

Table 6. SNP array data corresponding to EGFR/HER2/MET copy number in the indicated cell lines.

EGFR SNP array data

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
KYSE-520	6.718733	6.741493429	6.718733	6.660314	6.660314	6.718733	6.718733	6.718733	6.718733	6.44475	6.44475
OE21	6.595853	6.635474776	6.711752	6.711752	6.711752	6.595853	6.595853	6.542427	6.542427	6.538132	6.538132
H3255	6.410685	6.354872816	6.740301	6.60731	6.60731	6.60731	6.60731	6.740301	6.740301	6.410685	6.410685
NCI-H1573	6.040599	6.045958408	6.283974	6.045694	6.045694	6.040599	6.040599	6.040599	6.040599	5.823272	5.823272
HSC-2	5.94195	5.813092776	6.042633	5.94195	5.94195	6.042633	6.042633	6.042633	6.042633	5.94195	6.042633
HCC-827	5.624093	5.535087755	5.64778	5.64778	5.64778	5.749433	5.892709	5.778581	5.892709	5.778581	5.817453
Hs 852.T	5.523843	5.50758651	5.540834	5.540834	5.540834	5.523843	5.523843	5.523843	5.523843	5.428645	5.523843
KYSE-30	5.370001	5.336481449	5.504307	5.504307	5.504307	5.528331	5.528331	5.528331	5.528331	5.504307	5.428478
A431	5.251607	5.025420796	5.31399	5.31399	5.31399	5.401314	5.401314	5.251607	5.251607	4.952981	4.952981
BT-20	5.150625	5.128556102	5.198636	4.99139	4.99139	5.319849	5.319849	5.319849	5.319849	5.212884	5.212884
Ca9-22	4.671795	4.575755531	4.671795	4.671795	4.671795	4.893767	4.893767	4.893767	4.749979	4.671795	4.671795
NCI-H1568	4.362549	4.422823857	4.40939	4.508003	4.508003	4.621867	4.621867	4.621867	4.508003	4.621867	4.621867
PC-9	4.307727	4.364724714	4.41178	4.41178	4.41178	4.42986	4.42986	4.781583	4.42986	4.42986	4.621809
PNT1A	4.25082	4.194447653	4.258922	4.274299	4.274299	4.482283	4.482283	4.482283	4.258922	4.258236	4.258236
LUDLU-1	4.218594	4.141259612	4.269861	4.28757	4.28757	4.269861	4.269861	4.269861	4.075545	4.021817	4.021817
RERF-LC-MA	4.014391	3.983236102	3.874689	3.874689	3.874689	3.992777	4.064202	4	4	4	4
U373 MG	3.932433	3.789928306	3.999589	3.906136	3.906136	3.999589	4.048313	4.073112	4.048313	3.906136	4.048313
KYSE-450	3.869267	3.909038163	3.866752	3.866752	3.866752	3.869267	3.869267	3.917789	3.866752	3.815796	3.815796
KOSC-2 cl3-43	3.853418	3.66012851	3.871591	3.786779	3.786779	3.871591	4.047412	3.871591	3.853418	3.80363	3.80363
LU65	3.688177	3.603873857	3.688177	3.688177	3.688177	3.688177	3.719899	3.719899	3.688177	3.701103	3.701103
PC-14	3.68137	3.683669429	3.961857	3.961857	3.961857	3.961857	3.961857	3.961857	3.922973	3.906605	3.902885
NCI-H596	3.6606	3.72199549	3.92666	3.92666	3.92666	3.92666	3.92666	4.050986	3.92666	3.92666	3.653337
PC-3 [JPC-3]	3.647499	3.663076592	3.556417	3.556417	3.556417	3.655638	3.790605	3.790605	3.655638	3.767242	3.790605
647-V	3.633134	3.470668102	3.645379	3.645379	3.645379	3.645379	3.694535	3.645379	3.694535	3.694535	3.657458
HSC-3	3.570758	3.469518286	3.609322	3.609322	3.609322	3.635909	3.635909	3.635909	3.613514	3.613514	3.609322
CASKI	3.548381	3.565312755	3.518584	3.518584	3.512398	3.512398	3.536453	3.512398	3.512398	3.480862	3.480862
JHH-4	3.541971	3.443115082	3.72645	3.449344	3.601622	3.72645	3.72645	3.601622	3.601622	3.601622	3.655199
CAL-85-1	3.516086	3.438231571	3.629782	3.598705	3.598705	3.598705	3.629782	3.598705	3.598705	3.638944	3.638944
BT-549	3.510869	3.393711531	3.441616	3.441616	3.441616	3.489373	3.510869	3.495127	3.495127	3.510869	3.510869
HSC-4	3.450828	3.365350612	3.26703	3.26703	3.26703	3.287011	3.468548	3.480836	3.480836	3.480836	3.480836
NUGC-4	3.444932	3.439893429	3.581544	3.581544	3.581544	3.622942	3.622942	3.622942	3.622942	3.711979	3.743687
BICR10	3.436105	3.404563714	3.453907	3.267523	3.418784	3.453907	3.453907	3.418784	3.418784	3.418784	3.418784
KYSE-140	3.423342	3.430674939	3.431471	3.431471	3.431471	3.397922	3.397922	3.383845	3.383845	3.383845	3.500964
SNU-5	3.418943	3.358147245	3.465719	3.465719	3.465719	3.469476	3.469476	3.495591	3.469476	3.469476	3.420266
SK-MEL-3	3.41406	3.467373469	3.489846	3.524579	3.524579	3.549528	3.549528	3.549528	3.524579	3.549528	3.524579
KYSE-180	3.391391	3.375015286	3.582653	3.582653	3.582653	3.582653	3.582653	3.582653	3.306478	3.306478	3.204192
PNT2	3.387888	3.397246694	3.405847	3.562451	3.457522	3.405847	3.457522	3.578741	3.578741	3.457522	3.457522
G-361	3.372822	3.363805449	3.31356	3.316807	3.316807	3.372822	3.372822	3.372822	3.372822	3.372822	3.372822
KYSE-220	3.322587	3.368848184	3.244652	3.292362	3.292362	3.292362	3.292362	3.369054	3.369054	3.865377	3.865377
WM-115	3.272539	3.263409122	3.177676	3.179853	3.305193	3.393857	3.393857	3.436052	3.436052	3.436052	3.436052
KYSE-50	3.266165	3.222311694	3.209538	3.209538	3.209538	3.333031	3.333031	3.395734	3.333031	3.395734	3.395734
OSC-20	3.247106	3.134442612	2.979487	2.979487	3.108146	3.342224	3.342224	3.342224	3.132013	3.132013	3.132013

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
SAT	3.241407	3.153325755	3.206196	3.206196	3.210638	3.290546	3.290546	3.241407	3.241407	3.210638	3.210638
NCI-H1650	3.226752	3.218400714	3.83591	3.83591	3.83591	3.836849	3.836849	3.836849	3.83591	3.751073	3.518346
SK-CO-1	3.214665	3.265122184	3.502499	3.502499	3.502499	3.566612	3.652499	3.725649	3.566612	3.566612	3.566612
NCI-H1792	3.210818	3.195864449	3.37014	3.37014	3.37014	3.37014	3.691703	3.692449	3.691703	3.37014	3.229938
NCI-H1703	3.19572	3.149808551	3.16358	3.229891	3.229891	3.229891	3.229891	3.187826	3.16358	3.16358	3.16358
PCI 4B	3.171588	3.167475531	3.178856	3.237981	3.171588	3.237981	3.237981	3.237981	3.237981	3.171588	3.171588
NCI-H1563	3.156902	3.015406714	3.156902	3.156902	3.156902	3.241539	3.283369	3.301327	3.283369	3.283369	3.283369
NCI-H841	3.132091	3.048329694	3.132091	3.132091	3.132091	3.132091	3.132091	3.132091	3.132091	3.132091	3.132091
LOU-NH91	3.111632	3.083197041	3.091937	3.150625	3.150625	3.245861	3.383068	3.245861	3.217618	3.189097	3.189097
HuH-7	3.09566	2.977372102	3.007807	3.007807	3.007807	3.034269	3.09566	3.09566	3.034269	3.034269	3.034269
LC-1 sq	3.090563	2.950721367	3.175325	2.958327	3.090563	3.175325	3.191389	3.175325	3.090563	3.090563	3.090563
NCI-H522	3.088288	3.056570878	3.059922	3.059922	3.059922	3.059922	3.146846	3.169511	3.169511	3.169511	3.169511
201T	3.080531	2.957688367	3.09712	3.09712	3.09712	3.09712	3.162905	3.206871	3.206871	3.206871	3.09712
G-292 Clone A141B1	3.079211	3.077906061	3.101378	3.101378	3.101378	3.101378	3.101378	3.101378	3.021934	2.996675	2.996675
RPMI-7951	3.075119	3.038721531	3.041106	3.070971	3.070971	3.041106	3.084605	3.090128	3.090128	3.084605	3.084605
BEN	3.072106	3.005211204	3.072106	3.072106	3.072106	3.082038	3.160291	3.160291	3.160291	3.169925	3.189097
NCI-H2228	3.055512	3.015296918	3.055512	3.055512	3.055512	3.039728	3.055512	3.055512	3.039728	3.055512	3.324858
HCC-44	3.045739	2.911290592	3.05383	3.055292	3.055292	3.161227	3.222571	3.222571	3.161227	3.161227	3.161227
NCI-H1734	3.045608	3.014053653	2.97999	2.97999	3.084288	3.084288	3.084288	3.084288	3.049612	3.049612	3.049612
YKG-1	3.033795	2.986565571	3.418528	3.343058	3.343058	3.418528	3.418528	3.418528	3.418528	3.343058	3.343058
MOG-G-CCM	3.031173	3.081210755	3.170906	3.170906	3.170906	3.202471	3.232565	3.317986	3.202471	3.471859	3.520868
SK-BR-3	3.021924	2.964034673	3.227063	3.189097	3.189097	3.227063	3.328966	3.227063	3.227063	3.12143	3.12143
COLO 858	3.014686	3.064249245	3.109089	3.109089	3.167885	3.338335	3.338335	3.338335	3.267048	3.167885	3.167885
SCC-4	2.995215	3.022357388	3.394952	3.20894	3.394952	3.433174	3.609181	3.591725	3.521353	3.433174	3.433174
TE7	2.973552	2.907018959	3.036184	2.89907	2.89907	3.036184	3.036184	2.999687	2.999687	2.94022	2.999687
NCI-H1755	2.972533	2.915477939	3.024049	3.024049	3.072046	3.092628	3.092628	3.092628	3.072046	3.080074	3.092628
CaR-1	2.967637	2.952000245	2.94097	2.94097	2.948934	2.974674	2.974674	2.948934	2.94097	2.94097	2.948934
OVISe	2.964705	2.882252347	2.764106	2.764106	2.983398	3.035317	3.035317	2.983398	2.983398	2.952857	2.952857
Calu-3	2.960385	2.960519143	3.160928	3.160928	3.160928	3.142402	3.142402	3.142402	3.065731	3.065731	3.065731
NCI-H1623	2.952411	2.902007673	3.06155	3.034219	3.035015	3.034219	3.034219	3.034219	3.034219	3.034219	3.035015
BICR 78	2.950678	3.025212449	2.929577	2.929577	2.929577	3.230541	3.230541	3.230541	3.230541	3.445063	3.445063
LS 123	2.946317	2.916922939	3.045046	3.237154	3.237154	3.045046	3.237154	3.260578	3.237154	2.98877	2.979846
AN3CA	2.941199	2.922365857	2.62018	2.62018	2.653454	2.903978	2.903978	2.962557	2.962557	3.055782	3.055782
639-V	2.937969	2.807446408	2.937969	2.91964	2.937969	2.993201	2.993201	2.993201	2.981715	2.981715	2.981715
WM 266-4	2.930068	2.874593204	2.855266	2.855266	2.901953	3.057063	3.057063	3.11125	3.11125	3.054361	3.054361
SKN-3	2.928416	2.813150816	2.961006	2.961006	2.961006	3.125017	3.125017	3.125017	2.961006	2.961006	2.961006
NCI-H2085	2.928009	2.947168959	3.019797	3.019797	3.019797	3.019797	3.06164	3.175959	3.06164	3.113084	3.113084
RKN	2.92781	2.754522816	2.78781	2.78781	2.789445	3.049802	3.049802	3.049802	3.049802	3.049893	3.049893
MKN45	2.917551	2.898736	2.926439	2.926439	2.906593	3.00802	3.00802	3.145925	3.00802	2.898922	3.00802
KHOS/NP	2.902185	2.938604816	3.250765	3.250765	3.250765	3.292786	3.296351	3.296351	3.292786	3.250765	3.156554
SNU-16	2.899323	2.904760367	2.922701	2.922701	2.876703	2.876703	3.004233	3.03699	3.004233	3.004233	2.876703
HCE7	2.890214	2.745430184	2.486753	2.761115	2.832555	2.973748	2.978346	2.973748	2.973748	2.973748	2.973748
OCUG-1	2.888539	2.844002143	2.940095	2.940095	2.940095	2.970029	2.970029	2.940095	2.888539	2.888539	2.836196
NL20-TA	2.883583	2.820962367	2.542325	2.593667	2.542325	2.78182	2.876767	2.93947	2.93947	2.93947	2.93947
KYSE-70	2.866528	2.846896367	2.867169	2.867169	2.867169	3.02882	2.867169	2.867169	2.759153	2.740295	2.866528
MOG-G-UVW	2.865042	2.869016898	2.937552	3.087882	3.087882	3.087882	3.129781	3.263475	3.087882	2.859703	2.948934
NCI-H2342	2.860635	2.878060735	2.894629	2.894629	2.894629	2.894629	2.894629	2.894629	2.884125	2.860635	2.860635

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Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
ABC-1	2.855212	2.86351	2.811985	2.811985	2.811985	2.897802	2.897802	2.95017	2.95017	3.072106	3.140926
NCI-H650	2.854933	2.829436796	2.925675	2.925675	2.925675	2.925675	3.000187	3.142528	3.000187	2.925675	2.794719
FTC 133	2.854504	2.811006531	2.813128	2.813128	2.813128	2.863257	2.863257	2.863257	2.813128	2.75963	2.648858
MDST8	2.853753	2.828383551	2.853753	2.936228	2.925822	2.936228	2.936228	2.925822	2.913045	2.853753	2.853753
KYSE-150	2.852089	2.865999061	2.975281	2.975281	2.975281	2.966574	2.864069	2.864069	2.854032	2.854032	2.854032
SW 48	2.851531	2.832725265	2.859403	2.915252	2.915252	2.915252	2.915252	2.915252	2.896681	2.896681	2.896681
NCI-H2023	2.849425	2.815705388	2.836553	2.836553	2.86163	2.86163	2.86163	3.029658	2.86163	3.029658	3.029658
COLO-680N	2.846328	2.893982776	2.837675	2.837675	2.994213	3.063936	3.106593	3.295467	3.295467	3.578991	3.578991
KP-4	2.8349	2.716085265	2.976588	2.976588	2.976588	3.113457	3.113457	3.113457	2.976588	2.976588	2.976588
SW-1710	2.828271	2.801945388	3.005303	2.990623	3.005303	3.025081	3.025081	3.005303	3.025081	3.025081	3.025081
KATO III	2.813345	2.646816	2.991032	2.991032	2.991032	2.991032	3.003021	3.003021	2.991032	2.991032	2.991032
RERF-GC-1B	2.812758	2.764795	2.737425	2.737425	2.737425	2.871492	2.871492	2.871492	2.741277	2.741277	2.741277
CAL-29	2.811331	2.750759939	2.915294	2.915294	2.915294	3.045708	3.270877	3.270877	3.045708	2.915294	2.844175
Sci-1	2.804648	2.734233612	2.829592	2.829592	2.829592	2.845424	2.845424	2.845424	2.829592	2.81144	2.81144
NCI-H810	2.800201	2.807169837	2.58053	2.750768	2.750768	2.753248	2.753248	2.921559	2.921559	2.921559	2.753248
T.Tn	2.782865	2.72791102	2.796679	2.796679	2.782865	2.842624	2.842624	2.842624	2.782865	2.782865	2.782865
DOK	2.781798	2.813705898	3.154514	3.154514	3.090197	3.090197	3.090197	3.158389	3.090197	3.029083	2.992669
COLO 741	2.773798	2.771708367	2.771823	2.860753	2.860753	2.906066	2.906066	2.965542	2.906066	2.906066	2.906066
NCI-H1651	2.773677	2.673708429	2.773677	2.734753	2.773677	2.734753	2.773677	2.786129	2.786129	2.786129	2.786129
RWPE-1	2.768763	2.737865061	2.857614	2.788447	2.80242	2.788447	2.80242	2.788447	2.788447	2.788447	2.788447
KYSE-270	2.762888	2.689595388	2.767204	2.767204	2.839423	2.839423	2.839423	2.767204	2.839423	2.839423	2.839423
NCI-H196	2.748477	2.771432061	2.891337	2.982341	2.982341	2.982341	3.013204	3.013204	2.982341	2.982341	2.982341
SK-LU-1	2.741065	2.782619918	2.925497	3.011172	3.011172	3.240891	3.240891	3.256483	2.925497	2.925497	2.925497
NCI-H1693 (ship16)	2.73986	2.68732498	2.612233	2.73986	2.831701	2.73986	2.73986	2.831701	2.831701	2.826722	2.831701
NCI-H2135	2.738285	2.70914298	2.6742	2.564037	2.652971	2.564037	2.652971	2.652971	2.738285	2.738285	2.738285
VMRC-RCW	2.732449	2.653201673	3.060821	2.743508	2.865074	3.060821	3.060821	3.003877	2.865769	2.865769	2.865769
RERF-LC-Sq1	2.732214	2.740655163	2.767757	2.767757	2.767757	2.767757	2.959682	2.959682	2.767757	2.959682	2.959682
A375.S2	2.732202	2.769671592	2.730075	2.730075	2.730075	2.950606	2.964561	3.026759	2.964561	2.964561	2.950606
SW 1116	2.729671	2.775940694	2.992587	3.005242	3.005242	3.005242	3.005242	3.060481	3.060481	2.887298	2.887298
CAL-148	2.728069	2.636859184	2.596894	2.56399	2.596894	2.596894	2.811397	2.811397	2.811397	2.811397	2.891453
22RV1	2.727004	2.711221531	2.847447	2.847447	2.847447	2.847447	2.89983	2.89983	2.847447	2.89983	2.89983
IM-95	2.726768	2.714460102	2.833777	2.833777	2.833777	2.833777	2.833777	2.829516	2.829516	2.829516	2.829516
HT 115	2.724872	2.716082633	2.592551	2.592551	2.592551	2.724872	2.724872	2.592551	2.724872	2.592551	2.592551
NCI-H727	2.721212	2.703270735	2.80183	2.926523	2.926523	2.926523	2.926523	3.057193	2.926523	3.13694	2.926523
T.T	2.718233	2.706004306	2.687827	2.687827	2.687827	2.705201	2.705201	2.687827	2.705201	2.705201	2.705201
OE33	2.71355	2.719964857	2.875926	2.875926	2.875926	2.875926	2.875926	2.875926	2.837395	2.837395	2.740764
PLC/PRF/5	2.711645	2.673463837	2.541118	2.624064	2.656547	2.753014	2.656547	2.656547	2.656547	2.624064	2.656547
SAS	2.711081	2.608931592	3.10791	2.507132	2.704398	3.10791	3.10791	2.78525	2.78525	2.78525	2.883434
MKN74	2.706985	2.702259306	2.897369	2.897369	2.897369	2.95465	2.95465	2.975467	2.95465	2.897369	2.809501
HMCB	2.703291	2.678871612	2.698912	2.698912	2.910783	2.910783	2.910783	2.880916	2.880916	2.698912	2.490078
A2058	2.689237	2.609137531	2.536414	2.536414	2.625191	2.682503	2.930779	2.682503	2.930779	2.930779	2.932231
HCC-366	2.673229	2.556834857	2.708656	2.759275	2.759275	2.708656	2.83023	2.83023	2.708656	2.83023	2.83023
P31/FUJ	2.672578	2.625101571	2.793163	2.786745	2.786745	2.793163	2.793163	2.793163	2.786745	2.786745	2.78029
SW 1417	2.672327	2.65521902	2.807637	2.830392	2.830392	2.830392	2.933129	2.933129	2.830392	2.830392	2.814434
KMRC-1	2.666206	2.539772939	2.601747	2.374636	2.601747	2.666206	2.742515	2.742515	2.666206	2.601747	2.514323
OAW42	2.664958	2.659155429	2.633763	2.633763	2.633763	2.634261	2.70441	2.853506	2.759708	2.814912	2.814912
PE/CA-PJ15	2.662264	2.639397531	2.730814	2.730814	2.730814	2.730814	2.730814	2.730814	2.702872	2.662264	2.662264

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
NUGC-3	2.658408	2.663259469	2.628256	2.628256	2.710168	2.837405	2.837405	2.854998	2.854998	2.854998	2.837405
DMS 273	2.646371	2.609157347	2.814869	2.814869	2.814869	2.814869	2.890479	3.102323	2.814869	2.814869	2.759109
MKN1	2.642413	2.622494796	2.726903	2.726903	2.726903	2.803272	2.803272	2.803272	2.659877	2.659877	2.659877
RERF-LC-Ad2	2.634979	2.642198796	2.628105	2.628105	2.628105	2.767536	2.767536	2.767536	2.642171	2.642171	2.642171
TASK1	2.628593	2.500382918	2.533021	2.377245	2.533021	2.942739	2.942739	2.628593	2.628593	2.628593	2.729828
NCI-H23	2.62612	2.60171351	2.738743	2.738743	2.804058	2.804058	2.870991	2.952877	2.870991	2.804058	2.804058
KP-1N	2.623518	2.460675347	2.324974	2.324974	2.325065	2.678012	2.890458	2.678012	2.678012	2.678012	2.678012
NCI-H711	2.622425	2.586977082	2.399112	2.399112	2.58007	2.702985	2.702985	2.702985	2.648688	2.702985	2.702985
NCI-H661	2.621297	2.613435184	2.546877	2.546877	2.648216	2.648216	2.648216	2.648216	2.600904	2.648216	2.709604
EBC-1	2.62068	2.586313082	2.477574	2.477574	2.538066	2.666847	2.757023	2.757023	2.757023	2.757023	2.757023
JHH-6	2.616256	2.50211402	2.371619	2.228894	2.371619	2.808858	2.808858	2.752892	2.570852	2.570852	2.570852
NB69	2.609723	2.623469163	2.502446	2.609723	2.609723	2.609723	2.609723	2.609723	2.558847	2.558847	2.558847
SR	2.607445	2.643014612	2.788897	2.788897	2.788897	2.810405	3.132383	3.270246	3.132383	2.810405	2.810405
T84	2.604509	2.598727551	2.678526	2.671435	2.678526	2.712197	2.884454	2.712197	2.712197	2.671435	2.604509
HO-1-N-1	2.6041	2.427990837	2.556993	2.310372	2.556993	2.6041	2.727329	2.6041	2.6041	2.6041	2.625226
BALL-1	2.602836	2.603702612	2.388527	2.388527	2.473426	2.753936	2.753936	2.65773	2.65773	2.65773	2.65773
Nthy-ori 3-1	2.601196	2.610085694	2.37134	2.601196	2.601196	2.820574	2.820574	2.820574	2.671218	2.671218	2.671218
FTC 238	2.599685	2.571267122	2.460258	2.417234	2.460258	2.645045	2.645045	2.645045	2.417234	2.417234	2.599685
HuCCT1	2.595263	2.551327673	2.235536	2.227308	2.235536	2.742359	2.742359	2.742359	2.742359	2.742359	2.742359
HOS	2.591082	2.549625918	2.463319	2.463319	2.608742	2.672087	2.701515	2.701515	2.701515	2.694538	2.672087
NCI-H2170	2.587636	2.593833245	2.821442	2.821442	2.821442	2.581167	2.587636	2.826661	2.587636	2.587636	2.581167
Hs 936.T	2.583749	2.579530408	2.691758	2.743999	2.691758	2.571704	2.691758	2.739257	2.706782	2.739257	2.739257
NCI-H1793	2.582747	2.599235592	2.602953	2.605831	2.602953	2.573195	2.602953	2.602953	2.573195	2.796318	2.796318
MRK-nu-1	2.574874	2.526067612	2.483876	2.389763	2.483876	2.574874	2.574874	2.574874	2.574874	2.574874	2.587859
KHOS-312H	2.569064	2.573914143	2.780444	2.780444	2.780444	2.846436	2.924953	2.846436	2.846436	2.846436	2.709886
LCLC-103H	2.564998	2.510750082	2.624459	2.624459	2.624459	2.80349	2.80349	2.80349	2.720639	2.720639	2.720639
H69V	2.559905	2.697188429	2.791233	2.873464	2.873464	2.921265	2.921265	3.262218	2.873464	2.873464	2.873464
GCT	2.558752	2.501207816	2.478173	2.478173	2.478173	2.478173	2.478637	2.564014	2.478637	2.510651	2.510651
COLO 857	2.554101	2.516956816	2.60894	2.60894	2.60894	2.695422	2.695422	2.695422	2.60894	2.508547	2.508547
RERF-LC-MS	2.550014	2.554352776	2.453017	2.453017	2.655375	2.745904	2.745904	2.757023	2.655375	2.757023	2.757023
MKN7	2.538066	2.503053918	2.585563	2.585563	2.585563	2.585563	2.585563	2.585563	2.573762	2.573762	2.573762
A-427	2.534808	2.352632184	2.236281	2.064046	2.496985	2.573348	2.61572	2.573348	2.61572	2.496985	2.496985
T47D	2.533249	2.55140051	2.487338	2.449599	2.487338	2.541919	2.568306	2.568306	2.541919	2.541919	2.541919
BT-483	2.515964	2.554997571	2.642471	2.642471	2.642471	2.754181	2.754181	2.791409	2.754181	2.791409	2.604662
SKN	2.509067	2.411093816	2.222503	2.222503	2.576162	2.604966	2.773313	2.773313	2.773313	2.773313	2.773313
HCC202	2.504674	2.497351061	2.504674	2.504674	2.504674	2.604674	2.770288	2.770288	2.770288	2.729167	2.604673
OCUM-1	2.501923	2.496531714	2.609022	2.609022	2.609022	2.745904	2.745904	2.609022	2.573762	2.573762	2.609022
NCI-H2122	2.492184	2.453601163	2.469383	2.483351	2.483351	2.483351	2.617863	2.617863	2.483351	2.617863	2.617863
HT 29	2.48307	2.492675531	2.564061	2.564061	2.564061	2.689316	2.564061	2.480274	2.564061	2.362905	2.25788
huH-1	2.480336	2.400045388	2.58402	2.477488	2.477488	2.58402	2.658718	2.58402	2.58402	2.58402	2.58402
HCC-15	2.478943	2.454757265	2.460972	2.460972	2.460972	2.460972	2.478943	2.478943	2.478943	2.478943	2.478943
KHOS-240S	2.473022	2.427126306	2.648008	2.648008	2.648008	2.648008	2.659705	2.659705	2.648008	2.624087	2.46048
JIMT-1	2.467801	2.416077633	2.468733	2.273026	2.468733	2.284239	2.468733	2.473781	2.473781	2.473781	2.473781
KON	2.465788	2.393919245	2.600013	2.24814	2.649183	2.707968	2.707968	2.649183	2.603035	2.595451	2.595451
COLO 853	2.457194	2.42537798	2.383239	2.383239	2.435231	2.46515	2.46515	2.46515	2.435231	2.46515	2.53724
U-2 OS	2.453017	2.464540571	2.261862	2.261862	2.403268	2.561913	2.585563	2.585563	2.666847	2.700994	2.700994
ME-180	2.448612	2.296262408	2.375497	2.367621	2.375497	2.683015	2.693267	2.683015	2.650727	2.683015	2.683015

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
FU97	2.440152	2.394430122	2.104192	2.104192	2.528689	2.528689	2.60769	2.60769	2.528689	2.459828	2.459828
Calu-1	2.423993	2.393085122	2.324252	2.393302	2.417272	2.4839	2.5783	2.588071	2.5783	2.5783	2.5783
NCI-H2452	2.420291	2.388003878	2.362829	2.420291	2.420291	2.420291	2.420291	2.420291	2.420291	2.514239	2.514239
UMC-11	2.416248	2.343465367	2.802999	2.301574	2.623343	2.623343	2.701866	2.701866	2.701866	2.701866	2.701866
COLO 684	2.404058	2.400842408	2.585281	2.428759	2.428759	2.32184	2.550408	2.550408	2.550408	2.404058	2.404058
SVts-8	2.4032677	1.88929752	2.7680996	2.7680996	2.7901256	2.8228525	2.8872133	2.8872133	2.8228525	2.8228525	2.8872133
HEC-1	2.399539	2.336459061	2.145639	2.06111	2.06111	2.145639	2.465383	2.465383	2.465383	2.385323	2.616465
A-375	2.397768	2.300920306	2.208942	2.208942	2.208942	2.338028	2.420404	2.420404	2.420404	2.502141	2.502141
OUMS-23	2.396737	2.337490898	2.666206	2.462336	2.462336	2.577461	2.577461	2.577461	2.462336	2.462336	2.462336
EJ138	2.393516	2.434952878	2.683971	2.683971	2.683971	2.683971	2.683971	2.700168	2.683971	2.659659	2.659659
HO-1-u-1	2.39266	2.279093204	2.697292	2.068276	2.697292	2.697292	2.697292	2.697292	2.39266	2.388376	2.39266
HT55	2.391842	2.421966204	2.7445	2.770409	2.770409	2.770409	2.770409	2.770409	2.588871	2.533417	2.530346
NCI-H2347	2.388061	2.384280531	2.204886	2.53458	2.53458	2.53458	2.53458	2.53458	2.53458	2.53458	2.53458
RERF-LC-KJ	2.383365	2.438087082	2.664935	2.664935	2.664935	2.664935	2.736866	2.736866	2.664935	2.664935	2.664935
COR-L279	2.366668	2.336264245	2.473414	2.519642	2.519642	2.57726	2.685838	2.730724	2.730724	2.473414	2.590129
SW 1573	2.36377	2.233322878	2.317359	2.315963	2.315963	2.317359	2.374889	2.36377	2.36377	2.36377	2.36377
SW 900	2.356754	2.365797694	2.499934	2.499934	2.499934	2.441762	2.499934	2.499934	2.499934	2.441762	2.441762
HT 1376	2.351215	2.331488429	2.524024	2.56846	2.56846	2.56846	2.56846	2.395391	2.377663	2.377663	2.377663
HUP-T3	2.347342	2.241781898	2.312741	2.155293	2.312741	2.347342	2.428274	2.347342	2.428274	2.428274	2.428274
STC 1	2.34628	2.253264184	2.307312	2.307312	2.307312	2.35359	2.398622	2.398622	2.35359	2.35359	2.307312
NCI-H2286	2.341299	2.342265041	2.220807	2.220807	2.220807	2.311874	2.311874	2.311874	2.290037	2.290037	2.290037
VMRC-LCD	2.33985	2.341863837	2.352646	2.403268	2.403268	2.403268	2.44066	2.44066	2.465322	2.465322	2.643856
H69AR	2.338349	2.301531918	2.313453	2.356805	2.356805	2.356805	2.356805	2.478674	2.356805	2.356805	2.480726
VMRC-LCP	2.33487	2.236303122	2.419668	2.139019	2.32853	2.419668	2.445869	2.419668	2.445869	2.445869	2.445869
FLYA13	2.321944	2.322221102	2.595979	2.655294	2.655294	2.595979	2.595979	2.584833	2.4576	2.4576	2.4576
HUP-T4	2.308596	2.245841469	2.248246	2.248246	2.308596	2.452117	2.60541	2.452117	2.308596	2.308596	2.308596
COR-L 105	2.307818	2.293109898	2.532698	2.532698	2.18708	2.532698	2.532698	2.507664	2.47618	2.326598	2.326598
KMRC-20	2.302951	2.292869449	2.302951	2.302951	2.302951	2.370223	2.380952	2.370223	2.370223	2.302951	2.302951
NCI-H2081	2.295681	2.298436918	2.213956	2.213956	2.395089	2.395089	2.395089	2.491758	2.395089	2.491758	2.617456
VM-CUB1	2.293832	2.297199245	2.308441	2.618364	2.489336	2.489336	2.489336	2.618364	2.489336	2.403029	2.344732
BU25 TK	2.29009	2.329399041	2.4729	2.491685	2.496365	2.608052	2.608052	2.638276	2.608052	2.608052	2.608052
RCM-1	2.287897	2.188751082	2.249953	2.249953	2.319581	2.523122	2.555577	2.523122	2.319581	2.319581	2.319581
NCI-H2172	2.284971	2.277862041	2.284971	2.284971	2.284971	2.244245	2.284971	2.284971	2.284971	2.351752	2.351752
SK-NEP-1	2.283689	2.267003837	2.101561	2.057959	2.101561	2.101561	2.27481	2.101561	2.101561	2.101561	2.101561
D1.1	2.281505	2.266256898	2.093471	2.120455	2.120455	2.127241	2.127241	2.398446	2.398446	2.375801	2.375801
NCI-H2196	2.280615	2.285093878	2.283533	2.283533	2.283533	2.455285	2.455285	2.394737	2.455285	2.455285	2.394737
NCI-H1435	2.276856	2.193842163	2.170972	2.170972	2.170972	2.170972	2.240571	2.240571	2.240571	2.298246	2.298246
GRANTA-519	2.27561	2.147157163	2.087669	2.12228	2.129007	2.347419	2.359009	2.347419	2.347419	2.359009	2.413474
SK-MES	2.271936	2.264511735	2.083383	2.108336	2.232676	2.480983	2.480983	2.232676	2.232676	2.232676	2.232676
NCI-H1437	2.264798	2.262715878	2.295942	2.416747	2.416747	2.416747	2.417721	2.416747	2.393088	2.393088	2.393088
BFTC-905	2.259674	2.269937816	2.421089	2.096319	2.418183	2.421089	2.539694	2.539694	2.421089	2.418183	2.418183
DMS 153	2.258751	2.204225408	2.045716	2.045716	2.14218	2.14218	2.267074	2.267074	2.289711	2.507531	2.289711
Detroit 562	2.256824	2.266237776	2.427068	2.427068	2.484998	2.484998	2.427068	2.427068	2.253734	2.228054	2.228054
RT4	2.246948	2.22294298	2.38195	2.38195	2.38195	2.38195	2.405888	2.405888	2.38195	2.345052	2.38195
273T	2.238925	2.232078265	2.164192	2.171651	2.164192	2.269664	2.269664	2.282826	2.282826	2.282826	2.282826
SW 1463	2.238128	2.247130306	2.352377	2.352377	2.352377	2.417684	2.417684	2.288471	2.40639	2.40639	2.274574
H69	2.237583	2.252691306	2.237583	2.237583	2.237583	2.237583	2.237583	2.063214	2.061378	2.061378	2.061378

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
Hs 195.T	2.236919	2.20431902	1.974918	1.978002	1.991316	2.226788	2.503571	2.551683	2.427241	2.551683	2.575075
NCI-H1781	2.229587	2.225333224	2.20326	2.20326	2.229587	2.230879	2.357863	2.363032	2.363032	2.593749	2.593749
WIDR	2.223745	2.244019082	2.345858	2.345858	2.446586	2.451784	2.451784	2.451784	2.432761	2.335397	2.432761
SK-MEL-1	2.214036	2.254872041	2.593021	2.646106	2.646106	2.567915	2.567915	2.646106	2.567915	2.567915	2.567915
SUIT-2	2.212777	2.213396735	2.12033	2.12033	2.385234	2.462742	2.462742	2.385234	2.375763	2.375763	2.375763
WPE1-NB26	2.203354	2.140397327	2.342132	2.342132	2.342132	2.342132	2.364342	2.364342	2.342132	2.364342	2.364342
HCC-78	2.201821	2.246317673	2.124492	2.137768	2.137768	2.137768	2.354381	2.354381	2.354381	2.3738	2.562649
A549	2.201767	2.192650061	2.105569	2.141273	2.141273	2.253536	2.253536	2.301509	2.301509	2.301509	2.301509
LK-2	2.201363	2.149191898	2.201363	2.201363	2.201363	2.259753	2.266141	2.259753	2.094602	2.094602	2.083804
NCI-H2330	2.198132	2.180153082	2.088285	2.088285	2.088285	2.272724	2.366706	2.372707	2.367028	2.272724	2.366706
COLO 685	2.197742	2.163006286	2.883084	2.240014	2.240014	2.195289	2.240014	2.360028	2.360028	2.240014	2.360028
BICR22	2.184063	2.213391143	2.277157	2.277157	2.277157	2.379915	2.277157	2.277157	2.127751	2.127751	2.127751
MEWO	2.183021	2.180176102	2.124008	2.176337	2.176337	2.183021	2.183021	2.183021	2.183021	2.183021	2.183021
IHH-4	2.181084	2.159299041	2.326096	2.326096	2.326096	2.326096	2.326096	2.267456	2.267456	2.181084	2.181084
NCI-N87	2.180447	2.207232306	2.334677	2.334677	2.311848	2.341337	2.341337	2.341337	2.341337	2.238327	2.161493
KYSE-410	2.176975	2.200467918	2.320253	2.320253	2.320253	2.465678	2.465678	2.469616	2.274653	2.288106	2.288106
DMS 92	2.175971	2.099747673	2.144843	2.144843	2.175971	2.433468	2.433468	2.182343	2.175971	2.182343	2.182343
RO82-W-1	2.175428	2.149052918	2.302107	2.302107	2.302107	2.330576	2.330576	2.330576	2.330576	2.302107	2.226721
NCI-H82	2.170632	2.172188755	2.327036	2.327036	2.327036	2.327036	2.421663	2.488422	2.421663	2.277904	2.178562
C32	2.163402	2.076808429	1.849507	1.849507	2.146352	2.146352	2.146352	2.370287	2.125722	2.334536	2.334536
Hs 633T	2.159405	2.168496776	2.040955	2.172929	2.172929	2.172929	2.211745	2.211745	2.172929	2.172929	2.172929
NCI-H2405	2.148394	2.115157286	2.148394	2.151284	2.168414	2.151284	2.151284	2.168414	2.242163	2.168414	2.168414
Hs 895.T	2.141685	2.170353061	2.270676	2.270676	2.270676	2.270676	2.270676	2.364037	2.364037	2.364037	2.364037
MG 63	2.141685	2.148053612	1.905234	1.905234	2.018207	2.102925	2.102925	2.102925	2.018207	2.018207	2.102925
CAL-120	2.137493	1.99951251	2.140737	2.13664	2.137493	2.215269	2.243158	2.15269	2.168863	2.168863	2.168863
H33HJ-JA1	2.133418	2.156652918	2.164424	2.280929	2.280929	2.371835	2.426383	2.42912	2.42912	2.426383	2.426383
JSC-1	2.131158	2.157892816	2.212308	2.212308	2.287884	2.394611	2.476767	2.553089	2.476767	2.432177	2.432177
Hs 751.T	2.130414	2.105885837	2.00395	2.00395	2.015109	2.130414	2.344412	2.344412	2.130414	2.344412	2.344412
HMVII	2.1288	2.040526735	2.015769	2.015769	2.359086	2.359086	2.359086	2.383277	2.359086	2.245107	2.245107
fR2	2.124962	2.049262571	2.102354	2.102354	2.102354	2.118669	2.238912	2.238912	2.118669	2.118669	2.139019
COLO 201	2.124672	2.115652347	2.138703	2.138703	2.138703	2.208767	2.208767	2.208767	2.138703	2.208767	2.334947
NCI-H1915	2.119292	2.066165714	1.999986	2.105569	2.105569	2.21405	2.21405	2.21405	2.367519	2.21405	2.21405
IMR-90	2.117312	2.121663755	2.051588	1.892683	2.051588	2.525095	2.525095	2.423507	2.423507	2.423507	2.509708
RT112/84	2.115316	2.091732163	2.115316	2.115316	2.115316	2.35336	2.481325	2.481325	2.35336	2.35336	2.402916
Hs 894(E).Lu	2.11282	2.127957408	2.11282	2.262191	2.272501	2.262191	2.262191	2.272501	2.236773	2.236773	2.236773
LNCAp clone FGC	2.108322	2.093793735	2.032351	2.032351	2.059754	2.266719	2.266719	2.266719	2.108322	2.108322	2.108322
NCI-H1581	2.107682	2.072259918	2.147832	2.147832	2.147832	2.147832	2.11551	2.11551	2.111197	2.111197	2.111197
HCT 116	2.106237	2.085858327	2.106584	2.12423	2.12423	2.12423	2.12423	2.106584	2.055129	2.055129	2.042135
Hs 228.T	2.103719	2.11547302	2.076704	2.103719	2.076704	2.103719	2.226027	2.229014	2.226027	2.226027	2.229014
JURKAT EC-1	2.096947	2.077322082	2.096947	2.101254	2.101254	2.101254	2.096947	2.053444	2.096947	2.053444	2.096947
SK-N-BE(2)	2.096877	2.108693571	2.018135	2.018135	2.018135	2.018135	2.042718	2.042718	2.042718	2.158722	2.332968
LS180	2.093401	2.094016735	2.177654	2.177654	2.198335	2.198335	2.177654	2.177654	2.119015	2.070122	2.070122
CAL-51	2.092144	2.041400265	1.867583	1.871266	1.915502	2.202883	2.202883	2.185673	1.915502	2.185673	2.202883
SKG-IIIb	2.087795	1.906490816	2.072979	2.072979	2.087795	2.412624	2.412624	2.368941	2.087795	2.087795	2.087795
P116 T	2.084434	2.078256776	1.908914	2.009894	2.076353	2.076353	2.076353	2.151996	2.151996	2.009894	2.009894
Hs 417.Lu	2.082963	2.080747714	2.054988	2.054988	2.054988	2.082963	2.082963	2.082963	2.082963	2.29185	2.346984
WIL2 NS	2.081925	2.086687388	1.969631	1.969631	2.228521	2.24146	2.24146	2.24146	2.154161	2.24146	2.268021

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-2119130	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
COR-L95	2.081027	2.103553673	2.076198	2.168197	2.303146	2.303146	2.303146	2.385184	2.303146	2.208472	2.208472
SK-N-FI	2.08041	2.053238265	1.870829	2.029739	1.984551	2.093681	2.093681	2.093681	2.093681	2.093681	2.093681
WILCL	2.077982	2.059245714	1.977159	1.977159	2.007924	2.007924	2.027753	2.027753	2.027753	2.027753	2.26547
P32/ISH	2.07364	2.066828878	2.362689	1.960957	2.304185	2.362689	2.362689	2.304185	2.271306	2.108613	2.108613
EPLC-272H	2.069516	2.019263898	1.742631	1.681692	1.742631	2.269664	2.343272	2.269664	2.303211	2.303211	2.343272
A2780 ADR	2.066684	2.033308184	2.001832	2.031109	2.055143	2.194574	2.055143	2.194574	2.194574	2.055143	2.194574
C8166	2.062084	2.098756184	2.101226	2.101226	2.286212	2.286212	2.286212	2.390431	2.286212	2.232051	2.232051
Hs 908.T	2.060333	2.083260286	2.132178	2.132178	2.132178	2.073232	2.146599	2.146599	2.146599	2.146599	2.146599
NCI-H322	2.059372	2.052896837	2.085037	2.130634	2.130634	2.130662	2.173174	2.173174	2.173174	2.173174	2.130662
NCI-H1417	2.059316	2.01582951	2.205061	2.205061	2.205061	2.205061	2.222784	2.222784	2.690373	2.690373	2.754869
HBE4-E6/E7	2.058524	2.079743102	1.662684	1.944327	1.968274	2.180325	2.293024	2.180325	2.180325	2.066331	2.180325
NCI-H1522	2.058284	2.015362796	2.164261	2.16851	2.28902	2.16851	2.227654	2.227654	2.16851	2.16851	2.16851
LU99A	2.057138	2.072812837	2.071248	2.071248	2.085289	2.195222	2.195222	2.195222	2.195222	2.195222	2.195222
HCT-8	2.054634	2.041630286	2.022605	2.152872	2.133997	2.133997	2.133997	2.269244	2.269244	2.269244	2.269244
SBC-5	2.052538	2.032448408	1.938128	1.938128	2.091221	2.132123	2.132123	2.091221	2.068065	2.055426	2.055426
Ishikawa	2.04976	2.068361714	2.308389	2.308389	2.308389	2.308389	2.292607	2.308389	2.292607	2.216179	2.192389
U266B1	2.049136	2.080235408	2.044011	2.054096	2.044011	2.044011	2.054096	2.044011	2.044011	2.044011	2.044011
PA-TU-8988T	2.048497	1.999909714	1.994686	1.994686	1.994686	2.066444	2.14288	2.066444	2.14288	2.14288	2.175523
HRT-18	2.047547	2.010640898	2.135209	2.135209	2.135209	2.293271	2.293271	2.293271	2.293271	2.232503	2.232503
C170	2.047206	2.074730918	2.060997	2.075608	2.075608	2.075608	2.075608	2.093136	2.234512	2.093136	2.234512
J GAMMA 1	2.043016	2.015243286	2.191727	2.191727	2.191727	2.191727	2.191727	2.159623	2.043016	1.931599	2.043016
LS174T	2.040812	2.065228286	2.224052	2.224052	2.232157	2.232157	2.232157	2.250575	2.232157	2.172018	2.047887
NCI-H526	2.034746	1.991198633	1.696427	1.696427	1.743214	2.09206	2.09206	2.20131	2.09206	2.20131	2.20131
HLE	2.023206	1.98619998	2.19035	2.164301	2.164301	2.19035	2.19035	2.19035	2.164301	2.164152	2.164152
23132/87	2.014391	1.944262388	2.127006	2.168129	2.181707	2.208673	2.23539	2.23539	2.181707	2.181707	2.181707
Hs 834.T	2.012639	2.032355776	2.190147	2.190147	2.252941	2.252941	2.252941	2.266996	2.252941	2.252941	2.252941
Hs 891.T	2.010326	2.007325347	1.884045	1.912054	2.087879	2.087879	2.177979	2.177979	2.087879	2.177979	2.217531
Hs 839.T	2.004872	1.932453143	2.029153	2.029153	2.063214	2.307844	2.354777	2.307844	2.307844	2.307844	2.427404
SNU-1	2.004714	2.069788612	2.270295	2.270295	2.33001	2.36738	2.33001	2.270295	2.23016	2.219162	2.219162
EB1	2.000692	1.960849898	2.058439	2.093694	2.108516	2.108516	2.108516	2.108516	2.058439	2.058439	2.050752
NCI-H2009	1.988783	1.980017959	1.965983	1.965983	2.017074	2.114679	2.129669	2.168101	2.129669	2.129669	2.129669
NCI-H1048	1.986378	1.961069796	2.117838	2.104595	2.104595	2.098425	2.098425	2.098425	2.098425	2.098425	2.014305
A2780	1.985856	2.026895184	2.178847	2.248034	2.31993	2.31993	1.985856	1.985856	1.985856	1.897022	1.983013
NCI-H1666	1.985537	1.887812714	1.806077	1.806077	1.775051	1.61471	1.496922	1.444932	1.444932	1.496922	1.61471
G-402	1.983289	1.940407898	1.829022	1.834344	1.836406	1.836406	1.982011	1.961001	1.982011	1.995871	1.995871
JHH-7	1.981416	1.887399939	1.753258	1.753258	1.793288	2.077069	2.128703	2.077069	2.077069	1.793288	2.072247
OUS-11	1.981343	2.003307306	1.981343	1.981343	2.099025	2.160292	2.160292	2.099025	1.981343	2.099025	2.099025
AZ-521	1.978932	2.036937776	2.073021	2.073021	2.324858	2.324858	2.324858	2.324858	2.320653	2.320653	2.320653
K562/ADM	1.974715	1.825663102	1.848428	1.724244	1.995365	2.098146	2.110183	2.098146	2.098146	1.995365	1.995365
NCI-H889	1.974132	2.12430551	2.155019	2.155019	2.155019	2.628326	2.628326	2.659395	2.06224	2.06224	2.819173
OE19	1.971904	1.95641498	2.275243	2.145666	2.145666	2.275243	2.145666	1.974016	1.974016	1.878212	1.878212
HT 728.T	1.970593	1.957142551	1.971759	1.971759	2.021058	2.132247	2.155238	2.155238	2.370046	2.155238	2.155238
GA-10	1.967487	1.986767939	1.978293	1.978293	2.016457	1.978293	1.978293	1.978293	1.959173	1.959173	1.959173
Hs 737.T	1.962566	1.979357367	2.025338	1.902609	1.902609	2.025338	2.076043	1.902609	1.902609	1.902609	1.953739
NCI-H838	1.962434	1.928405	2.040855	2.186079	2.040855	2.040855	2.186079	2.226881	2.040855	2.040855	2.101951
NCI-H1155	1.96068	1.972872878	1.96068	1.988928	1.988928	2.010527	2.052212	2.052212	2.052212	2.052212	2.010527
A2780cis	1.955571	1.949626163	1.847546	1.981736	2.116952	2.186945	2.116952	2.186945	2.186945	2.116952	2.116952

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
G-401	1.952639	1.91027649	1.894599	1.894599	1.986276	2.027539	2.119472	2.027539	1.986276	1.985247	1.985247
MS-1-L	1.952346	1.982695653	2.25656	2.134107	2.1996	2.25656	2.25656	2.25656	2.1996	2.1996	2.1996
Hs 746T	1.949352	1.95437602	1.996969	1.996969	2.123055	1.996969	1.996969	2.123055	2.198833	2.198833	2.198833
SW 13	1.948559	1.919654286	1.979441	1.979441	1.979441	1.979441	2.094574	2.094574	2.094574	2.094574	2.094574
AGS	1.947164	1.943232163	1.990781	1.990781	2.010613	2.010613	2.010613	2.010613	2.010613	2.010613	2.02322
NCI-H2171	1.942958	1.950063061	1.940927	1.940927	2.223665	2.173174	2.173174	2.173174	2.173174	2.350219	2.37913
Hs 573.T	1.930985	1.953761673	1.847637	1.847637	1.928191	2.063595	2.082444	2.082444	2.082444	2.082444	2.082444
HT1080	1.927481	1.964792755	2.029924	2.029924	2.069009	2.069009	2.069009	1.927481	1.927481	1.67104	1.581778
COR-L 23/CPR	1.924772	1.992882388	1.807758	1.976417	1.976417	2.272895	2.272895	2.298376	2.272895	2.272895	2.272895
HGC-27	1.923706	1.861182878	1.884451	1.884451	1.950571	1.950571	1.962449	1.962449	1.950571	1.950571	1.958792
SBC-3	1.922283	1.87430849	1.985305	1.956875	1.985305	1.999062	1.999062	1.999062	1.956875	1.956875	1.956875
COLO 792	1.915651	1.882537204	1.855838	2.08428	2.08428	2.118461	2.118461	2.106668	2.08428	2.029739	2.029739
H-EMC-SS	1.912559	1.953568061	1.951803	1.951803	1.951803	2.342503	2.347905	2.347905	2.342503	2.347905	2.347905
J82	1.910432	1.781440857	1.762499	1.520425	1.762499	1.867191	1.920742	1.920742	1.920742	2.128552	2.128552
Hs 600.T	1.905368	1.931164163	1.864153	1.864153	2.00516	2.00516	2.052127	2.00516	1.864153	2.00516	2.052127
Hs 781.T	1.904757	1.954014347	1.802077	1.802077	1.898472	1.898472	2.012395	2.081518	2.012395	2.413911	2.413911
BPH-1	1.894345	1.916480796	1.782916	1.782916	1.856172	1.98471	2.044366	2.044366	1.98471	2.044366	2.044366
COR-L23	1.892174	1.897527306	1.928013	1.928013	1.928013	2.06135	2.06135	2.06135	2.06135	2.153241	2.220459
A673	1.889162	1.913707939	1.717012	1.833534	1.833534	1.833534	1.733832	1.733832	1.733832	1.717012	1.717012
COLO 205	1.884661	1.92261249	1.951891	2.01274	2.01274	2.132468	2.132468	1.951891	1.874433	1.865876	1.865876
RMG-I	1.866707	1.822054367	1.866707	1.866707	1.891305	2.23939	2.23939	2.175849	1.91773	1.91773	1.91773
WI-26 VA4	1.854321	1.881021184	1.818701	1.929803	1.929803	1.99577	2.108766	2.108766	2.108766	2.008456	2.008456
Eos-HL-60	1.853456	1.859891449	2.186701	2.186701	2.186701	2.186701	2.186701	2.186701	1.970782	1.932876	1.848245
GP5D	1.85253	1.869270245	1.972458	1.805614	1.972458	2.040898	2.040898	1.805614	1.733737	1.733737	1.733737
MES-SA/Dx-5	1.846861	1.889437184	2.305353	2.305353	2.305353	2.305353	2.3466	2.305353	2.305353	2.305353	1.974962
SBC-2	1.845355	1.869572347	1.805305	1.746445	1.805305	1.805305	1.805305	1.880078	1.988175	2.052269	2.130166
MCAS	1.842675	1.692514	1.842675	1.842675	1.878664	1.929596	1.929596	1.878664	1.878664	1.878664	1.878664
Ishikawa (Heraklio) 02 ER-	1.833641	1.871376959	2.233448	2.233448	2.233448	2.233448	2.073429	2.073429	1.907603	1.870271	1.870271
MOLT-4F	1.831195	1.671263408	1.729684	1.729684	1.859749	1.955542	1.967282	1.955542	1.955542	1.955542	1.967282
HARA	1.828838	1.811220184	1.867719	1.858779	1.856384	1.858779	1.949147	1.858779	1.858779	1.858779	1.924757
NCI-H1944	1.820237	1.805162857	1.611014	1.735809	2.029339	2.029339	2.029339	2.029339	2.001212	1.993689	1.993689
BICR 31	1.819485	1.843385122	2.01944	1.941913	1.941913	1.941913	1.871055	1.841335	1.795472	1.814335	1.814335
D341 Med	1.819101	1.797722388	1.819101	1.894135	1.904727	1.904727	1.894135	1.894135	1.819101	1.819101	1.819101
KHM-3S	1.79781	1.971451857	1.714258	1.668791	1.668791	1.714258	2.359328	2.735357	3.248595	3.020628	3.248595
COR-L88	1.789457	1.778430939	1.789457	1.845659	1.845659	1.944327	1.944327	2.139528	1.944327	1.857779	1.944327
NCI-H358	1.786631	1.775809122	2.034461	2.034461	2.034461	1.985508	2.034461	1.985508	1.985508	1.985508	1.985508
1321N	1.785668	1.814321673	2.104359	2.104359	2.104359	2.190107	2.190107	2.190107	2.019296	1.863563	2.072683
KYSE-510	1.717203	1.727553184	1.853988	1.853988	1.853988	1.853988	1.811703	1.811703	1.747926	1.811703	1.811703
TCO-1	1.702926	1.634383041	1.371826	1.394676	1.449981	1.613308	1.619487	1.613308	1.619487	1.619487	1.619487
NCI-H441	1.692289	1.722444694	1.666444	1.666444	1.713191	1.873604	1.873604	1.873604	1.713191	1.872669	1.872669
SNU-398	1.644383	1.610436694	1.812534	1.812534	1.812534	1.878152	1.812534	1.812534	1.665391	1.665391	1.665391
NCI-H1299	1.610634	1.607423367	1.582261	1.662474	1.672575	1.672575	1.672575	1.672575	1.683222	1.683222	1.719669
ASH-3	1.3333496	1.282063418	0.9883396	0.9883396	1.4151733	1.4205219	1.4964415	1.4964415	1.4964415	1.4964415	1.4964415
MCF7	1.150625	1.234855673	1.427392	1.427392	1.374121	1.427392	1.374121	1.319849	1.283092	1.283092	1.283092
SKOV3	1.0107454	1.027430129	1.0822168	1.0874075	1.2565393	1.2565393	1.0822168	0.9882676	1.0822168	1.2565393	1.2565393
ChaGo-K-1	0.9705445	0.907140933	0.9705445	1.0860414	1.0860414	1.0860414	0.9705445	0.9705445	1.13213	1.13213	1.13213
A373 C6	0.869123	0.901690388	0.8781185	0.8714279	0.8714279	0.8714279	0.8442507	0.8095883	0.8442507	0.8095883	0.7818635

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
MES-SA	0.7528253	0.784296504	0.9228057	0.9228057	0.9228057	0.9228057	1.0695177	1.0695177	0.8519282	0.8519282	0.8519282
OAW28	0.7357038	0.746945755	0.798911	0.8649355	0.798911	0.7909373	0.6828103	0.6828103	0.9794969	0.9794969	0.9794969
NCI-H460	0.6484461	0.569243312	0.3147084	0.3777133	0.3777133	0.3777133	0.3897879	0.3897879	0.7146986	0.7146986	0.7146986

B. ERBB2 SNP array data

ProbeSetID	Mean	SNP_A-2138860	SNP_A-4199416
Affy_SNP.ID		7268231	11276532
dbSNP		rs4252612	rs4252643
Chromosome		17	17
Position		35117234	35130293
Strand		+	-
Cytoband		q12	q12
KYSE-410	7.406145	7.40414	7.40815
Calu-3	6.984205	6.783041	7.185369
NCI-N87	6.5425605	6.565467	6.519654
NCI-H2170	6.4934345	6.063419	6.92345
MKN7	6.382685	6.355836	6.409534
VMRC-LCP	6.1759995	6.244285	6.107714
OE33	5.930246	5.930246	5.930246
OE19	5.611658	5.611658	5.611658
SK-BR-3	5.537203	5.501213	5.573193
HCC202	5.4161105	5.509775	5.322446
HO-1-u-1	5.2524505	5.300472	5.204429
JIMT-1	5.205902	5.205902	5.205902
NCI-H1693 (ship16)	4.5363085	4.374288	4.698329
SKOV3	4.210977	4.286284	4.13567
RMG-I	3.8286955	3.847051	3.81034
BICR 31	3.446782	3.528013	3.365551
NCI-H1563	3.438787	3.438787	3.438787
KYSE-50	2.899275	2.901678	2.896872
NUGC-4	2.8919245	2.833679	2.95017
NCI-H522	2.881096	2.878906	2.883286
REF-GC-1B	2.8742675	2.960582	2.787953
SK-MEL-1	2.8493985	2.831863	2.866934
COLO-680N	2.843106	2.860389	2.825823
T.Tn	2.819216	2.819216	2.819216
LU65	2.811984	2.81414	2.809828
NCI-H2081	2.78139	2.78139	2.78139
A549	2.7641	2.729223	2.798977
LK-2	2.741504	2.744667	2.738341
NCI-H2009	2.709708	2.732214	2.687202
COLO 201	2.698706	2.729694	2.667718
COLO 792	2.697088	2.697088	2.697088
KHM-3S	2.6923705	2.69346	2.691281
SNU-5	2.677204	2.620494	2.733914

C. MET SNP array data

ProbeSetID	Median	Mean	SNP_A-2060488	SNP_A-1990987	SNP_A-4250633	SNP_A-2119519
Affy_SNP.ID			6070863	2625703	11264768	6981648
dbSNP			rs38841	rs10487353	rs38847	rs2299433
Chromosome			7	7	7	7
Position			115913877	115917887	115919277	115924829
Strand			+	-	+	+
Cytoband			q31.2	q31.2	q31.2	q31.2
Hs 746T	6.353536	6.179734786	6.476262	6.483741	6.476262	6.476262
MKN45	5.9993945	6.047987857	5.954915	5.935075	5.935075	5.935075
EBC-1	5.798258	5.561305071	5.775051	5.821465	5.821465	5.821465
SNU-5	5.634547	5.730620286	6.112681	6.112681	5.634547	5.634547
NCI-H1573	5.58455	5.582480786	5.730309	5.644351	5.644351	5.644351
OE33	5.221678	5.108011429	5.374789	5.322405	5.322405	5.322405
OCUG-1	4.686753	4.565070857	4.686372	4.687134	4.687134	4.687134
VMRC-RCW	4.47436	4.442329857	3.999351	4.023979	4.586229	4.586229
SK-BR-3	4.078277	4.040958214	4.1339	4.078277	4.078277	4.078277
C32	4.062967	4.063081143	4.322447	4.322447	4.095586	4.095586
Detroit 562	3.7587865	3.724144071	3.736275	3.593785	3.593785	3.736275
NCI-H2342	3.6937315	3.6740345	3.881318	3.881318	3.881318	3.881318
FTC 133	3.5743185	3.465995857	3.647539	3.614116	3.614116	3.614116
PNT1A	3.507132	3.520597643	3.590536	3.545874	3.486409	3.545874
KMRC-20	3.427928	3.426840643	3.572808	3.504144	3.492471	3.492471
COLO 858	3.374802	3.354036786	3.374802	3.374802	3.374802	3.374802
NUGC-4	3.3327075	3.311047214	2.918863	2.98114	3.217618	3.400879
YKG-1	3.2886055	3.2031295	3.267634	3.309577	3.309577	3.312846
H3255	3.215802	3.251627286	3.522366	3.437622	3.413821	3.413821
SK-CO-1	3.1729975	3.118538429	3.351289	3.305964	3.292095	3.292095
LOU-NH91	3.160291	3.155499429	3.292325	3.198636	3.160291	3.292325
SK-MEL-3	3.159654	3.225826786	3.319119	3.319119	3.159654	3.319119
NCI-H2170	3.153093	3.076933786	3.016807	3.016807	3.016807	3.153093
G-361	3.1326005	3.145947357	3.352589	3.352589	3.352589	3.352589
SK-NEP-1	3.094811	3.065128357	2.875521	2.904969	3.116878	3.147903
HCC-44	3.087111	3.039358857	2.763962	2.763962	2.994182	3.043408
WM 266-4	3.082038	3.087722714	3.38531	3.082038	3.082038	3.082038
NCI-H441	3.077623	3.082637643	3.077623	3.077623	3.077623	3.199245
P31/FUJ	3.074602	3.084332929	2.648757	2.84764	3.074602	3.262888
MOG-G-CCM	3.072613	2.931386286	3.242317	3.072613	3.072613	3.072613
SW 1417	3.069209	3.029638286	3.069209	3.069209	3.069209	3.069209
COLO 853	3.0386255	3.034300357	2.975415	2.975415	2.975415	3.029699
NCI-H2085	3.016188	2.965419071	3.016188	3.016188	3.016188	3.016188

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
D341 Med	2.671545	2.666515	2.676575		NCI-H23	2.9948565	2.981778143	3.198321	3.127195	2.996522	3.127195
NCI-H2135	2.6496145	2.617956	2.681273		A2058	2.971587	2.894833286	3.088328	3.088328	3.037372	3.037372
UMC-11	2.641747	2.607421	2.676073		LUDLU-1	2.9644875	2.931987	2.804811	2.804811	2.804811	2.954671
STC 1	2.629892	2.629892	2.629892		COLO 741	2.963187	2.964942	2.963187	2.903029	2.7736	2.903029
KATO III	2.6271375	2.562506	2.691769		KP-4	2.9485235	2.916517429	2.642656	2.749701	3.117642	3.117642
HMVII	2.592692	2.592692	2.592692		SR	2.9181915	2.909131286	3.001675	2.889578	2.829007	2.889578
COLO 205	2.573715	2.573715	2.573715		WM-115	2.913138	2.927133857	3.129917	2.814586	2.814586	3.01169
JHH-4	2.5730025	2.536773	2.609232		NCI-H810	2.908847	2.833355214	3.020364	2.908847	2.908847	2.933108
HuCCT1	2.557023	2.609069	2.504977		SBC-5	2.8988535	2.9239435	2.980914	2.980914	3.115644	3.115644
SK-LU-1	2.552743	2.552743	2.552743		MKN1	2.896216	2.875414929	2.91091	2.91091	3.014514	3.014514
WI-26 VA4	2.549048	2.549048	2.549048		Sci-1	2.872665	2.896049571	3.176132	3.014808	2.872665	2.891581
HCC-44	2.546363	2.546363	2.546363		NCI-H1755	2.868482	2.868767071	2.961833	2.868482	2.868482	2.868482
KYSE-450	2.5447545	2.566279	2.52323		HCC202	2.867372	2.873158429	3.066449	2.999513	2.793963	2.999513
MS-1-L	2.5130265	2.593702	2.432351		LCLC-103H	2.86148	2.901389929	2.804244	2.804244	2.822462	3.018571
201T	2.491139	2.465481	2.516797		KMRC-1	2.8593295	2.785395357	2.79199	3.042845	2.942573	2.942573
NCI-H810	2.488556	2.488556	2.488556		DMS 273	2.855502	2.845410929	3.329949	2.736252	2.736252	2.855502
OAW42	2.4695	2.442368	2.496632		SNU-16	2.847565	2.799459929	2.639629	2.639629	2.639629	2.66286
NCI-H650	2.455322	2.455322	2.455322		KATO III	2.840764	2.839600857	2.83463	2.83463	2.846898	2.846898
NCI-H1666	2.4525955	2.403268	2.501923		COLO 857	2.818847	2.852647143	2.818847	2.818847	2.818847	2.818847
NCI-H2122	2.449908	2.449908	2.449908		22RV1	2.8179675	2.825521071	2.743274	2.717423	2.717423	2.743274
MOG-G-UVW	2.449451	2.449451	2.449451		JSC-1	2.808346	2.929921429	3.341663	3.341663	3.028245	3.028245
NCI-H1581	2.439793	2.439793	2.439793		U373 MG	2.7962715	2.773538929	2.855051	2.889885	2.889885	2.889885
OVISE	2.425734	2.45409	2.397378		U266B1	2.794051	2.850709643	2.789072	2.789072	2.789072	2.789072
A2780cis	2.42403	2.42403	2.42403		JHH-4	2.793842	2.699646929	2.618259	2.672715	2.793842	2.862679
JHH-7	2.4203845	2.432475	2.408294		COLO 792	2.784492	2.840433	2.99227	2.784492	2.784492	2.99227
SW-1710	2.4193445	2.369551	2.469138		MRK-nu-1	2.7834295	2.745245143	2.597645	2.622227	2.7575	2.826527
FTC 238	2.4169605	2.441613	2.392308		CAL-148	2.769583	2.757274714	2.864123	2.864123	2.739413	2.864123
NCI-H1755	2.414511	2.414511	2.414511		COLO 684	2.766928	2.727863643	2.708114	2.708114	2.766928	2.766928
COR-L279	2.4071345	2.428934	2.385335		PC-14	2.7653415	2.741458214	2.544226	2.544226	2.544226	2.621913
U373 MG	2.403481	2.403481	2.403481		NCI-H841	2.758824	2.786344929	3.22942	3.22942	3.22942	3.22942
COLO 685	2.398208	2.398208	2.398208		BICR 78	2.750902	2.755712929	2.86377	2.831571	2.766706	2.831571
NCI-H23	2.389082	2.389082	2.389082		RERF-LC-KJ	2.750535	2.782063286	2.978089	2.849264	2.750535	2.849264
KP-4	2.3784035	2.27237	2.484437		JHH-6	2.7420105	2.649656214	2.726454	2.757567	2.757567	2.846317
LUDLU-1	2.377726	2.377726	2.377726		SW 48	2.736453	2.781257071	2.722447	2.722447	2.722447	2.736453
SNU-1	2.3758135	2.345525	2.406102		NCI-H196	2.722144	2.627415786	2.326482	2.326482	2.326482	2.62778
A373 C6	2.3655755	2.366173	2.364978		BALL-1	2.721345	2.6989785	2.598724	2.811048	2.825368	2.825368
YKG-1	2.3630035	2.327435	2.398572		NCI-H889	2.718772	2.7394705	2.820291	2.820291	2.646544	2.820291
NCI-H838	2.361989	2.361989	2.361989		NCI-H2009	2.707223	2.677878929	2.712716	2.718694	2.70173	2.718694
KYSE-180	2.35133	2.35133	2.35133		SK-LU-1	2.687566	2.605813143	2.92203	2.811495	2.687566	2.687566
NCI-H460	2.344501	2.344501	2.344501		HEC-1	2.6858805	2.5141325	2.73463	2.73463	2.751825	2.751825
SW 1116	2.335792	2.413849	2.257735		H69AR	2.679096	2.667231214	2.679096	2.679096	2.658729	2.679096
VMRC-LCD	2.3268825	2.352646	2.301119		OCUM-1	2.666847	2.602363143	2.573762	2.573762	2.666847	2.678275
OCUG-1	2.3204395	2.31401	2.326869		KHOS-312H	2.645598	2.433654786	1.801104	1.801104	1.947972	2.00742
BPH-1	2.319775	2.319775	2.319775		IM-95	2.644987	2.6314285	2.661105	2.644987	2.556993	2.644987
MCAS	2.3140655	2.321234	2.306897		OAW42	2.642609	2.645555786	3.119033	2.907857	2.65734	2.65734
NCI-H2342	2.310903	2.310903	2.310903		MDST8	2.624947	2.635751143	2.772143	2.734037	2.619959	2.733321
PE/CA-PJ15	2.310125	2.310125	2.310125		HT 115	2.623534	2.595708071	2.515264	2.515264	2.629857	2.629857

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
NCI-H1781	2.309879	2.309879	2.309879		HMGB	2.6161475	2.754683929	3.212003	3.137835	3.137835	3.137835
MOLT-4F	2.309737	2.309737	2.309737		RERF-LC-MS	2.614851	2.641049286	2.597317	2.597317	2.597317	2.609022
A-427	2.305198	2.267982	2.342414		FTC 238	2.6090915	2.576791643	2.719413	2.719413	2.624575	2.624575
WM 266-4	2.2927635	2.286839	2.298688		KHOS-240S	2.593009	2.750341429	2.295434	2.295434	2.295434	2.406979
NB69	2.284069	2.284069	2.284069		AN3CA	2.591834	2.596447143	2.17138	2.17138	2.17138	2.591834
HRT-18	2.277852	2.277852	2.277852		NCI-H1651	2.5754805	2.6328635	2.921894	2.909468	2.909468	2.909468
G-361	2.2738	2.2738	2.2738		1321N	2.572272	2.536160643	2.743876	2.705031	2.695751	2.695751
SK-CO-1	2.2704025	2.216166	2.324639		LU65	2.5719265	2.571867357	2.876831	2.876831	2.876831	2.876831
EJ138	2.263008	2.263008	2.263008		HARA	2.571444	2.569916214	2.753503	2.732113	2.732113	2.732113
HT 115	2.2587945	2.241009	2.27658		RCM-1	2.569842	2.519454429	2.229627	2.265877	2.220098	2.436967
CASKI	2.257669	2.257669	2.257669		HuH-7	2.566611	2.553114071	2.566611	2.566611	2.566611	2.566611
MKN45	2.2487385	2.253483	2.243994		WIDR	2.566433	2.566277143	2.687884	2.667615	2.667615	2.667615
Sci-1	2.246948	2.246948	2.246948		UMC-11	2.564286	2.5420845	2.564286	2.45749	2.564286	2.587377
COLO 853	2.242959	2.242959	2.242959		huH-1	2.563592	2.536612786	2.204994	2.374725	2.564737	2.58587
MEWO	2.240239	2.240239	2.240239		HCC-827	2.561117	2.416543071	2.394334	2.561117	2.750924	2.784206
P32/ISH	2.237343	2.198577	2.276109		NCI-H2081	2.5595745	2.579073857	2.53049	2.588659	2.512729	2.658167
U266B1	2.2343765	2.23838	2.230373		SW 13	2.54389	2.553215357	2.469407	2.469407	2.469407	2.473646
HEC-1	2.233717	2.172847	2.294587		KYSE-140	2.54327	2.546991071	2.511267	2.511267	2.511267	2.54327
COLO 857	2.232437	2.232437	2.232437		LC-1 sq	2.536981	2.6609255	2.408357	2.431258	2.450858	2.559917
KYSE-510	2.228081	2.228081	2.228081		KYSE-410	2.536665	2.595577714	2.443408	2.443408	2.443408	2.457785
FLYA13	2.22508	2.22508	2.22508		CAL-85-1	2.5314445	2.538900429	2.814923	2.814923	2.814923	2.814923
fr2	2.221007	2.221007	2.221007		HCC-366	2.5302205	2.335572857	2.597023	2.597023	2.733478	2.733478
KHOS/NP	2.220299	2.220299	2.220299		SW 1463	2.527391	2.533177857	2.84117	2.474148	2.474148	2.527391
DOK	2.217758	2.217758	2.217758		MOLT-4F	2.527259	2.464744857	2.509768	2.538569	2.538569	2.64501
MKN1	2.217196	2.217196	2.217196		SW-1710	2.52394	2.505554929	2.506526	2.667992	2.52394	2.667992
COR-L 105	2.2156705	2.221822	2.209519		HCC-15	2.5105875	2.505424286	2.525672	2.525672	2.495503	2.495503
SNU-16	2.215644	2.215644	2.215644		PLC/PRF/5	2.49792	2.456491357	2.245372	2.284199	2.49792	2.505933
Detroit 562	2.2156035	2.188513	2.242694		A375.S2	2.495648	2.555792429	2.800967	2.639201	2.495648	2.639201
SK-MES	2.21513	2.225454	2.204806		COR-L279	2.488386	2.366638286	2.588906	2.588906	2.529565	2.529565
SBC-5	2.214679	2.214679	2.214679		COLO 685	2.486619	2.523850357	2.600001	2.600001	2.486619	2.851198
Calu-1	2.211678	2.211678	2.211678		FLYA13	2.482021	2.434360714	2.518738	2.518738	2.526838	2.526838
NCI-H661	2.209774	2.209774	2.209774		JHH-7	2.476583	2.507523714	2.622157	2.622157	2.622157	2.622157
RO82-W-1	2.203892	2.203892	2.203892		NCI-H1623	2.469236	2.424541714	2.348774	2.348774	2.271254	2.348774
NCI-H2085	2.201606	2.201606	2.201606		HGC-27	2.468389	2.442615429	2.468389	2.468389	2.468389	2.468389
NCI-H1734	2.201175	2.201175	2.201175		NCI-H2347	2.456307	2.456934214	2.425251	2.425251	2.425251	2.487363
H33HJ-JA1	2.197728	2.197728	2.197728		SKN	2.444542	2.398401571	2.688384	2.688384	2.477048	2.675114
NCI-H2405	2.196812	2.196812	2.196812		VMRC-LCP	2.444231	2.318382643	1.967939	2.548881	2.548881	2.563919
C170	2.194628	2.194628	2.194628		SAT	2.439669	2.372732786	1.970505	2.05088	2.439669	2.446993
GRANTA-519	2.1942145	2.132991	2.255438		CaR-1	2.439296	2.479265357	2.393655	2.393655	2.484937	2.629637
D1.1	2.188067	2.188067	2.188067		SW 900	2.437711	2.447624571	2.614799	2.614799	2.438046	2.438046
HCC-78	2.186268	2.186268	2.186268		Nthy-ori 3-1	2.43466	2.420241214	2.567926	2.567926	2.567926	2.567926
NCI-H2347	2.185971	2.185971	2.185971		NCI-H526	2.434635	2.449528357	2.489129	2.489129	2.434635	2.489129
GA-10	2.180718	2.180718	2.180718		HUP-T3	2.434586	2.438426929	2.280013	2.281152	2.281152	2.586082
HARA	2.1764785	2.251633	2.101324		BT-483	2.430152	2.444356357	2.370871	2.370871	2.370871	2.430152
JSC-1	2.171044	2.117395	2.224693		NCI-H2330	2.429767	2.341269929	2.454977	2.454977	2.429767	2.454977
Hs 936.T	2.170917	2.170917	2.170917		NCI-H711	2.427935	2.495495643	2.416498	2.416498	2.416498	2.439372
MES-SA	2.166931	2.166931	2.166931		NCI-H661	2.423557	2.370422	2.179769	2.179769	2.179769	2.192888

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
BU25 TK	2.1653915	2.15905	2.171733		EJ138	2.422922	2.410984429	2.456726	2.435504	2.435504	2.435504
PNT1A	2.164697	2.164697	2.164697		HOS	2.419543	2.516425357	2.175659	2.114082	2.114082	2.175659
ChaGo-K-1	2.161288	2.161288	2.161288		KHM-3S	2.416223	2.379781714	2.416223	2.470192	2.416223	2.470192
MKN74	2.1582565	2.181002	2.135511		ChaGo-K-1	2.4103015	2.364104214	2.530214	2.498794	2.459705	2.459705
T47D	2.154945	2.139789	2.170101		H69V	2.404434	2.262584786	2.198173	2.596765	2.198173	2.479114
EBC-1	2.154486	2.154486	2.154486		Hs 936.T	2.401059	2.443547429	2.422012	2.422012	2.422012	2.401059
NCI-H82	2.151055	2.14321	2.1589		201T	2.386913	2.374132286	2.682514	2.386913	2.338016	2.386913
A2780 ADR	2.1489815	2.156113	2.14185		SUIT-2	2.386862	2.354446929	2.242163	2.242163	2.31595	2.386862
RERF-LC-Sq1	2.1489525	2.191349	2.106556		NCI-H1792	2.3837015	2.387611786	2.647674	2.647674	2.647674	2.647674
RT4	2.143004	2.143004	2.143004		NCI-H460	2.366694	2.393000786	2.472642	2.438851	2.438851	2.472642
HSC-3	2.142811	2.142811	2.142811		SNU-398	2.36574	2.305664	2.389246	2.389246	2.389246	2.389246
WPE1-NB26	2.141273	2.141273	2.141273		COR-L95	2.365452	2.352801929	2.565722	2.461255	2.336154	2.39475
KMRC-1	2.1410335	2.049164	2.232903		SK-MES	2.365297	2.242536286	2.357327	2.373267	2.373267	2.400745
NCI-H441	2.138112	2.097268	2.178956		BFTC-905	2.363592	2.474462286	2.296046	2.742192	2.742192	2.742192
NCI-H1573	2.133341	2.168006	2.098676		EB1	2.3578555	2.347330071	2.336745	2.336745	2.336745	2.336745
RERF-LC-KJ	2.130386	2.130386	2.130386		CASKI	2.3549885	2.326869286	2.434089	2.403029	2.434089	2.434089
K562/ADM	2.1287835	1.959582	2.297985		HO-1-N-1	2.353416	2.264480214	2.397303	2.397303	2.412549	2.57071
OSC-20	2.1246775	2.088858	2.160497		PC-3 [JPC-3]	2.352646	2.394180286	2.390695	2.352646	2.352646	2.352646
SBC-3	2.124022	2.124022	2.124022		TE7	2.351203	2.332805286	2.419905	2.419905	2.419905	2.419905
RT112/84	2.1236595	2.151558	2.095761		TASK1	2.346099	2.287112	2.450525	2.567073	2.567073	2.567073
BFTC-905	2.121332	2.119029	2.123635		NCI-H1734	2.3429835	2.330537429	2.142166	2.142166	2.38555	2.421227
KHOS-240S	2.120358	2.120358	2.120358		HRT-18	2.3415235	2.327379071	2.53151	2.528845	2.473757	2.473757
HCE7	2.116809	2.060771	2.172847		RT112/84	2.339449	2.333729	2.504105	2.34381	2.504105	2.504105
NCI-H1522	2.114374	2.114374	2.114374		CAL-29	2.3384115	2.358125429	2.120939	2.390431	2.464216	2.464216
LNCaP clone FGC	2.11103	2.11103	2.11103		IMR-90	2.3380535	2.219999857	2.511207	2.623332	2.373584	2.373584
SBC-2	2.1074895	2.183142	2.031837		RO82-W-1	2.3379215	2.342567071	2.657811	2.498794	2.484534	2.484534
NCI-H1703	2.103677	2.103677	2.103677		NCI-H1581	2.335088	2.321850643	2.391061	2.335088	2.335088	2.391061
LS174T	2.100028	2.100028	2.100028		RT4	2.334754	2.227424	2.399816	2.389057	2.399816	2.399816
NCI-H2023	2.099554	2.099554	2.099554		HT 29	2.3175745	2.391541143	2.7243	2.7243	2.550503	2.550503
NCI-H2171	2.097728	2.097728	2.097728		KON	2.313129	2.282338857	2.323969	2.450463	2.450463	2.450463
WIL2 NS	2.094127	2.094127	2.094127		BT-20	2.3070205	2.272815214	2.415786	2.365385	2.365385	2.415786
CaR-1	2.091204	1.993544	2.188864		BU25 TK	2.295543	2.256701214	2.080256	2.080256	2.080256	2.355759
HLE	2.090341	2.090341	2.090341		RPMI-7951	2.295004	2.364569143	2.702635	2.695603	2.695603	2.695603
HUP-T4	2.089963	2.089963	2.089963		IHH-4	2.276764	2.226101286	1.996449	2.040514	2.083916	2.19475
NL20-TA	2.087809	2.087809	2.087809		fR2	2.267548	2.151334929	2.539443	2.475654	2.475654	2.475654
RERF-LC-MA	2.085289	2.085289	2.085289		CAL-51	2.264522	2.214289857	2.27835	2.27835	2.373495	2.373495
LCLC-103H	2.084448	2.084448	2.084448		COR-L 105	2.26032	2.257736357	2.248405	2.248405	2.248405	2.26032
KYSE-270	2.08285	2.08285	2.08285		LK-2	2.2595615	2.266311786	2.024236	2.024236	2.236374	2.278048
NCI-H2286	2.0828265	2.033406	2.132247		FU97	2.254841	2.294969643	2.173065	2.173065	2.51822	2.51822
COR-L95	2.082528	2.082528	2.082528		MOG-G-UVW	2.253695	2.213891643	2.486692	2.44834	2.44834	2.44834
KYSE-140	2.0820655	2.091082	2.073049		NCI-H2452	2.243583	2.280705929	2.188134	2.188134	2.17574	2.188134
EB1	2.0820005	2.070474	2.093527		HuCCT1	2.236575	2.2690245	2.246683	2.246683	2.36092	2.36092
BICR22	2.080242	2.080242	2.080242		NCI-H1417	2.234991	2.250631214	2.30286	2.068643	2.234991	2.234991
SKN-3	2.07914	2.072711	2.085569		Hs 633T	2.231598	2.162600714	2.312314	2.239337	2.231598	2.239337
RERF-LC-Ad2	2.075355	2.075355	2.075355		A-375	2.225076	2.229321929	2.179864	2.191309	2.179864	2.191309
C8166	2.073969	2.034119	2.113819		NCI-H1650	2.217624	2.248877857	2.552422	2.513949	2.513949	2.513949
PC-9	2.067108	2.002437	2.131779		KP-1N	2.2119685	2.252706	2.293454	2.502893	2.502893	2.530754

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-2119130	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
SW 1573	2.0585865	1.997604	2.119569		NCI-H2135	2.211716	2.137554071	1.972968	1.972968	1.972968	1.972968
SK-N-BE(2)	2.052679	2.052679	2.052679		HT55	2.210316	2.257225071	2.486193	2.418233	2.336373	2.336373
AZ-521	2.0481285	2.163293	1.932964		NCI-H1781	2.209344	2.233240357	2.209344	2.209344	2.209344	2.553077
NCI-H526	2.0465215	1.977144	2.115899		RERF-GC-1B	2.20256	2.1819985	2.419768	2.352288	2.419768	2.419768
A2780	2.043954	2.043954	2.043954		NCI-H727	2.198523	2.193538286	2.192591	2.192591	2.192591	2.192591
LU99A	2.042959	2.042959	2.042959		647-V	2.1977405	2.124148286	2.217624	2.278729	2.306923	2.306923
CAL-85-1	2.0392425	1.811318	2.267167		NCI-H2023	2.1952225	2.153250214	2.30147	2.248418	2.248418	2.30147
KYSE-30	2.03482	2.024436	2.045204		Calu-3	2.1951075	2.130328	1.907052	1.907052	1.907052	2.123884
GCT	2.034148	2.034148	2.034148		H69	2.194696	2.240359286	2.20811	2.194696	2.194696	2.20811
SUIT-2	2.032365	2.032365	2.032365		RWPE-1	2.188864	2.265799714	2.188864	2.188864	2.514432	2.514432
Eos-HL-60	2.031894	2.031894	2.031894		VMRC-LCD	2.187831	2.183962286	2.248656	2.127006	2.113167	2.248656
NCI-H1417	2.029264	2.074441	1.984087		T84	2.1878105	2.100313071	2.205397	2.205397	2.311136	2.311136
IM-95	2.0284025	2.030767	2.026038		SW 1116	2.1815105	2.224802929	2.66545	2.379561	2.379561	2.379561
HGC-27	2.028153	2.028153	2.028153		A373 C6	2.180846	2.13408	1.937818	1.937818	1.937818	2.102132
SAT	2.0273125	1.924832	2.129793		COLO 201	2.177844	2.169826429	2.06135	2.06135	2.06135	2.177844
NCI-H196	2.02525	2.041253	2.009247		KHOS/NP	2.177559	2.109167571	1.656028	1.656028	1.754421	1.755096
IHH-4	2.01792	2.01792	2.01792		Hs 891.T	2.163893	2.164133643	2.163893	2.163893	2.163893	2.163893
Hs 633T	2.016787	2.016787	2.016787		PA-TU-8988T	2.153719	2.009981643	2.318186	2.202117	2.194493	2.202117
KMRC-20	2.015023	1.957138	2.072908		BICR22	2.151134	2.171526714	2.151134	2.151134	2.151134	2.261598
U-2 OS	2.014391	2.014391	2.014391		MES-SA/Dx-5	2.1505655	2.082107357	2.318935	2.274207	2.149901	2.15123
Hs 573.T	2.009837	2.009837	2.009837		SK-MEL-1	2.149079	2.110367643	2.27124	2.076015	2.076015	2.222143
SR	2.008005	1.992618	2.023392		KYSE-70	2.1414645	2.233324643	2.442443	2.384919	2.384919	2.384919
BALL-1	2.006312	2.006312	2.006312		A673	2.141122	2.176495	2.522616	2.387026	2.535215	2.535215
MG 63	2.005434	2.005434	2.005434		D1.1	2.1368305	2.111685429	2.277328	1.996333	1.936093	1.936093
NCI-H2330	2.00372	2.00372	2.00372		NCI-H1793	2.1344265	2.102215571	1.899847	1.899847	1.899847	2.169285
Hs 891.T	2.0034095	2.010182	1.996637		OE19	2.130235	2.074995929	2.17176	2.17176	2.130235	2.130235
NCI-H1435	1.999668	1.999668	1.999668		WILCL	2.129159	2.044889429	2.361989	2.165228	2.159664	2.159664
DMS 273	1.995235	1.995235	1.995235		HCC-78	2.127448	2.050012071	1.790031	1.790031	1.790031	2.253008
MES-SA/Dx-5	1.994888	1.994888	1.994888		BEN	2.1264775	2.121312286	2.23539	2.23539	2.326997	2.326997
Hs 852.T	1.99217	1.99217	1.99217		Calu-1	2.1217	2.1226085	2.42104	2.229107	2.1217	2.229107
Ishikawa	1.989319	1.989319	1.989319		SK-N-BE(2)	2.121547	2.108961643	2.092381	2.092381	2.053982	2.092381
COLO 741	1.988015	1.988015	1.988015		HUP-T4	2.119334	2.171675643	2.225253	2.229134	2.577367	2.577367
CAL-29	1.985363	1.985363	1.985363		MCF7	2.113167	2.055814	2.390695	2.113167	2.113167	2.390695
HCT-8	1.9822545	1.932743	2.031766		Ishikawa	2.107752	2.118464643	2.107752	2.107752	2.107752	2.107752
NCI-H1048	1.981663	1.981663	1.981663		Hs 195.T	2.096249	2.024884714	2.096249	2.096249	2.096249	2.233288
VM-CUB1	1.976048	1.944474	2.007622		HT 1376	2.09322	2.115561	2.09322	2.09322	2.09322	2.09322
AGS	1.975846	2.05452	1.897172		COLO 205	2.091641	2.077503714	2.090788	2.090788	2.091641	2.091641
SKG-IIIb	1.9745945	1.903459	2.04573		NCI-H1437	2.0894955	2.049091214	2.038336	2.16953	2.16953	2.354037
COLO 858	1.974016	1.974016	1.974016		A2780	2.086899	2.094334714	2.086899	2.086899	1.942458	2.086899
KYSE-150	1.969281	1.969281	1.969281		A549	2.086801	2.091357429	2.296202	2.107599	2.107599	2.296202
Nthy-ori 3-1	1.964961	1.964961	1.964961		NCI-H1522	2.0813295	2.082866929	1.954516	1.954516	1.846785	1.954516
WM-115	1.964202	1.964202	1.964202		G-401	2.081199	2.069127643	2.1186	2.1186	2.1186	2.1186
SW 13	1.963009	1.955644	1.970374		PC-9	2.081056	2.04861	2.081056	2.081056	2.081056	2.081056
P116 T	1.96163	1.96163	1.96163		GRANTA-519	2.080312	2.073158786	2.080312	2.080312	2.080312	2.203542
huH-1	1.960943	1.960943	1.960943		639-V	2.0775215	2.092880714	2.565876	2.302016	2.302016	2.302016
SK-NEP-1	1.954531	1.954531	1.954531		A2780 ADR	2.072101	2.119415643	2.261585	2.261585	2.158422	2.158422
Hs 417.Lu	1.952082	1.952082	1.952082		A-427	2.0718575	1.988570429	2.122972	2.142358	2.142358	2.142358

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
Hs 737.T	1.95195	1.95195	1.95195		NB69	2.068502	2.1822505	2.381306	2.381306	2.007665	2.068502
WILCL	1.9502925	1.975428	1.925157		NCI-H2286	2.0673985	2.1009515	2.259529	2.319039	2.319039	2.319039
A-375	1.949	1.904369	1.993631		SBC-2	2.064498	2.035906571	2.06674	2.06674	2.06674	2.06674
LC-1 sq	1.9466415	1.923084	1.970199		HCT-8	2.063637	2.079253214	2.063637	2.063637	2.063637	2.172807
NUGC-3	1.9372555	1.781096	2.093415		Hs 895.T	2.063143	2.007857071	1.751529	1.821435	1.821435	1.968843
PNT2	1.934278	1.934278	1.934278		K562/ADM	2.0600015	2.028466857	2.140998	2.194332	2.272777	2.272777
BICR 78	1.932758	1.932758	1.932758		Hs 228.T	2.059343	1.972181429	2.107627	2.107627	2.107627	2.107627
273T	1.9230275	1.918042	1.928013		STC 1	2.057322	2.039048643	1.819807	1.819807	2.057322	2.142221
PLC/PRF/5	1.921557	1.921557	1.921557		LNCAp clone F	2.052311	2.0496905	2.05309	2.043286	1.974161	2.043286
MDST8	1.919812	1.878769	1.960855		C8166	2.047845	2.077550643	2.462644	2.138112	2.078207	2.078207
NCI-H1651	1.917938	1.917938	1.917938		DMS 92	2.046383	2.1337025	2.09717	2.040258	2.040258	2.060001
COR-L23	1.91724	1.91724	1.91724		NCI-H2196	2.0449175	2.0819925	2.264772	2.264772	2.264772	2.275453
HT 29	1.912529	1.905785	1.919273		HMVII	2.044125	2.034169357	1.94612	1.94612	1.94612	2.00124
J GAMMA 1	1.9121195	1.905144	1.919095		Hs 417.Lu	2.038564	1.999182143	1.942649	1.942649	1.942649	2.038564
Hs 908.T	1.905785	1.905785	1.905785		H-EMC-SS	2.035751	1.955006429	2.224947	2.224947	2.155183	2.155183
P31/FUJ	1.9031855	1.908437	1.897934		H33HJ-JA1	2.034119	2.084017429	2.034119	2.034119	2.018809	2.034119
RCM-1	1.901952	1.901952	1.901952		HCT 116	2.032336	2.007684857	2.032336	2.032336	2.032336	2.032336
Hs 834.T	1.901758	1.901758	1.901758		GA-10	2.032023	2.036001071	1.890646	1.890646	2.135732	2.140573
SW 48	1.899384	1.899384	1.899384		NCI-H1666	2.028711	1.809154286	2.028711	2.028711	2.028711	2.140779
DMS 153	1.8988655	1.888787	1.908944		LS174T	2.0282345	2.047543429	2.241009	2.065105	2.036456	2.065105
HOS	1.891313	1.89165	1.890976		Hs 908.T	2.023521	2.037572429	2.023521	2.023521	2.023521	2.023521
RWPE-1	1.8909905	1.891335	1.890646		C170	2.0207925	2.020573143	2.262231	2.256758	2.070234	2.070234
BICR10	1.887453	1.887453	1.887453		J82	2.017933	1.879096571	2.10472	2.112654	2.112654	2.112654
Hs 895.T	1.887108	1.887108	1.887108		A2780cis	2.013759	2.007294429	2.242787	2.019583	1.876647	2.125305
WIDR	1.886012	1.886012	1.886012		OUS-11	2.013329	1.962009286	2.027082	2.09915	2.179281	2.179281
A2058	1.885864	1.805058	1.96667		KYSE-150	2.013228	1.975970357	1.908408	1.908408	1.908408	1.909971
JURKAT EC-1	1.8828705	1.838512	1.927229		COR-L23	2.011432	1.975228	2.011432	2.011432	2.011432	2.011432
T.T	1.882799	1.882799	1.882799		MS-1-L	2.010826	1.994067571	2.072402	2.072402	2.091878	2.091878
OUS-11	1.881809	1.848413	1.915205		SKG-IIIb	2.0046715	2.042940786	1.989029	2.048427	2.048427	2.214733
SK-MEL-3	1.878393	1.878393	1.878393		P32/ISH	2.0034955	2.0459235	2.016529	2.096347	2.096347	2.199681
SW 1417	1.8775645	1.875608	1.879521		NCI-H1693 (shij	1.98761	2.0918535	2.372925	2.294561	2.294561	2.294561
EPLC-272H	1.877083	1.877083	1.877083		Hs 781.T	1.987479	1.985078143	2.146297	1.936521	1.936521	2.010671
NCI-H1623	1.875367	1.875367	1.875367		Hs 600.T	1.986741	1.980923429	2.36405	2.36405	2.36405	2.36405
SVts-8	1.8724745	1.985537	1.759412		Eos-HL-60	1.978554	1.970614857	1.978554	1.978554	2.077617	2.077617
Hs 228.T	1.870225	1.870225	1.870225		P116 T	1.975907	1.965617857	2.116148	2.116148	1.918294	1.918294
NCI-H1437	1.869365	1.869365	1.869365		273T	1.975733	1.9910575	2.260333	2.157985	2.157985	2.260333
647-V	1.8670015	1.85549	1.878513		NCI-H2172	1.974554	1.894567643	2.095817	2.052269	2.052269	2.052269
HCT 116	1.866904	1.866904	1.866904		KYSE-220	1.9735615	2.036028571	1.96575	1.96575	1.96575	1.96575
NCI-H2228	1.859855	1.859855	1.859855		Hs 852.T	1.9655375	1.933576714	1.760728	1.760728	1.760728	1.843924
PC-3 [JPC-3]	1.8587765	1.790605	1.926948		AZ-521	1.963764	1.947559071	1.923261	1.923261	1.923261	2.004267
NCI-H322	1.8566565	1.877686	1.835627		NCI-H650	1.959056	2.0152965	2.375345	2.111266	1.959056	2.375345
23132/87	1.851999	1.851999	1.851999		23132/87	1.947479	1.731576571	2.071248	2.042959	2.042959	2.071248
BT-549	1.8516385	1.806524	1.896753		RKN	1.9466865	1.924271214	1.683528	1.798552	1.955761	1.955761
G-402	1.837503	1.696379	1.978627		HBE4-E6/E7	1.9431555	1.874076643	2.008197	1.960665	2.140435	2.140435
FTC 133	1.835536	1.835536	1.835536		NCI-H1299	1.940986	2.004233071	2.367354	2.147407	1.940986	1.940986
HUP-T3	1.829175	1.829175	1.829175		Hs 573.T	1.939159	1.989307	2.028196	1.939159	1.928339	1.939159
NCI-H889	1.8165655	1.77093	1.862201		Hs 737.T	1.933247	1.870785571	2.106403	2.027839	2.027839	2.106403

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
22RV1	1.811395	1.811395	1.811395		NCI-H2171	1.929906	1.968905857	1.929906	1.929906	1.884451	1.929906
H3255	1.8106385	1.80728	1.813997		G-402	1.928679	1.990094571	2.09111	2.09111	1.892803	2.09111
1321N	1.810554	1.84896	1.772148		MEWO	1.923217	1.942139714	1.916022	1.916022	1.916022	1.923217
CAL-51	1.804006	1.781594	1.826418		CAL-120	1.921765	1.839351571	1.669342	1.680451	1.669342	1.680451
Hs 751.T	1.803885	1.803885	1.803885		AGS	1.916557	1.935016	1.916557	1.916557	1.916557	1.916557
G-292 Clone A141B1	1.803766	1.786957	1.820575		WIL2 NS	1.913585	1.902580429	1.976082	1.913585	1.913585	1.938246
NCI-H711	1.802367	1.748335	1.856399		COR-L 23/CPR	1.906858	1.921375	2.095649	1.878273	1.878273	1.906858
BT-483	1.8011375	1.795797	1.806478		GP5D	1.904712	1.889742357	1.924491	1.924491	1.924491	1.97726
HT 728.T	1.794948	1.744602	1.845294		MCAS	1.9018165	1.772718	1.673513	1.679194	1.890226	1.913407
COLO 684	1.7927315	1.805722	1.779741		COLO-680N	1.901206	1.887987286	1.952449	1.952449	1.901206	1.901206
LS180	1.788976	1.788976	1.788976		LS180	1.90016	1.869492143	2.134672	1.908586	1.90016	1.908586
SK-N-FI	1.787812	1.787812	1.787812		DMS 153	1.896005	1.889634286	1.947149	1.896005	1.896005	1.947149
KYSE-220	1.785009	1.779165	1.790853		SBC-3	1.894659	1.732911643	1.968318	1.894659	1.966567	1.966567
KHOS-312H	1.7823285	1.796649	1.768008		HT 728.T	1.885993	1.887304143	2.034718	1.991996	1.991996	2.034718
Hs 781.T	1.77707	1.758157	1.795983		SW 1573	1.884211	1.9192265	1.745957	1.884211	1.884211	1.985537
HSC-4	1.772512	1.71386	1.831164		MKN74	1.88246	1.874296286	1.869244	1.869244	1.869244	1.895676
NCI-H2172	1.769977	1.769977	1.769977		SAS	1.877083	1.973357714	1.853168	1.986392	2.189161	2.189161
A375.S2	1.7648095	1.749909	1.77971		MKN7	1.874679	1.813077	2.014391	1.956391	2.014391	2.014391
KP-1N	1.763482	1.642588	1.884376		RMG-I	1.8729565	1.807113286	1.957724	1.957724	1.839397	1.971336
PC-14	1.761731	1.761731	1.761731		SVts-8	1.867145	1.834762286	1.912113	1.867145	1.912113	1.912113
ME-180	1.758172	1.758172	1.758172		Hs 894(E).Lu	1.861021	1.853384071	1.827261	1.827261	1.800532	1.827261
G-401	1.7564935	1.674078	1.838909		Hs 751.T	1.859582	1.909508786	2.402402	2.30451	1.859582	1.859582
DMS 92	1.754586	1.702926	1.806246		SK-N-FI	1.850343	1.853057286	2.166182	1.889252	1.889252	2.065034
HO-1-N-1	1.7542175	1.833259	1.675176		NCI-H82	1.848263	1.859927214	2.079681	1.895901	1.895901	2.079681
GP5D	1.7540665	1.742252	1.765881		J GAMMA 1	1.847911	1.855349929	2.23919	2.020672	1.991678	1.991678
SCC-4	1.750302	1.750302	1.750302		HSC-4	1.845019	1.863611571	1.6889	1.82016	1.869878	1.98146
VMRC-RCW	1.748681	1.748681	1.748681		BT-549	1.8412445	1.874990357	2.06279	2.048143	1.948251	1.948251
RERF-LC-MS	1.743687	1.743687	1.743687		KYSE-30	1.834833	1.828907786	1.958046	1.958046	1.958046	1.958046
SKN	1.739092	1.693588	1.784596		Ca9-22	1.834222	1.854555929	1.792761	1.834222	1.838024	1.933422
ABC-1	1.7333545	1.851999	1.61471		NCI-H2405	1.8317355	1.822290714	2.121921	1.762828	1.762828	1.890137
Hs 894(E).Lu	1.731201	1.59834	1.864062		SNU-1	1.829466	1.857600857	2.04837	1.890646	1.890646	2.04837
KOSC-2 cl3-43	1.72599	1.72599	1.72599		BPH-1	1.822954	1.868315286	2.148065	1.968143	1.959012	1.959012
SAS	1.7234445	1.668565	1.778324		NCI-H1944	1.8136765	1.7780645	1.932433	1.879792	1.841654	1.879792
Hs 195.T	1.723323	1.723323	1.723323		HSC-2	1.810627	1.845119571	1.621988	1.915592	1.915592	1.915592
HT1080	1.722561	1.722561	1.722561		HICR 31	1.8103235	1.791867214	1.849811	1.870814	1.870814	1.870814
NCI-H1650	1.722053	1.722053	1.722053		KYSE-50	1.799248	1.779096786	1.799496	1.799496	1.753698	1.799248
Hs 839.T	1.7179875	1.709811	1.726164		NCI-H1915	1.785119	1.8673545	2.223692	2.223692	2.223692	2.223692
LOU-NH91	1.715981	1.564817	1.867145		D341 Med	1.772413	1.698846643	1.807218	1.807218	1.807218	1.807218
C32	1.715628	1.715628	1.715628		PE/CA-PJ15	1.7722435	1.805475286	2.05868	1.858749	1.858749	1.858749
NCI-H1792	1.711836	1.711836	1.711836		KYSE-510	1.7609535	1.770406429	1.666946	1.666946	1.666946	1.711868
A431	1.7067945	1.678049	1.73554		KOSC-2 cl3-43	1.756854	1.890756214	1.694086	1.723466	1.756854	2.187147
PCI 4B	1.7057235	1.682223	1.729224		OUMS-23	1.7526995	1.740450643	2.109655	2.107238	2.107238	2.107238
NCI-H1155	1.703022	1.703022	1.703022		HT1080	1.7494585	1.814138786	1.993616	1.993616	2.050696	1.993616
HBE4-E6/E7	1.6954085	1.695032	1.695785		JURKAT EC-1	1.748791	1.785515643	2.176785	1.875096	1.839748	1.825913
NCI-H2452	1.6915975	1.622169	1.761026		COR-L88	1.746288	1.664343214	1.867855	1.85743	1.846101	1.846101
NCI-H358	1.683913	1.681933	1.685893		DOK	1.744822	1.792563571	2.054761	1.939159	1.888383	1.816934
NCI-H1944	1.6768705	1.67587	1.677871		NCI-H838	1.736552	1.716605071	2.050993	1.849249	1.849249	1.849249

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-225288	SNP_A-201261	SNP_A-211913	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
OAW28	1.6684815	1.648754	1.688209		NCI-H358	1.7251345	1.758348429	2.04259	1.843055	1.843055	2.04259
Ishikawa (Heraklio) 02 ER-	1.6645965	1.64084	1.688353		NCI-H1048	1.72264	1.709694714	1.823951	1.818102	1.799032	1.818102
HMCB	1.6525275	1.61532	1.689735		ME-180	1.712807	1.552682714	1.606851	1.826204	1.818763	1.826204
SW 1463	1.643089	1.630052	1.656126		A431	1.711788	1.7580915	1.555795	1.834329	2.055964	2.055964
HT55	1.640859	1.538866	1.742852		Hs 834.T	1.7024475	1.764145929	2.131186	2.002394	1.876572	2.002394
AN3CA	1.640333	1.640333	1.640333		LS 123	1.7023875	1.629909786	1.390687	1.47482	1.47482	1.718126
OE21	1.634574	1.593245	1.675903		NCI-H1703	1.697225	1.7182055	1.979441	1.940058	1.940058	1.979441
TASK1	1.631984	1.610981	1.652987		VM-CUB1	1.691614	1.709047143	1.952654	1.918754	1.793939	1.793939
KYSE-520	1.631298	1.631298	1.631298		KYSE-180	1.691582	1.566332714	1.892788	1.742079	1.742079	1.742079
OCUM-1	1.6309645	1.663754	1.598175		NCI-H1435	1.68673	1.649101643	1.68673	1.68673	1.68673	1.688434
Ca9-22	1.621495	1.621495	1.621495		KYSE-450	1.670808	1.687648357	1.878739	1.741479	1.687051	1.741479
Hs 600.T	1.6113275	1.612549	1.610106		ABC-1	1.6635725	1.665212857	1.631151	1.631151	1.631151	1.695994
SW 900	1.60389	1.60389	1.60389		Ishikawa (Heraklio) 02 ER-	1.660806	1.820513929	2.312314	2.312301	2.272803	2.272803
HCC-366	1.581444	1.581444	1.581444		NUGC-3	1.6583775	1.754460357	1.621594	1.621594	1.621594	1.630331
BEN	1.580338	1.496922	1.663754		G-292 Clone A1	1.632151	1.819873429	2.284879	2.142015	2.142015	2.142015
Hs 746T	1.5696785	1.542825	1.596532		U-2 OS	1.621524	1.550534214	1.727877	1.727877	1.727877	1.743687
HSC-2	1.56527	1.56527	1.56527		WI-26 VA4	1.612912	1.722526357	2.075495	1.818271	2.075495	2.075495
NCI-H1793	1.556046	1.511365	1.600727		HSC-3	1.60895	1.720568786	1.544295	1.579158	1.589722	1.900101
H-EMC-SS	1.554018	1.543974	1.564062		NCI-H1155	1.606156	1.617121857	1.789892	1.758392	1.606156	1.758392
COR-L 23/CPR	1.550807	1.513073	1.588541		HLE	1.602328	1.656423071	2.075861	1.865906	1.533309	1.865906
BT-20	1.5224575	1.496922	1.547993		TCO-1	1.5967445	1.629325714	1.786429	1.786429	1.783383	1.783383
IMR-90	1.5117595	1.630872	1.392647		NCI-H2122	1.5953045	1.601759429	1.501436	1.501436	1.501436	1.612302
SNU-398	1.497936	1.497936	1.497936		HO-1-u-1	1.584022	1.6465835	1.563542	1.604502	1.703646	1.878258
RPMI-7951	1.470438	1.46945	1.471426		OVISE	1.5777555	1.471765714	1.559427	1.625276	1.625276	1.633642
HCC-827	1.4694065	1.322405	1.616408		JIMT-1	1.569711	1.572775786	1.814166	1.569711	1.569711	1.569711
J82	1.4686905	1.544058	1.393323		GCT	1.559225	1.531814	1.559225	1.559225	1.559225	1.559225
KYSE-70	1.468114	1.468114	1.468114		HCE7	1.5537375	1.6148405	2.067093	1.764879	1.729082	1.764879
NCI-H841	1.4585645	1.448165	1.468964		EPLC-272H	1.5372735	1.428109571	1.554667	1.596516	1.596516	1.700815
JHH-6	1.4487305	1.473573	1.423888		SKN-3	1.5054765	1.490851429	1.524664	1.524664	1.619915	1.619915
HCC-15	1.41844	1.41844	1.41844		NCI-H596	1.50135	1.428170929	1.424804	1.424804	1.50135	1.50135
NCI-H1568	1.416904	1.416904	1.416904		NL20-TA	1.48882015	1.493631193	2.0274962	1.7099382	1.7099382	1.7099382
LS 123	1.4056445	1.284534	1.526755		Hs 839.T	1.486995	1.6536245	1.321401	1.911132	1.911132	2.101073
639-V	1.3860315	1.329038	1.443025		KYSE-270	1.4817775	1.453861071	1.420345	1.333095	1.420345	1.481061
T84	1.3761225	1.264388	1.487857		ASH-3	1.45072175	1.241679157	1.4380864	1.5032531	1.5032531	1.5032531
RKN	1.3741605	1.295513	1.452808		MG 63	1.3840415	1.342891357	1.5996	1.5996	1.5996	1.5996
NCI-H727	1.3719525	1.291864	1.452041		NCI-H522	1.3650465	1.311362857	1.581811	1.581811	1.581811	1.581811
CAL-148	1.35256	1.355801	1.349319		T.T	1.336256	1.387053214	1.508527	1.482494	1.482494	1.482494
HT 1376	1.346498	1.346498	1.346498		T47D	1.322646	1.289987429	1.32811	1.32811	1.32811	1.334058
TE7	1.309042	1.30467	1.313414		SCC-4	1.322629	1.363823	1.218148	1.478901	1.478901	1.478901
CAL-120	1.2995455	1.274516	1.324575		KYSE-520	1.295384	1.345186857	1.526449	1.468964	1.295384	1.468964
MOG-G-CCM	1.286604	1.286604	1.286604		RERF-LC-Sq1	1.2943443	1.204703271	1.5180389	1.5180389	1.4561719	1.4561719
H69V	1.256774	1.078268	1.43528		PCI 4B	1.28546145	1.178102836	1.4498418	1.2974278	1.4498418	1.2974278
HuH-7	1.2508275	1.235518	1.266137		NCI-H1568	1.254747	1.440961214	1.91577	1.794636	1.794636	1.794636
MRK-nu-1	1.2431475	1.227554	1.258741		OSC-20	1.25329	1.312805929	1.292123	1.113359	1.113359	1.495961
KON	1.19488	1.161622	1.228138		RERF-LC-MA	1.2352369	1.109684443	1.4273916	1.4273916	1.4273916	1.4273916
OUMS-23	1.159545	1.096972	1.222118		BICR10	1.225724	1.223030214	1.247097	1.208238	1.208238	1.225724
TCO-1	1.151825	1.151825	1.151825		OE21	1.2121205	1.190524143	1.228044	1.228044	1.228044	1.228044

ProbeSetID	Median	Mean	SNP_A-1989988	SNP_A-206887	SNP_A-2252883	SNP_A-2012613	SNP_A-2119130	SNP_A-2222805	SNP_A-2101835	SNP_A-1989990	SNP_A-4212396
Affy_SNP.ID			2568889	6195985	9040198	4941856	6973989	8566536	6695364	2568944	2568952
dbSNP			rs763317	rs12538489	rs12718939	rs7785313	rs7789965	rs759167	rs729969	rs723527	rs10488140
Chromosome			7	7	7	7	7	7	7	7	7
Position			54869406	54874045	54879529	54887304	54887957	54901993	54902416	54909081	54912597
Strand			+	+	+	-	-	+	-	+	-
Cytoband			p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
NCI-H1299	1.136823	1.136823	1.136823		RERF-LC-Ad2	1.196541	1.216891214	1.105384	1.105384	1.105384	1.129069
MCF7	1.111632	1.111632	1.111632		SKOV3	1.0113754	1.005652571	0.9638997	0.9685444	0.9685444	0.9685444
NCI-H596	1.082276	1.082276	1.082276		T.Tn	0.9614609	0.92826555	0.9614609	0.9614609	1.0114567	1.019878
COR-L88	1.0207988	0.9901213	1.0514763		OAW28	0.9538827	0.988743093	0.7158246	0.6369229	0.7158246	0.9460151
NCI-H2196	0.9397781	0.8724516	1.0071046		NCI-H1563	0.947366	0.914825664	1.0949194	1.0865166	1.0949194	1.0949194
PA-TU-8988T	0.92865655	0.9271194	0.9301937		PNT2	0.90216975	0.829820071	1.0213973	0.9739229	0.9739229	0.914327
ASH-3	0.86861705	0.7290104	1.0082237		NCI-N87	0.8192706	0.81994745	0.7912006	0.7372462	0.7912006	0.8192706
H69	0.8183582	0.8183582	0.8183582		NCI-H2228	0.7844841	0.692114007	0.813008	0.813008	0.813008	0.813008
NCI-H1915	0.59179185	0.5044195	0.6791642		MES-SA	0.6931083	0.711113443	0.6931083	0.6931083	0.5848332	0.6931083
A673	0.5400225	0.5222913	0.5577537		WPE1-NB26	0.60873945	0.59515735	0.7047713	0.7047713	0.7826116	0.7826116
H69AR	0.44263765	0.4036942	0.4815811		NCI-H322	0.5817452	0.548923157	0.7016281	0.6703033	0.6703033	0.6703033
FU97	0.38777175	0.3312451	0.4442984		LU99A	0.112633415	0.132148134	0.27500705	0.27500705	0.40326772	0.40326772

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301045	SNP_A-2050945	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
KYSE-520	6.44475	6.96283	6.96283	6.96283	6.96283	6.96283	6.96283	6.911036	6.66341	6.810133	6.810133
OE21	6.538132	6.538132	6.538132	6.542427	6.542427	6.703002	6.763342	6.763342	6.763342	6.763342	6.763342
H3255	6.410685	6.410685	6.410685	6.489619	6.489619	6.725333	6.725333	6.489619	6.410685	6.436391	6.387916
NCI-H1573	5.669633	5.823272	5.823272	5.756219	5.823272	6.228164	6.228164	5.756219	5.756219	5.797845	5.797845
HSC-2	6.042633	6.082549	6.082549	6.082549	6.082549	6.082549	6.082549	6.07028	5.862791	5.862791	5.862791
HCC-827	5.817453	5.778581	5.778581	5.778581	5.817453	5.778581	5.718564	5.624093	5.624093	5.592099	5.592099
Hs 852.T	5.523843	5.523843	5.523843	5.483529	5.483529	5.483529	5.483529	5.394872	5.394872	5.320132	5.225252
KYSE-30	5.428478	5.528331	5.528331	5.370001	5.370001	5.370001	5.370001	5.22546	5.370001	5.370001	5.370001
A431	4.952981	5.251607	5.251607	5.251607	5.449759	5.319466	5.406551	5.406551	5.406551	5.412951	5.412951
BT-20	5.328966	5.396437	5.396437	5.396437	5.396437	5.396437	5.396437	5.212884	5.212884	5.165112	5.165112
Ca9-22	4.749979	4.940741	4.940741	4.749979	4.749979	4.749979	4.801972	4.749979	4.64452	4.64452	4.64452
NCI-H1568	4.676529	4.676529	4.676529	4.676529	4.676529	4.843599	4.843599	4.676529	4.676529	4.345007	4.345007
PC-9	4.636894	4.636894	4.636894	4.636894	4.636894	4.636894	4.636894	4.621809	4.621809	4.411873	4.277177
PNT1A	4.258236	4.267817	4.267817	4.178325	4.25082	4.25082	4.25082	4.178325	4.178325	4.25082	4.25082
LUDLU-1	4.021817	4.021817	3.977944	3.977944	4.302562	4.33679	4.33679	4.302562	4.302562	4.302562	4.302562
RERF-LC-MA	4.064202	4.168129	4.168129	4.168129	4.168129	4.140779	4.168129	4.168129	4.140779	4.085289	4.078277
U373 MG	4.048313	4.048313	4.048313	4.048313	4.048313	4.048313	4.19363	3.932433	4.19363	4.19363	3.932433
KYSE-450	3.815796	3.815796	3.815796	3.799071	3.799071	3.799071	3.799071	3.791148	3.799071	3.799071	3.839512
KOSC-2 cl3-43	3.853418	3.853418	3.853418	3.853418	4.047412	3.853418	3.853418	3.853418	3.891889	3.891889	3.891889
LU65	3.719899	3.767734	3.751521	3.719899	3.719899	3.719899	3.742142	3.719327	3.719327	3.719327	3.742142
PC-14	3.906605	3.906605	3.902885	3.735516	3.735516	3.735516	3.735516	3.68137	3.68137	3.68137	3.68137
NCI-H596	3.653337	3.653337	3.592797	3.592797	3.645819	3.653337	3.645819	3.592797	3.645819	3.553951	3.645819
PC-3 [JPC-3]	3.798351	3.806077	3.806077	3.806077	3.806077	3.806077	3.806077	3.806077	3.806077	3.806077	3.759412
647-V	3.657458	3.657458	3.657458	3.657458	3.694535	3.694535	3.665173	3.665173	3.657458	3.633134	3.633134
HSC-3	3.613514	3.743829	3.743829	3.613514	3.613514	3.574781	3.574781	3.405058	3.534462	3.278307	3.278307
CASKI	3.469614	3.480862	3.536453	3.536453	3.574606	3.536453	3.497627	3.574606	3.497627	3.574606	3.574606
JHH-4	3.655199	3.655199	3.655199	3.655199	3.655199	3.541971	3.655199	3.655199	3.655199	3.541971	3.541971
CAL-85-1	3.638944	3.566452	3.566452	3.566452	3.566452	3.638944	3.566452	3.516086	3.516086	3.469467	3.469467
BT-549	3.511322	3.512381	3.512381	3.511322	3.512381	3.518653	3.541819	3.511322	3.511322	3.406628	3.541819
HSC-4	3.480836	3.480836	3.480836	3.480836	3.724958	3.480836	3.724958	3.724958	3.450828	3.253103	3.450828
NUGC-4	3.889717	3.889717	3.889717	3.671848	3.671848	3.671848	3.671848	3.598175	3.671848	3.556417	3.444932
BICR10	3.498425	3.498425	3.498425	3.565496	3.565496	3.565496	3.565496	3.565496	3.562107	3.498425	3.562107
KYSE-140	3.6205	3.500964	3.609801	3.500964	3.500964	3.500964	3.500964	3.423342	3.500964	3.423342	3.423342
SNU-5	3.495591	3.495591	3.420266	3.331158	3.420266	3.420266	3.461767	3.331158	3.331158	3.331158	3.418943
SK-MEL-3	3.569896	3.649552	3.569896	3.569896	3.671889	3.671889	3.671889	3.569896	3.671889	3.671889	3.671889
KYSE-180	3.306478	3.514559	3.514559	3.391391	3.391391	3.391391	3.400959	3.391391	3.391391	3.226931	3.391391
PNT2	3.457522	3.578741	3.578741	3.457522	3.430399	3.430399	3.42269	3.344443	3.285994	3.285994	3.285994
G-361	3.684526	3.684526	3.684526	3.504307	3.504307	3.684526	3.684526	3.504307	3.504307	3.419261	3.419261
KYSE-220	3.865377	3.440758	3.440758	3.440758	3.440758	3.578799	3.578799	3.440758	3.278242	3.150044	3.150044
WM-115	3.477138	3.578849	3.578849	3.477138	3.522324	3.522324	3.522324	3.477138	3.477138	3.304945	3.272539
KYSE-50	3.407955	3.407955	3.407955	3.324356	3.324356	3.324356	3.324356	3.237999	3.237999	3.178913	3.178913
OSC-20	3.270497	3.270497	3.270497	3.132013	3.270497	3.270497	3.270497	3.247106	3.270497	3.247106	3.366962

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301043	SNP_A-2050943	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
SAT	3.241407	3.241407	3.241407	3.210638	3.241407	3.241407	3.371414	3.371414	3.371414	3.371414	3.499189
NCI-H1650	3.518346	3.498253	3.484961	3.411823	3.410682	3.410682	3.280761	3.26993	3.26993	3.250625	3.226752
SK-CO-1	3.566612	3.579901	3.579901	3.566612	3.400036	3.400036	3.400036	3.31009	3.31009	3.298624	3.298624
NCI-H1792	3.37014	3.37014	3.210818	3.064305	3.210818	3.210818	3.097603	3.017324	3.037221	3.097603	3.210818
NCI-H1703	3.091284	3.187826	3.254784	3.187826	3.254784	3.210372	3.343755	3.343755	3.343755	3.343755	3.343755
PCI 4B	3.171588	3.235594	3.196797	3.196797	3.196797	3.139215	3.139215	3.139215	3.139215	3.139215	3.139215
NCI-H1563	3.301327	3.309338	3.309338	3.191313	3.191313	3.191313	3.191313	3.165517	3.165517	2.929295	3.165517
NCI-H841	3.132091	3.132091	3.052188	2.892354	3.052188	3.154253	3.198607	3.198607	2.747096	2.747096	2.865897
LOU-NH91	3.217618	3.189097	3.189097	3.150625	3.189097	3.189097	3.189097	3.150625	3.131194	3.091937	3.150625
HuH-7	3.09566	3.058333	3.058333	3.09566	3.174125	3.156129	3.174125	3.156129	3.248015	3.156129	3.248015
LC-1 sq	3.090563	3.090563	3.090563	3.090563	3.081959	3.081959	3.081959	3.081959	3.13976	2.967719	3.244118
NCI-H522	3.146846	3.146846	3.146846	2.843185	3.045447	3.146846	3.343293	3.25499	3.045447	3.045447	3.25499
201T	3.09712	3.09712	2.902449	2.876777	3.138729	3.206871	3.138729	2.876777	2.876777	2.876777	2.926805
G-292 Clone A141B1	2.996675	2.996675	2.996675	2.996675	3.068352	3.178798	3.246704	3.178798	3.178798	3.178798	3.246704
RPMI-7951	3.084605	3.090128	3.090128	3.084605	3.075119	3.090128	3.155249	3.075119	2.930215	2.930215	2.889567
BEN	3.310704	3.310704	3.310704	3.310704	3.310704	3.189097	3.310704	3.169925	3.091937	3.072106	3.072106
NCI-H2228	3.387308	3.387308	3.387308	3.244455	3.244455	3.387308	3.244455	2.920468	2.920468	2.871503	2.871503
HCC-44	3.222571	3.232207	3.222571	3.086408	3.086408	3.086408	3.086408	3.034713	3.045739	3.034713	3.045739
NCI-H1734	3.084288	3.084288	3.084288	3.065023	3.065023	3.409576	3.409576	3.065023	3.065023	3.050554	3.065023
YKG-1	3.343058	3.216663	3.216663	3.176103	3.176103	3.176103	3.216663	3.176103	3.176103	3.06135	3.033795
MOG-G-CCM	3.539163	3.539163	3.539163	3.471859	3.471859	3.471859	3.208959	3.059132	3.059132	2.86254	2.86254
SK-BR-3	3.12143	3.082038	3.082038	3.042101	3.12143	3.12143	3.042101	3.021924	3.021924	3.021924	3.021924
COLO 858	3.167885	3.267048	3.267048	3.167885	3.129868	3.129868	3.129868	3.086051	3.086051	3.086051	3.086051
SCC-4	3.521353	3.521353	3.521353	3.521353	3.521353	3.401146	3.277575	2.968483	2.968483	2.91732	2.968483
TE7	2.999687	2.999687	2.999687	2.999687	3.060421	3.060421	3.072852	3.072852	3.067934	3.067934	3.072852
NCI-H1755	3.092628	3.092628	3.092628	3.080074	2.974829	2.974829	2.974829	2.972533	2.972533	2.78226	2.78226
CaR-1	2.974674	2.948934	2.948934	2.918339	2.918339	2.918339	3.096893	3.096893	3.140528	3.140528	3.140528
OVI5E	2.983398	2.983398	2.983398	2.952857	2.952857	2.952857	2.978038	2.76086	2.978038	2.978038	2.98711
Calu-3	3.065731	3.142402	3.081483	3.065731	3.065731	3.065731	3.033321	2.960385	2.960385	2.960385	2.960385
NCI-H1623	3.035015	3.035015	2.952411	3.035015	2.952411	3.07115	3.07115	3.07115	3.082603	2.793064	2.793064
BICR 78	3.445063	3.445063	3.445063	3.254168	3.254168	3.203602	3.203602	3.103987	3.203602	3.103987	3.091729
LS 123	2.98877	2.98877	3.021833	2.98877	2.98877	3.021833	2.940199	2.940199	2.940199	2.940199	2.946317
AN3CA	3.209319	3.209319	3.098264	3.098264	3.113153	3.276333	3.276333	3.113153	3.113153	3.098264	3.098264
639-V	2.993201	2.981715	2.981715	2.981715	2.93198	2.93198	2.93198	2.93198	3.000034	3.000034	3.045247
WM 266-4	3.117231	3.117231	3.054361	3.054361	3.054361	3.054361	3.054361	2.932147	3.054361	2.932147	2.932147
SKN-3	3.089119	3.089119	3.089119	2.928416	2.928416	2.928416	3.029406	2.928416	3.029406	2.928416	3.029406
NCI-H2085	3.113084	3.113084	3.113084	3.113084	3.113084	3.175959	3.113084	2.956287	2.956287	2.783821	2.783821
RKN	3.073538	3.049893	3.049893	3.049802	3.073538	2.946472	2.946472	2.946472	2.946472	2.798058	2.946472
MKN45	3.029497	3.029497	3.029497	2.992608	2.992608	2.745771	2.992608	2.745771	2.917551	2.917551	2.917551
KHOS/NP	3.250765	3.296351	3.296351	3.061939	3.061939	3.018986	3.015112	2.902185	2.902185	3.015112	3.015112
SNU-16	3.01634	3.005893	3.005893	2.945984	3.005893	3.005893	2.945984	2.899323	2.899323	2.899323	2.945984
HCE7	2.973748	3.024403	3.024403	3.024403	3.024403	2.903451	2.903451	3.024403	2.934736	2.903451	2.934736
OCUG-1	2.836196	2.836196	2.836196	2.836196	2.866464	2.866464	3.05363	3.05363	3.08614	3.05363	3.08614
NL20-TA	2.93947	2.93947	2.93947	2.883583	2.93947	2.910762	2.910762	2.883583	2.883583	2.883583	2.910762
KYSE-70	2.867169	2.866528	2.866528	2.866528	2.866528	2.866528	2.970544	2.866528	2.866528	2.895285	2.970544
MOG-G-UVVW	3.087882	3.087882	3.042735	2.952794	2.952794	2.948934	2.865042	2.865042	2.865042	2.865042	2.865042
NCI-H2342	2.894629	2.894629	2.860635	2.797719	2.797719	2.860635	3.12489	2.860635	2.860635	2.797719	2.797719

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301043	SNP_A-2050942	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
ABC-1	3.150625	3.140926	3.111632	3.072106	3.072106	3.072106	3.072106	2.939772	2.897802	2.712288	2.712288
NCI-H650	2.925675	2.911215	2.858761	2.794719	2.858761	2.858761	2.819032	2.794719	2.794719	2.819032	2.858761
FTC 133	2.75963	2.886184	2.886184	2.886184	2.886184	2.727228	2.816935	2.816935	2.816935	2.816935	2.854504
MDST8	2.853753	2.803163	2.803163	2.803163	2.803163	2.803163	2.803163	2.832804	2.832804	2.863203	2.887913
KYSE-150	2.854032	2.852089	2.852089	2.820802	2.820802	2.852089	2.852089	2.820802	2.699953	2.595075	2.699953
SW 48	2.915252	2.896681	2.753003	2.753003	2.866763	2.753003	2.753003	2.67929	2.67929	2.67929	2.67929
NCI-H2023	3.029658	3.06171	3.06171	2.997217	2.849425	2.997217	2.997217	2.849425	2.849425	2.849425	2.741009
COLO-680N	3.578991	3.535174	3.535174	3.152154	3.152154	3.106593	3.079558	2.881362	2.881362	2.881362	2.881362
KP-4	3.113457	2.889514	2.889514	2.858707	2.8349	2.8349	2.8349	2.830803	2.8349	2.830803	2.8349
SW-1710	3.088862	3.07501	3.07501	2.828271	2.828271	2.701334	2.527199	2.527199	2.527199	2.527199	2.828271
KATO III	2.991032	3.003021	3.040995	2.905454	2.813345	2.813345	2.813345	2.728663	2.728663	2.589047	2.589047
RERF-GC-1B	2.829895	2.871492	2.829895	2.741277	2.741277	2.741277	2.829895	2.741277	2.646094	2.646094	2.646094
CAL-29	2.844175	2.790488	2.790488	2.790488	2.844175	2.811331	2.811331	2.790488	2.811331	2.811331	2.811331
Sci-1	2.81144	2.845424	2.804648	2.727654	2.727654	2.727654	2.804648	2.727654	2.727654	2.727654	2.727654
NCI-H810	3.1006	2.929305	2.929305	2.929305	3.039144	3.039144	3.039144	2.929305	2.929305	2.800201	2.800201
T.Tn	2.803403	2.917436	2.803403	2.803403	2.803403	2.803403	2.803403	2.648515	2.803403	2.648515	2.648515
DOK	2.992669	2.992669	2.707821	2.707821	2.707821	2.805252	2.67187	2.67187	2.67187	2.67187	2.849264
COLO 741	2.91316	2.91316	2.91316	2.824728	2.91316	2.91316	2.91316	2.824728	2.824728	2.808684	2.808684
NCI-H1651	2.786129	2.786129	2.786129	2.786129	2.778219	2.795332	2.814695	2.778219	2.732046	2.631318	2.631318
RWPE-1	2.80242	2.80242	2.80242	2.80242	2.925539	2.966925	2.925539	2.925539	2.766043	2.66291	2.766043
KYSE-270	2.844789	2.839423	2.839423	2.762888	2.742594	2.742594	2.762888	2.641385	2.641385	2.578489	2.641385
NCI-H196	3.013204	3.076123	3.076123	2.647766	2.787766	3.080223	2.787766	2.736006	2.736006	2.729414	2.736006
SK-LU-1	2.980021	2.980021	2.980021	2.937094	2.937094	2.937094	2.937094	2.818695	2.818695	2.771823	2.766452
NCI-H1693 (ship16)	2.831701	2.831701	2.831701	2.826722	2.688384	2.826722	2.826722	2.688384	2.641512	2.54591	2.490456
NCI-H2135	2.743898	2.801294	2.743898	2.801294	2.801294	2.826267	2.826267	2.801294	2.739424	2.683915	2.739424
VMRC-RCW	3.003877	2.910899	2.910899	2.910899	2.910899	2.878991	2.878991	2.732449	2.732449	2.70904	2.70904
RERF-LC-Sq1	2.985706	3.022673	3.039667	3.022673	3.022673	3.039667	3.022673	2.632094	2.732214	2.732214	2.732214
A375.S2	3.030668	2.950606	3.030668	2.89832	2.89832	2.89832	2.884985	2.640507	2.640507	2.570793	2.640507
SW 1116	3.060481	3.047184	3.047184	2.887298	2.887298	2.887298	2.691292	2.606509	2.606509	2.578677	2.578677
CAL-148	3.103396	2.972317	2.972317	2.972317	2.972317	2.972317	2.972317	2.972317	2.86971	2.823568	2.728069
22RV1	2.917677	2.917677	2.917677	2.751691	2.751691	2.578418	2.570591	2.541728	2.541728	2.541728	2.570591
IM-95	2.860356	2.829516	2.860356	2.829516	2.829516	2.829516	2.829516	2.825975	2.811756	2.789939	2.726768
HT 115	2.676803	2.743084	2.743084	2.72651	2.72651	2.72651	2.72651	2.72651	2.676803	2.72651	2.72651
NCI-H727	3.13694	2.984609	2.984609	2.773754	2.773754	2.773754	2.721212	2.535743	2.721212	2.721212	2.721212
T.T	2.79774	2.79774	2.79774	2.79774	2.79774	2.864925	2.864925	2.77169	2.77169	2.765135	2.765135
OE33	2.740764	2.726006	2.726006	2.672418	2.726006	2.635835	2.635835	2.635835	2.635835	2.635835	2.71355
PLC/PRF/5	2.658546	2.729492	2.75821	2.729492	2.729492	2.729492	2.75821	2.75821	2.729492	2.75821	2.780598
SAS	2.883434	2.829938	2.829938	2.829938	2.829938	2.78525	2.829938	2.829938	2.829938	2.770354	2.770354
MKN74	2.717288	2.717288	2.809501	2.71947	2.71947	2.71947	2.735871	2.71947	2.71947	2.614064	2.638692
HMCB	2.343707	2.343707	2.270676	2.270676	2.270676	2.343707	2.343707	2.810558	2.343707	2.664935	2.664935
A2058	2.98587	2.932231	2.932231	2.932231	2.932231	2.931113	2.931113	2.931113	2.926732	2.802048	2.802048
HCC-366	2.947958	2.947958	2.947958	2.754436	2.754436	2.754436	2.745926	2.694957	2.694957	2.676061	2.676061
P31/FUJ	2.78029	2.695229	2.695229	2.695229	2.695229	2.695229	2.698595	2.695229	2.695229	2.672578	2.672578
SW 1417	2.814434	2.814434	2.814434	2.775243	2.775243	2.775243	2.762046	2.709818	2.706353	2.706353	2.706353
KMRC-1	2.666206	2.666206	2.666206	2.666206	2.703912	2.673538	2.673538	2.555006	2.673538	2.673538	2.703912
OAW42	2.814912	2.759708	2.759708	2.69617	2.69617	2.69617	2.69617	2.639675	2.556862	2.548929	2.556862
PE/CA-PJ15	2.662264	2.662264	2.662264	2.602075	2.602075	2.602075	2.662264	2.662264	2.602075	2.602075	2.602075

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301043	SNP_A-2050943	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989994
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
NUGC-3	2.837405	2.817467	2.817467	2.817467	2.755391	2.755391	2.81328	2.755391	2.578146	2.520268	2.520268
DMS 273	2.810612	2.814869	2.810612	2.759109	2.757711	2.810612	2.753225	2.676952	2.676952	2.651855	2.651855
MKN1	2.797554	2.797554	2.797554	2.686998	2.749545	2.686998	2.749545	2.686998	2.686998	2.642413	2.642413
RERF-LC-Ad2	2.86254	2.755491	2.755491	2.723222	2.723222	2.642667	2.642667	2.554791	2.642667	2.554791	2.554791
TASK1	2.942739	2.729828	2.729828	2.729828	2.729828	2.655076	2.729828	2.729828	2.729828	2.655076	2.655076
NCI-H23	2.751325	2.640172	2.640172	2.62612	2.618341	2.62612	2.62612	2.618341	2.618341	2.553791	2.590847
KP-1N	2.890458	2.890458	2.890458	2.890458	2.890458	2.634724	2.634724	2.634724	2.623518	2.60679	2.623518
NCI-H711	2.775431	2.775431	2.775431	2.702985	2.648688	2.648688	2.775431	2.618259	2.618259	2.618259	2.622425
NCI-H661	2.839876	2.709604	2.709604	2.709604	2.702759	2.702759	2.702759	2.642402	2.642402	2.642402	2.621297
EBC-1	2.757023	2.757023	2.757023	2.734742	2.734742	2.734742	2.734742	2.689657	2.632291	2.62068	2.62068
JHH-6	2.661449	2.661449	2.661449	2.661449	2.616256	2.616256	2.616256	2.616256	2.661449	2.616256	2.779299
NB69	2.609723	2.840501	2.840501	2.708351	2.708351	2.840501	2.840501	2.708351	2.708351	2.708351	2.708351
SR	2.744132	2.810405	2.744132	2.720684	2.720684	2.720684	2.691145	2.662424	2.662424	2.66083	2.66083
T84	2.604509	2.643925	2.643925	2.643925	2.643925	2.631886	2.604509	2.631886	2.604509	2.631886	2.643925
HO-1-N-1	2.627502	2.627502	2.627502	2.627502	2.803468	2.803468	2.803468	2.803468	2.77179	2.77179	2.803468
BALL-1	2.753936	2.753936	2.753936	2.753936	2.65773	2.65773	2.790268	2.790268	2.880693	2.532925	2.880693
Nthy-ori 3-1	2.861887	2.811745	2.811745	2.671218	2.671218	2.671218	2.810122	2.671218	2.671218	2.612139	2.612139
FTC 238	2.645045	2.599685	2.645045	2.599685	2.599685	2.599685	2.599685	2.4464	2.599685	2.4464	2.4464
HuCC1	2.764659	2.742359	2.742359	2.742359	2.721436	2.721436	2.721436	2.700847	2.700847	2.595263	2.700847
HOS	2.694538	2.701515	2.72192	2.72192	2.72192	2.72192	2.694538	2.493096	2.493096	2.493096	2.52382
NCI-H2170	2.587636	2.587636	2.587636	2.587636	2.587636	2.825108	2.825108	2.450488	2.450821	2.450488	2.450821
Hs 936.T	2.739257	2.739257	2.739257	2.706782	2.706782	2.739257	2.706782	2.441861	2.414761	2.318884	2.318884
NCI-H1793	2.9002	2.9002	2.629938	2.582747	2.582747	2.629938	2.582747	2.397793	2.450315	2.450315	2.526093
MRK-nu-1	2.719807	2.587859	2.587859	2.587859	2.587859	2.587859	2.719807	2.587859	2.58007	2.577449	2.577449
KHOS-312H	2.846436	2.981171	2.846436	2.817185	2.817185	2.801251	2.569064	2.479847	2.479847	2.569064	2.584645
LCLC-103H	2.80349	2.808335	2.720639	2.607854	2.670486	2.670486	2.670486	2.607854	2.607854	2.543019	2.564998
H69V	2.873464	3.262218	3.36689	3.262218	3.262218	3.186182	3.186182	2.767624	2.767624	2.559905	2.559905
GCT	2.564014	2.564014	2.607164	2.510651	2.607164	2.672738	2.672738	2.607164	2.607164	2.607164	2.607164
COLO 857	2.695422	2.717771	2.717771	2.607971	2.607971	2.577945	2.607971	2.577945	2.577945	2.577945	2.577945
RERF-LC-MS	2.822852	2.757023	2.757023	2.712288	2.712288	2.712288	2.712288	2.632291	2.632291	2.550014	2.550014
MKN7	2.585563	2.585563	2.573762	2.477574	2.573762	2.597317	2.597317	2.477574	2.597317	2.477574	2.42825
A-427	2.68388	2.66153	2.66153	2.66153	2.66153	2.558871	2.534808	2.534808	2.534808	2.534808	2.534808
T47D	2.568306	2.659625	2.659625	2.659625	2.659625	2.577367	2.443767	2.443767	2.443767	2.443767	2.510155
BT-483	2.592633	2.515964	2.515964	2.515964	2.509575	2.509575	2.502857	2.495977	2.495977	2.401599	2.445374
SKN	2.827037	2.778385	2.778385	2.773313	2.773313	2.773313	2.752747	2.53615	2.53615	2.362409	2.53615
HCC202	2.604673	2.604673	2.604673	2.604673	2.592868	2.461845	2.445856	2.39203	2.384451	2.384451	2.368726
OCUM-1	2.62068	2.62068	2.62068	2.573762	2.573762	2.573762	2.573762	2.477574	2.62068	2.390695	2.378068
NCI-H2122	2.703042	2.617863	2.617863	2.617863	2.617863	2.492184	2.492184	2.438715	2.492184	2.492184	2.505158
HT 29	2.337246	2.480274	2.581439	2.477097	2.477097	2.477097	2.477097	2.477097	2.477097	2.477097	2.581439
huH-1	2.658718	2.507265	2.507265	2.507265	2.507265	2.480336	2.480336	2.480336	2.480336	2.419942	2.419942
HCC-15	2.487058	2.487058	2.487058	2.487058	2.578205	2.593925	2.611848	2.593925	2.578205	2.570994	2.570994
KHOS-240S	2.624087	2.69312	2.641974	2.624087	2.624087	2.554315	2.432761	2.351458	2.351458	2.432761	2.554315
JIMT-1	2.473781	2.467801	2.467801	2.467801	2.512548	2.467801	2.467801	2.467801	2.458241	2.467801	2.512548
KON	2.595451	2.595451	2.595451	2.595451	2.463553	2.463553	2.586717	2.332814	2.586717	2.332814	2.586717
COLO 853	2.624901	2.53724	2.605223	2.624901	2.624901	2.605223	2.605223	2.421598	2.516326	2.516326	2.457194
U-2 OS	2.757023	2.700994	2.700994	2.700994	2.700994	2.700994	2.700994	2.643856	2.453017	2.453017	2.453017
ME-180	2.693267	2.693267	2.693267	2.693267	2.650727	2.506357	2.506357	2.506357	2.448612	2.442938	2.448612

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301043	SNP_A-2050942	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
FU97	2.459828	2.459828	2.459828	2.440152	2.440152	2.440152	2.440152	2.440152	2.440152	2.585999	2.675708
Calu-1	2.5783	2.5783	2.528088	2.4839	2.528088	2.528088	2.528088	2.399313	2.399313	2.436149	2.528088
NCI-H2452	2.912677	2.945329	2.945329	2.527775	2.527775	2.527775	2.527775	2.476767	2.476767	2.471895	2.471895
UMC-11	2.701866	2.623343	2.623343	2.623343	2.619004	2.58521	2.457428	2.457428	2.182736	2.167067	1.938157
COLO 684	2.404058	2.404058	2.310035	2.179417	2.371416	2.404058	2.371416	2.371416	2.317165	2.317165	2.317165
SVts-8	2.8872133	2.8872133	2.8872133	2.8228525	2.844466	2.844466	2.844466	2.7791336	2.7791336	2.7791336	2.7791336
HEC-1	2.616465	2.430301	2.430301	2.430301	2.430301	2.430301	2.430301	2.422174	2.422174	2.422174	2.399539
A-375	2.592656	2.502141	2.502141	2.397768	2.592656	2.397768	2.397768	2.390305	2.397768	2.397768	2.598911
OUMS-23	2.462336	2.462336	2.462336	2.34103	2.34103	2.34103	2.549609	2.316907	2.549609	2.316907	2.611953
EJ138	2.656961	2.700168	2.705958	2.606884	2.606884	2.606884	2.399853	2.375953	2.375953	2.366312	2.366312
HO-1-u-1	2.39266	2.39266	2.39266	2.39266	2.39266	2.39266	2.395492	2.39266	2.395492	2.395492	2.443841
HT55	2.530346	2.530346	2.533417	2.454521	2.391842	2.391842	2.391842	2.391842	2.391842	2.391842	2.391842
NCI-H2347	2.53458	2.53458	2.534568	2.534568	2.613551	2.613551	2.613551	2.613551	2.483705	2.249212	2.249212
REFL-LC-KJ	2.736866	2.736866	2.736866	2.602707	2.602707	2.602707	2.602707	2.488033	2.488033	2.488033	2.602707
COR-L279	2.730724	2.590129	2.590129	2.413361	2.413361	2.413361	2.413361	2.413361	2.413361	2.4376	2.413361
SW 1573	2.374889	2.36377	2.36377	2.36377	2.374889	2.36377	2.36377	2.157425	2.157425	2.157425	2.41606
SW 900	2.52834	2.528004	2.528004	2.486168	2.486168	2.528004	2.528004	2.486168	2.387657	2.319439	2.486168
HT 1376	2.487631	2.477182	2.377663	2.362001	2.377663	2.377663	2.477182	2.477182	2.477182	2.477182	2.477182
HUP-T3	2.529121	2.529121	2.529121	2.529121	2.464892	2.428274	2.464892	2.464892	2.464892	2.200381	2.464892
STC 1	2.35359	2.35359	2.356945	2.356945	2.356945	2.393126	2.387316	2.356945	2.356945	2.356945	2.356945
NCI-H2286	2.311874	2.290037	2.311874	2.311874	2.311874	2.410647	2.410647	2.341299	2.341299	2.341299	2.341299
VMRC-LCD	2.643856	2.585563	2.585563	2.465322	2.465322	2.465322	2.585563	2.585563	2.585563	2.33985	2.33985
H69AR	2.480726	2.480726	2.551766	2.480726	2.480726	2.551766	2.551766	2.300534	2.300534	2.118904	2.065076
VMRC-LCP	2.76178	2.435529	2.435529	2.435529	2.435529	2.389145	2.389145	2.389145	2.380218	2.33487	2.33487
FLYA13	2.515542	2.4576	2.4576	2.415211	2.415211	2.415211	2.415211	2.415211	2.397931	2.274574	2.274574
HUP-T4	2.452117	2.421351	2.421351	2.308596	2.421351	2.260623	2.260623	2.260623	2.421351	2.40023	2.421351
COR-L 105	2.341952	2.341952	2.341952	2.326598	2.326598	2.321698	2.317592	2.315278	2.306832	2.236002	2.315278
KMRC-20	2.302951	2.235191	2.235191	2.182858	2.235191	2.235191	2.235191	2.366757	2.366757	2.366757	2.43173
NCI-H2081	2.617456	2.529457	2.529457	2.491758	2.363108	2.363108	2.409371	2.345065	2.345065	2.317514	2.295681
VM-CUB1	2.261282	2.344732	2.344732	2.320149	2.182276	2.182276	2.182276	2.182276	2.182276	2.182276	2.320149
BU25 TK	2.496365	2.496365	2.496365	2.21729	2.14288	2.14288	2.171339	2.171339	2.21729	2.171339	2.21729
RCM-1	2.523122	2.523122	2.523122	2.319581	2.290351	2.287897	2.287897	2.287897	2.287897	2.146105	2.287897
NCI-H2172	2.351752	2.262178	2.229294	2.229294	2.229294	2.262178	2.262178	2.229294	2.229294	2.216554	2.216554
SK-NEP-1	2.283689	2.283689	2.283689	2.283689	2.283689	2.378903	2.283689	2.283689	2.283689	2.283153	2.283689
D1.1	2.375801	2.375801	2.506539	2.375801	2.099415	2.099415	2.114554	2.114554	2.124299	2.114554	2.114554
NCI-H2196	2.394737	2.381381	2.381381	2.279764	2.280615	2.280615	2.280615	2.279764	2.280615	2.280615	2.280615
NCI-H1435	2.322486	2.322486	2.325245	2.322486	2.325245	2.325245	2.325245	2.276947	2.276947	2.276947	2.276947
GRANTA-519	2.713561	2.413474	2.413474	2.150394	2.444422	2.150394	2.150394	2.27561	2.31256	2.27561	2.27561
SK-MES	2.486278	2.486278	2.508499	2.486278	2.486278	2.362205	2.379902	2.379902	2.379902	2.362205	2.357519
NCI-H1437	2.264798	2.264798	2.264798	2.120496	2.120496	2.264798	2.256917	2.264798	2.264798	2.304795	2.304795
BFTC-905	2.418183	2.259674	2.259674	2.259674	2.259674	2.259674	2.259674	2.259674	2.283193	2.306314	2.350002
DMS 153	2.52028	2.289711	2.289711	2.491283	2.324549	2.324549	2.324549	2.324549	2.324549	2.03882	2.03882
Detroit 562	2.228054	2.256824	2.256824	2.239549	2.226267	2.239549	2.256824	2.256824	2.256824	2.256824	2.256824
RT4	2.382456	2.382456	2.382456	2.345052	2.345052	2.345052	2.382456	2.375561	2.375561	2.375561	2.338118
273T	2.186728	2.186728	2.186728	2.174695	2.186728	2.186728	2.241686	2.241686	2.241686	2.241686	2.208539
SW 1463	2.40639	2.40639	2.40639	2.329058	2.329058	2.329058	2.329058	2.238128	2.329058	2.238128	2.329058
H69	2.063214	2.151435	2.160074	2.160074	2.160074	2.151435	2.160074	2.289007	2.294118	2.294118	2.294118

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301043	SNP_A-2050943	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989994
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
Hs 195.T	2.684495	2.684495	2.684495	2.684495	2.714316	2.684495	2.434399	2.37101	2.37101	2.37101	2.37101
NCI-H1781	2.593749	2.417422	2.417422	2.417422	2.417422	2.363032	2.268626	2.198496	2.109016	2.103329	2.103329
WIDR	2.451784	2.451784	2.451784	2.432761	2.234699	2.234699	2.229947	2.229947	2.229947	2.229947	2.229947
SK-MEL-1	2.567915	2.567915	2.26086	2.257748	2.190472	2.257748	2.257748	2.190472	2.190472	2.190472	2.049902
SUIT-2	2.462742	2.462742	2.462742	2.375763	2.357225	2.212777	2.212777	2.212777	2.212777	2.153378	2.153378
WPE1-NB26	2.364342	2.364342	2.294639	2.273459	2.273459	2.294639	2.294639	2.273459	2.273459	2.104707	2.273459
HCC-78	2.656788	2.656788	2.656788	2.616582	2.656788	2.656788	2.616582	2.594888	2.594888	2.47794	2.47794
A549	2.301509	2.301509	2.301509	2.19754	2.19754	2.19754	2.201767	2.201767	2.201767	2.201767	2.201767
LK-2	2.259753	2.259753	2.266141	2.083804	2.266141	2.083804	2.083804	2.274391	2.274391	2.274391	2.298519
NCI-H2330	2.377207	2.377207	2.398308	2.398308	2.398308	2.377207	2.198132	2.198132	2.198132	2.198132	2.198132
COLO 685	2.360028	2.310333	2.310333	2.310333	2.21145	2.310333	2.21145	2.08013	2.07098	2.1798351	2.1798351
BICR22	2.379915	2.379915	2.446351	2.446351	2.446351	2.277943	2.277943	2.076409	2.108224	2.086451	2.086451
MEWO	2.309996	2.309996	2.38406	2.38406	2.38406	2.38406	2.38406	2.309996	2.309996	2.38406	2.298337
IHH-4	2.181084	2.181084	2.181084	2.162106	2.181084	2.181084	2.181084	2.301691	2.058849	2.190917	2.301691
NCI-N87	2.161493	2.161493	2.221328	2.161493	2.161493	2.161493	2.094672	2.094672	2.094672	2.094672	2.011404
KYSE-410	2.298454	2.288106	2.274653	2.252717	2.226761	2.226761	2.216206	2.186417	2.186417	2.176907	2.176907
DMS 92	2.513393	2.513393	2.607526	2.513393	2.513393	2.182343	2.224373	2.398936	2.398936	2.292229	2.224373
RO82-W-1	2.302107	2.453214	2.249927	2.226721	2.226721	2.222009	2.222009	2.222009	2.199237	2.199237	2.175428
NCI-H82	2.170632	2.170632	2.166386	2.07122	2.060672	2.060672	2.060672	2.060672	2.060672	2.060672	2.150066
C32	2.334536	2.223905	2.150216	2.125722	2.125722	2.150216	2.171556	2.171556	2.171556	2.171556	2.183697
Hs 633T	2.211745	2.26789	2.26789	2.211745	2.26789	2.26789	2.26789	2.26789	2.26789	2.255636	2.255636
NCI-H2405	2.214545	2.214545	2.214545	2.168414	2.122446	2.122446	2.118683	2.118683	2.071515	2.071515	2.132537
Hs 895.T	2.364037	2.364037	2.141685	2.141685	2.142777	2.141685	2.188107	2.114277	2.114277	2.114277	2.114277
MG 63	2.118254	2.175347	2.269874	2.269874	2.269874	2.175347	2.118254	2.118254	2.118254	2.269874	2.269874
CAL-120	2.168863	2.168863	2.168863	2.190458	2.215269	2.190458	2.190458	2.222517	2.222517	2.299286	2.299286
H33HJ-JA1	2.426383	2.440028	2.440028	2.31745	2.31745	2.31745	2.309853	2.309853	2.309853	2.297556	2.288041
JSC-1	2.439508	2.439508	2.439508	2.432177	2.394611	2.230693	2.131158	2.131158	2.131158	2.126771	2.126771
Hs 751.T	2.552124	2.344412	2.344412	2.225494	2.225494	2.225494	2.02931	1.934809	1.934809	1.888652	1.934809
HMVII	2.245107	2.383277	2.245107	2.208579	1.965911	1.965911	2.208579	2.208579	2.208579	2.208579	1.959539
fR2	2.238912	2.238912	2.215376	2.139019	2.139019	2.139019	2.139019	2.139019	2.139019	2.139019	2.129366
COLO 201	2.382355	2.382355	2.382355	2.283127	2.283127	2.283127	2.382355	2.283127	2.283127	2.124672	2.124672
NCI-H1915	2.416622	2.416622	2.377156	2.362116	2.345922	2.362116	2.362116	2.345922	2.315006	2.315006	2.345922
IMR-90	2.525095	2.525095	2.525095	2.525095	2.509708	2.423507	2.509708	2.509708	2.394661	2.290298	2.290298
RT112/84	2.439235	2.402916	2.402916	2.380522	2.380522	2.380522	2.209988	2.209988	2.31379	2.31379	2.209988
Hs 894(E).Lu	2.236773	2.161997	2.047121	2.047121	2.047121	2.047121	2.047121	2.047121	2.047121	2.121091	2.126412
LNCaP clone FGC	2.156577	2.156577	2.156577	2.108322	2.156577	2.108322	2.108322	2.108322	1.956875	1.956875	1.916809
NCI-H1581	2.111197	2.111197	2.111197	2.057152	2.050965	2.057152	2.057152	1.960797	1.960797	2.057152	2.163811
HCT 116	2.032465	2.032465	2.042135	2.042135	2.055129	2.098676	2.098676	2.146407	2.146407	2.146407	2.146407
Hs 228.T	2.331837	2.331837	2.376561	2.331837	2.331837	2.331837	2.331837	2.31626	2.31626	2.18635	2.18635
JURKAT EC-1	2.096947	2.179796	2.179796	2.069755	1.983941	1.983941	2.069755	2.127751	2.069755	2.069755	2.069755
SK-N-BE(2)	2.345602	2.332968	2.332968	2.332968	2.332968	2.332968	2.332968	2.304834	2.304834	2.210619	2.209197
LS180	2.070122	2.070122	2.119015	1.926623	1.926623	1.926623	2.127986	2.130304	2.130304	2.20404	2.20404
CAL-51	2.202883	2.185673	2.202883	2.185673	2.077996	2.077996	2.077996	2.23539	2.23539	2.172399	2.172399
SKG-IIIb	2.195936	2.087795	2.087795	2.028596	2.028596	2.028596	2.028596	2.028596	2.150476	2.028596	2.150476
P116 T	2.151996	2.17608	2.18612	2.17608	2.18612	2.18612	2.21153	2.21153	2.21153	2.21153	2.21153
Hs 417.Lu	2.346984	2.29185	2.346984	2.177138	2.177138	2.346984	2.346984	2.177138	2.177138	2.177138	2.177138
WIL2 NS	2.268021	2.24146	2.24146	2.000173	1.936727	1.936727	2.000173	2.000173	2.000173	2.000173	1.936727

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896871	SNP_A-4301043	SNP_A-2050943	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
COR-L95	2.208472	2.208472	2.100794	2.084308	2.010944	2.010944	2.010944	2.010944	2.010944	2.010944	2.100794
SK-N-FI	2.175509	2.209774	2.30997	2.209774	2.209774	2.175509	2.223385	2.175509	2.175509	2.094211	2.08041
WILCL	2.323492	2.323492	2.323492	2.26547	2.26547	2.323492	2.091417	2.091417	2.091417	1.945738	1.945738
P32/ISH	2.108613	2.066444	2.07364	2.07364	2.07364	2.07364	2.07364	2.107724	2.107724	2.102647	2.107724
EPLC-272H	2.430338	2.343272	2.343272	2.343272	2.303211	2.212375	2.177911	2.177911	2.177911	2.069516	2.069516
A2780 ADR	2.194574	2.194574	2.194574	2.066684	1.90331	1.90331	2.066684	2.066684	2.066684	2.066684	2.066684
C8166	2.232051	2.232051	2.189026	1.94781	1.94781	1.978976	1.94781	1.94781	1.94781	1.946106	1.94781
Hs 908.T	2.146599	2.073232	2.031809	2.014893	2.014893	2.031809	2.031809	2.014893	1.975122	1.975122	2.014893
NCI-H322	2.173174	2.173174	2.241407	2.228627	2.228627	2.173174	2.20178	2.196327	2.196327	2.196327	2.196327
NCI-H1417	2.754869	2.690373	2.690373	2.690373	2.405111	2.405111	2.405111	2.174573	1.824426	1.824426	1.817518
HBE4-E6/E7	2.293024	2.180325	2.180325	2.150627	2.150627	2.150627	2.451512	2.451512	2.451512	2.451512	2.451512
NCI-H1522	2.16851	2.050511	1.939999	1.939999	1.92	1.939999	1.939999	1.980646	1.939999	1.939999	1.980646
LU99A	2.275007	2.275007	2.275007	2.275007	2.378068	2.275007	2.403268	2.168129	2.154486	2.154486	2.140779
HCT-8	2.269244	2.267035	2.133997	2.054634	2.054634	2.054634	2.054634	2.054634	2.054634	2.054634	2.114263
SBC-5	2.068065	2.055426	2.055426	1.915309	1.915309	1.915309	2.055426	2.055426	2.055426	1.952654	1.952654
Ishikawa	2.192389	2.068868	2.068868	2.068868	1.89608	1.89608	2.107224	2.107224	2.150353	2.150353	2.208888
U266B1	2.054096	2.054096	2.054096	2.067248	2.067248	2.141466	2.141466	2.142757	2.19363	2.141466	2.19363
PA-TU-8988T	2.175523	2.14288	2.14288	2.14288	2.14288	2.048497	2.048497	2.048497	2.048497	2.048497	2.048497
HRT-18	2.293271	2.232503	2.232503	2.205598	2.205598	2.205598	2.196435	2.196435	2.196435	2.196435	2.134203
C170	2.234512	2.234512	2.286578	2.234512	2.093136	2.047107	2.021331	2.021331	1.9244	2.021331	2.021331
J GAMMA 1	2.043016	2.043016	2.043016	2.040471	1.756101	1.740579	1.756101	1.756101	1.756101	2.031609	1.982055
LS174T	2.040812	2.172018	2.04259	2.040812	2.040812	2.040812	1.939027	1.939027	1.939027	2.040812	1.958778
NCI-H526	2.203018	2.20131	2.203018	2.203018	2.09206	2.210833	2.210833	2.084924	2.084924	2.049476	2.049476
HLE	2.10607	2.10607	2.10607	2.10607	2.072768	1.945841	1.945841	1.945841	1.951627	1.945841	1.951627
23132/87	2.208673	2.168129	2.071248	2.028711	2.014391	2.014391	2.014391	2.014391	2.014391	2.014391	2.014391
Hs 834.T	2.085177	2.252941	2.055766	2.023879	1.973812	1.973812	1.973812	1.973812	1.973812	1.973812	2.012639
Hs 891.T	2.217531	2.32451	2.32451	2.231212	2.231212	2.32451	2.320356	2.231212	2.231212	2.194345	2.076676
Hs 839.T	2.800671	2.427404	2.427404	2.427404	2.427404	2.159009	2.159009	2.159009	2.159009	2.049263	2.049263
SNU-1	2.23016	2.219162	2.219162	2.219162	2.195235	1.91311	1.91311	1.91311	1.91311	1.976388	2.004714
EB1	2.050752	2.058439	2.050752	2.004555	1.832953	1.732882	1.732882	1.732882	1.732882	1.892354	2.000692
NCI-H2009	2.168101	2.129669	2.129669	2.086073	2.040599	2.040599	2.040599	1.988783	2.040599	1.988783	2.040599
NCI-H1048	2.014305	1.986378	1.986378	1.8791	1.986378	1.986378	2.092787	1.986378	1.8791	1.794357	1.8791
A2780	1.985856	1.985856	1.985856	1.985856	1.983013	1.983013	1.983013	1.983013	1.983013	1.983013	2.063764
NCI-H1666	1.806077	1.743687	1.61471	1.743687	1.743687	1.743687	1.897202	2.057138	2.057138	2.042959	2.042959
G-402	2.057308	1.995871	2.057308	1.995871	1.995871	1.995871	1.995871	2.095286	2.095286	2.095286	2.130069
JHH-7	2.077069	2.072247	2.072247	2.072247	2.063736	2.010239	2.010239	2.063736	2.010239	2.010239	2.010239
OUS-11	2.160292	2.099025	2.014046	1.950439	1.942002	1.942002	1.942002	1.942002	1.942002	1.942002	1.942002
AZ-521	2.320653	2.29939	2.118987	1.978932	1.967151	1.967151	1.978932	1.978932	2.051631	1.978932	1.978932
K562/ADM	1.995365	1.995365	1.995365	1.995365	1.990187	1.990187	1.990187	1.990187	1.974715	1.900414	1.974715
NCI-H889	2.846005	2.846005	2.846005	2.283689	2.283689	2.283689	2.343259	2.283689	2.283689	2.19278	2.143443
OE19	1.878212	1.878212	1.971904	1.971904	1.974016	1.971904	1.878212	1.971904	1.907946	1.971904	1.971904
HT 728.T	2.155238	2.132247	2.155238	2.08243	1.988102	1.67453	1.67453	1.784341	1.67453	1.67453	2.023406
GA-10	1.822095	1.822095	1.774973	1.715516	1.715516	1.715516	1.774973	1.774973	1.774973	2.197297	2.197297
Hs 737.T	1.953739	1.953739	2.012941	2.012941	2.012941	1.953739	1.948647	1.953739	1.948647	1.934145	1.934145
NCI-H838	2.187972	2.101951	2.10351	2.101951	2.101951	2.101951	2.076156	2.007967	2.007967	1.98355	1.98355
NCI-H1155	2.010527	1.969077	1.909197	1.899458	1.802108	1.746776	1.746776	1.899458	1.746776	1.802108	1.899458
A2780cis	2.116952	1.972181	1.972181	1.955571	1.955571	1.955571	1.972181	1.972181	1.955571	1.955571	1.955571

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301043	SNP_A-2050943	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
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Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
G-401	1.986276	1.985247	1.985247	1.952639	1.952639	1.952639	1.952639	1.985247	1.952639	2.011131	2.023163
MS-1-L	2.1996	2.134107	2.134107	1.952346	1.952346	1.952346	2.055186	1.952346	2.055186	1.952346	2.055186
Hs 746T	2.198833	2.055426	2.198833	2.055426	1.949352	1.929315	1.929315	1.830353	1.830353	1.830353	1.830353
SW 13	1.979441	1.979441	1.743419	1.625769	1.625769	1.985073	1.985073	1.985073	1.964757	1.76455	1.791706
AGS	2.02322	2.02322	2.02322	2.02322	2.02322	2.031495	2.039987	2.039987	2.039987	2.031495	1.886192
NCI-H2171	2.37913	2.350219	2.350219	2.173174	2.002653	2.002653	2.002653	1.96429	1.949015	1.942958	1.942958
Hs 573.T	2.408031	2.408031	2.408031	2.392093	2.392093	2.126536	2.126536	1.836788	1.942973	1.930985	1.930985
HT1080	1.581778	1.67104	1.67104	1.67104	1.67104	1.67104	1.833534	1.859203	1.833534	1.859203	2.068981
COR-L 23/CPR	2.093681	1.924772	1.982795	1.924772	1.982795	1.924772	1.982795	1.924772	1.924772	1.924772	1.924772
HGC-27	1.962449	1.958792	1.958792	1.950571	1.895482	1.958792	1.958792	1.958792	1.958792	1.958792	2.035117
SBC-3	1.956875	1.956875	1.70526	1.70526	1.70526	1.70526	1.804857	1.804857	1.829007	1.804857	1.804857
COLO 792	2.106668	2.029739	2.029739	2.021302	2.021302	1.951143	1.915651	1.915651	1.915651	1.915651	1.915651
H-EMC-SS	2.48257	2.48257	2.48257	2.351611	2.351611	2.331001	2.232743	2.145117	2.232743	2.145117	2.145117
J82	2.160155	2.160155	2.160155	2.160155	2.160155	1.920742	2.144514	1.910432	1.910432	1.646031	1.910432
Hs 600.T	2.155798	2.052127	2.236533	2.236533	2.236533	1.80691	1.80691	1.80691	1.805151	1.805151	1.805151
Hs 781.T	2.413911	2.366922	2.366922	2.254342	2.254342	2.254342	2.169884	2.000707	2.000707	1.904757	1.904757
BPH-1	2.044366	1.977435	1.977435	1.924743	1.827384	1.827384	1.783523	1.724815	1.783523	1.724815	1.724815
COR-L23	2.293037	2.220459	2.220459	2.220459	1.914982	1.914982	2.08897	1.914982	1.914982	1.914982	1.914982
A673	1.733832	1.773896	1.773896	1.733832	1.733832	1.773896	1.831929	1.889162	1.773896	1.889162	2.036741
COLO 205	1.874433	1.874433	1.973113	1.874433	1.874433	1.874433	1.973113	2.109197	2.109197	2.109197	1.973113
RMG-I	2.175849	2.090481	2.090481	2.090481	1.91773	1.91773	2.090481	1.784191	2.090481	1.784191	2.090481
WI-26 VA4	2.008456	1.949499	1.949499	1.887483	1.887483	1.845735	1.845735	1.845735	1.845735	1.845735	1.845735
Eos-HL-60	1.890511	1.890511	1.890511	1.848245	1.853456	1.853456	1.853456	1.853456	1.853456	1.890511	1.890511
GP5D	2.040898	1.710672	2.040898	2.040898	1.710672	1.689044	1.689044	1.689044	1.689044	1.689044	1.689044
MES-SA/Dx-5	1.974962	1.773225	1.773225	1.773225	1.773225	1.773225	1.773225	1.773225	1.773225	1.846861	1.846861
SBC-2	2.130166	2.052269	2.052269	2.052269	1.823383	1.823383	1.823383	1.783321	1.783321	1.74312	1.583328
MCAS	1.878664	1.863139	1.878664	1.863139	1.863139	1.851908	1.863139	1.863139	1.87627	1.863139	1.87627
Ishikawa (Heraklio) 02 ER-	1.833641	1.833641	1.833641	1.749767	1.740138	1.678887	1.678887	1.678887	1.678887	1.749767	1.815412
MOLT-4F	1.969849	1.967282	1.969849	1.969849	1.969849	1.753038	1.753038	1.790279	1.831195	1.831195	1.831195
HARA	1.973157	1.924757	1.924757	1.924757	1.973157	1.924757	1.924757	1.973157	1.924757	1.770243	1.770243
NCI-H1944	1.993689	1.993689	2.001212	1.993689	1.993689	1.820237	1.993689	1.997416	1.997416	2.021961	1.997416
BICR 31	1.814335	1.795472	1.795472	1.806879	1.806879	1.795472	1.795472	1.795472	1.806879	1.819485	1.819485
D341 Med	1.757482	1.751183	1.780333	1.751183	1.648493	1.751183	1.780333	1.780333	1.780333	1.751183	1.751183
KHM-3S	3.248595	3.020628	3.020628	3.020628	2.735357	2.542792	2.476094	2.06953	2.038379	1.84173	1.79781
COR-L88	1.857779	1.697148	1.851133	1.851133	1.851133	1.697148	1.697148	1.697148	1.697148	1.622745	1.622745
NCI-H358	1.72821	1.72821	1.72821	1.664289	1.664289	1.664289	1.707353	1.707353	1.707353	1.72821	1.72821
1321N	2.072683	2.072683	1.863563	1.863563	1.83497	1.83497	2.011978	2.011978	2.011978	2.011978	1.941531
KYSE-510	1.811703	1.748902	1.748902	1.723006	1.573025	1.573025	1.723006	1.723006	1.723006	1.671897	1.671897
TCO-1	1.770946	1.702926	1.75178	1.75178	1.75178	1.75178	1.770946	1.85987	1.831378	1.85987	1.893866
NCI-H441	1.873604	1.872669	1.872669	1.661874	1.661874	1.661874	1.661874	1.519743	1.68422	1.648591	1.648591
SNU-398	1.767843	1.669519	1.669519	1.665391	1.661955	1.60187	1.661955	1.661955	1.661955	1.60187	1.669519
NCI-H1299	1.683222	1.683222	1.683222	1.641722	1.599517	1.641722	1.683222	1.683222	1.641722	1.589006	1.531765
ASH-3	1.4964415	1.4151733	1.4151733	1.4151733	1.331404	1.240009	1.240009	1.240009	1.240009	1.331404	1.456886
MCF7	1.283092	1.208143	1.208143	1.208143	1.150625	1.091937	1.091937	1.150625	1.091937	1.150625	1.150625
SKOV3	1.3220215	1.3220215	1.3597303	1.3597303	1.3597303	1.3597303	1.3597303	1.3650747	1.3597303	1.3597303	1.3597303
ChaGo-K-1	1.13213	0.9315729	0.9315729	0.3994263	0.3994263	0.3994263	0.3994263	0.3994263	0.3819881	0.3819881	0.7489773
A373 C6	0.7818635	0.7818635	0.7818635	0.7818635	0.8095883	0.7818635	0.7818635	0.8458438	0.7818635	0.8458438	0.8999572

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301045	SNP_A-2050945	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989997
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
MES-SA	0.9603825	0.8519282	0.8519282	0.8519282	0.7528253	0.9024908	0.9024908	0.7528253	0.7528253	0.7528253	0.9024908
OAW28	0.7155319	0.6828103	0.6828103	0.6516708	0.6156497	0.6156497	0.6516708	0.6516708	0.6516708	0.7155319	0.7432961
NCI-H460	0.7146986	0.4828994	0.4828994	0.4149362	0.3897879	0.3897879	0.4149362	0.4149362	0.3347929	0.3347929	0.4179333

SNP_A-4199624	SNP_A-2101390	SNP_A-2253557	ProbeSetID	SNP_A-2096988	SNP_A-2194428	SNP_A-1990988	SNP_A-4249755	SNP_A-1930452	SNP_A-2210006	SNP_A-2034119
11314429	6689767	9048083	Affy_SNP.ID	6625899	8104571	2625759	11152768	12275185	8339875	5678614
rs2896191	rs38852	rs38857	dbSNP	rs10215153	rs2237717	rs2283053	rs6978135	rs41735	rs41736	rs6566
7	7	7	Chromosome	7	7	7	7	7	7	7
115927995	115948476	115959364	Position	115993082	115999338	116020970	116024313	116029367	116029719	116032369
-	-	-	Strand	-	-	+	-	-	+	+
q31.2	q31.2	q31.2	Cytoband	q31.2	q31.2	q31.2	q31.2	q31.2	q31.2	q31.2
6.483741	6.483741	6.353536	Hs 746T	6.353536	6.285127	6.285127	5.589738	5.589738	5.589738	5.589738
6.572275	6.572275	6.043874	MKN45	6.14231	6.14231	5.849256	5.652385	5.652385	6.14231	6.14231
5.821465	5.821465	5.821465	EBC-1	5.821465	5.444932	5.179527	5.169925	5.179527	5.179527	5.179527
6.112681	6.112681	5.634547	SNU-5	5.634547	5.634547	5.582103	5.480071	5.582103	5.480474	5.480474
5.644351	5.644351	5.472339	NCI-H1573	5.737824	5.498854	5.498854	5.472339	5.498854	5.524749	5.498854
5.322405	5.322405	5.260284	OE33	5.183072	4.85022	4.85022	4.83089	4.85022	4.85022	4.85022
4.833896	4.833896	4.833896	OCUG-1	4.833896	4.487783	4.216219	4.216219	4.302471	4.302471	4.302471
4.665404	4.665404	4.665404	VMRC-RCW	4.641322	4.641322	4.362491	4.313745	4.347246	4.347246	4.347246
4.078277	4.078277	4.078277	SK-BR-3	4.078277	4.057138	3.782838	3.782838	4.057138	4.057138	4.154486
4.062967	4.033007	4.033007	C32	4.062967	4.062967	3.910254	3.910254	4.062967	3.95434	3.95434
3.781298	3.781298	3.736275	Detroit 562	3.781298	3.781298	3.781298	3.681808	3.781298	3.781298	3.590728
3.881318	3.881318	3.736046	NCI-H2342	3.651417	3.430943	3.430943	3.430943	3.51867	3.51867	3.430943
3.614116	3.614116	3.614116	FTC 133	3.534521	3.331986	3.331986	3.164621	3.164621	3.331986	3.331986
3.486409	3.486409	3.507132	PNT1A	3.545874	3.507132	3.507132	3.497738	3.507132	3.507132	3.567584
3.44082	3.427928	3.427928	KMRC-20	3.427928	3.427928	3.427928	3.363546	3.363546	3.363546	3.242777
3.570984	3.570984	3.570984	COLO 858	3.560813	3.560813	3.368309	3.102875	3.102875	3.046291	3.002379
3.436175	3.436175	3.400879	NUGC-4	3.514046	3.245861	3.245861	3.514046	3.514046	3.264536	3.264536
3.312846	3.312846	3.312846	YKG-1	3.309577	3.198226	2.984106	2.984106	3.12276	3.12276	2.984106
3.296618	3.215802	3.215802	H3255	3.215802	3.215802	3.047856	3.047856	3.215802	3.131906	3.131906
3.292095	3.259627	3.258172	SK-CO-1	3.087823	2.752303	2.752303	2.752303	3.087823	3.087823	3.087823
3.292325	3.160291	3.169925	LOU-NH91	3.169925	3.160291	3.072106	3.052138	3.052138	3.052138	3.052138
3.159654	3.159654	3.319119	SK-MEL-3	3.409408	3.319119	3.127722	3.127722	3.140722	3.140722	3.140722
3.153093	3.153093	3.153093	NCI-H2170	3.153093	3.153093	3.153093	2.957374	3.153093	2.946711	2.897823
3.352589	3.352589	3.25889	G-361	3.006311	2.928197	2.928197	2.928197	2.97487	2.97487	2.928197
3.147903	3.147903	3.147903	SK-NEP-1	3.116878	3.094811	2.939616	3.094811	3.094811	3.040945	3.040945
3.111966	3.111966	3.111966	HCC-44	3.198578	3.111966	3.087111	3.087111	3.181942	3.087111	2.895793
3.082038	3.082038	3.013387	WM 266-4	3.126037	2.940897	2.940897	2.940897	3.017466	3.375385	3.077652
3.199245	3.199245	3.199245	NCI-H441	3.077623	3.008041	3.008041	3.008041	3.059222	3.059222	2.906888
3.262888	3.262888	3.074602	P31/FUJ	2.9714	2.910215	2.9714	3.028255	3.074602	3.34875	3.441774
3.228383	3.228383	3.072613	MOG-G-CCM	3.072613	2.633473	2.633473	2.633473	2.721684	2.721684	2.633473
3.069209	3.069209	3.069209	SW 1417	3.069209	3.069209	2.932534	2.932534	3.072842	2.932534	2.921611
3.087635	3.087635	3.087635	COLO 853	3.044162	3.033089	3.044162	3.044162	3.044162	3.033089	3.01853
3.016188	2.907836	2.907836	NCI-H2085	3.016188	3.016188	3.016188	2.851037	3.039134	2.851037	2.829483

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301043	SNP_A-2050943	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
3.127195	2.909826	3.073827	NCI-H23	3.073827	2.993191	2.569111	2.569111	2.993191	2.993191	2.993191	2.993191
3.037372	3.037372	3.037372	A2058	2.905802	2.691156	2.691156	2.691156	2.746862	2.746862	2.691156	2.691156
2.979754	2.979754	2.979754	LUDLU-1	2.979754	2.974304	2.974304	2.974304	2.954671	2.974304	2.954671	2.927444
2.903029	2.7736	2.903029	COLO 741	2.963187	3.070583	3.070583	3.070583	3.070583	3.070583	3.070583	3.070583
3.117642	3.117642	3.081493	KP-4	3.081493	2.774791	2.54315	2.774791	3.081493	2.815554	2.815554	2.815554
2.889578	2.829007	2.889578	SR	2.946805	2.946805	2.946805	2.829007	2.946805	2.946805	2.946805	2.946805
3.129917	3.01169	2.814586	WM-115	2.814586	2.774074	2.814586	2.814586	3.01169	3.01169	3.01169	3.01169
2.933108	2.933108	2.908847	NCI-H810	2.908847	2.789785	2.706556	2.706556	2.706556	2.651222	2.651222	2.651222
3.115644	3.115644	3.115644	SBC-5	2.764028	2.708363	2.708363	2.764028	2.816793	2.816793	2.816793	2.816793
2.91091	2.896216	2.896216	MKN1	2.896216	2.871353	2.78681	2.78681	2.78681	2.78681	2.78681	2.78681
2.872665	2.872665	2.891581	Sci-1	2.891581	2.882043	2.855191	2.758209	2.855191	2.855191	2.855191	2.855191
2.961833	2.868482	2.805641	NCI-H1755	2.822516	2.805641	2.822516	2.822516	2.868482	2.995317	2.822516	2.822516
2.999513	2.793963	2.793963	HCC202	2.867372	2.793963	2.665782	2.665782	2.867372	2.885484	3.031586	3.031586
3.018571	3.018571	3.018571	LCLC-103H	3.018571	2.86148	2.86148	2.804244	2.853485	2.86148	2.86148	2.853485
2.942573	2.942573	2.942573	KMRC-1	2.926669	2.747141	2.590564	2.558051	2.558051	2.550586	2.516773	2.516773
2.855502	2.736252	2.855502	DMS 273	2.855502	2.736252	2.736252	2.736252	2.855502	2.905391	2.905391	2.905391
2.786547	2.639629	2.786547	SNU-16	2.908583	2.908583	2.918055	2.918055	2.918055	2.918055	2.908583	2.908583
2.846898	2.83463	2.829202	KATO III	2.83463	2.846898	2.846898	2.846898	2.846898	2.829202	2.829202	2.829202
3.002807	3.002807	3.002807	COLO 857	2.887542	2.887542	2.887542	2.771425	2.771425	2.738464	2.709311	2.709311
2.743274	2.717423	2.892661	22RV1	2.892661	2.717423	2.892661	2.944539	2.94536	2.94536	2.944539	2.944539
3.028245	2.861513	2.808346	JSC-1	2.808346	2.79267	2.79267	2.79267	2.800977	2.800977	2.79267	2.79267
2.937938	2.937938	2.889885	U373 MG	2.737492	2.65281	2.700576	2.700576	2.700576	2.700576	2.346472	2.346472
2.789072	2.789072	2.794051	U266B1	2.893698	2.893698	2.955883	2.955883	2.955883	2.932356	2.794051	2.794051
2.959889	2.959889	2.959889	JHH-4	2.862679	2.793842	2.379383	2.379383	2.793842	2.379383	2.379383	2.379383
2.99227	2.839736	2.839736	COLO 792	2.839736	2.784492	2.783019	2.783019	2.784492	2.783019	2.783019	2.783019
2.932001	2.932001	2.932001	MRK-nu-1	2.826527	2.7575	2.448043	2.448043	2.786558	2.786558	2.780301	2.780301
2.864123	2.864123	2.739413	CAL-148	2.739413	2.647051	2.483546	2.647051	2.799753	2.685838	2.799753	2.799753
2.766928	2.766928	2.767226	COLO 684	2.813661	2.767226	2.766928	2.641824	2.689634	2.641824	2.617828	2.617828
2.621913	2.621913	2.856735	PC-14	3.075477	2.856735	2.856735	2.848898	2.856735	2.848898	2.681785	2.681785
3.22942	3.111711	2.945755	NCI-H841	2.571893	2.272146	2.272146	2.272146	2.571893	2.571893	2.272146	2.272146
2.766706	2.766706	2.750902	BICR 78	2.750902	2.748488	2.688919	2.688919	2.717951	2.688919	2.717951	2.717951
2.849264	2.849264	2.750535	RERF-LC-KJ	2.750535	2.750535	2.739737	2.612653	2.739737	2.739737	2.739737	2.739737
2.85965	2.85965	2.846317	JHH-6	2.846317	2.651659	2.139404	2.139404	2.651659	2.506611	2.506611	2.506611
2.736453	2.736453	2.840145	SW 48	3.015224	2.840145	2.840145	2.840145	2.840145	2.672475	2.672475	2.672475
2.741511	2.741511	2.741511	NCI-H196	2.741511	2.591528	2.722144	2.722144	2.861063	2.722144	2.591528	2.591528
2.825368	2.825368	2.825368	BALL-1	2.811048	2.514371	2.514371	2.514371	2.631642	2.631642	2.631642	2.631642
2.75277	2.75277	2.75277	NCI-H889	2.718772	2.703924	2.703924	2.703924	2.718772	2.718772	2.718772	2.718772
2.718694	2.718694	2.718694	NCI-H2009	2.718694	2.668633	2.581391	2.581391	2.668633	2.626527	2.63712	2.63712
2.687566	2.687566	2.687566	SK-LU-1	2.698674	2.488958	2.478234	2.188013	2.478234	2.488958	2.488958	2.488958
2.755014	2.755014	2.751825	HEC-1	2.524878	1.942708	1.942708	1.942708	2.524878	2.637131	2.448081	2.448081
2.679096	2.679096	2.859971	H69AR	2.859971	2.679096	2.617945	2.431208	2.720785	2.559026	2.559026	2.559026
2.678275	2.678275	2.678275	OCUM-1	2.666847	2.573762	2.326997	2.326997	2.700994	2.700994	2.609022	2.609022
2.187648	2.354955	2.645598	KHOS-312H	2.645598	2.746383	2.746383	2.753492	2.776159	2.776159	2.881192	2.881192
2.644987	2.644987	2.644987	IM-95	2.644987	2.571964	2.571964	2.571964	2.571964	2.707155	2.756968	2.756968
2.65734	2.642609	2.589694	OAW42	2.642609	2.642609	2.508281	2.501596	2.508281	2.501596	2.501596	2.501596
2.733321	2.619959	2.624947	MDST8	2.733321	2.709626	2.624947	2.545539	2.545539	2.386988	2.516869	2.516869
2.629857	2.515264	2.629857	HT 115	2.629857	2.617211	2.452437	2.617211	2.670383	2.670383	2.617211	2.617211

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301043	SNP_A-2050943	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
3.137835	2.94173	2.778649	HMCB	2.453646	2.422423	2.422423	2.422423	2.453646	2.453646	2.453646	
2.609022	2.597317	2.609022	RERF-LC-MS	2.62068	2.678275	2.678275	2.712288	2.712288	2.678275	2.678275	
2.638114	2.624575	2.593608	FTC 238	2.570319	2.422112	2.422112	2.422112	2.593608	2.624575	2.475972	
2.410835	2.435119	2.593009	KHOS-240S	2.593009	2.957861	3.179527	3.179527	3.184364	3.339124	3.339124	
2.591834	2.591834	2.865267	AN3CA	2.932983	2.865267	2.865267	2.591834	2.865267	2.591834	2.482899	
2.71866	2.71866	2.640842	NCI-H1651	2.510119	2.400318	2.400318	2.400318	2.510119	2.510119	2.400318	
2.695751	2.578394	2.572272	1321N	2.572272	2.572272	2.322782	2.322782	2.343105	2.343105	2.343105	
2.876831	2.876831	2.876831	LU65	2.267022	2.266746	2.266746	2.266746	2.267022	2.267022	2.267022	
2.632778	2.571444	2.571444	HARA	2.571444	2.533441	2.249172	2.249172	2.533441	2.545205	2.571444	
2.708126	2.708126	2.708126	RCM-1	2.708126	2.642044	2.265877	2.642044	2.642044	2.49764	2.49764	
2.566611	2.602918	2.602918	HuH-7	2.439012	2.439012	2.439012	2.590917	2.602918	2.602918	2.590917	
2.575063	2.566433	2.566433	WIDR	2.566433	2.466782	2.466782	2.466782	2.498005	2.566433	2.498005	
2.664408	2.587377	2.587377	UMC-11	2.587377	2.564286	2.564286	2.564286	2.45749	2.469027	2.36983	
2.662	2.662	2.662	huH-1	2.58587	2.562447	2.374725	2.562447	2.58587	2.562447	2.562447	
2.784206	2.784206	2.784206	HCC-827	2.750924	2.561117	1.60523	1.60523	2.539515	1.963194	1.963194	
2.670143	2.658167	2.512729	NCI-H2081	2.512729	2.326315	2.326315	2.512729	2.765954	2.765954	2.765954	
2.473646	2.469407	2.473646	SW 13	2.614134	2.614134	2.621541	2.621541	2.739424	2.621541	2.614134	
2.548738	2.548738	2.54327	KYSE-140	2.54327	2.532626	2.54327	2.54327	2.617176	2.617176	2.54327	
3.036526	3.036526	3.036526	LC-1 sq	3.036526	2.514045	2.450858	2.514045	2.631735	2.514045	2.631735	
2.457785	2.457785	2.457785	KYSE-410	2.615545	2.615545	2.87114	2.87114	2.716669	2.615545	2.615545	
2.650278	2.650278	2.650278	CAL-85-1	2.245266	2.245266	2.245266	2.360449	2.412611	2.412611	2.412611	
2.733478	2.733478	2.733478	HCC-366	2.463418	1.915592	1.855399	1.855399	1.915592	1.915592	1.915592	
2.527391	2.527391	2.527391	SW 1463	2.532146	2.527391	2.45712	2.45712	2.527391	2.532146	2.532146	
2.64501	2.64501	2.527259	MOLT-4F	2.527259	2.178454	2.178454	2.221635	2.527259	2.412086	2.412086	
2.52394	2.634851	2.634851	SW-1710	2.52394	2.52394	2.25726	2.25726	2.451759	2.451759	2.451759	
2.581108	2.581108	2.581108	HCC-15	2.581108	2.54875	2.495503	2.478539	2.478539	2.430264	2.277563	
2.505933	2.505933	2.504589	PLC/PRF/5	2.504589	2.504589	2.49792	2.49792	2.49792	2.464769	2.373293	
2.639201	2.495648	2.495648	A375.S2	2.650462	2.650462	2.46693	2.446932	2.46693	2.446932	2.446932	
2.490127	2.490127	2.488386	COR-L279	2.488386	2.320989	1.827506	1.827506	2.320989	2.320989	2.320989	
2.869369	2.486619	2.37347	COLO 685	2.486619	2.486619	2.486619	2.486619	2.486619	2.486619	2.146914	
2.526838	2.518738	2.482021	FLYA13	2.296163	2.252082	2.252082	2.348966	2.482021	2.482021	2.348966	
2.622157	2.476583	2.451031	JHH-7	2.476583	2.409296	2.409296	2.476583	2.476583	2.409296	2.409296	
2.348774	2.348774	2.503996	NCI-H1623	2.504189	2.504189	2.504189	2.469236	2.504189	2.469236	2.469236	
2.512451	2.468389	2.468389	HGC-27	2.512451	2.468389	2.406866	2.406866	2.464351	2.406866	2.208042	
2.541058	2.541058	2.541058	NCI-H2347	2.56923	2.541058	2.541058	2.393743	2.393743	2.34912	2.222837	
2.675114	2.675114	2.477048	SKN	2.412036	2.377207	2.000058	2.000058	2.377207	2.027425	2.027425	
2.658569	2.658569	2.658569	VMRC-LCP	2.563919	2.339581	2.14218	2.14218	2.14218	1.760995	1.760995	
2.474014	2.580896	2.580896	SAT	2.474014	2.446993	2.439669	2.439669	2.439669	2.217196	2.217196	
2.86641	2.86641	2.484937	CaR-1	2.393478	2.221515	2.218641	2.218641	2.393478	2.484937	2.659384	
2.438046	2.437376	2.437376	SW 900	2.438046	2.438046	2.437376	2.365804	2.437376	2.365804	2.365804	
2.529889	2.43466	2.43466	Nthy-ori 3-1	2.43466	2.227121	2.227121	2.227121	2.43466	2.43466	2.227121	
2.489129	2.434635	2.620669	NCI-H526	2.620669	2.323711	2.323711	2.323711	2.424093	2.406954	2.424093	
2.586082	2.586082	2.586082	HUP-T3	2.586082	2.434586	2.434586	2.434586	2.434586	2.313453	2.313453	
2.430152	2.430152	2.430152	BT-483	2.475776	2.475776	2.475776	2.475776	2.578736	2.475776	2.430152	
2.454977	2.454977	2.429767	NCI-H2330	2.429767	2.358067	2.155375	2.155375	2.319052	2.112862	2.112862	
2.439372	2.300963	2.300963	NCI-H711	2.300963	2.300963	2.576658	2.683493	2.781566	2.781566	2.781566	
2.423557	2.423557	2.518015	NCI-H661	2.576386	2.518015	2.518015	2.456147	2.456147	2.281937	2.281937	

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896871	SNP_A-4301043	SNP_A-2050943	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
2.456726	2.435504	2.422922	EJ138	2.422922	2.390053	2.368091	2.368091	2.390053	2.368091	2.368091	
2.188472	2.188472	2.419543	HOS	2.419543	2.695388	2.855395	2.970915	2.970915	2.970915	2.970915	
2.470192	2.416223	2.30399	KHM-3S	2.470192	2.470192	2.470192	2.30399	2.30399	2.30399	2.30399	
2.459705	2.437996	2.382607	ChaGo-K-1	2.437996	2.382607	2.209626	2.209626	2.209626	2.209626	2.209626	
2.479114	2.479114	2.446536	H69V	2.446536	2.404434	2.404434	1.781485	2.101909	1.8302	1.8302	
2.401059	2.297361	2.297361	Hs 936.T	2.401059	2.297361	2.401059	2.401059	2.694175	2.732404	2.619671	
2.338016	2.338016	2.386913	201T	2.477415	2.477415	1.996738	2.477415	2.477415	2.477415	2.477415	
2.386862	2.386862	2.386862	SUIT-2	2.386862	2.386862	2.355695	2.386862	2.386862	2.355695	2.355695	
2.647674	2.545754	2.461193	NCI-H1792	2.30621	2.161575	2.161575	2.161575	2.161575	2.119541	2.109197	
2.438851	2.422959	2.366694	NCI-H460	2.355529	2.355529	2.355529	2.355529	2.366694	2.366694	2.295017	
2.403506	2.403506	2.403506	SNU-398	2.342234	2.17388	2.17388	2.17388	2.342234	2.17388	2.131806	
2.39475	2.336154	2.39475	COR-L95	2.461255	2.39475	2.29572	2.202749	2.29572	2.202749	2.202749	
2.400745	2.373267	2.373267	SK-MES	2.373267	2.036371	2.036371	2.036371	2.087081	2.087081	2.087081	
2.742192	2.742192	2.742192	BFTC-905	2.363592	2.363592	2.363592	2.206297	2.206297	2.194952	2.194952	
2.392408	2.392408	2.392408	EB1	2.392408	2.392408	2.378966	2.302315	2.378966	2.302315	2.191039	
2.403029	2.403029	2.389221	CASKI	2.320756	2.320756	2.153843	2.153843	2.153843	2.286277	2.286277	
2.57071	2.57071	2.539694	HO-1-N-1	2.309529	2.1506	1.673448	1.673448	2.309529	2.063595	2.063595	
2.352646	2.352646	2.352646	PC-3 [JPC-3]	2.352646	2.390695	2.261862	2.261862	2.573762	2.585563	2.585563	
2.419905	2.419905	2.416011	TE7	2.286395	2.239031	2.193293	2.193293	2.239031	2.286395	2.286395	
2.799228	2.799228	2.567073	TASK1	2.241673	1.951701	1.826909	1.826909	1.951701	1.951701	1.951701	
2.479578	2.479578	2.479578	NCI-H1734	2.393428	2.143841	2.300417	2.300417	2.393428	2.300417	2.265733	
2.473757	2.37366	2.37366	HRT-18	2.309387	2.23229	2.118309	2.118309	2.23229	2.225467	2.118309	
2.541225	2.541225	2.541225	RT112/84	2.335088	2.098983	2.098983	2.098983	2.230693	2.230693	2.098983	
2.480592	2.480592	2.390431	CAL-29	2.341401	2.335422	2.335422	2.335422	2.335422	2.269625	2.269625	
2.373584	2.373584	2.399477	IMR-90	2.302523	2.260069	2.157711	1.856976	1.856976	1.760415	1.856976	
2.497155	2.484534	2.484534	RO82-W-1	2.191309	2.146269	2.146269	2.146269	2.191309	2.191309	2.191309	
2.335088	2.283833	2.279725	NCI-H1581	2.279725	2.279725	2.066078	2.066078	2.477684	2.507991	2.477684	
2.399816	2.381394	2.364635	RT4	2.304873	1.956054	1.956054	1.956054	2.15532	2.15532	1.965911	
2.7243	2.426732	2.339324	HT 29	2.295825	2.152886	2.152886	2.152886	2.152886	2.23842	2.295825	
2.611287	2.611287	2.611287	KON	2.302289	2.09763	1.986494	1.986494	2.09763	1.986494	1.986494	
2.415786	2.415786	2.415786	BT-20	2.248656	2.195222	2.113167	2.113167	2.113167	2.113167	2.113167	
2.355759	2.355759	2.355759	BU25 TK	2.355759	2.313143	2.313143	2.277943	2.277943	2.263337	2.128745	
2.637258	2.542278	2.295004	RPMI-7951	2.295004	2.09083	2.09083	2.09083	2.09083	2.09083	2.09083	
2.276764	2.276764	2.327603	IHH-4	2.327603	2.327603	2.276764	2.276764	2.276764	2.24158	2.24158	
2.440648	2.361976	2.267548	fr2	2.239668	1.576837	1.576837	1.576837	2.267548	2.267548	1.576837	
2.314152	2.309387	2.309387	CAL-51	2.250694	1.973332	1.973332	2.063863	2.167407	2.167407	2.167407	
2.26032	2.26032	2.26032	COR-L 105	2.410134	2.26032	2.26032	2.26032	2.26032	2.1852	2.1852	
2.461587	2.461587	2.461587	LK-2	2.278048	2.230466	2.230466	2.241075	2.27979	2.27979	2.241075	
2.55616	2.51822	2.51822	FU97	2.372266	2.336617	2.089362	2.089362	2.089362	2.088718	2.088718	
2.430984	2.354164	2.253695	MOG-G-UVW	2.253695	1.719542	1.719542	1.719542	2.237424	2.237424	2.236759	
2.243583	2.243583	2.391124	NCI-H2452	2.445053	2.500262	2.500262	2.445053	2.445053	2.049278	1.92649	
2.591564	2.591564	2.343144	HuCC1	2.226467	2.209264	2.169938	2.169938	2.169938	2.03966	2.03966	
2.234991	2.068643	2.068643	NCI-H1417	2.234991	2.234991	2.234991	2.234991	2.234991	2.56006	2.56006	
2.239337	2.231598	2.231598	Hs 633T	2.231598	2.158667	2.017992	2.017992	2.158667	2.017992	1.948383	
2.191309	2.340812	2.340812	A-375	2.326753	2.326753	2.260438	2.258843	2.258843	2.081799	2.081799	
2.513949	2.220072	2.217624	NCI-H1650	2.217624	2.128524	1.990926	1.990926	2.128524	1.990926	1.990926	
2.502893	2.502893	2.319491	KP-1N	2.130483	2.055978	1.82323	2.055978	2.130483	2.055978	2.130483	

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896871	SNP_A-4301043	SNP_A-2050943	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
2.237716	2.237716	2.237716	NCI-H2135	2.237716	2.237716	2.236068	2.187364	2.236068	2.187364	1.998441	
2.174545	2.174545	2.486193	HT55	2.502493	2.155019	2.079245	2.079245	2.079245	2.047362	2.246087	
2.553077	2.553077	2.411936	NCI-H1781	2.411936	2.086045	2.086045	2.086045	2.086045	1.905025	1.905025	
2.419768	2.311719	2.311719	RERF-GC-1B	2.093401	1.883415	1.883415	1.883415	2.049845	2.049845	2.049845	
2.198523	2.192591	2.198523	NCI-H727	2.198712	2.198712	2.198712	2.198712	2.198712	2.198523	2.157452	
2.306923	2.306923	2.306923	647-V	2.177857	1.81158	1.81158	1.81158	2.085373	2.085373	1.923765	
2.248418	2.230426	2.230426	NCI-H2023	2.160019	1.9641	1.9641	1.9641	2.160019	2.160019	1.9641	
2.207438	2.123884	2.207438	Calu-3	2.230186	2.230186	2.230186	2.230186	2.230186	2.182777	2.107085	
2.512403	2.512403	2.512403	H69	2.262125	2.170523	2.170523	2.076324	2.093164	2.093164	2.156386	
2.514432	2.514432	2.514432	VMRC-LCD	2.21204	2.123138	2.062565	2.123138	2.123138	2.123138	2.064724	
2.248656	2.113167	2.113167	VMRC-LCD	2.390695	2.113167	2	2.113167	2.248656	2.248656	2.248656	
2.311291	2.311136	2.285088	T84	1.759537	1.717028	1.717028	1.759537	2.170224	2.170224	2.170224	
2.302925	2.302925	2.18781	SW 1116	2.175211	2.12246	2.12246	2.116079	2.116079	2.116079	1.78108	
2.182411	2.182411	2.210069	A373 C6	2.214331	2.214331	2.210069	2.179281	2.179281	2.179281	2.210069	
2.177844	2.177844	2.177844	COLO 201	2.177844	2.281492	2.281492	2.177844	2.281492	2.177844	2.104136	
1.99123	2.171203	2.177559	KHOS/NP	2.177559	2.211892	2.211892	2.307416	2.479505	2.479505	2.499012	
2.171189	2.163893	2.112903	Hs 891.T	2.138813	2.138813	2.171189	2.15808	2.218734	2.210605	2.15808	
2.194493	2.194493	2.194493	PA-TU-8988T	2.112945	1.987015	1.638094	1.638094	1.987015	1.638094	1.638094	
2.311874	2.311874	2.261598	BICR22	2.23406	2.159459	2.140339	2.042277	2.042277	2.042277	2.140339	
2.15123	2.149901	2.15123	MES-SA/Dx-5	2.15123	2.15123	1.759835	1.759835	1.794775	2.092982	2.092982	
2.076015	2.024394	2.024394	SK-MEL-1	2.222143	2.222143	1.81615	1.81615	2.222143	2.238101	2.238101	
2.343336	2.343336	2.14347	KYSE-70	2.112057	2.112057	2.112057	2.112057	2.139459	2.139459	2.112057	
2.342439	2.189702	2.141122	A673	2.035502	1.851133	1.851133	1.851133	2.035502	2.141122	2.05207	
2.277328	2.277328	2.338067	D1.1	2.382355	2.382355	2.338067	1.831608	1.831608	1.831608	1.927425	
2.230693	2.230693	2.230693	NCI-H1793	2.321453	2.230693	2.230693	2.099568	2.099568	2.062762	1.825376	
2.17176	2.17176	2.17176	OE19	2.130235	1.972108	1.972108	1.972108	1.963632	1.960241	1.960241	
2.154349	2.129159	2.125128	WILCL	2.125128	1.728194	1.728194	1.728194	2.129159	2.129159	1.805243	
2.043357	2.043357	2.127448	HCC-78	2.127448	2.043357	2.127448	2.145735	2.145735	2.145735	2.127448	
2.326997	2.23539	2.181707	BEN	2.071248	2.057138	1.743687	1.743687	2.071248	2.071248	2.071248	
2.229107	2.1217	2.229107	Calu-1	2.278913	1.993833	1.967268	1.967268	1.993833	1.967268	1.967268	
2.140147	2.092381	2.092381	SK-N-BE(2)	2.121547	2.121547	2.121547	2.121547	2.140147	2.121547	2.121547	
2.578028	2.578028	2.577367	HUP-T4	2.013415	1.861323	1.801104	1.801104	1.861323	1.861323	1.861323	
2.390695	2.113167	2.014391	MCF7	1.806077	1.647499	1.647499	1.647499	1.806077	2.127006	2.573762	
2.107752	2.072641	2.180284	Ishikawa	2.228787	2.180284	2.083509	2.083509	2.136805	2.083509	2.070417	
2.27279	2.27279	2.27279	Hs 195.T	2.21409	2.102368	2.036898	1.676532	1.676532	1.625029	1.676532	
2.369068	2.09322	2.161411	HT 1376	2.263851	2.161411	2.09322	2.050596	2.050596	2.042092	1.959509	
2.106695	2.091641	2.106695	COLO 205	2.163156	2.163156	2.106695	2.091641	2.091641	1.982447	1.816427	
2.354037	2.16953	2.140655	NCI-H1437	2.140655	1.811256	1.811256	1.811256	1.965049	1.876075	1.876075	
2.086899	2.086899	2.086899	A2780	2.182587	2.192213	2.192213	2.032536	2.192213	2.032536	2.032536	
2.296202	2.086801	2.086801	A549	2.086801	2.029196	2.008413	2.008413	2.008413	2.003129	1.857233	
1.970986	1.954516	1.970986	NCI-H1522	2.233994	2.233994	2.233994	2.233994	2.233994	2.191673	2.191673	
2.152311	2.1186	2.1186	G-401	2.043798	2.043798	1.976228	2.043798	2.043798	1.976228	1.976228	
2.081757	2.081757	2.081757	PC-9	2.081757	2.081056	2.038151	2.038151	2.038151	1.97694	1.856839	
2.225467	2.225467	2.225467	GRANTA-519	2.203542	2.080312	1.923898	1.923898	1.923898	1.923898	1.923898	
2.302016	2.115483	2.115483	639-V	2.03956	1.794186	1.794186	1.794186	2.03956	2.03956	1.794186	
2.056813	2.041125	2.056813	A2780 ADR	2.158422	2.244418	2.056813	2.056813	2.087389	2.056813	2.016386	
2.164329	2.142358	2.122972	A-427	2.020743	1.917789	1.609611	1.609611	1.917789	1.892369	1.892369	

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301043	SNP_A-2050943	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
2.068502	2.068502	2.381306	NB69	2.381306	2.437872	2.068502	2.02282	2.068502	2.068502	2.146914	
2.234366	2.234366	2.175917	NCI-H2286	1.95888	1.905502	1.905502	1.905502	1.95888	1.95888	1.95888	
2.06674	2.064498	2.064498	SBC-2	2.064498	2.064498	1.9822	1.9822	1.9822	1.9822	1.9822	
2.172807	2.063637	2.063637	HCT-8	2.063637	1.885292	1.885292	2.172807	2.207868	2.172807	2.058043	
2.063143	2.063143	2.083243	Hs 895.T	2.108752	2.083243	2.083243	2.07874	2.07874	2.063143	2.041367	
2.272777	2.194332	2.140998	K562/ADM	1.979005	1.537089	1.537089	1.919347	1.979005	1.979005	1.979005	
2.145968	2.145968	2.107627	Hs 228.T	2.011059	1.828333	1.70395	1.70395	1.828333	1.828333	1.876511	
2.145542	2.142221	2.142221	STC 1	2.142221	2.057322	1.97614	1.97614	2.057322	2.057322	2.011073	
2.043286	2.043286	2.052311	LNcaP clone Fc	2.05309	2.052311	2.052311	2.052311	2.052311	2.077799	2.102828	
2.034276	1.905055	2.034276	C8166	2.078207	2.034276	2.034276	2.034276	2.078207	2.047845	2.047845	
2.406001	2.383176	2.383176	DMS 92	2.383176	2.046383	2.040258	2.040258	2.040258	1.519079	2.046383	
2.264772	2.115746	2.115746	NCI-H2196	1.95349	1.95349	1.799125	1.95349	1.974089	1.974089	1.974089	
2.044125	2.00124	1.94612	HMVII	2.044125	2.044125	2.044125	2.044125	2.100962	2.268862	2.100962	
1.942649	1.868942	2.038564	Hs 417.Lu	2.242813	2.076577	2.076577	2.038564	2.076577	2.038564	1.722212	
2.231465	2.231465	2.155183	H-EMC-SS	1.916319	1.676112	1.676112	1.676112	1.685475	1.676112	1.685475	
2.227908	2.018809	2.254184	H33HJ-JA1	2.285219	2.254184	2.254184	1.934145	1.94615	1.934145	1.94615	
2.032336	2.023578	2.023578	HCT 116	2.032336	2.023578	2.032336	2.032336	2.023578	1.877294	1.877294	
2.140573	2.135732	2.135732	GA-10	2.135732	2.032023	2.007233	2.007233	2.032023	2.007233	1.812904	
2.181707	2.140779	2.140779	NCI-H1666	2.028711	1.598175	1.427392	1.427392	1.427392	1.427392	1.301529	
2.065105	2.003042	2.065105	LS174T	2.013013	2.003042	2.003042	2.003042	2.013013	2.020013	2.069516	
2.023521	2.017562	2.017562	Hs 908.T	2.017562	2.017562	2.072247	2.072247	2.072247	2.103858	2.017562	
2.117326	2.070234	1.971351	C170	1.906247	1.862806	1.862806	1.906247	1.907112	1.907112	2.117326	
2.10472	2.10472	2.082977	J82	1.952889	1.808868	1.500407	1.500407	1.808868	1.500407	1.500407	
2.013759	1.876647	2.013759	A2780cis	2.125305	2.013759	1.876647	1.876647	2.013759	2.013759	2.013759	
2.09915	2.027082	2.013329	OUS-11	2.013329	1.583595	1.583595	1.915814	1.915814	1.915814	1.915814	
2.013228	2.013228	2.013228	KYSE-150	2.110086	2.110086	2.110086	2.110086	2.110086	1.786119	1.652157	
1.883024	2.049689	2.049689	COR-L23	2.049689	1.883024	1.883024	1.883024	2.022061	2.021216	1.883024	
2.138772	2.138772	2.091878	MS-1-L	1.94925	1.856506	1.782729	1.782729	1.94925	1.94925	1.94925	
2.46888	2.46888	2.46888	SKG-IIIb	2.020314	1.712362	1.675983	1.712362	1.924298	1.924298	1.924298	
2.199681	2.096347	2.096347	P32/ISH	1.990462	1.990462	1.948001	1.990462	1.990462	1.941339	1.990462	
2.146187	1.98732	1.98732	NCI-H1693 (shij)	1.98732	1.983434	1.98732	1.98761	1.98761	1.98761	1.98761	
2.010671	2.010671	1.987479	Hs 781.T	2.010671	1.987479	1.936521	1.936521	1.987479	1.987479	1.906113	
2.36405	2.36405	2.107001	Hs 600.T	1.866481	1.736046	1.315481	1.315481	1.736046	1.736046	1.736046	
2.077617	1.978554	1.978554	Eos-HL-60	1.978554	1.947134	1.898427	1.898427	1.947134	1.898427	1.873438	
1.918294	1.918294	1.918294	P116 T	2.03352	2.03352	2.03352	2.03352	2.03352	1.763642	1.763642	
2.260333	1.879055	1.797547	273T	1.773412	1.773412	1.773412	1.773412	1.975733	2.05612	1.975733	
1.978089	1.974554	1.730286	NCI-H2172	1.730286	1.730286	1.730286	1.974554	1.974554	1.730286	1.718142	
1.96575	1.96575	1.96575	KYSE-220	2.108322	2.108322	2.108322	2.108322	2.221167	2.108322	1.981373	
1.89907	1.843924	1.933009	Hs 852.T	2.005103	1.998066	1.998066	2.005103	2.049207	2.163211	2.049207	
1.923261	1.923261	2.004267	AZ-521	2.027682	2.004267	2.004267	2.004267	2.004267	1.798119	1.798119	
2.375345	1.959056	1.959056	NCI-H650	2.066402	2.066402	1.710911	1.710911	1.848352	1.848352	1.848352	
2.071248	2.042959	1.851999	23132/87	1.479696	1.011782	1.011782	1.011782	1.663754	2.140779	1.727877	
1.955761	2.098969	2.098969	RKN	1.955761	1.955761	1.937612	1.937612	1.937612	1.834069	1.834069	
2.140435	1.960665	1.960665	HBE4-E6/E7	1.925646	1.586077	1.586077	1.586077	1.747233	1.747233	1.747233	
2.147407	1.940986	1.940986	NCI-H1299	2.147407	1.940986	1.852819	1.852819	2.147407	1.852819	1.838894	
2.028196	1.939159	2.037439	Hs 573.T	2.165337	2.229787	2.229787	1.846435	1.846435	1.846435	1.846435	
2.106403	2.027839	1.962975	Hs 737.T	1.903519	1.903205	1.561981	1.561981	1.561981	1.561981	1.770649	

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896877	SNP_A-4301045	SNP_A-2050945	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
1.998615	1.998615	1.785264	NCI-H2171	1.785264	1.752425	1.752425	1.752425	2.35516	2.35516	2.35516	
2.09111	2.09111	2.09111	G-402	1.928679	1.928679	1.892803	1.928679	1.928679	1.907171	1.907171	
1.923217	1.916022	1.923217	MEWO	1.923217	1.870029	1.923217	1.923217	1.984174	2.148189	1.984174	1.923217
1.93568	1.93568	1.935208	CAL-120	1.935208	1.921765	1.700735	1.921765	1.921765	1.921765	1.921765	1.921765
1.922002	1.916557	1.916557	AGS	1.922002	1.916557	1.916557	1.916557	1.945444	1.945444	2.008686	2.01419
1.913585	1.786585	1.786585	WIL2 NS	1.786585	1.786585	1.786585	1.880514	1.938246	2.114679	2.114679	2.114679
1.906858	1.906858	1.906858	COR-L 23/CPR	1.906858	1.878273	1.878273	1.878273	1.906858	2.06423	1.906858	1.906858
1.97726	1.924491	1.904712	GP5D	1.904712	1.872609	1.834008	1.872609	1.872609	1.705069	1.837581	
1.927969	1.927969	1.927969	MCAS	1.913407	1.673513	1.399441	1.399441	1.913407	1.927969	1.650627	
1.901206	1.901206	1.901206	COLO-680N	1.901206	1.901206	1.867946	1.857688	1.857688	1.81758	1.81758	
1.908586	1.90016	1.908586	LS180	1.908586	1.90016	1.776532	1.776532	1.747248	1.747248	1.747248	
1.896005	1.896005	1.849234	DMS 153	1.837261	1.800996	1.800996	1.800996	1.837261	2.024909	2.024909	
1.966567	1.894659	1.894659	SBC-3	1.894659	1.430996	1.430996	1.488029	1.488029	1.488029	1.488029	
2.034718	1.991996	1.77999	HT 728.T	1.991996	1.77999	1.77999	1.77999	1.77999	1.67018	1.77999	
2.20713	2.20713	2.20713	SW 1573	1.926504	1.926504	1.884211	1.777124	1.777124	1.679274	1.777124	
1.869244	1.802788	1.869244	MKN74	1.895676	1.910105	1.910105	1.910105	1.910105	1.910105	1.749263	
2.189161	2.189161	2.189161	SAS	1.877083	1.736978	1.736978	1.859355	1.877083	1.877083	1.877083	
2.014391	1.882213	1.882213	MKN7	1.867145	1.496922	1.496922	1.496922	1.759412	1.743687	1.743687	
1.979441	1.979441	1.979441	RMG-I	1.674514	1.623534	1.540745	1.540745	1.674514	1.674514	1.906516	
1.912113	1.912113	1.867145	SVts-8	1.679919	1.679919	1.679919	1.679919	1.867145	2.014391	1.790605	
1.883174	1.827261	1.861021	Hs 894(E).Lu	1.861021	1.861021	1.883174	1.883174	1.883174	1.861021	1.861021	
1.859582	1.859582	1.859582	Hs 751.T	1.922343	1.878528	1.878528	1.789038	1.789038	1.735413	1.735413	
1.889252	1.889252	1.850343	SK-N-FI	1.850343	1.741779	1.741779	1.741779	1.742599	1.742978	1.742978	
2.079681	1.864016	1.83251	NCI-H82	2.036841	1.83251	1.584161	1.584161	1.83251	1.78444	1.656987	
1.991678	1.987523	1.847911	J GAMMA 1	1.847911	1.697372	1.697308	1.697308	1.65489	1.65489	1.65489	
1.98146	2.003777	2.003777	HSC-4	2.003777	1.961835	1.716392	1.813181	1.813181	1.716392	1.716392	
1.948251	1.942487	1.860809	BT-549	1.82168	1.784269	1.772132	1.772132	1.784269	1.772132	1.784269	
1.958046	1.834833	1.834833	KYSE-30	1.834833	1.715246	1.715246	1.715246	1.724498	1.724498	1.715246	
1.933422	1.933422	1.933422	Ca9-22	1.933422	1.834222	1.800455	1.800455	1.834222	1.800455	1.761857	
1.890137	1.890137	1.890137	NCI-H2405	1.773334	1.659001	1.659001	1.659001	1.773334	1.890137	1.890137	
1.890646	1.829466	1.829466	SNU-1	1.829466	1.772601	1.772601	1.772601	1.772601	1.829466	1.829466	
1.957241	1.890976	1.822954	BPH-1	1.822954	1.667432	1.792125	1.792125	1.792125	1.792125	1.792125	
1.879792	1.841654	1.785699	NCI-H1944	1.785699	1.785699	1.548246	1.548246	1.785699	1.548246	1.850252	
2.180352	2.180352	1.869018	HSC-2	1.869018	1.752236	1.677613	1.752236	1.752236	1.677613	1.752236	
1.929906	1.956787	1.929906	BICR 31	1.770836	1.726735	1.726735	1.770836	1.726735	1.542706	1.542706	
1.753698	1.799248	1.799248	KYSE-50	1.834604	1.799248	1.799248	1.723625	1.799248	1.723625	1.723625	
2.223692	2.121907	1.985493	NCI-H1915	1.584745	1.526041	1.526041	1.526041	1.584745	1.584745	1.584745	
1.807218	1.772413	1.772413	D341 Med	1.772413	1.571754	1.571754	1.571754	1.571754	1.571754	1.571754	
1.778293	1.766194	1.778293	PE/CA-PJ15	1.778293	1.766194	1.754892	1.754892	1.754892	1.754892	1.754892	
1.711868	1.711868	1.711868	KYSE-510	1.810039	1.810039	1.865423	1.865423	1.873453	1.873453	1.83955	
2.238168	2.238168	2.187147	KOSC-2 cl3-43	1.9272	1.756854	1.749263	1.756854	1.756854	1.749263	1.749263	
2.107238	2.107238	2.107238	OUMS-23	1.398161	1.340194	1.340194	1.340194	1.398161	1.398161	1.398161	
1.795395	1.795395	1.795395	HT1080	1.716312	1.709252	1.709252	1.709252	1.716312	1.716312	1.716312	
1.748791	1.748791	1.839748	JURKAT EC-1	2.026982	1.73293	1.73293	1.633969	1.541303	1.541303	1.73293	
1.85743	1.846101	1.746288	COR-L88	1.746288	1.662522	1.436193	1.436193	1.436193	1.358055	1.358055	
1.777514	1.744822	1.744822	DOR	1.744822	1.707448	1.735445	1.735445	1.735445	1.735445	1.735445	
1.849249	1.799743	1.736552	NCI-H838	1.736552	1.386175	1.386175	1.386175	1.736552	1.708279	1.708279	

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896871	SNP_A-4301043	SNP_A-2050943	SNP_A-4194333	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy_SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
2.04259	1.843055	1.833977	NCI-H358	1.616292	1.601406	1.601406	1.601406	1.616292	1.544582	1.544582	
1.799032	1.799032	1.72264	NCI-H1048	1.72264	1.582111	1.582111	1.582111	1.72264	1.582111	1.582111	
1.832067	1.832067	1.832067	ME-180	1.818763	1.152715	1.152715	1.152715	1.283	1.283	1.236571	1.236571
2.055964	1.934809	1.934809	A431	1.711788	1.429045	1.429045	1.480405	1.711788	1.711788	1.711788	1.711788
1.798831	1.655427	1.655427	Hs 834.T	1.655427	1.588674	1.588674	1.588674	1.655427	1.749468	1.749468	1.749468
1.766147	1.718126	1.718126	LS 123	1.766147	1.718126	1.718126	1.686649	1.686649	1.491094	1.491094	1.491094
1.979441	1.940058	1.849158	NCI-H1703	1.545292	1.545292	1.496768	1.496768	1.496768	1.369566	1.496768	1.496768
1.793939	1.793939	1.691614	VM-CUB1	1.691614	1.583095	1.583095	1.581944	1.583095	1.583095	1.583095	1.581944
1.742079	1.742079	1.691582	KYSE-180	1.691582	1.115124	1.115124	1.115124	1.691582	1.691582	1.213775	1.213775
1.688434	1.68673	1.688434	NCI-H1435	1.688434	1.688434	1.642016	1.597578	1.597578	1.463583	1.597578	1.597578
1.687051	1.654565	1.62544	KYSE-450	1.62544	1.62544	1.62133	1.62133	1.654565	1.731584	1.731584	1.731584
1.695994	1.695994	1.695994	ABC-1	1.821465	1.631151	1.531069	1.821465	1.821465	1.427392	1.581544	1.581544
2.272803	1.809793	1.739901	Ishikawa (Herak	1.581711	1.485461	1.485461	1.485461	1.485461	1.485461	1.485461	1.485461
1.630331	1.621594	1.630331	NUGC-3	1.894255	1.894255	1.894255	1.938629	1.938629	1.938629	1.938629	1.686424
2.142015	1.862549	1.632151	G-292 Clone A1	1.632151	1.586493	1.586493	1.586493	1.632151	1.553404	1.553404	1.553404
1.743687	1.743687	1.711979	U-2 OS	1.531069	1.531069	1.091937	1.091937	1.444932	1.444932	1.444932	1.444932
2.075495	1.777872	1.612912	WI-26 VA4	1.612912	1.475702	1.475702	1.475702	1.612912	1.475702	1.475702	1.475702
1.946282	1.950923	1.950923	HSC-3	1.946282	1.900101	1.579158	1.60895	1.60895	1.491559	1.491559	1.491559
1.606156	1.596084	1.606156	NCI-H1155	1.606156	1.880484	1.549157	1.549157	1.549157	1.361765	1.422602	1.422602
1.533309	2.054025	2.054025	HLE	1.824073	1.671347	1.502773	1.33404	1.33404	1.207269	1.33404	1.33404
1.783383	1.642882	1.642882	TCO-1	1.500115	1.500115	1.500115	1.550607	1.550607	1.500115	1.500115	1.500115
1.805614	1.805614	1.805614	NCI-H2122	1.612302	1.612302	1.612302	1.491989	1.578307	1.491989	1.491989	1.491989
1.896948	1.896948	1.896948	HO-1-u-1	1.878258	1.499806	1.422073	1.483547	1.483547	1.422073	1.422073	1.422073
1.681482	1.625276	1.625276	OVISE	1.596084	1.350078	1.056803	1.350078	1.350078	1.262972	1.262972	1.262972
1.560923	1.569711	1.569711	JIMT-1	1.569711	1.569711	1.560923	1.560923	1.560923	1.486513	1.486513	1.486513
1.559225	1.559225	1.559225	GCT	1.634182	1.559225	1.425808	1.425808	1.634182	1.425808	1.425808	1.425808
1.764879	1.575099	1.575099	HCE7	1.532376	1.532376	1.467941	1.467941	1.467941	1.449091	1.449091	1.449091
1.700815	1.700815	1.700815	EPLC-272H	1.51988	1.229118	1.078447	1.078447	1.229118	1.078447	1.229118	1.229118
1.619915	1.619915	1.619915	SKN-3	1.486289	1.469727	1.383551	1.383551	1.383551	1.308174	1.308174	1.308174
1.540136	1.540136	1.532783	NCI-H596	1.540136	1.532783	1.532783	1.277556	1.325049	1.277556	1.277556	1.043167
1.6904903	1.657718	1.5713857	NL20-TA	1.4062546	1.2869552	1.2869552	1.2869552	1.4062546	1.2417358	0.9188213	0.9188213
2.101073	2.101073	1.911132	Hs 839.T	1.486995	1.349446	1.161179	1.349446	1.479333	1.479333	1.486995	1.486995
1.481061	1.420345	1.420345	KYSE-270	1.482494	1.482494	1.482494	1.482494	1.482494	1.482494	1.482494	1.482494
1.5332242	1.5332242	1.5332242	ASH-3	1.4633571	1.0416993	0.6029178	0.6029178	1.0416993	1.0416993	1.0416993	1.0416993
1.571386	1.494878	1.477639	MG 63	1.290444	1.105502	1.105502	1.105502	1.105502	1.072862	1.072862	1.072862
1.581811	1.40831	1.379672	NCI-H522	1.350421	1.051937	1.051937	1.051937	1.051937	1.051937	1.051937	1.051937
1.482494	1.482494	1.336256	T.T	1.336256	1.297796	1.297796	1.297796	1.336256	1.297796	1.297796	1.297796
1.334058	1.334058	1.334058	T47D	1.317182	1.288082	1.125496	1.24014	1.288082	1.24014	1.24014	1.24014
1.591883	1.591883	1.591883	SCC-4	1.331222	1.314036	1.259058	1.259058	1.259058	1.120295	1.120295	1.120295
1.468964	1.295384	1.271063	KYSE-520	1.295384	1.295384	1.189498	1.189498	1.295384	1.323572	1.448724	1.448724
1.4561719	1.4070343	1.4070343	RERF-LC-Sq1	1.1816543	0.8110917	0.8110917	0.8110917	1.1065244	1.1065244	0.8192055	0.8192055
1.2974278	1.2974278	1.2974278	PCI 4B	1.2734951	1.2734951	1.1000873	0.9104992	0.9104992	0.8192706	0.8192706	0.8192706
1.794636	1.399085	1.254747	NCI-H1568	1.254747	1.135441	1.135441	1.135441	1.254747	1.254747	1.254747	1.254747
1.495961	1.495961	1.495961	OSC-20	1.486272	1.318041	1.214457	1.214457	1.214457	1.214457	1.214457	1.214457
1.4273916	1.3198491	1.3198491	RERF-LC-MA	1.1506247	0.6438562	0.4775737	0.4775737	1.1506247	0.9293365	0.9293365	0.9293365
1.246048	1.246048	1.246048	BICR10	1.246048	1.246048	1.225724	1.189919	1.189919	1.189919	1.189919	1.189919
1.228044	1.228044	1.228044	OE21	1.196197	1.188734	1.128425	1.150412	1.150412	1.128425	1.128425	1.128425

ProbeSetID	SNP_A-2135346	SNP_A-4200418	SNP_A-2166333	SNP_A-1952822	SNP_A-1896874	SNP_A-4301045	SNP_A-2050942	SNP_A-4194332	SNP_A-2259746	SNP_A-4230238	SNP_A-1989992
Affy.SNP.ID	7217859	11481086	7664895	12588794	11787387	9465412	5926612	10390310	9140266	7909743	2569009
dbSNP	rs17586344	rs17586365	rs17172432	rs4140770	rs6960438	rs17172438	rs10244108	rs7785013	rs759169	rs3735061	rs2877260
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54914607	54914995	54915526	54915685	54916117	54925746	54926546	54926876	54928845	54938076	54945838
Strand	-	+	+	-	+	+	+	+	-	-	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
1.129069	1.129069	1.15668	RERF-LC-Ad2	1.236402	1.236402	1.236402	1.497214	1.497214	1.236402	1.236402	
0.9685444	0.9685444	1.0113754	SKOV3	1.0113754	1.0113754	1.0113754	1.0426041	1.0426041	1.0426041	1.0992004	
1.019878	1.019878	1.019878	T.Tn	1.019878	0.9251308	0.7779219	0.7779219	0.9251308	0.7779219	0.7779219	
0.9617503	1.0602814	1.0602814	OAW28	0.9460151	0.9460151	0.9460151	1.0147267	1.1441107	1.1593644	1.5892559	
1.0865166	1.052959	0.947366	NCI-H1563	0.947366	0.8467476	0.5995443	0.5995443	0.8467476	0.8467476	0.6627455	
0.8900125	0.8900125	0.914327	PNT2	0.9777086	0.9777086	0.6220993	0.5957445	0.6220993	0.6220993	0.6220993	
0.8192706	0.7912006	0.8192706	NCI-N87	1.0172224	0.8192706	0.8192706	0.8192706	0.8192706	0.8192706	0.7970291	
0.7998296	0.7998296	0.7998296	NCI-H2228	0.7691386	0.5840321	0.5840321	0.5840321	0.5840321	0.5038623	0.428946	
0.6931083	0.6931083	0.764117	MES-SA	0.7979483	0.7979483	0.764117	0.6839943	0.7291	0.6839943	0.6839943	
0.6945376	0.6945376	0.6945376	WPE1-NB26	0.5229413	0.3940197	0.3940197	0.3940197	0.5229413	0.5229413	0.5229413	
0.6227272	0.5817452	0.5817452	NCI-H322	0.5817452	0.5637644	0.4553836	0.4005065	0.4005065	0.3921312	0.3921312	
0.27500705	0.16812853	0.16812853	LU99A	0.0571383	-0.05829269	-0.05829269	-0.05829269	0	0	0	

ProbeSetID	SNP_A-1989993	SNP_A-2075433	SNP_A-1867443	SNP_A-2145133	SNP_A-2236084	SNP_A-2196993	SNP_A-1989993	SNP_A-2212413	SNP_A-1958322	SNP_A-2024856	SNP_A-1959701
Affy_SNP.ID	2569010	6293987	11350852	7356959	8796070	8136360	2569034	8389173	12670418	5521926	12690481
dbSNP	rs2110290	rs12671550	rs7803660	rs7796139	rs980653	rs13244925	rs10488142	rs17172446	rs6947594	rs12666347	rs17172448
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54946977	54947884	54948280	54950085	54960099	54966465	54970156	54976807	54977506	54980139	54981429
Strand	-	+	+	+	+	+	-	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
KYSE-520	6.810133	6.810133	6.810133	6.911036	7.123581	6.911036	6.810133	6.810133	7.022481	7.077002	7.022481
OE21	6.763342	6.763342	6.763342	7.025238	7.12099	7.025238	7.025238	7.025238	6.676592	7.12099	6.674109
H3255	6.387916	6.387916	6.387916	6.387916	6.436391	6.071772	6.071772	6.436391	6.76763	6.76763	6.76763
NCI-H1573	6.228164	6.228164	6.228164	6.271808	6.393148	6.271808	6.271808	6.393148	6.651435	6.651435	6.651435
HSC-2	5.862791	5.862791	5.862791	6.07028	6.164015	6.07028	6.056095	6.056095	6.056095	6.305957	6.056095
HCC-827	5.592099	5.592099	5.592099	5.890594	5.929655	5.940205	5.940205	5.940205	5.940205	6.067734	5.940205
Hs 852.T	5.225252	5.095803	5.225252	5.225252	5.320132	5.320132	5.320132	5.380848	5.689587	5.69901	5.69901
KYSE-30	5.370001	5.370001	5.370001	5.543344	5.561594	5.543344	5.543344	5.561594	5.660296	5.682268	5.682268
A431	5.412951	5.406551	5.406551	5.412951	5.412951	5.412951	5.412951	5.412951	5.403763	5.403763	5.311354
BT-20	5.165112	5.165112	5.165112	5.165112	5.255214	4.98627	4.98627	5.03707	5.03707	5.255214	5.03707
Ca9-22	4.64452	4.447982	4.607766	4.64452	4.801972	4.801972	4.607766	4.894021	5.002894	5.077503	5.002894
NCI-H1568	4.345007	4.234798	4.234798	4.234798	4.299728	4.299728	4.299728	4.300957	4.40944	4.639796	4.639796
PC-9	4.277177	4.198772	4.198772	4.198772	4.198772	4.130393	4.130393	4.198772	4.228008	4.228008	4.533135
PNT1A	4.25082	4.19421	4.19421	4.25082	4.325935	4.325935	4.325935	4.355007	4.423277	4.433431	4.433431
LUDLU-1	4.302562	4.302562	4.302562	4.302562	4.328124	4.218594	4.218594	4.328124	4.328124	4.385474	4.342817
RERF-LC-MA	4.078277	4.078277	4.078277	4.085289	4.085289	4.078277	4.014391	4.078277	4.078277	4.228734	4.228734
U373 MG	3.932433	3.932433	3.987465	4.19363	4.19363	3.999538	3.987465	3.987465	3.987465	3.987465	3.987465
KYSE-450	3.839512	3.839512	3.848588	3.850594	3.891358	3.850594	3.850594	3.891358	3.891358	4.196933	4.196933
KOSC-2 cl3-43	3.891889	3.891889	3.891889	3.891889	3.891889	3.96648	3.891889	3.891889	3.836376	3.96648	3.836376
LU65	3.742142	3.719327	3.519079	3.519079	3.742142	3.519079	3.445273	3.519079	3.763611	3.844304	3.763611
PC-14	3.68137	3.54121	3.522732	3.522732	3.54121	3.54121	3.54121	3.752228	3.782379	3.8063	3.782379
NCI-H596	3.805698	3.645819	3.645819	3.645819	3.805698	3.607008	3.607008	3.6606	3.6606	3.805698	3.846534
PC-3 [JPC-3]	3.759412	3.573193	3.573193	3.556417	3.556417	3.453662	3.383068	3.453662	3.556417	3.581544	3.564817
647-V	3.633134	3.633134	3.633134	3.665173	3.633134	3.665173	3.633134	3.633134	3.61471	3.69285	3.544312
HSC-3	3.278307	3.278307	3.534462	3.536572	3.689422	3.749374	3.689422	3.689422	3.749374	3.750396	3.749374
CASKI	3.576586	3.576586	3.576586	3.576586	3.576586	3.723045	3.723045	3.723045	3.723045	3.760595	3.723045
JHH-4	3.541971	3.541971	3.541971	3.896589	3.896589	3.915621	3.896589	3.896589	3.34333	3.915621	3.34333
CAL-85-1	3.469467	3.516086	3.516086	3.66219	3.66219	3.66219	3.66219	3.66219	3.676831	3.719764	3.676831
BT-549	3.573703	3.573703	3.576227	3.603244	3.605214	3.605214	3.576227	3.576227	3.605214	3.621248	3.605214
HSC-4	3.450828	3.450828	3.450828	3.724958	3.450828	3.822233	3.450828	3.450828	3.265616	3.822233	3.265616
NUGC-4	3.444932	3.365147	3.365147	3.365147	3.365147	3.338054	3.255214	3.338054	3.338054	3.365147	3.365147
BICR10	3.562107	3.162741	3.162741	3.162741	3.315289	3.315289	3.315289	3.451125	3.451125	3.549798	3.451125
KYSE-140	3.423342	3.55894	3.414961	3.380047	3.55894	3.380047	3.380047	3.55894	3.55894	3.567433	3.55894
SNU-5	3.418943	3.331158	3.384649	3.418943	3.418943	3.384649	3.281381	3.384649	3.418943	3.508005	3.508005
SK-MEL-3	3.671889	3.342804	3.342804	3.342804	3.342804	3.329676	3.329676	3.342804	3.342804	3.792404	3.792404
KYSE-180	3.391391	3.391391	3.350747	3.400959	3.44654	3.400959	3.350747	3.383399	3.44654	3.456599	3.44654
PNT2	3.344443	3.344443	3.344443	3.344443	3.344443	3.387888	3.387888	3.43292	3.596798	3.609776	3.596798
G-361	3.419261	3.415262	3.415262	3.415262	3.419261	3.415262	3.366944	3.415262	3.43471	3.580101	3.580101
KYSE-220	3.150044	3.150044	3.228393	3.228393	3.278242	3.278242	3.278242	3.278242	3.309952	3.322587	3.416851
WM-115	3.272539	3.272539	3.272539	3.217533	3.217533	3.217533	3.125027	3.33799	3.389698	3.47088	3.47088
KYSE-50	3.178913	2.985901	2.861266	2.861266	2.985901	2.985901	2.861266	2.985901	3.272484	3.644571	3.644571
OSC-20	3.366962	3.247106	3.247106	3.247106	3.247106	3.247106	3.074493	3.254485	3.254485	3.451482	3.254485

ProbeSetID	SNP_A-1989993	SNP_A-2075433	SNP_A-1867443	SNP_A-2145133	SNP_A-2236083	SNP_A-2196993	SNP_A-1989993	SNP_A-2212413	SNP_A-1958323	SNP_A-2024853	SNP_A-1959703
Affy_SNP.ID	2569010	6293987	11350852	7356959	8796070	8136360	2569034	8389173	12670418	5521926	12690481
dbSNP	rs2110290	rs12671550	rs7803660	rs7796139	rs980653	rs13244925	rs10488142	rs17172446	rs6947594	rs12666347	rs17172448
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54946977	54947884	54948280	54950085	54960099	54966465	54970156	54976807	54977506	54980139	54981429
Strand	-	+	+	+	+	+	-	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
SAT	3.499189	3.499189	3.499189	3.613984	3.613984	3.499189	3.410532	3.410532	3.410532	3.646178	3.410532
NCI-H1650	3.250625	3.250625	3.226752	3.250625	3.226752	3.226752	3.21881	3.226752	3.226752	3.286336	3.286336
SK-CO-1	3.298624	3.214665	3.083634	3.083634	3.083634	3.083634	3.083634	3.104548	3.312425	3.359676	3.312425
NCI-H1792	3.210818	3.210818	3.097603	3.307588	3.307588	3.326225	3.326225	3.326225	3.495952	3.466622	3.466622
NCI-H1703	3.350656	3.350656	3.31151	3.350656	3.350656	3.355215	3.350656	3.350656	3.350656	3.350656	3.31151
PCI 4B	3.196797	3.196797	3.139215	3.139215	3.216227	3.493305	3.493305	3.389636	3.493305	3.493305	3.447126
NCI-H1563	3.165517	3.165517	2.929295	2.904189	2.994152	2.904189	2.786788	2.904189	2.994152	3.065561	2.994152
NCI-H841	3.198607	3.198607	3.198607	3.258508	3.258508	3.258508	3.210031	3.26769	3.26769	3.339659	3.339659
LOU-NH91	3.150625	3.236477	3.236477	3.255214	3.236477	3.111632	3.042101	3.111632	3.111632	3.255214	3.111632
HuH-7	3.248015	3.248015	3.248015	3.156129	3.14575	3.145741	3.132062	3.145741	3.132062	3.145741	3.132062
LC-1 sq	3.244118	3.244118	3.244118	3.270998	3.270998	3.317127	3.270998	3.270998	3.317127	3.325277	3.317127
NCI-H522	3.343293	3.25499	3.25499	3.343293	3.343293	3.315966	3.25499	3.315966	3.346851	3.315966	3.23984
201T	2.996777	2.996777	2.996777	3.080531	3.080531	3.080531	3.080531	3.080531	3.209689	3.219482	3.209689
G-292 Clone A141B1	3.246704	3.178798	3.068352	3.068352	3.273746	3.346851	3.189576	3.288949	3.288949	3.288949	3.111819
RPMI-7951	2.930215	2.930215	2.889567	3.075119	3.237276	3.237276	3.136163	3.237276	3.251775	3.27814	3.27814
BEN	3.072106	3.072106	3.072106	2.801076	3.072106	3.072106	2.801076	2.801076	2.801076	3.208143	2.918863
NCI-H2228	2.871503	2.854193	2.807179	2.807179	2.807179	2.786316	2.786316	2.807179	3.14985	3.203498	3.203498
HCC-44	3.045739	3.042494	3.045739	3.045739	3.05395	3.045739	3.042494	3.045739	3.042494	3.05395	3.042494
NCI-H1734	3.065023	3.050554	3.050554	2.890712	2.890712	2.890712	2.890712	2.993099	2.993099	2.993099	2.851112
YKG-1	3.033795	3.033795	3.033795	3.146245	3.364257	3.146245	3.033795	3.033795	3.033795	3.146245	3.146245
MOG-G-CCM	2.86254	2.849092	2.828661	2.849092	3.031173	2.961564	2.849092	2.961564	3.031173	3.031173	3.195196
SK-BR-3	3.021924	3.021924	3.021924	3.111632	3.011782	2.970854	2.970854	2.970854	2.970854	3.111632	2.970854
COLO 858	3.086051	2.99048	2.884603	3.086051	3.259441	3.014686	2.884603	2.901488	3.014686	3.117069	3.014686
SCC-4	2.968483	2.968483	2.968483	3.075656	2.968483	3.075656	3.075656	3.281205	3.075656	3.281205	3.075656
TE7	3.072852	3.067934	3.067934	3.072852	3.067934	2.973552	2.922271	2.922271	2.922271	2.922271	2.922271
NCI-H1755	2.972533	2.932106	2.932106	2.972533	2.988022	2.972533	2.932106	2.932106	2.988022	3.056232	3.110788
CaR-1	3.140528	3.140528	3.140528	3.168405	3.168405	3.060541	3.046502	3.060541	3.046502	3.060541	3.046502
OVISe	2.98711	2.978038	2.978038	2.978038	3.062548	2.964705	2.872409	2.964705	2.964705	2.973583	2.964705
Calu-3	2.968905	2.960385	2.947605	2.947605	2.947605	2.947605	2.947605	2.968905	3.045146	3.340448	3.045146
NCI-H1623	2.864433	2.864433	2.864433	2.864433	2.966213	2.864433	2.864433	2.966213	2.966213	3.001706	3.001706
BICR 78	3.091729	2.98919	2.950678	2.950678	2.950678	2.950678	2.950678	2.950678	2.950678	3.082782	3.082782
LS 123	2.946317	2.751925	2.743541	2.751925	2.946317	2.946317	2.946317	2.946317	3.095048	3.233732	3.233732
AN3CA	3.098264	3.059462	3.006993	2.929357	2.923445	2.923445	2.923445	2.929357	2.929357	3.010512	3.010512
639-V	3.045247	3.05393	3.05393	3.05393	3.05393	3.045247	3.000034	2.939334	2.939334	2.939334	2.923518
WM 266-4	2.932147	2.930068	2.743898	2.930068	2.758188	2.758188	2.758188	2.764571	2.764571	2.934027	2.764571
SKN-3	3.029406	2.845994	2.845994	3.029406	3.029406	3.034188	2.770619	2.782018	2.782018	3.034188	2.782018
NCI-H2085	2.956287	2.783821	2.719346	2.719346	2.783821	2.769006	2.769006	2.783821	2.975518	3.280992	2.975518
RKN	2.946472	2.798058	2.92781	3.062369	3.062369	3.087655	3.062369	3.062369	2.92781	3.062369	2.92781
MKN45	2.917551	2.915472	2.915472	2.917551	2.93364	2.93364	2.917551	2.917551	2.93364	2.972131	2.93364
KHOS/NP	2.898552	2.898552	2.89373	2.79313	2.79313	2.79313	2.68205	3.009302	3.065113	3.102904	3.009302
SNU-16	2.945984	2.899323	2.899323	2.899323	2.899323	2.898964	2.803598	2.898964	3.06163	3.06163	3.0794
HCE7	2.934736	2.934736	2.934736	3.025829	3.025829	3.025829	2.934736	2.890214	2.832436	2.890214	2.7579
OCUG-1	3.08614	3.08614	3.08614	3.104705	3.08614	2.981715	2.885473	2.885473	2.836887	2.885473	2.836887
NL20-TA	3.008478	2.960892	2.960892	3.008478	3.008478	3.044443	3.008478	3.008478	3.050063	3.226394	3.226394
KYSE-70	2.982228	2.982228	2.895285	2.855137	2.895285	2.855137	2.788359	2.855137	2.895285	2.920657	2.855137
MOG-G-UVW	2.865042	2.808771	2.77446	2.77446	2.77446	2.808771	2.808771	2.808771	2.890246	3.032303	2.890246
NCI-H2342	2.797719	2.797719	2.748554	2.913392	2.985368	2.913392	2.86797	2.913392	2.985368	3.114487	3.114487

ProbeSetID	SNP_A-1989993	SNP_A-2075433	SNP_A-1867443	SNP_A-2145133	SNP_A-2236083	SNP_A-2196993	SNP_A-1989999	SNP_A-2212413	SNP_A-1958323	SNP_A-2024853	SNP_A-1959703
Affy_SNP.ID	2569010	6293987	11350852	7356959	8796070	8136360	2569034	8389173	12670418	5521926	12690481
dbSNP	rs2110290	rs12671550	rs7803660	rs7796139	rs980653	rs13244925	rs10488142	rs17172446	rs6947594	rs12666347	rs17172448
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54946977	54947884	54948280	54950085	54960099	54966465	54970156	54976807	54977506	54980139	54981429
Strand	-	+	+	+	+	+	-	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
ABC-1	2.712288	2.501923	2.712288	2.501923	2.501923	2.501923	2.501923	2.585563	2.929337	3.12143	2.929337
NCI-H650	2.858761	2.858761	2.819032	3.033231	3.033231	3.108333	3.108333	3.108333	3.108333	3.15666	3.108333
FTC 133	2.854504	2.854504	2.854504	2.854504	2.930674	2.930674	2.930674	2.930674	2.901161	2.930674	3.037151
MDST8	2.901552	2.901552	2.901552	2.891538	2.891538	2.891538	2.891538	2.891538	2.891538	2.891538	2.891538
KYSE-150	2.699953	2.820802	2.853313	2.853313	2.976979	2.853313	2.830263	2.853313	2.976979	3.058522	2.976979
SW 48	2.67929	2.67929	2.718244	2.718244	2.718244	2.773247	2.773247	2.785657	3.015924	3.064594	3.064594
NCI-H2023	2.849425	2.660853	2.660853	2.662482	2.662482	2.662482	2.662482	2.894005	2.981808	2.99363	2.99363
COLO-680N	2.881362	2.881362	2.846328	2.824089	2.824089	2.846328	2.846328	2.846328	2.846328	2.846328	2.824089
KP-4	2.8349	2.915294	2.915294	2.915294	3.017151	2.915294	2.881022	2.881022	2.881022	2.915294	2.881022
SW-1710	2.828271	2.828271	2.828271	2.911509	2.945558	2.945558	2.945558	2.945558	2.945558	2.945558	2.911509
KATO III	2.976341	2.969411	2.969411	2.976341	3.047355	2.976341	2.969411	2.969411	2.976341	2.976341	2.969411
RERF-GC-1B	2.646094	2.646094	2.864561	2.864561	2.905275	2.864561	2.798813	2.798813	2.812758	2.864561	2.812758
CAL-29	2.811331	2.905222	2.905222	2.905222	2.834447	3.048188	2.834447	2.834447	2.834447	2.778671	2.834447
Sci-1	2.727654	2.525828	2.525828	2.702691	2.816032	2.816032	2.702691	2.816032	2.816032	2.936228	2.936228
NCI-H810	2.800201	2.800201	2.800201	2.800201	2.800201	2.866635	2.866635	2.870222	2.870222	2.887754	2.958016
T.Tn	2.648515	2.648515	2.648515	2.648515	2.834522	2.826181	2.826181	2.826181	2.826181	2.871151	2.871151
DOK	2.86147	2.849264	2.849264	2.849264	2.755414	2.755414	2.755414	2.755414	2.86147	2.862593	2.912571
COLO 741	2.808684	2.808684	2.735782	2.6317	2.6317	2.6317	2.6317	2.654984	2.654984	2.735782	2.654984
NCI-H1651	2.732046	2.732046	2.732046	2.814292	2.814292	2.787206	2.642702	2.787206	2.787425	2.814292	2.787425
RWPE-1	2.766043	2.766043	2.766043	2.766043	2.766043	2.708904	2.708904	2.708904	2.768763	2.781853	2.768763
KYSE-270	2.641385	2.762888	2.808357	2.934997	2.994284	2.934997	2.808357	2.808357	2.934997	2.934997	2.905929
NCI-H196	2.736006	2.729414	2.736006	2.787766	2.875531	2.736006	2.729414	2.855963	2.875531	3.047636	2.875531
SK-LU-1	2.766452	2.766452	2.741065	2.64471	2.64471	2.64471	2.64471	2.741065	2.741065	2.84104	2.84104
NCI-H1693 (ship16)	2.490456	2.480372	2.480372	2.480372	2.490456	2.490456	2.480372	2.625598	2.625598	2.757767	2.775541
NCI-H2135	2.739424	2.683915	2.683915	2.683915	2.683915	2.683915	2.683915	2.739424	2.807975	2.810307	2.810307
VMRC-RCW	2.70904	2.70904	2.70904	2.70904	2.70904	2.732449	2.5876	2.846339	2.846339	2.897305	2.846339
RERF-LC-Sq1	2.732214	2.664625	2.664625	2.664625	2.664625	2.631295	2.588083	2.664625	2.664625	2.664625	2.67573
A375.S2	2.640507	2.640507	2.640507	2.732202	2.89428	2.732202	2.732202	2.89428	2.963817	3.04182	3.04182
SW 1116	2.578677	2.578677	2.465444	2.495928	2.578677	2.578677	2.578677	2.636934	2.682628	2.728136	2.910699
CAL-148	2.728069	2.728069	2.590999	2.728069	2.728069	2.728069	2.640091	2.728069	2.922303	2.948374	2.948374
22RV1	2.570591	2.570591	2.570591	2.570591	2.680441	2.680441	2.680441	2.788216	2.859307	2.859307	2.788216
IM-95	2.726768	2.71284	2.71284	2.548464	2.548464	2.470988	2.467064	2.470988	2.470988	2.564571	2.720706
HT 115	2.752992	2.752992	2.752992	2.783217	2.752992	2.606907	2.752992	2.888136	2.888136	2.894883	2.888136
NCI-H727	2.721212	2.688134	2.636206	2.688134	2.768608	2.688134	2.688134	2.768608	2.808128	2.808128	2.808128
T.T	2.765135	2.724491	2.616559	2.616559	2.724491	2.646752	2.616559	2.616559	2.646752	2.724491	2.646752
OE33	2.71355	2.813356	2.813356	2.974191	2.813356	2.974191	2.974191	2.813356	2.927695	2.927695	2.927695
PLC/PRF/5	2.791804	2.791804	2.780598	2.791804	2.791804	2.791804	2.780598	2.723031	2.780598	2.834663	2.723031
SAS	2.770354	2.770354	2.730892	2.730892	2.706455	2.867372	2.640114	2.640114	2.563539	2.640114	2.550694
MKN74	2.71947	2.638692	2.638692	2.638692	2.734463	2.638692	2.624157	2.624157	2.638692	2.734463	2.734463
HMCB	2.810558	2.810558	2.810558	2.837513	2.810558	2.810558	2.703291	2.703291	2.703291	3.046953	3.046953
A2058	2.802048	2.926732	2.926732	2.931113	2.931113	2.926732	2.689237	2.689237	2.6922	2.861331	2.6922
HCC-366	2.676061	2.673229	2.501184	2.673229	2.501184	2.501184	2.450513	2.673229	2.673229	2.760617	2.34381
P31/FUJ	2.672578	2.672578	2.672578	2.672578	2.672578	2.672578	2.447636	2.544966	2.523266	2.544966	2.523266
SW 1417	2.709818	2.706353	2.667168	2.667168	2.667168	2.665496	2.53452	2.665496	2.672327	2.672327	2.672327
KMRC-1	2.703912	2.703912	2.703912	2.982526	2.982526	2.982526	2.685098	2.685098	2.685098	2.818901	2.685098
OAW42	2.639675	2.639675	2.639675	2.548929	2.684119	2.664958	2.539072	2.664958	2.664958	2.684119	2.684119
PE/CA-PJ15	2.602075	2.618352	2.618352	2.750212	2.750235	2.750235	2.750212	2.750235	2.773809	2.989917	2.989917

ProbeSetID	SNP_A-1989993	SNP_A-2075433	SNP_A-1867443	SNP_A-2145133	SNP_A-2236083	SNP_A-2196993	SNP_A-1989993	SNP_A-2212413	SNP_A-1958323	SNP_A-2024853	SNP_A-1959703
Affy_SNP.ID	2569010	6293987	11350852	7356959	8796070	8136360	2569034	8389173	12670418	5521926	12690481
dbSNP	rs2110290	rs12671550	rs7803660	rs7796139	rs980653	rs13244925	rs10488142	rs17172446	rs6947594	rs12666347	rs17172448
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54946977	54947884	54948280	54950085	54960099	54966465	54970156	54976807	54977506	54980139	54981429
Strand	-	+	+	+	+	+	-	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
NUGC-3	2.520268	2.520268	2.658408	2.658408	2.784162	2.658408	2.358704	2.358704	2.658408	2.784162	2.784162
DMS 273	2.651855	2.596038	2.479432	2.596038	2.479432	2.432525	2.410622	2.410622	2.432525	2.803818	2.432525
MKN1	2.642413	2.642413	2.650727	2.628987	2.628987	2.628987	2.536078	2.536078	2.628987	2.650727	2.650727
RERF-LC-Ad2	2.554791	2.503269	2.503269	2.503269	2.554791	2.554791	2.503269	2.609326	2.503269	2.609326	2.634979
TASK1	2.655076	2.772209	2.772209	2.917509	2.772209	2.917509	2.772209	2.772209	2.647455	2.920521	2.569751
NCI-H23	2.590847	2.590847	2.590847	2.773534	2.784261	2.808346	2.773534	2.773534	2.787623	2.787623	2.787623
KP-1N	2.623518	2.623518	2.623518	2.623518	2.623518	2.643498	2.623518	2.623518	2.584539	2.643498	2.271359
NCI-H711	2.622425	2.622425	2.622425	2.695184	2.740764	2.695184	2.622425	2.622425	2.695184	2.695184	2.695184
NCI-H661	2.621297	2.642402	2.583113	2.583113	2.583113	2.585352	2.585352	2.585352	2.647144	2.669251	2.669251
EBC-1	2.62068	2.403268	2.378068	2.378068	2.42825	2.248656	2.181707	2.248656	2.42825	2.573762	2.573762
JHH-6	2.779299	2.616256	2.616256	2.779299	2.779299	2.76178	2.642355	2.642355	2.642355	2.76178	2.642355
NB69	2.699919	2.699919	2.689168	2.689168	2.771381	2.771381	2.771381	2.771381	2.908541	2.931249	2.931249
SR	2.66083	2.584221	2.584221	2.607445	2.584221	2.539168	2.539168	2.584221	2.584221	2.607445	2.539168
T84	2.888762	2.631886	2.571408	2.571408	2.57326	2.571408	2.571408	2.57326	2.491003	2.491003	2.428822
HO-1-N-1	2.803468	2.803468	2.803468	2.803468	2.77179	2.742058	2.643568	2.643568	2.563124	2.643568	2.563124
BALL-1	2.880693	2.530202	2.609933	2.609933	2.609933	2.609933	2.588918	2.588918	2.588918	2.609933	2.588918
Nthy-ori 3-1	2.612139	2.612139	2.553458	2.553458	2.777624	2.777624	2.553458	2.553458	2.777624	2.803567	2.803567
FTC 238	2.4464	2.610295	2.464597	2.610295	2.709581	2.709581	2.709581	2.709581	2.762467	2.803217	2.762467
HuCCT1	2.700847	2.820899	2.820899	2.820899	2.820899	2.631828	2.595263	2.436273	2.436273	2.631828	2.436273
HOS	2.52382	2.52382	2.461168	2.52382	2.591082	2.591082	2.591082	2.591082	2.688646	2.688646	2.662103
NCI-H2170	2.450821	2.450821	2.450821	2.450821	2.653983	2.653983	2.653983	2.672121	2.760118	2.822158	2.760118
Hs 936.T	2.318884	2.318884	2.376169	2.414761	2.477415	2.477415	2.477415	2.618352	2.662046	2.662046	2.662046
NCI-H1793	2.526093	2.526093	2.450315	2.478588	2.526093	2.526093	2.526093	2.59349	2.686361	2.939439	2.686361
MRK-nu-1	2.577449	2.577449	2.577449	2.577449	2.577449	2.492123	2.468905	2.492123	2.492123	2.692303	2.492123
KHOS-312H	2.479847	2.479847	2.467813	2.443321	2.443321	2.443321	2.333239	2.584645	2.708284	2.746929	2.708284
LCLC-103H	2.564998	2.543019	2.564998	2.564998	2.564998	2.543019	2.469591	2.543019	2.564998	2.615696	2.615696
H69V	2.559905	2.512645	2.353756	2.512645	2.559905	2.559905	2.559905	2.512645	2.685656	3.005221	3.005221
GCT	2.607164	2.57446	2.566172	2.566172	2.566172	2.566172	2.53827	2.566172	2.57446	2.619295	2.619295
COLO 857	2.577945	2.561936	2.554101	2.554101	2.577945	2.554101	2.537048	2.537048	2.554101	2.592715	2.554101
RERF-LC-MS	2.550014	2.514021	2.514021	2.465322	2.465322	2.42825	2.415786	2.42825	2.42825	2.62068	2.42825
MKN7	2.42825	2.42825	2.42825	2.42825	2.42825	2.42825	2.378068	2.514021	2.514021	2.538066	2.538066
A-427	2.534808	2.534808	2.534808	2.534808	2.558871	2.640033	2.640033	2.709017	2.640033	2.709017	2.640033
T47D	2.510155	2.510155	2.510155	2.510155	2.510155	2.533249	2.533249	2.5755	2.5755	2.533249	2.503281
BT-483	2.495977	2.495977	2.445374	2.445374	2.495977	2.495977	2.445374	2.590541	2.590541	2.590541	2.590541
SKN	2.53615	2.53615	2.53615	2.53615	2.752747	2.521268	2.509067	2.509067	2.509067	2.521268	2.509067
HCC202	2.368726	2.368726	2.368726	2.445856	2.445856	2.558586	2.558586	2.631492	2.558586	2.631492	2.658408
OCUM-1	2.378068	2.378068	2.378068	2.378068	2.390695	2.378068	2.288093	2.378068	2.378068	2.501923	2.501923
NCI-H2122	2.505158	2.505158	2.560499	2.560499	2.560499	2.560499	2.560499	2.6412	2.560499	2.54407	2.53476
HT 29	2.728484	2.581439	2.477097	2.728484	2.45166	2.45166	2.522147	2.522147	2.522147	2.621641	2.522147
huH-1	2.419942	2.480336	2.480336	2.908268	2.908268	2.908268	2.885229	2.885229	2.57843	2.885229	2.57843
HCC-15	2.570994	2.570994	2.570994	2.570994	2.559477	2.341068	2.338426	2.338426	2.341068	2.559477	2.341068
KHOS-240S	2.473022	2.473022	2.432761	2.432761	2.473022	2.568839	2.568839	2.568839	2.568839	2.568839	2.568839
JIMT-1	2.512548	2.724536	2.724536	2.724536	2.756934	2.756934	2.724536	2.480543	2.558431	2.480543	2.480543
KON	2.586717	2.332814	2.266351	2.586717	2.586717	2.586717	2.465788	2.465788	2.465788	2.628581	2.465788
COLO 853	2.457194	2.457194	2.457194	2.444113	2.457194	2.457194	2.457194	2.457194	2.444113	2.444113	2.439805
U-2 OS	2.453017	2.477574	2.514021	2.514021	2.477574	2.44066	2.44066	2.44066	2.514021	2.514021	2.477574
ME-180	2.448612	2.448612	2.448612	2.506357	2.448612	2.663112	2.448612	2.448612	2.253708	2.663112	1.958807

ProbeSetID	SNP_A-1989993	SNP_A-2075433	SNP_A-1867443	SNP_A-2145133	SNP_A-2236083	SNP_A-2196993	SNP_A-1989993	SNP_A-2212413	SNP_A-1958323	SNP_A-2024853	SNP_A-1959703
Affy_SNP.ID	2569010	6293987	11350852	7356959	8796070	8136360	2569034	8389173	12670418	5521926	12690481
dbSNP	rs2110290	rs12671550	rs7803660	rs7796139	rs980653	rs13244925	rs10488142	rs17172446	rs6947594	rs12666347	rs17172448
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54946977	54947884	54948280	54950085	54960099	54966465	54970156	54976807	54977506	54980139	54981429
Strand	-	+	+	+	+	+	-	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
FU97	2.675708	2.697847	2.697847	2.710777	2.710777	2.710777	2.697847	2.585999	2.585999	2.710777	2.522135
Calu-1	2.564903	2.564903	2.564903	2.568282	2.564903	2.436149	2.423993	2.436149	2.436149	2.573892	2.423993
NCI-H2452	2.471895	2.427814	2.23931	2.427814	2.427814	2.427814	2.427814	2.427814	2.427814	2.427814	2.427814
UMC-11	1.938157	1.938157	1.938157	1.938157	2.167067	2.261901	2.261901	2.416872	2.428859	2.489482	2.489482
COLO 684	2.338093	2.317165	2.338093	2.371416	2.545086	2.545086	2.545086	2.545086	2.570994	2.752581	2.570994
SVts-8	2.844466	2.7122876	2.7122876	2.4530171	2.4032677	2.2220626	2.1952216	1.3741214	0.9708537	0.9708537	0.9293365
HEC-1	2.399539	2.399539	2.399539	2.399539	2.422174	2.422174	2.399539	2.354994	2.354994	2.399539	2.354994
A-375	2.598911	2.608029	2.608029	2.608029	2.598911	2.598911	2.585045	2.585045	2.57368	2.57368	2.399791
OUMS-23	2.611953	2.611953	2.611953	2.72614	2.72614	2.611953	2.382607	2.396737	2.396737	2.611953	2.396737
EJ138	2.375953	2.375953	2.35813	2.375953	2.393516	2.393516	2.393516	2.393516	2.472704	2.472704	2.491661
HO-1-u-1	2.443841	2.540448	2.540448	2.540448	2.540448	2.484412	2.403218	2.403218	2.403218	2.484412	2.403218
HT55	2.460062	2.391842	2.349784	2.391842	2.460062	2.460062	2.460062	2.558527	2.564239	2.564239	2.564239
NCI-H2347	2.249212	2.195815	2.195815	2.195815	2.195815	2.195815	2.195815	2.249212	2.347662	2.398333	2.398333
REFR-LC-KJ	2.602707	2.488033	2.343964	2.343964	2.317643	2.317643	2.293402	2.317643	2.383365	2.535982	2.383365
COR-L279	2.413361	2.366668	2.366668	2.366668	2.366668	2.364978	2.358844	2.358844	2.364978	2.364978	2.358844
SW 1573	2.41606	2.427839	2.427839	2.438566	2.473034	2.566623	2.566623	2.566623	2.473034	2.566623	2.473034
SW 900	2.486168	2.319439	2.319439	2.355734	2.355734	2.319439	2.315006	2.315006	2.355734	2.356754	2.355734
HT 1376	2.477182	2.351215	2.351215	2.351215	2.351215	2.350538	2.313635	2.226974	2.313635	2.350538	2.313635
HUP-T3	2.464892	2.464892	2.464892	2.71578	2.71578	2.732807	2.732807	2.732807	2.212563	2.732807	2.212563
STC 1	2.356945	2.34628	2.34628	2.34628	2.34628	2.34628	2.163252	2.34628	2.390632	2.401071	2.401071
NCI-H2286	2.341299	2.352007	2.352007	2.179945	2.179945	2.179945	2.179945	2.352007	2.500298	2.500298	2.500298
VMRC-LCD	2.33985	2.33985	2.301119	2.261862	2.301119	2.288093	2.261862	2.261862	2.288093	2.301119	2.288093
H69AR	2.065076	1.994946	1.994946	1.994946	2.02478	2.02478	1.912842	2.02478	2.065076	2.338336	2.547151
VMRC-LCP	2.33487	2.33487	2.238646	2.33487	2.33487	2.33487	2.238646	2.33487	2.33487	2.554327	2.091864
FLYA13	2.274574	2.274574	2.274574	2.274574	2.275807	2.275807	2.275807	2.275807	2.321944	2.405061	2.405061
HUP-T4	2.421351	2.554518	2.554518	2.554518	2.405312	2.554518	2.405312	2.405312	2.40023	2.405312	2.390153
COR-L 105	2.315278	2.226814	2.190769	2.175252	2.190769	2.190769	2.172331	2.172331	2.226814	2.307818	2.307818
KMRC-20	2.43173	2.43173	2.43173	2.480702	2.480702	2.43173	2.323259	2.30787	2.214398	2.30787	2.214398
NCI-H2081	2.295681	2.295681	2.295681	2.295681	2.179945	2.160811	2.125432	2.160811	2.179945	2.400218	2.400218
VM-CUB1	2.374383	2.320149	2.175292	2.293832	2.427964	2.427964	2.427964	2.293832	2.293832	2.427964	2.293832
BU25 TK	2.23834	2.23834	2.23834	2.302796	2.302796	2.312055	2.312055	2.312055	2.312055	2.345205	2.345205
RCM-1	2.287897	2.287897	2.146105	2.146105	2.146105	2.393843	2.146105	2.146105	2.393843	2.416048	2.393843
NCI-H2172	2.262178	2.341145	2.423931	2.423931	2.423931	2.374763	2.341145	2.374763	2.374763	2.474381	2.374763
SK-NEP-1	2.450525	2.450525	2.450525	2.580813	2.580813	2.561758	2.561758	2.561758	2.54419	2.561758	2.54419
D1.1	2.124299	2.124299	2.114554	2.124299	2.366592	2.366592	2.480714	2.366592	2.366592	2.480714	2.366592
NCI-H2196	2.280615	2.35799	2.35799	2.35799	2.362307	2.362307	2.35799	2.35799	2.356448	2.356448	2.356448
NCI-H1435	2.276947	2.276947	2.276856	2.276947	2.276947	2.364825	2.364825	2.364825	2.364825	2.364825	2.364825
GRANTA-519	2.27561	2.31256	2.31256	2.398585	2.31256	2.394322	2.31256	2.31256	2.394322	2.398585	2.394322
SK-MES	2.357519	2.271936	2.268061	2.271936	2.271936	2.271936	2.268061	2.23402	2.252717	2.268061	2.252717
NCI-H1437	2.39072	2.304795	2.304795	2.39072	2.39072	2.519136	2.519136	2.519136	2.519136	2.519136	2.110822
BFTC-905	2.350002	2.518364	2.518364	2.565508	2.565508	2.565508	2.306314	2.283193	2.266746	2.266746	2.13069
DMS 153	2.03882	2.313078	2.313078	1.964421	1.964421	1.964421	1.964076	2.09869	2.058934	2.09869	2.124008
Detroit 562	2.376818	2.277328	2.277328	2.277328	2.376818	2.277328	2.277328	2.261335	2.277328	2.281858	2.277328
RT4	2.338118	2.246948	2.145117	2.145117	2.246948	2.246948	2.150764	2.145117	2.150764	2.246948	2.150764
273T	2.241686	2.241686	2.241686	2.324033	2.440846	2.324033	2.224973	2.224973	2.42958	2.440846	2.440846
SW 1463	2.329058	2.238128	2.238128	2.051985	2.051985	2.051985	2.011619	2.051985	2.051985	2.283101	2.219791
H69	2.382633	2.503874	2.503874	2.503874	2.503874	2.574401	2.503874	2.361951	2.361951	2.361951	2.282643

ProbeSetID	SNP_A-1989993	SNP_A-2075433	SNP_A-1867443	SNP_A-2145133	SNP_A-2236083	SNP_A-2196993	SNP_A-1989999	SNP_A-2212413	SNP_A-1958323	SNP_A-2024853	SNP_A-1959703
Affy_SNP.ID	2569010	6293987	11350852	7356959	8796070	8136360	2569034	8389173	12670418	5521926	12690481
dbSNP	rs2110290	rs12671550	rs7803660	rs7796139	rs980653	rs13244925	rs10488142	rs17172446	rs6947594	rs12666347	rs17172448
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54946977	54947884	54948280	54950085	54960099	54966465	54970156	54976807	54977506	54980139	54981429
Strand	-	+	+	+	+	+	-	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
Hs 195.T	2.37101	2.272173	2.272173	2.272173	2.272173	2.272173	2.249688	2.167012	2.167012	2.167012	1.957812
NCI-H1781	2.103329	2.041495	2.041495	2.041495	2.103329	1.947458	1.922921	1.947458	2.090858	2.160961	2.249503
WIDR	2.229947	2.229947	2.229947	2.136172	2.229947	2.136172	2.136172	2.136172	2.145378	2.148888	2.145378
SK-MEL-1	2.049902	2.049902	2.049902	2.050497	2.057845	2.057845	2.057845	2.077841	2.214036	2.214036	2.25162
SUIT-2	2.153378	2.153378	2.153378	2.153378	2.153378	2.323079	2.142248	2.323079	2.34995	2.35799	2.34995
WPE1-NB26	2.273459	2.273459	2.273459	2.144019	2.396448	2.144019	2.104707	1.954838	2.144019	2.25774	2.144019
HCC-78	2.47794	2.408832	2.244404	2.244404	2.244404	2.23273	2.146105	2.117118	2.146105	2.146105	2.117118
A549	2.313712	2.201767	2.297192	2.297192	2.313712	2.313712	2.297192	2.297192	2.313712	2.316894	2.316894
LK-2	2.298519	2.321415	2.308985	2.308985	2.308985	2.308985	2.274391	2.260926	2.260926	2.308985	2.260926
NCI-H2330	2.198132	2.031038	2.00408	2.00408	2.031038	2.031038	2.059824	1.994122	2.059824	2.223799	2.059824
COLO 685	2.07098	1.798351	2.07098	2.07098	2.08013	2.08013	2.08013	2.197742	2.342926	2.417621	2.342926
BICR22	2.086451	2.108224	2.086451	2.086451	2.086451	2.108224	2.127751	2.127751	2.127751	2.382645	2.355632
MEWO	2.298337	2.19278	2.133804	2.133804	2.143978	2.19278	2.143978	2.19278	2.198496	2.198496	2.198496
IHH-4	2.301691	2.301691	2.301691	2.301691	2.190917	2.190917	2.164669	2.08407	2.164669	2.164669	2.08407
NCI-N87	2.192578	2.180447	2.180447	2.180447	2.180447	2.063172	2.063172	2.063172	2.063172	2.180447	2.180447
KYSE-410	2.176907	2.176907	2.176907	2.176975	2.216206	2.176975	2.176907	2.175075	2.176975	2.247584	2.247584
DMS 92	2.224373	2.224373	2.133391	2.206378	2.206378	2.224373	2.206378	2.138524	2.138524	2.138524	2.138524
RO82-W-1	2.175428	2.174926	2.127834	2.127834	2.127834	2.127834	2.127834	2.174926	2.174926	2.174926	2.127834
NCI-H82	2.173975	2.150066	2.173975	2.2636	2.372316	2.372316	2.372316	2.294522	2.372316	2.407643	2.407643
C32	2.202224	2.183697	2.183697	2.202224	2.212563	2.212563	2.202224	2.183697	2.258698	2.374395	2.258698
Hs 633T	2.255636	2.203408	1.983289	2.063383	2.063383	2.159405	2.159405	2.159405	2.255636	2.321143	2.159405
NCI-H2405	2.132537	2.132537	2.071515	2.132537	2.167026	2.219697	2.219697	2.369665	2.369665	2.369665	2.369665
Hs 895.T	2.188107	2.188107	2.188107	2.114277	2.188107	2.034476	1.995987	2.034476	2.207586	2.058708	2.283075
MG 63	2.269874	2.234007	2.045701	2.10312	2.10312	2.234007	2.34232	2.234007	2.234007	2.234007	2.141685
CAL-120	2.299286	2.299286	2.299286	2.299286	1.917448	2.299286	2.215001	1.917448	1.917448	1.917448	1.781158
H33HJ-JA1	2.297556	2.288041	2.133418	2.133418	2.133418	2.133418	2.092633	2.092633	2.092633	2.092633	1.788557
JSC-1	2.126771	2.126771	2.021989	2.021989	2.126771	2.144829	2.365105	2.144829	2.144829	2.365105	2.144829
Hs 751.T	1.934809	1.888652	1.888652	1.888652	1.970651	1.801737	1.800563	1.970651	2.140559	2.146283	2.248312
HMVII	2.208579	1.891964	1.712521	1.891964	2.1288	2.1288	1.891964	1.891964	2.1288	2.132688	2.132688
fR2	2.129366	2.099889	2.124962	2.124962	2.124962	2.124962	2.124962	2.088229	2.124962	2.130276	2.130276
COLO 201	2.124672	2.124672	2.131379	2.131379	2.131379	2.059443	2.059443	2.03021	2.080649	2.131379	2.080649
NCI-H1915	2.315006	2.203058	2.119292	2.119292	2.119292	2.119292	2.032821	1.986407	2.032821	2.119292	1.986407
IMR-90	2.290298	2.268442	2.268442	2.268442	2.268442	2.268442	2.091613	2.055101	2.055101	2.117312	2.055101
RT112/84	2.209988	2.111169	1.934426	1.934426	2.111169	2.004339	1.952082	1.952082	1.952082	2.004339	2.004339
Hs 894(E).Lu	2.156427	2.156427	2.20283	2.210525	2.210525	2.20283	2.156427	2.105666	2.156427	2.156427	2.105666
LNCAp clone FGC	1.915933	1.915933	1.916809	1.956875	1.915933	1.956875	1.956875	1.962829	1.962829	1.962829	2.012495
NCI-H1581	2.306897	2.163811	2.107682	2.27359	2.306897	2.27359	2.107682	2.107682	2.27359	2.27359	2.107682
HCT 116	2.149107	2.149107	2.149107	2.149107	2.114623	2.114623	2.114623	2.106237	2.106237	2.106237	2.106237
Hs 228.T	2.18635	2.112543	1.997041	1.997041	1.997041	1.997041	1.981619	1.981619	1.981619	2.017834	1.981619
JURKAT EC-1	2.069755	2.069755	1.983941	2.125128	2.126757	2.127751	2.327384	2.126757	2.126757	2.252056	2.126757
SK-N-BE(2)	2.209197	2.209197	1.974278	1.974278	2.113542	1.974278	1.952067	1.974278	1.999293	2.113542	1.999293
LS180	2.205289	2.20404	2.130304	2.20404	2.20404	2.205289	2.153843	2.075285	2.153843	2.153843	2.153843
CAL-51	2.23539	2.23539	2.172399	2.300066	2.300066	2.300066	2.172399	1.999697	2.172399	2.300066	1.999697
SKG-IIIb	2.150476	2.150476	2.150476	2.150476	2.162816	2.254316	2.136695	2.136695	2.136695	2.243026	2.136695
P116 T	2.194911	2.18612	2.084434	2.194911	2.084434	2.194911	2.084434	2.084434	2.084434	2.194911	2.084434
Hs 417.Lu	2.235363	2.235363	2.036001	2.235363	2.235363	1.914507	1.870512	1.870512	2.235363	2.235363	1.870512
WIL2 NS	2.000173	2.071417	2.071417	2.071417	2.071417	1.986581	1.986581	1.986581	2.071417	2.071417	2.096361

ProbeSetID	SNP_A-1989993	SNP_A-2075433	SNP_A-1867443	SNP_A-2145133	SNP_A-2236083	SNP_A-2196993	SNP_A-1989993	SNP_A-2212413	SNP_A-1958323	SNP_A-2024853	SNP_A-1959703
Affy_SNP.ID	2569010	6293987	11350852	7356959	8796070	8136360	2569034	8389173	12670418	5521926	12690481
dbSNP	rs2110290	rs12671550	rs7803660	rs7796139	rs980653	rs13244925	rs10488142	rs17172446	rs6947594	rs12666347	rs17172448
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54946977	54947884	54948280	54950085	54960099	54966465	54970156	54976807	54977506	54980139	54981429
Strand	-	+	+	+	+	+	-	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
COR-L95	2.202991	2.010944	1.924817	2.202991	2.202991	2.135856	2.088746	1.984406	2.088746	2.088746	2.081027
SK-N-FI	2.08041	2.08041	1.993428	1.993428	1.993428	2.08041	1.984319	1.984319	2.050625	2.204429	2.204429
WILCL	1.945738	1.911682	1.826326	1.826326	1.826326	1.826326	2.077982	2.077982	2.085765	2.143553	2.085765
P32/ISH	2.107724	2.107724	2.107724	2.107724	2.102647	2.102647	2.061887	1.925379	1.925379	2.061887	1.925379
EPLC-272H	2.069516	2.177911	1.857612	2.123843	2.123843	2.177911	2.123843	2.123843	2.053374	2.123843	2.053374
A2780 ADR	2.108363	2.108363	2.108363	2.157028	2.157028	2.134438	2.134438	2.108363	2.108363	2.134438	2.134438
C8166	1.978976	1.866828	1.962931	1.978976	2.062084	2.062084	2.062084	2.062084	2.206552	2.216808	2.206552
Hs 908.T	2.048583	2.109377	2.109377	2.117617	2.183007	2.183007	2.117617	2.117617	2.183007	2.183007	2.265877
NCI-H322	2.196327	2.14897	2.059372	2.010742	2.059372	2.059372	2.034404	2.034404	2.034404	2.059372	2.034404
NCI-H1417	1.817518	1.474335	1.454848	1.454848	1.415368	1.415368	1.415368	1.415368	1.421703	1.421703	1.480059
HBE4-E6/E7	2.451512	2.451512	2.451512	2.541596	2.451512	2.16919	2.058524	2.058524	2.058524	2.058524	1.829757
NCI-H1522	1.989608	1.989608	2.078866	2.078866	2.180081	2.180081	2.180081	2.205948	2.301626	2.352045	2.301626
LU99A	2.140779	2.028711	2.028711	2.028711	1.836772	1.836772	1.836772	1.836772	1.867145	1.941707	1.867145
HCT-8	2.114263	2.135388	2.135388	2.234685	2.234685	2.340337	2.234685	2.114263	2.114263	1.881867	1.839107
SBC-5	1.952654	1.952654	2.017031	2.15383	2.139377	2.017031	1.952654	2.017031	2.017031	2.078263	2.078263
Ishikawa	2.202816	2.202816	2.202816	2.202816	2.208888	2.202816	2.04976	2.025609	2.025609	2.04976	2.025609
U266B1	2.19363	2.19363	2.19363	2.17783	2.17783	2.17783	2.17783	2.045446	2.045446	2.17783	2.048909
PA-TU-8988T	2.048497	2.07933	2.07933	2.119375	2.119375	2.119375	2.07933	2.07933	2.07933	2.07933	2.074722
HRT-18	2.134203	2.134203	1.931679	1.931679	1.950805	1.950805	1.950805	1.910551	1.950805	1.950805	1.950805
C170	2.021331	2.047206	2.021331	2.047206	2.047206	2.248167	2.349899	2.248167	2.248167	2.137452	2.102535
J GAMMA 1	2.031609	2.031609	1.982055	1.982055	1.982055	2.031609	2.031609	1.982055	1.982055	1.982055	1.905606
LS174T	2.04259	2.04259	1.958778	1.994917	1.958778	1.994917	2.298961	1.994917	2.298961	2.315122	2.315122
NCI-H526	2.049476	2.049476	2.034746	1.887858	1.884511	1.861944	1.861944	1.884511	1.884511	1.884511	1.884511
HLE	1.951627	1.951627	1.951627	2.212603	2.226427	2.250257	2.226427	2.226427	2.226427	2.273892	2.226427
23132/87	2.014391	2.042959	2.042959	2.042959	2.042959	1.727877	1.727877	1.727877	1.836772	2.042959	1.836772
Hs 834.T	2.012639	1.973812	1.973812	2.012639	2.162502	2.058637	2.012639	2.058637	2.162502	2.162502	2.058637
Hs 891.T	2.076676	2.010326	2.010326	1.955966	1.955966	1.866118	1.800918	1.800918	1.866118	1.866118	1.866118
Hs 839.T	2.049263	2.049263	1.913541	1.913541	1.913541	1.913541	1.770477	1.913541	1.913541	2.004872	1.913541
SNU-1	2.004714	2.004714	2.004714	2.004714	2.004714	2.115649	2.004714	2.004714	2.004714	2.115649	2.115649
EB1	2.000692	1.892354	1.864243	1.892354	1.892354	2.000692	1.892354	1.864243	1.892354	2.060559	1.864243
NCI-H2009	2.040599	2.06207	2.06207	2.06207	2.065387	2.06207	1.988783	1.950189	1.950189	2.065387	1.950189
NCI-H1048	1.8791	1.794357	2.045929	2.045929	1.831547	1.831547	1.629511	1.831547	1.897142	2.025137	2.025137
A2780	2.075411	2.075411	2.075411	2.075411	2.075411	2.05694	2.075411	2.05694	2.05694	2.075411	2.05694
NCI-H1666	2.042959	2.042959	2.042959	2.042959	2.042959	2.042959	2.042959	1.912113	1.61471	1.61471	1.61471
G-402	2.130069	2.159159	2.159159	2.159159	2.159159	2.159159	2.14749	1.983289	1.983289	2.117464	1.894943
JHH-7	2.010239	2.063736	2.063736	2.063736	1.981416	2.223278	1.981416	1.874568	1.874568	1.874568	1.845644
OUS-11	1.942002	1.908661	1.942002	1.942002	1.880755	1.863502	1.863502	1.880755	1.880755	1.880755	2.049675
AZ-521	2.051631	1.978932	1.967151	1.967151	2.051631	1.973972	1.803437	1.803437	1.803437	1.973972	2.033064
K562/ADM	1.974715	1.974715	1.974715	1.974715	2.218628	2.000837	1.974715	1.783787	2.000837	2.049178	2.000837
NCI-H889	2.143443	1.980051	1.974132	1.887648	1.974132	1.928723	1.800053	1.928723	1.928723	1.974132	1.942267
OE19	1.971904	1.984565	1.989999	1.989999	1.989999	2.039902	2.059514	2.039902	2.039902	2.059514	2.059514
HT 728.T	2.08243	2.023406	1.79076	1.79076	1.79076	1.900571	1.79076	1.79076	1.88089	1.88089	1.79076
GA-10	2.283663	2.341363	2.341363	2.341363	2.283663	2.283663	2.283663	2.283663	1.949998	1.949998	1.929448
Hs 737.T	1.948647	2.033763	1.961542	1.961542	1.934145	1.961542	2.14986	1.961542	1.962566	2.14986	2.099011
NCI-H838	1.98355	1.965137	1.796138	1.965137	1.796138	1.796138	1.793009	1.796138	1.796138	1.962434	1.793009
NCI-H1155	1.899458	2.198065	2.198065	2.198065	2.198065	2.198065	2.191255	2.008471	2.191255	2.191255	2.008471
A2780cis	1.972181	1.994715	1.994715	1.994715	1.994715	1.994715	1.882814	1.881596	1.882814	1.994715	1.994715

ProbeSetID	SNP_A-1989993	SNP_A-2075433	SNP_A-1867443	SNP_A-2145133	SNP_A-2236085	SNP_A-2196993	SNP_A-1989999	SNP_A-2212413	SNP_A-1958322	SNP_A-2024856	SNP_A-1959701
Affy_SNP.ID	2569010	6293987	11350852	7356959	8796070	8136360	2569034	8389173	12670418	5521926	12690481
dbSNP	rs2110290	rs12671550	rs7803660	rs7796139	rs980653	rs13244925	rs10488142	rs17172446	rs6947594	rs12666347	rs17172448
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54946977	54947884	54948280	54950085	54960099	54966465	54970156	54976807	54977506	54980139	54981429
Strand	-	+	+	+	+	+	-	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
G-401	2.023163	2.071543	2.071543	2.071543	2.071543	2.085247	2.071543	1.856036	1.856036	1.856036	1.828823
MS-1-L	2.055186	2.197674	2.197674	2.197674	2.349171	2.197674	1.945518	1.945518	1.941045	1.945518	1.941045
Hs 746T	1.810763	1.810763	1.810763	1.830353	1.830353	1.932861	1.932861	2.066825	2.076971	2.138744	2.204873
SW 13	1.964757	1.791706	1.791706	1.964757	1.791706	1.791706	1.783196	1.783196	1.791706	2.076226	2.076226
AGS	2.031495	1.886192	1.83549	1.83549	1.83549	1.83549	1.685684	1.685684	1.83549	1.947164	1.947164
NCI-H2171	1.942958	1.890197	1.890197	1.825484	1.75585	1.713398	1.585294	1.713398	1.75585	1.825484	1.825484
Hs 573.T	1.930985	1.836788	1.836788	1.836788	1.860975	1.860975	1.930985	1.930985	1.930985	1.976402	1.860975
HT1080	2.162611	2.211557	2.211557	2.323814	2.211557	2.439496	2.439496	2.211557	2.211557	2.439496	2.439496
COR-L 23/CPR	1.982795	2.276318	2.276318	2.276318	1.776189	2.186485	1.918754	1.918754	2.186485	2.186485	1.918754
HGC-27	2.035117	2.035117	1.875246	1.856612	2.031409	1.923706	1.785761	1.761355	1.785761	1.923706	1.785761
SBC-3	1.829007	1.804857	1.829007	1.804857	1.801166	1.739001	1.698077	1.698077	1.698077	1.801166	1.922283
COLO 792	1.915651	1.915651	1.879702	1.879702	1.915651	2.070093	2.070093	1.965297	1.965297	1.965297	1.965297
H-EMC-SS	2.145117	1.96664	1.96664	1.844365	1.823506	1.844365	1.823506	1.823506	1.823506	1.844365	1.844365
J82	1.910432	1.910432	1.910432	1.910432	1.965283	2.042519	1.965283	1.965283	1.965283	2.042519	1.965283
Hs 600.T	1.805151	1.805151	1.805151	1.805151	1.737958	1.905368	1.737958	1.905368	1.905368	2.002754	2.002754
Hs 781.T	1.904757	1.904757	1.873332	1.873332	1.836711	1.781843	1.780473	1.836711	1.933319	2.032679	2.032679
BPH-1	1.700079	1.700079	1.686698	1.700079	1.700079	1.700079	1.700079	1.795766	1.795766	2.167693	2.244272
COR-L23	1.914982	1.914982	1.914982	1.897919	1.826694	1.826694	1.774536	1.774536	1.774536	1.774536	1.774536
A673	2.074005	2.152585	2.152585	2.152585	2.074005	2.074005	2.074005	1.97335	2.074005	2.152585	1.865483
COLO 205	1.973113	1.899249	1.884661	1.884661	1.899249	1.899249	1.899249	1.884661	1.899249	2.025366	2.025366
RMG-I	2.090481	1.784191	1.761794	1.761794	2.256362	1.758486	1.737184	1.758486	2.040656	1.758486	1.758486
WI-26 VA4	1.854321	1.955937	1.854321	1.854321	1.854321	1.955937	1.854321	1.854321	1.854321	1.854321	1.783087
Eos-HL-60	1.891695	1.981823	1.853456	1.853456	1.796386	1.981823	1.796386	1.796386	1.796386	1.796386	1.796386
GP5D	1.689044	1.76123	1.76123	2.010958	2.010958	2.012151	2.012151	2.012151	2.010958	2.010958	2.021245
MES-SA/Dx-5	1.996304	1.996304	1.992777	1.992777	1.893746	1.992777	1.893746	1.893746	1.893746	1.874252	1.893746
SBC-2	1.74312	1.583328	1.74312	1.814704	1.814704	1.663106	1.663106	1.814704	1.814704	1.953666	1.959275
MCAS	1.87627	1.880634	1.87627	1.87627	1.833045	1.880634	1.833045	1.833045	1.809916	1.833045	1.729192
Ishikawa (Heraklio) 02 ER-	1.899578	1.815412	1.677436	1.815412	2.155334	2.215523	2.15722	2.155334	2.155334	2.15722	2.155334
MOLT-4F	1.831195	1.831195	1.831195	1.831195	1.831195	1.831791	1.831195	1.831195	1.830904	1.830904	1.34409
HARA	1.770243	1.903146	1.903146	1.903146	1.681192	1.786491	1.681192	1.602715	1.681192	1.786491	1.681192
NCI-H1944	2.021961	2.112335	1.997416	1.997416	1.810579	1.997416	1.810579	1.810579	1.810579	1.810579	1.747863
BICR 31	1.819485	1.902564	1.902564	1.902564	1.902564	1.924535	1.924535	1.924535	1.924535	1.924535	1.784285
D341 Med	1.780333	1.780333	1.840297	1.864878	1.864878	1.864878	1.864878	1.846435	1.864878	1.864878	1.846435
KHM-3S	1.79781	1.79781	1.445684	1.420275	1.445684	1.445684	1.420275	1.37396	1.445684	1.79781	1.445684
COR-L88	1.653898	1.653898	1.622745	1.577822	1.577822	1.622745	1.622745	1.577822	1.622745	1.653898	1.653898
NCI-H358	1.843771	1.843771	1.707353	1.786631	1.843771	1.94415	1.843771	1.786631	1.831287	1.831287	1.831287
1321N	2.011978	1.941531	1.941531	1.763408	1.763408	1.755253	1.755253	1.755253	1.763408	1.785668	1.763408
KYSE-510	1.671897	1.63672	1.671897	1.717203	1.744617	1.717203	1.671897	1.671897	1.717203	1.904876	1.904876
TCO-1	1.893866	1.954076	1.954076	1.954076	1.954076	2.115441	1.954076	1.831378	1.817073	1.817073	1.762546
NCI-H441	1.648591	1.648591	1.648591	1.68422	1.776251	1.776251	1.776251	1.776251	1.776251	1.844426	1.844426
SNU-398	1.669519	1.60187	1.327199	1.274998	1.327199	1.418246	1.418246	1.418246	1.528998	1.746808	1.656516
NCI-H1299	1.589006	1.589006	1.589006	1.589006	1.665326	1.665326	1.531765	1.504572	1.531765	1.665326	1.665326
ASH-3	1.456886	1.6048663	1.6048663	1.6675451	1.6675451	1.6675451	1.6048663	1.6048663	1.4974205	1.4974205	1.3502046
MCF7	1.356144	1.356144	1.264536	1.091937	1.150625	1.150625	1.150625	1.150625	1.150625	1.150625	1.150625
SKOV3	1.3597303	1.3161213	1.1122604	1.1122604	1.0107454	1.0107454	1.0107454	0.8360128	0.7764232	0.7764232	0.7176477
ChaGo-K-1	0.9027652	1.0011962	0.9027652	0.9027652	0.7489773	1.0011962	1.0011962	0.7489773	1.0011962	1.0011962	1.0011962
A373 C6	0.8999572	1.085051	0.8999572	1.0501332	1.0501332	1.1413339	1.1413339	1.0501332	1.0501332	1.0501332	1.0501332

ProbeSetID	SNP_A-198999	SNP_A-207543	SNP_A-186744	SNP_A-214513	SNP_A-223608	SNP_A-219699	SNP_A-198999	SNP_A-221241	SNP_A-195832	SNP_A-202485	SNP_A-195970
Affy.SNP.ID	2569010	6293987	11350852	7356959	8796070	8136360	2569034	8389173	12670418	5521926	12690481
dbSNP	rs2110290	rs12671550	rs7803660	rs7796139	rs980653	rs13244925	rs10488142	rs17172446	rs6947594	rs12666347	rs17172448
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54946977	54947884	54948280	54950085	54960099	54966465	54970156	54976807	54977506	54980139	54981429
Strand	-	+	+	+	+	+	-	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
MES-SA	0.7528253	0.7528253	0.7280241	0.6639377	0.6639377	0.6639377	0.6483078	0.4327235	0.6235642	0.6235642	0.6235642
OAW28	0.8076479	0.8753291	0.8076479	0.8076479	0.8076479	0.8753291	0.8753291	0.8646361	0.8753291	0.8753291	0.8646361
NCI-H460	0.4179333	0.6484461	0.4179333	0.6484461	0.6484461	0.7269926	0.7269926	0.6484461	0.7269926	0.7269926	0.7269926

ProbeSetID	SNP_A-2082958	SNP_A-1989993	SNP_A-1989996	SNP_A-1989999	SNP_A-2099333	SNP_A-4303186	SNP_A-4246316	SNP_A-4273285	SNP_A-2183844	SNP_A-4252184	SNP_A-2234157
Affy_SNP.ID	6417094	2569063	2569064	2569066	6658895	9748516	10659347	5794037	7934783	11486260	8763615
dbSNP	rs4947984	rs2075109	rs2075110	rs11506105	rs11536635	rs7781264	rs2241055	rs6968014	rs845558	rs845559	rs7795728
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54985525	54993112	54993368	54994386	55003224	55005049	55014670	55021651	55021797	55022169	55024300
Strand	+	+	-	-	-	-	+	+	-	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
KYSE-520	6.795414	6.638875	6.605428	6.605428	6.605428	6.605428	6.765417	6.605428	6.541713	6.541713	6.605428
OE21	6.598788	6.517906	6.517906	6.517906	6.517906	6.496708	6.598788	6.496708	6.425512	6.425512	6.598788
H3255	6.711301	6.068365	6.068365	6.068365	6.068365	6.059432	6.178403	6.059432	5.719868	5.719868	6.059432
NCI-H1573	6.499318	6.185886	6.185886	6.185886	5.776193	5.776193	5.937855	5.776193	5.776193	5.776193	5.776193
HSC-2	5.640284	5.637055	5.637055	5.637055	5.637055	5.277746	5.472184	5.472184	5.273699	5.273699	5.273699
HCC-827	5.505771	5.484077	5.006087	5.006087	5.006087	4.554309	5.006087	5.006087	5.01294	5.01294	5.01294
Hs 852.T	5.689587	5.689587	5.689587	5.689587	5.608417	5.608417	5.69901	5.608417	5.564213	5.564213	5.564213
KYSE-30	5.660296	5.161531	5.045703	5.045703	5.028042	5.028042	5.055517	5.028042	5.028042	5.028042	5.028042
A431	5.164804	5.000207	4.849715	4.702284	4.606726	4.322911	4.459508	4.362625	4.362625	4.362625	4.362625
BT-20	4.98627	4.892512	4.892512	4.892512	4.892512	5.03707	5.150625	5.150625	4.892512	4.892512	4.849844
Ca9-22	4.894021	4.894021	4.894021	4.894021	3.939181	3.862511	3.862511	3.862511	3.862511	3.862511	3.862511
NCI-H1568	4.330074	4.330074	4.639796	4.468095	4.362549	4.362549	4.468095	4.362549	4.330074	4.330074	4.330074
PC-9	4.228008	4.228008	4.161384	4.161384	4.159569	4.161384	4.50813	4.161384	4.161384	4.307727	4.161384
PNT1A	4.423277	4.423277	4.423277	4.391829	3.911622	3.909822	3.911622	3.909822	3.909822	3.909822	3.877783
LUDLU-1	4.328124	4.218594	4.218594	4.038422	3.958441	3.937206	3.958441	3.937206	3.937206	3.937206	3.936454
RERF-LC-MA	3.821465	3.798351	3.798351	3.798351	3.798351	3.759412	3.821465	3.759412	3.759412	3.759412	3.655638
U373 MG	3.914574	3.89688	3.86716	3.785303	3.633584	3.624002	3.624002	3.359145	3.332595	3.332595	3.184345
KYSE-450	3.891358	3.891358	3.881491	3.881491	3.881491	3.881491	4.009916	4.009916	3.971591	3.971591	3.971591
KOSC-2 cl3-43	3.627747	3.44257	3.44257	3.44257	3.44257	3.014909	3.014909	3.266909	3.223591	3.223591	3.223591
LU65	3.519079	3.519079	3.398615	3.398615	3.398615	3.507654	3.543247	3.507654	3.428992	3.428992	3.428992
PC-14	3.752228	3.752228	3.752228	3.721957	3.468929	3.468929	3.476991	3.476991	3.476991	3.476991	3.476991
NCI-H596	3.6606	3.6606	3.6606	3.6606	3.631725	3.6606	3.966662	3.966662	3.631725	3.631725	3.477121
PC-3 [JPC-3]	3.556417	3.556417	3.556417	3.564817	3.564817	3.647499	3.647499	3.647499	3.647499	3.647499	3.647499
647-V	3.495411	2.892311	3.052298	3.052298	3.052298	2.75709	3.052298	3.052298	3.070951	3.070951	3.070951
HSC-3	3.689422	3.570758	3.570758	3.570758	3.483098	3.083277	3.083277	3.026608	3.026608	3.026608	3.026608
CASKI	3.548381	3.534216	3.548381	3.534216	3.4255	3.4255	3.534216	3.534216	3.534216	3.534216	3.593694
JHH-4	3.241267	3.141577	3.141577	3.141577	3.141577	2.937333	2.937333	2.937333	3.023907	3.023907	3.023907
CAL-85-1	3.461593	3.406325	3.406325	3.406325	3.461593	3.178683	3.178683	3.170723	2.800803	2.800803	2.800803
BT-549	3.576227	3.418528	3.086527	3.086527	3.086527	2.981832	3.086527	3.183837	3.097396	3.097396	3.097396
HSC-4	3.227337	3.003306	3.227337	3.227337	3.227337	2.991564	2.991564	2.977606	2.977606	2.977606	2.977606
NUGC-4	3.338054	3.255214	3.101801	3.101801	3.101801	3.101801	3.101801	3.101801	3.101801	3.101801	3.101801
BICR10	3.315289	3.315289	3.315289	3.315289	3.436105	3.252159	3.436105	3.367842	3.252159	3.252159	3.088981
KYSE-140	3.380047	3.141975	3.269967	3.269967	3.269967	3.296535	3.34572	3.34572	3.296535	3.296535	3.296535
SNU-5	3.429942	3.429942	3.429942	3.429942	3.056182	3.120892	3.293975	3.176976	3.176976	3.176976	3.120892
SK-MEL-3	3.329676	3.329676	3.408168	3.408168	3.282028	3.408168	3.41406	3.408168	3.249072	3.249072	3.249072
KYSE-180	3.383399	3.350747	3.344869	3.344869	2.9046	2.846382	3.344869	3.344869	3.298136	3.298136	3.298136
PNT2	3.43292	3.387888	3.387888	3.345122	3.324621	3.321656	3.324621	3.321656	3.321656	3.321656	3.234785
G-361	3.415262	3.366944	3.366944	3.366944	3.366944	3.158167	3.457678	3.158167	3.116153	3.116153	3.073389
KYSE-220	3.322587	3.322587	3.487512	3.322587	3.322587	3.446925	3.446925	3.446925	3.336992	3.336992	3.291717
WM-115	3.389698	3.33799	3.088981	3.335003	3.015894	2.899323	3.015894	3.015894	2.899323	2.899323	2.899323
KYSE-50	3.311885	3.311885	3.442913	3.311885	3.311885	3.311885	3.311885	3.266165	3.066828	3.066828	2.931312
OSC-20	3.074493	3.037513	3.254485	3.455998	3.455998	3.455998	3.254485	2.432376	2.432376	2.432376	2.432376

ProbeSetID	SNP_A-2082958	SNP_A-1989999	SNP_A-1989999	SNP_A-1989999	SNP_A-2099333	SNP_A-4303186	SNP_A-4246316	SNP_A-4273285	SNP_A-2183844	SNP_A-4252184	SNP_A-2234157
Affy_SNP.ID	6417094	2569063	2569064	2569066	6658895	9748516	10659347	5794037	7934783	11486260	8763615
dbSNP	rs4947984	rs2075109	rs2075110	rs11506105	rs11536635	rs7781264	rs2241055	rs6968014	rs845558	rs845559	rs7795728
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54985525	54993112	54993368	54994386	55003224	55005049	55014670	55021651	55021797	55022169	55024300
Strand	+	+	-	-	-	-	+	+	-	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
SAT	3.163339	3.143383	3.163339	3.163339	3.410532	2.852776	2.852776	2.402816	2.402816	2.402816	2.614624
NCI-H1650	3.21881	2.950222	2.833398	2.811614	2.811614	2.630843	2.728585	2.728585	2.728585	2.728585	2.728585
SK-CO-1	3.104548	3.104548	3.104548	3.104548	3.069468	3.064365	3.069468	3.064365	3.013955	3.013955	3.064365
NCI-H1792	3.326225	3.038701	2.860977	2.860977	2.860977	2.646902	2.860977	3.006524	3.006524	3.006524	3.006524
NCI-H1703	3.229891	3.19572	2.881341	2.881341	2.881341	2.429356	2.881341	2.893624	2.893624	2.893624	2.893624
PCI 4B	3.216227	3.168299	3.168299	2.937677	2.937677	2.895391	2.937677	2.971565	2.937677	2.937677	2.971565
NCI-H1563	2.786788	2.733679	2.733679	2.733679	2.733679	2.515928	2.733679	2.733679	2.733679	2.733679	2.515928
NCI-H841	3.21368	3.210031	3.210031	2.978593	2.978593	2.727049	2.79542	2.79542	2.753814	2.753814	2.753814
LOU-NH91	3.011782	2.897802	2.822852	2.822852	2.822852	2.822852	2.897802	2.897802	2.897802	2.897802	2.897802
HuH-7	3.106544	2.925215	2.925215	2.925215	3.145741	2.925215	2.925215	2.379725	2.379725	2.379725	2.379725
LC-1 sq	3.238882	2.60998	2.887532	2.887532	2.887532	2.485559	2.485559	2.213795	2.213795	2.213795	2.446536
NCI-H522	3.196711	3.088288	3.088288	2.781324	2.781324	2.610902	2.781324	2.781324	2.610902	2.610902	2.781324
201T	3.209689	2.810754	2.810754	2.529997	2.529997	2.493619	2.529997	2.529997	2.529997	2.529997	2.53985
G-292 Clone A141B1	3.079211	3.079211	3.111819	3.111819	3.111819	2.868771	2.868771	2.868771	2.868771	2.868771	2.868771
RPMI-7951	3.251775	3.237276	3.237276	2.970936	2.788633	2.785943	2.788633	2.788633	2.788633	2.788633	2.788633
BEN	2.801076	2.597317	2.573762	2.597317	2.597317	2.597317	2.918863	2.970854	2.918863	2.918863	2.876586
NCI-H2228	3.161854	3.161854	3.203498	3.161854	2.925508	3.161854	3.3276	3.161854	2.925508	2.925508	2.580554
HCC-44	2.863192	2.543891	2.473463	2.543891	2.543891	2.449649	2.543891	2.543891	2.543891	2.543891	2.449649
NCI-H1734	2.851112	2.851112	2.974304	2.993099	2.993099	2.993099	3.045608	3.045608	3.045608	3.045608	3.045608
YKG-1	2.821648	2.654811	2.654811	2.654811	2.654811	2.629834	2.629834	2.431245	2.431245	2.431245	2.431245
MOG-G-CCM	2.961564	2.961564	2.961564	2.917898	2.917898	2.8508	2.917898	2.917898	2.8508	2.8508	2.814401
SK-BR-3	2.811985	2.62068	2.62068	2.62068	2.62068	2.62068	2.757023	2.757023	2.757023	2.757023	2.757023
COLO 858	3.008681	3.008681	3.008681	3.008681	2.972317	2.901488	2.972317	2.972317	2.972317	2.972317	2.916039
SCC-4	2.995215	2.925539	2.995215	2.995215	3.340602	2.995215	2.995215	2.185984	2.158654	2.158654	2.185984
TE7	2.858032	2.823764	2.823764	2.823764	2.823764	2.7065	2.7065	2.474528	2.474528	2.474528	2.498442
NCI-H1755	3.056232	3.056232	2.680236	2.623436	2.623436	2.623436	2.680236	2.829916	2.778252	2.778252	2.623436
CaR-1	2.836844	2.836844	3.024707	3.024707	3.024707	2.438009	2.438009	2.438009	2.724468	2.724468	2.724468
OVISE	2.872409	2.872409	2.964705	2.973583	2.973583	2.973583	2.973583	2.530334	2.530334	2.530334	2.530334
Calu-3	2.947605	2.89851	2.89851	2.89851	2.741489	2.674258	2.741489	2.741489	2.674258	2.674258	2.741489
NCI-H1623	2.966213	2.820053	3.001706	2.820053	2.820053	2.820053	3.001706	2.820053	2.667981	2.667981	2.667981
BICR 78	2.950678	2.950678	2.782249	2.756535	2.756535	2.662057	2.756535	2.782249	2.756535	2.756535	2.756535
LS 123	3.095048	3.095048	3.095048	2.882053	2.882053	2.854172	2.854172	2.759863	2.604931	2.604931	2.516157
AN3CA	2.941199	2.929357	2.941199	2.941199	3.010512	2.941199	2.941199	2.795069	2.712254	2.712254	2.429381
639-V	2.805324	2.549549	2.805324	2.805324	2.8107	2.8107	2.8107	2.53712	1.990462	1.990462	1.990462
WM 266-4	2.758188	2.758188	2.764571	2.764571	3.039859	3.039859	3.078645	3.039859	2.515373	2.515373	2.480336
SKN-3	2.770619	2.770619	2.770619	2.770619	2.782018	2.532985	2.532985	2.250244	2.250244	2.250244	2.250244
NCI-H2085	2.928009	2.928009	2.928009	2.808422	2.808422	2.928009	3.022977	2.928009	2.903862	2.903862	2.808422
RKN	2.708848	2.560487	2.560487	2.560487	2.560487	2.098202	2.098202	2.063101	2.063101	2.063101	2.063101
MKN45	2.915472	2.701515	2.701515	2.701515	2.670681	2.680521	2.701515	2.864037	2.864037	2.864037	2.86907
KHOS/NP	3.009302	2.705314	2.705314	2.705314	2.705314	2.705314	2.748688	2.705314	2.651613	2.651613	2.651613
SNU-16	2.898964	2.898964	2.889705	2.780158	2.715431	2.780158	2.848673	2.848673	2.780158	2.780158	2.766308
HCE7	2.656444	2.634898	2.471822	2.471822	2.471822	2.3488	2.292985	2.292985	2.292985	2.292985	2.292985
OCUG-1	2.742382	2.742382	2.836887	2.836887	2.836887	2.318845	2.314269	2.268889	2.314269	2.314269	2.314269
NL20-TA	2.960892	2.581733	2.581733	2.581733	2.581733	2.549597	2.574117	2.574117	2.549597	2.549597	2.574117
KYSE-70	2.788359	2.788359	2.720448	2.720448	2.720448	2.720448	2.806601	2.806601	2.720448	2.720448	2.746283
MOG-G-UJVV	2.713888	2.703788	2.890246	2.890246	3.032303	2.844337	2.844337	2.734518	2.598888	2.598888	2.598888
NCI-H2342	2.985368	2.86797	2.86797	2.86797	2.86797	2.847511	2.847511	2.847511	2.832199	2.832199	2.832199

ProbeSetID	SNP_A-2082958	SNP_A-1989999	SNP_A-1989999	SNP_A-1989999	SNP_A-2099333	SNP_A-4303186	SNP_A-4246316	SNP_A-4273285	SNP_A-2183844	SNP_A-4252184	SNP_A-2234151
Affy_SNP.ID	6417094	2569063	2569064	2569066	6658895	9748516	10659347	5794037	7934783	11486260	8763615
dbSNP	rs4947984	rs2075109	rs2075110	rs11506105	rs11536635	rs7781264	rs2241055	rs6968014	rs845558	rs845559	rs7795728
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54985525	54993112	54993368	54994386	55003224	55005049	55014670	55021651	55021797	55022169	55024300
Strand	+	+	-	-	-	-	+	+	-	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
ABC-1	2.855212	2.855212	2.585563	2.855212	2.723537	2.723537	2.855212	2.855212	2.855212	2.855212	2.723537
NCI-H650	2.854933	2.833355	2.833355	2.629671	2.629671	2.589377	2.589377	2.589377	2.589377	2.589377	2.589377
FTC 133	2.930674	2.901161	2.901161	2.853721	2.548702	2.498599	2.548702	2.548702	2.548702	2.548702	2.569538
MDST8	2.800475	2.779563	2.779563	2.741869	2.679495	2.659843	2.679495	2.741869	2.659843	2.659843	2.636425
KYSE-150	2.853313	2.830263	2.830133	2.830263	2.830133	2.762855	2.830133	2.830133	2.762855	2.762855	2.762855
SW 48	3.064594	3.064594	3.064594	3.01044	2.773247	2.851531	2.851531	2.851531	2.767149	2.767149	2.759752
NCI-H2023	2.981808	2.981808	2.99363	2.981808	2.543019	2.480177	2.53615	2.53615	2.53615	2.53615	2.543019
COLO-680N	2.625644	2.602602	2.536845	2.536845	2.625644	2.536845	2.625644	2.726073	2.536845	2.536845	2.640357
KP-4	2.638114	2.596308	2.638114	2.638114	2.638114	2.08156	2.08156	2.048398	2.048398	2.048398	2.048398
SW-1710	2.828065	2.703528	2.80065	2.80065	2.80065	2.486302	2.486302	2.437476	2.437476	2.437476	2.437476
KATO III	2.75679	2.669823	2.510953	2.105082	1.878288	1.796866	1.796866	1.796866	1.768602	1.768602	1.796866
RERF-GC-1B	2.798813	2.667946	2.474883	2.474883	2.474883	2.474883	2.50145	2.812758	2.825715	2.825715	2.825715
CAL-29	2.691451	2.512451	2.691451	2.750591	2.750591	2.512451	1.87743	2.706624	2.141685	2.141685	2.141685
Sci-1	2.816032	2.816032	2.936228	2.953033	2.936228	2.54205	2.936228	2.54205	2.42104	2.42104	2.54205
NCI-H810	2.887754	2.870222	2.870222	2.866635	2.810111	2.708667	2.810111	2.708667	2.686315	2.686315	2.538342
T.Tn	2.826181	2.714374	2.807637	2.719391	2.489591	2.489591	2.667878	2.667878	2.523736	2.523736	2.523736
DOK	2.862593	2.862593	2.862593	2.755414	2.512101	2.512101	2.655651	2.655651	2.610155	2.610155	2.610155
COLO 741	2.654351	2.6317	2.654351	2.654351	2.654351	2.609221	2.654351	2.654351	2.654351	2.654351	2.609221
NCI-H1651	2.787206	2.642702	2.426433	2.426433	2.426433	2.354866	2.354866	2.354866	2.354866	2.354866	2.354866
RWPE-1	2.708904	2.708092	2.376409	2.376409	2.376409	2.455729	2.639594	2.536462	2.536462	2.536462	2.536462
KYSE-270	2.808357	2.641732	2.436942	2.436942	2.436942	2.360792	2.426495	2.426495	2.426495	2.426495	2.448476
NCI-H196	2.855963	2.855963	2.632094	2.630657	2.630657	2.630657	2.632094	2.630657	2.517641	2.517641	2.303925
SK-LU-1	2.711792	2.64471	2.711792	2.711792	2.711792	2.711792	2.720189	2.711792	2.439235	2.439235	2.455716
NCI-H1693 (ship16)	2.757767	2.757767	2.757767	2.749322	2.757767	2.749322	2.749322	2.749322	2.600704	2.600704	2.479835
NCI-H2135	2.58594	2.58594	2.810307	2.810307	2.846638	2.810307	2.824208	2.824208	2.630657	2.630657	2.630657
VMRC-RCW	2.814999	2.320911	2.814999	2.814999	2.814999	2.27149	2.27149	2.27149	1.817042	1.817042	1.884856
RERF-LC-Sq1	2.588083	2.588083	2.588083	2.740429	2.815673	2.815673	2.815673	2.740429	2.740429	2.740429	2.480897
A375.S2	2.963817	2.732202	2.963817	2.80195	2.80195	2.80195	2.80195	2.602051	2.569905	2.569905	2.531246
SW 1116	2.910699	2.910699	2.910699	2.910699	2.910699	2.784624	2.928594	2.784624	2.635731	2.635731	2.635731
CAL-148	2.922303	2.640091	2.948374	2.907341	2.948374	2.907341	2.907341	1.89433	1.89433	1.89433	1.642784
22RV1	2.680441	2.499218	2.468524	2.468524	2.468524	2.356792	2.727004	2.727004	2.588024	2.588024	2.588024
IM-95	2.637686	2.637686	2.564571	2.564571	2.564571	2.637686	2.720706	2.637686	2.720706	2.720706	2.637686
HT 115	2.888136	2.828488	2.828488	2.621192	2.621192	2.621192	2.621192	2.621192	2.609548	2.609548	2.621192
NCI-H727	2.636206	2.636206	2.661507	2.372152	2.372152	2.253232	2.626979	2.626979	2.253232	2.253232	2.253232
T.T	2.616559	2.44165	2.646752	2.646752	2.646752	2.744389	2.744389	2.718233	2.718233	2.718233	2.718233
OE33	2.663055	2.503426	2.503426	2.503426	2.503426	2.48992	2.503426	2.672932	2.503426	2.503426	2.559775
PLC/PRF/5	2.71534	2.71534	2.618969	2.590482	2.590482	2.42185	2.42185	2.590482	2.42185	2.42185	2.590482
SAS	2.142619	2.082654	2.082654	2.142619	2.142619	2.082654	2.040926	2.040926	2.345589	2.345589	2.345589
MKN74	2.624157	2.532901	2.46075	2.46075	2.46075	2.366757	2.46075	2.652753	2.623959	2.623959	2.623959
HMCB	2.703291	2.6165	2.703291	2.703291	2.776699	2.6165	2.776699	2.776699	2.776699	2.776699	2.776699
A2058	2.689237	2.6649	2.32992	2.32992	2.32992	2.32992	2.32992	2.32992	1.961133	1.961133	2.016357
HCC-366	2.335936	2.335936	2.34381	2.34381	2.599826	2.34381	2.34381	2.205961	2.205961	2.205961	2.205961
P31/FUJ	2.447636	2.447636	2.523266	2.544966	2.544966	2.523266	2.523266	2.465395	2.434213	2.434213	2.434213
SW 1417	2.665496	2.665496	2.672327	2.672327	2.665496	2.448328	2.663571	2.448328	2.328401	2.328401	2.328401
KMRC-1	2.569147	2.403268	2.365791	2.365791	2.365791	2.355657	2.259331	1.919125	1.919125	1.919125	2.233142
OAW42	2.684119	2.664958	2.664958	2.686076	2.664958	2.540041	2.75861	2.761082	2.75861	2.75861	2.540041
PE/CA-PJ15	2.773809	2.773809	2.989917	2.773809	2.773809	2.66584	2.66584	2.471062	2.35957	2.35957	2.35957

ProbeSetID	SNP_A-2082958	SNP_A-1989994	SNP_A-1989996	SNP_A-1989999	SNP_A-2099333	SNP_A-4303186	SNP_A-4246318	SNP_A-4273285	SNP_A-2183844	SNP_A-4252184	SNP_A-2234157
Affy_SNP.ID	6417094	2569063	2569064	2569066	6658895	9748516	10659347	5794037	7934783	11486260	8763615
dbSNP	rs4947984	rs2075109	rs2075110	rs11506105	rs11536635	rs7781264	rs2241055	rs6968014	rs845558	rs845559	rs7795728
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54985525	54993112	54993368	54994386	55003224	55005049	55014670	55021651	55021797	55022169	55024300
Strand	+	+	-	-	-	-	+	+	-	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
NUGC-3	2.658408	2.447154	2.447154	2.447154	2.375649	2.447154	2.560321	2.560321	2.560321	2.560321	2.560321
DMS 273	2.410622	2.324497	2.410622	2.410622	2.441353	2.441353	2.441353	2.441353	2.353233	2.353233	2.353233
MKN1	2.628987	2.471087	2.471087	2.471087	2.471087	2.367519	2.503256	2.503256	2.503256	2.503256	2.503256
RERF-LC-Ad2	2.609326	2.609326	2.634979	2.634979	2.634979	2.609326	2.609326	2.634979	2.634979	2.634979	2.522424
TASK1	2.398233	2.219162	2.398233	2.398233	2.398233	1.949822	1.949822	1.985566	1.985566	1.985566	1.985566
NCI-H23	2.737593	2.525852	2.737593	2.525852	2.525852	2.157493	2.157493	2.157493	2.157493	2.157493	2.414686
KP-1N	2.230799	2.013314	2.230799	2.230799	2.230799	2.013314	2.013314	2.013314	2.062621	2.062621	2.062621
NCI-H711	2.617164	2.558003	2.446734	2.446734	2.446734	2.345385	2.345385	2.345385	2.345385	2.345385	2.345385
NCI-H661	2.610003	2.585352	2.610003	2.669251	2.631295	2.610003	2.631295	2.631295	2.610003	2.610003	2.610003
EBC-1	2.42825	2.42825	2.42825	2.573762	2.655375	2.790126	2.790126	2.655375	2.609022	2.609022	2.465322
JHH-6	2.505364	2.293558	2.642355	2.642355	2.76178	2.293558	2.293558	2.054832	2.054832	2.054832	2.054832
NB69	2.931249	2.498733	2.498733	2.498733	2.498733	2.498733	2.498733	2.299377	2.299377	2.299377	2.299377
SR	2.50202	2.374155	2.50202	2.462951	2.50202	2.50202	2.646717	2.50202	2.462951	2.462951	2.403356
T84	2.339452	2.339452	2.428822	2.485876	2.485876	2.428822	2.485876	2.529241	2.529241	2.529241	2.566326
HO-1-N-1	2.522111	2.342606	2.522111	2.522111	2.563124	2.02611	1.655004	1.712729	1.712729	1.712729	1.712729
BALL-1	2.490139	2.490139	2.490139	2.588918	2.668919	2.490139	2.490139	2.368865	2.368865	2.368865	2.368865
Nthy-ori 3-1	2.418819	2.410547	2.335088	2.410547	2.410547	2.335088	2.410547	2.493947	2.493947	2.493947	2.493947
FTC 238	2.610295	2.606884	2.606884	2.594289	2.51256	2.395743	2.489348	2.508995	2.508995	2.508995	2.508995
HuGCT1	2.395441	2.325194	2.325194	2.436273	2.631828	2.325194	2.325194	2.437847	2.31344	2.31344	2.31344
HOS	2.608765	2.608765	2.608765	2.452018	2.340132	2.318935	2.340132	2.340132	2.340132	2.318935	2.318935
NCI-H2170	2.672121	2.672121	2.672121	2.672121	2.61579	2.55905	2.61579	2.55905	2.446388	2.446388	2.446388
Hs 936.T	2.618352	2.618352	2.600353	2.593232	2.583749	2.518533	2.583749	2.567737	2.518533	2.518533	2.484876
NCI-H1793	2.59349	2.539945	2.59349	2.59349	2.59349	2.723548	2.723548	2.539945	2.519979	2.519979	2.519979
MRK-nu-1	2.468905	2.360398	2.492123	2.692303	2.708588	2.708588	2.708588	2.413086	2.286447	2.286447	2.286447
KHOS-312H	2.708284	2.350692	2.350692	2.350692	2.350692	2.350692	2.390015	2.277249	2.277249	2.277249	2.277249
LCLC-103H	2.469591	2.318974	2.222316	2.318974	2.318974	2.222316	2.265022	2.222316	2.222316	2.222316	2.222316
H69V	2.367341	2.367341	2.869176	2.464241	2.464241	2.431481	2.43786	2.43786	2.431481	2.431481	2.431481
GCT	2.558752	2.558752	2.619295	2.619295	2.654558	2.558752	2.558752	2.320782	2.269506	2.269506	2.236108
COLO 857	2.537048	2.460258	2.37975	2.37975	2.354369	2.268968	2.354369	2.354369	2.354369	2.354369	2.354369
RERF-LC-MS	2.415786	2.390695	2.140779	2.390695	2.390695	2.390695	2.390695	2.390695	2.390695	2.390695	2.365385
MKN7	2.514021	2.378068	2.403268	2.514021	2.632291	2.585563	2.585563	2.403268	2.403268	2.403268	2.403268
A-427	2.427491	1.903787	1.903787	2.640033	2.677077	1.903787	1.64727	1.64727	1.64727	1.64727	1.762107
T47D	2.503281	2.465653	2.503281	2.503281	2.520774	2.465653	2.520774	2.520774	2.520774	2.586611	2.602215
BT-483	2.621227	2.550729	2.600482	2.550729	2.550729	2.46386	2.488228	2.488228	2.46386	2.488228	2.488228
SKN	2.437785	2.437785	2.437785	2.437785	2.437785	2.369538	2.369538	1.822156	1.780613	1.780613	1.780613
HCC202	2.631492	2.536665	2.536665	2.252571	2.252571	2.219831	2.252571	2.252571	2.219831	2.219831	2.252571
OCUM-1	2.378068	2.378068	2.378068	2.44066	2.44066	2.44066	2.44066	2.44066	2.44066	2.44066	2.44066
NCI-H2122	2.213728	2.213728	2.213728	2.213728	2.53476	2.53476	2.53476	2.414186	2.22956	2.22956	2.22956
HT 29	2.48307	2.522147	2.522147	2.522147	2.522147	2.48307	2.48307	2.48307	2.408332	2.408332	2.48307
huH-1	2.307546	1.87764	1.87764	1.87764	1.87764	1.808159	1.774177	1.87764	2.066148	2.066148	2.066148
HCC-15	2.338426	2.312185	2.312185	2.338426	2.341068	2.268889	2.552886	2.304055	2.304055	2.304055	2.304055
KHOS-240S	2.568839	2.201754	2.201754	2.071783	2.071783	2.040556	2.071783	2.071783	2.040556	2.040556	2.040556
JIMT-1	2.476718	2.146982	2.419418	2.419418	2.419418	2.146982	2.03305	2.219697	2.147805	2.147805	2.19723
KON	2.266351	2.266351	2.465788	2.465788	2.56183	1.981329	1.981329	1.85467	1.640333	1.640333	1.85467
COLO 853	2.327113	2.124907	2.124907	2.21646	2.21646	2.124907	2.21646	2.253919	2.298154	2.298154	2.253919
U-2 OS	2.365385	2.248656	2.248656	2.261862	2.365385	2.365385	2.365385	2.365385	2.314087	2.314087	2.261862
ME-180	1.8093	1.78052	1.8093	1.8093	1.8093	1.78052	1.571051	1.571051	1.900086	1.900086	1.900086

ProbeSetID	SNP_A-2082958	SNP_A-1989999	SNP_A-1989999	SNP_A-1989999	SNP_A-2099333	SNP_A-4303186	SNP_A-4246311	SNP_A-4273285	SNP_A-2183844	SNP_A-4252184	SNP_A-2234151
Affy_SNP.ID	6417094	2569063	2569064	2569066	6658895	9748516	10659347	5794037	7934783	11486260	8763615
dbSNP	rs4947984	rs2075109	rs2075110	rs11506105	rs11536635	rs7781264	rs2241055	rs6968014	rs845558	rs845559	rs7795728
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54985525	54993112	54993368	54994386	55003224	55005049	55014670	55021651	55021797	55022169	55024300
Strand	+	+	-	-	-	-	+	+	-	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
FU97	2.284592	2.201256	2.201256	2.201256	2.201256	2.159773	2.159773	2.159773	2.052679	2.052679	2.052679
Calu-1	2.402741	2.093904	2.402741	2.402741	2.402741	2.093904	2.093904	2.093904	2.046113	2.046113	2.170006
NCI-H2452	2.30473	2.204389	2.30473	2.204389	2.204389	2.204389	2.204389	2.30473	2.204389	2.204389	1.940146
UMC-11	2.477794	2.477794	2.477794	2.428859	2.416248	2.330422	2.330422	2.198348	2.157616	2.138346	1.960387
COLO 684	2.545086	2.545086	2.521798	2.478417	2.378156	2.099731	2.212563	2.212563	2.212563	2.212563	2.212563
SVts-8	0.6438562	0.6438562	0.6668475	0.6668475	0.6668475	0.6668475	0.5737623	0.6668475	0.6668475	0.6668475	0.689657
HEC-1	2.036841	2.003907	2.036841	2.446413	2.446413	2.036841	1.892309	2.320859	2.320859	2.320859	2.320859
A-375	2.271043	1.985899	2.271043	2.271043	2.57368	1.848048	1.832082	1.832082	1.832082	1.832082	1.832082
OUMS-23	2.382607	2.27216	2.382607	2.382607	2.396737	2.396737	2.396737	1.685491	1.685491	1.685491	1.685491
EJ138	2.472704	2.451747	2.451747	2.300768	2.164656	2.107279	2.164656	2.164656	2.164656	2.164656	2.164656
HO-1-u-1	2.379206	2.198119	2.379206	2.379206	2.403218	1.935813	1.776501	1.920771	1.689687	1.689687	1.689687
HT55	2.308778	2.308778	2.471442	2.308778	2.243901	2.062254	2.243901	2.243901	2.062254	2.062254	2.243901
NCI-H2347	2.347662	2.347662	2.398333	2.398333	2.398333	2.347662	2.522604	2.522604	2.337708	2.337708	1.940205
REFR-LC-KJ	2.293402	2.237331	2.155088	2.155088	2.383365	2.205061	2.205061	2.205061	2.205061	2.205061	2.205061
COR-L279	2.250257	2.147695	2.250257	2.250257	2.250257	2.147695	2.147695	2.136998	1.848489	1.848489	2.014118
SW 1573	2.425039	2.136833	2.204362	2.204362	2.204362	1.761042	1.685443	1.891665	1.685443	1.685443	1.883685
SW 900	2.315006	2.092242	2.092242	2.092242	2.489251	2.489251	2.489251	2.452733	2.092242	2.092242	2.034048
HT 1376	2.226974	2.220285	2.220285	2.220285	2.159418	2.059429	2.019196	2.019196	2.019196	2.019196	2.059429
HUP-T3	2.136007	1.887438	2.136007	2.136007	2.212563	1.779741	1.46426	1.539662	1.434263	1.434263	1.539662
STC 1	2.390632	2.390632	2.077238	1.970534	2.077238	1.970534	2.077238	2.017892	1.970534	1.970534	1.922135
NCI-H2286	2.389901	2.389901	2.389901	2.460455	2.460455	2.456837	2.460455	2.456837	2.389901	2.389901	2.110016
VMRC-LCD	2.261862	2.222063	2.222063	2.222063	2.222063	2.181707	2.222063	2.222063	2.085289	2.085289	2
H69AR	2.338336	2.338336	2.393642	2.393642	2.393642	2.338349	2.393642	2.393642	2.338349	2.338349	2.260438
VMRC-LCP	2.019139	2.019139	2.091864	2.091864	2.324768	2.091864	2.091864	2.033007	1.831011	1.831011	1.831011
FLYA13	2.303614	2.275807	2.303614	2.382418	2.405061	2.382418	2.382418	2.303614	2.023249	2.023249	1.931325
HUP-T4	2.084266	1.859521	1.859521	1.859521	1.859521	1.756493	1.298753	1.859521	1.972064	1.972064	1.972064
COR-L 105	2.172331	2.172331	2.172331	2.307818	2.172331	2.307818	2.398786	2.398786	2.132468	2.132468	2.123635
KMRC-20	2.161029	2.12376	2.161029	2.161029	2.16531	2.161029	2.12376	2.161029	2.16531	2.16531	2.16531
NCI-H2081	2.160811	2.125432	2.114277	2.114277	2.125432	2.114277	2.211008	2.211008	2.114277	2.114277	2.092801
VM-CUB1	2.293832	2.293832	2.293832	2.228214	2.165746	2.165746	2.078375	2.165746	2.165746	2.012366	2.165746
BU25 TK	2.266101	2.088858	2.266101	2.266101	2.29009	2.266101	2.282905	2.29009	2.282905	2.282905	2.282905
RCM-1	2.142921	1.738195	1.738195	2.393843	2.706409	1.738195	1.459253	1.848869	1.842065	1.842065	1.842065
NCI-H2172	2.341145	2.060305	2.060305	2.027139	2.060305	2.027139	2.333971	2.333971	2.230693	2.230693	2.230693
SK-NEP-1	2.189648	2.182939	2.189648	2.189648	2.244683	2.182939	2.182939	1.941163	1.941163	1.941163	1.941163
D1.1	2.334831	2.334831	2.366592	2.334831	2.334831	2.282225	2.282225	2.282225	2.281505	2.279843	2.281505
NCI-H2196	2.191376	2.097839	2.097839	2.097839	2.097839	2.097839	2.140394	2.140394	2.140394	2.140394	2.140394
NCI-H1435	2.360117	2.175835	2.175835	2.01706	2.01706	1.898427	2.01706	2.01706	1.898427	1.898427	1.875367
GRANTA-519	2.192335	1.980893	1.980893	1.864773	1.864773	1.804641	1.804641	1.864773	1.804641	1.804641	1.804641
SK-MES	2.148997	2.010786	1.966086	1.966086	1.966086	1.946091	1.966086	2.252717	2.0965	2.0965	2.0965
NCI-H1437	1.980298	1.896708	1.980298	1.980298	1.980298	1.849036	1.980298	2.152078	2.152078	2.152078	2.248458
BFTC-905	2.127475	2.113125	2.127475	2.127475	2.13069	2.13069	2.113125	2.243861	2.133832	2.133832	2.133832
DMS 153	2.124008	2.09869	2.124008	2.227508	2.258751	2.258751	2.325928	2.325928	2.325928	2.296853	2.258751
Detroit 562	2.261335	2.244285	2.244285	2.244285	2.243874	2.134727	2.134727	2.209197	2.134727	2.134727	2.134727
RT4	2.145117	2.145117	2.150764	2.150764	2.150764	2.137438	2.069854	2.069854	1.959495	1.959495	1.959495
273T	2.42958	2.224973	2.403719	2.241182	2.224973	2.224973	2.118821	2.118821	1.980661	1.980661	2.118821
SW 1463	2.051985	2.051985	2.051985	2.219791	2.173568	2.173568	2.213447	2.173568	2.173568	2.173568	2.148696
H69	2.282643	2.143498	2.143498	2.282643	2.282643	2.282643	2.143498	2.302705	2.302705	2.156017	2.302705

ProbeSetID	SNP_A-2082958	SNP_A-1989996	SNP_A-1989996	SNP_A-1989996	SNP_A-209933	SNP_A-4303186	SNP_A-424631	SNP_A-4273285	SNP_A-218384	SNP_A-4252184	SNP_A-2234157
Affy_SNP.ID	6417094	2569063	2569064	2569066	6658895	9748516	10659347	5794037	7934783	11486260	8763615
dbSNP	rs4947984	rs2075109	rs2075110	rs11506105	rs11536635	rs7781264	rs2241055	rs6968014	rs845558	rs845559	rs7795728
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54985525	54993112	54993368	54994386	55003224	55005049	55014670	55021651	55021797	55022169	55024300
Strand	+	+	-	-	-	-	+	+	-	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
Hs 195.T	1.842141	1.839763	1.755441	1.842141	1.842141	1.755441	1.755441	1.842141	1.842141	1.842141	1.842141
NCI-H1781	2.090858	2.171801	2.249503	2.282709	2.282709	2.282709	2.299546	2.299546	2.282709	2.282709	2.171801
WDR	2.136172	2.145378	2.145378	2.148888	2.148888	2.160155	2.172046	2.162829	2.160155	2.112043	2.160155
SK-MEL-1	2.283846	2.283846	2.374737	2.374737	2.407041	2.407041	2.407041	2.374737	2.207385	2.207385	2.177993
SUIT-2	2.323079	1.83929	2.323079	2.35799	2.362192	2.362192	2.362192	1.794279	1.794279	1.794279	1.838055
WPE1-NB26	1.954838	1.785342	1.781874	1.785342	1.785342	1.781874	1.781874	1.895594	1.74069	1.74069	1.761136
HCC-78	2.075762	2.044111	1.971802	1.971802	1.971802	1.808575	1.971802	1.867009	1.867009	1.867009	1.867009
A549	2.297192	2.297192	2.162666	2.162666	2.085975	1.867463	2.035032	2.035032	1.867463	1.867463	1.995698
LK-2	2.101574	2.049802	1.90939	1.90939	1.90939	1.906426	1.8475	1.906426	1.906426	1.906426	1.906426
NCI-H2330	1.994122	1.994122	2.059824	2.156509	2.059824	2.059824	2.156509	2.224546	2.156509	2.156509	2.156509
COLO 685	2.197742	2.197742	2.197742	2.197742	2.072571	2.035944	2.035944	2.035944	1.894674	1.894674	1.907961
BICR22	2.127751	2.065528	2.065528	2.065528	2.127751	2.127751	2.184063	2.289111	2.289111	2.289111	2.184063
MEWO	2.143978	2.143978	2.198496	2.143978	2.198496	2.142468	2.142468	2.142468	2.005635	2.005635	2.005635
IHH-4	2.063877	1.96667	2.063877	2.063877	2.063877	2.08407	1.452093	1.888173	1.888173	1.888173	2.100181
NCI-N87	2.163416	2.137204	2.137204	2.137204	2.163416	2.247822	2.247822	2.247822	2.173119	2.173119	2.201538
KYSE-410	2.176975	2.165909	2.134727	2.134727	2.134727	2.134727	2.134727	2.134727	1.860597	1.860597	2.064399
DMS 92	2.044863	2.003677	2.044863	2.003677	2.003677	1.640382	1.640382	1.640382	1.640382	1.640382	1.640382
RO82-WV-1	2.108461	2.108461	2.108461	2.191943	2.271661	2.271661	2.191943	2.010139	1.625309	1.625309	2.004411
NCI-H82	2.372316	2.372316	2.294522	2.213983	2.213983	1.966903	1.966903	1.966903	1.787439	1.787439	1.966903
C32	2.163402	2.163402	2.258698	2.163402	2.163402	1.72291	1.610156	1.610156	1.593611	1.593611	1.610156
Hs 633T	2.063383	2.063383	2.132495	2.159405	2.321143	2.132495	2.132495	2.124078	2.104526	2.104526	2.104526
NCI-H2405	2.219697	2.033506	2.033506	1.929167	1.929167	1.889522	1.802139	1.802139	1.802139	1.802139	1.889522
Hs 895.T	2.058708	2.058708	2.058708	2.058708	2.058708	2.058708	2.092898	2.092898	2.058708	2.092898	2.04094
MG 63	2.10312	2.0761	2.0761	2.0761	2.141685	2.0761	2.0761	2.141685	2.179905	2.0761	2.179905
CAL-120	1.651636	1.480595	1.480595	1.651636	1.994527	1.441853	1.039667	1.441853	1.447204	1.447204	1.892219
H33HJ-JA1	1.672042	1.788557	2.02641	2.02641	2.02641	2.0133	2.0133	2.006183	2.006183	2.006183	2.006183
JSC-1	1.968114	1.968114	1.968114	1.968114	1.968114	1.794837	1.794837	1.794837	1.794837	1.794837	1.836162
Hs 751.T	2.248312	2.248312	2.36148	2.36148	2.36148	2.36148	2.36148	2.251805	2.239629	2.239629	2.215215
HMVII	2.1288	2.132688	2.150079	2.150079	2.150079	1.917002	1.917002	1.836803	1.594375	1.594375	1.72518
fR2	2.124962	2.124962	2.130276	2.1316	2.137933	2.1316	2.042789	1.614545	1.582378	1.582378	1.614545
COLO 201	2.03021	1.954926	1.939425	1.954926	1.954926	1.954926	1.939425	1.954926	1.874146	1.874146	1.874146
NCI-H1915	1.986407	1.751246	1.751246	1.751246	1.751246	1.560419	1.502105	1.560419	1.560419	1.560419	1.751246
IMR-90	1.981126	1.973535	1.981126	2.055101	2.117312	2.055101	2.055101	1.459358	1.459358	1.459358	1.459358
RT112/84	1.952082	1.952082	2.004339	2.137314	2.137314	2.137314	2.137314	1.808945	1.739238	1.739238	1.560503
Hs 894(E).Lu	1.985073	1.985073	1.985073	1.985073	2.077378	2.077378	2.105666	2.177572	2.077378	2.077378	2.077378
LNcaP clone FGC	2.016802	2.016802	2.119776	2.119776	2.172209	2.172209	2.160045	2.216045	2.216045	2.216045	2.119776
NCI-H1581	1.994079	1.971875	1.971875	1.971875	1.971875	1.84785	1.828654	1.84785	1.828654	1.828654	1.84785
HCT 116	2.103148	1.9972	1.9972	1.958602	1.9972	1.943562	1.943562	1.9972	2.071572	2.071572	1.967691
Hs 228.T	1.981619	1.981619	1.981619	1.98191	1.98191	1.981619	1.98191	2.017834	1.971088	1.971088	1.944797
JURKAT EC-1	2.125128	2.126757	2.126757	2.117409	2.117409	2.000563	2.000563	2.117409	2.117409	2.000563	2.117409
SK-N-BE(2)	1.99298	1.99298	1.99298	1.99298	1.99298	1.99298	1.99298	2.096877	1.99298	1.99298	1.832189
LS180	1.970126	1.937184	2.029767	2.029767	2.093401	2.029767	2.029767	2.029767	2.020786	2.020786	2.020786
CAL-51	1.954091	1.767977	1.819224	1.819224	1.999697	1.819224	1.50411	1.819224	1.524272	1.524272	1.772086
SKG-IIIb	1.808344	1.520135	1.520135	2.243026	2.243026	1.520135	1.350855	1.21404	1.213036	1.213036	1.21404
P116 T	2.084434	2.084434	1.954208	1.938216	1.954208	1.938216	1.938216	1.938216	1.870482	1.870482	2.025981
Hs 417.Lu	1.821619	1.821619	1.870512	1.870512	1.880845	1.880845	1.952229	2.173934	1.952229	1.952229	1.952229
WIL2 NS	2.071417	1.986581	2.096361	2.096361	2.096361	2.096361	2.086241	2.086241	2.081925	2.081925	1.962288

ProbeSetID	SNP_A-2082958	SNP_A-1989993	SNP_A-1989994	SNP_A-1989995	SNP_A-2099333	SNP_A-4303186	SNP_A-4246311	SNP_A-4273285	SNP_A-2183844	SNP_A-4252184	SNP_A-2234151
Affy_SNP.ID	6417094	2569063	2569064	2569066	6658895	9748516	10659347	5794037	7934783	11486260	8763615
dbSNP	rs4947984	rs2075109	rs2075110	rs11506105	rs11536635	rs7781264	rs2241055	rs6968014	rs845558	rs845559	rs7795728
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54985525	54993112	54993368	54994386	55003224	55005049	55014670	55021651	55021797	55022169	55024300
Strand	+	+	-	-	-	-	+	+	-	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
COR-L95	2.081027	2.081027	2.081027	2.060291	2.060291	1.990549	2.060291	2.060291	2.037368	2.037368	2.060291
SK-N-FI	2.050625	1.997373	1.997373	1.997373	1.997373	1.905447	1.997373	1.997373	1.774692	1.774692	1.750428
WILCL	2.077982	2.085765	2.085765	2.085765	2.085765	2.193414	1.973637	1.973637	1.867493	1.867493	1.867493
P32/ISH	1.894001	1.765302	1.765302	1.925379	2.003403	1.925379	1.925379	2.003403	1.986001	1.986001	1.986001
EPLC-272H	2.009147	1.829573	1.979383	1.979383	1.979383	1.829573	1.610255	1.558065	1.50738	1.50738	1.610255
A2780 ADR	2.037056	2.134438	2.134438	1.860218	1.860218	1.797051	1.797051	1.860218	1.786165	1.786165	1.786165
C8166	2.062084	2.062084	2.206552	2.206552	2.063778	2.057591	2.057591	2.057591	1.97582	1.97582	1.97582
Hs 908.T	2.117617	2.060333	2.060333	2.008255	2.008255	1.973244	2.008255	2.060333	2.008255	2.060333	1.988754
NCI-H322	2.010742	2.010742	2.034404	2.034404	2.034404	2.012438	1.821527	1.821527	1.846024	1.846024	1.846024
NCI-H1417	1.80098	2.335461	2.447525	2.447525	2.447525	2.335461	2.335461	2.1312	2.059316	1.80098	1.742315
HBE4-E6/E7	1.810193	1.810193	1.829757	1.829757	1.921172	1.829757	1.829757	1.829757	1.805521	1.805521	1.805521
NCI-H1522	2.205948	2.205948	2.022147	2.058284	1.946943	1.826985	1.67921	1.67921	1.391008	1.391008	1.67921
LU99A	1.836772	1.836772	1.867145	1.941707	2.014391	1.941707	2.057138	2.057138	2.057138	2.057138	2.057138
HCT-8	1.839107	1.743886	1.839107	1.839107	1.87377	1.613819	1.613819	1.87377	1.87377	1.949998	1.949998
SBC-5	2.017031	2.017031	2.078263	2.078263	2.052538	2.052538	2.052538	2.052538	1.961557	1.961557	1.961557
Ishikawa	1.963413	1.763846	1.763846	1.934573	1.934573	1.772242	1.772242	1.934573	1.934573	1.934573	1.972341
U266B1	2.045446	2.045446	2.048909	2.048909	2.049136	2.048909	2.049136	2.049136	2.048909	2.048909	2.049136
PA-TU-8988T	1.947605	1.830369	1.947605	1.947605	2.033278	1.830369	1.830369	1.651734	1.651734	1.651734	1.651734
HRT-18	1.910551	1.751686	1.950805	2.047547	2.117644	2.047547	1.751686	1.745358	1.668613	1.668613	1.668613
C170	2.047206	2.045034	2.045034	1.932418	2.045034	1.932418	1.932418	2.045034	1.932418	1.932418	1.932418
J GAMMA 1	1.905606	2.065147	2.151394	2.065147	2.065147	2.065147	2.044622	2.065147	2.065147	2.044622	2.065147
LS174T	2.002466	1.994917	2.002466	2.071389	2.071389	2.002466	2.002466	2.071389	2.002466	2.002466	2.002466
NCI-H526	1.861944	1.829007	1.861944	1.829007	1.829007	1.829007	1.829007	1.829007	1.829007	1.829007	1.930024
HLE	2.149353	2.023206	2.023206	2.023206	2.023206	1.800331	1.486272	1.486272	1.486272	1.486272	1.617495
23132/87	1.727877	1.679919	1.61471	1.61471	1.61471	1.679919	1.61471	1.61471	1.598175	1.598175	1.61471
Hs 834.T	1.806246	1.806246	1.838878	1.838878	1.806246	1.793009	1.793009	1.793009	1.739032	1.793009	1.793009
Hs 891.T	1.86506	1.811472	1.800918	1.770352	1.770352	1.770352	1.811472	1.86506	1.811472	1.811472	1.810424
Hs 839.T	1.770477	1.603029	1.770477	2.004872	2.039959	2.004872	2.004872	1.603029	1.166692	1.166692	1.166692
SNU-1	1.955849	1.955849	1.934234	1.934234	1.934234	1.903161	1.835444	1.835444	1.903161	1.903161	1.960753
EB1	1.839794	1.839794	1.863245	1.779975	1.863245	1.779975	1.8569	1.863245	1.863245	1.863245	2.052226
NCI-H2009	1.865226	1.839123	1.839123	1.820222	1.820222	1.820222	1.839123	1.835337	1.820222	1.820222	1.78136
NCI-H1048	1.897142	1.897142	1.897142	1.897142	2.025137	2.025137	2.025137	2.145817	1.822202	1.822202	1.770024
A2780	1.986639	1.986639	1.986639	1.98053	1.972341	1.972341	1.972341	1.98053	1.972341	1.947913	1.947913
NCI-H1666	1.61471	1.985537	1.985537	1.985537	1.985537	1.985537	1.985537	1.985537	1.985537	1.985537	1.985537
G-402	1.845187	1.716121	1.845187	1.894943	2.050639	1.894943	1.894943	1.630052	1.622482	1.622482	1.630052
JHH-7	1.763783	1.763783	1.763783	1.763783	1.849492	1.634166	1.280372	1.634166	1.280372	1.280372	1.661971
OUS-11	1.880755	1.880755	1.880755	2.074708	2.111516	2.111516	2.111516	2.111516	2.095105	2.095105	1.688964
AZ-521	1.973972	1.973972	2.033064	2.033064	1.973972	1.814335	1.814335	1.814335	1.814335	1.814335	1.793722
K562/ADM	1.518499	1.45558	1.45558	2.000837	2.000837	1.45558	1.45558	1.45558	1.33344	1.33344	1.33344
NCI-H889	1.928723	1.928723	1.942267	1.942267	1.942267	1.942267	1.972603	1.942267	1.910581	1.910581	1.828073
OE19	2.039902	1.797794	1.797794	1.797794	1.797794	1.774473	1.774473	1.797794	1.797794	1.797794	1.797794
HT 728.T	1.790419	1.790419	1.829267	1.829267	1.88089	1.829267	1.88089	1.88089	1.829267	1.829267	1.970593
GA-10	1.928176	1.928176	1.928176	1.949998	2.031609	2.031609	1.949998	2.031609	2.031609	1.823199	1.98587
Hs 737.T	1.962566	1.962566	1.962566	1.962566	1.962566	1.962566	1.962566	2.034404	2.034404	1.949191	1.949191
NCI-H838	1.764003	1.764003	1.764003	1.758925	1.758925	1.713621	1.713621	1.713621	1.713621	1.713621	1.758925
NCI-H1155	1.944371	1.775737	1.934957	1.775737	1.775737	1.704381	1.929477	1.929477	1.929477	1.929477	1.929477
A2780cis	1.878769	1.721544	1.782278	1.782278	1.782278	1.721544	1.721544	1.763768	1.721544	1.721544	1.763768

ProbeSetID	SNP_A-2082958	SNP_A-1989996	SNP_A-1989996	SNP_A-1989996	SNP_A-2099333	SNP_A-4303186	SNP_A-4246313	SNP_A-4273285	SNP_A-2183844	SNP_A-4252184	SNP_A-2234157
Affy_SNP.ID	6417094	2569063	2569064	2569066	6658895	9748516	10659347	5794037	7934783	11486260	8763615
dbSNP	rs4947984	rs2075109	rs2075110	rs11506105	rs11536635	rs7781264	rs2241055	rs6968014	rs845558	rs845559	rs7795728
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54985525	54993112	54993368	54994386	55003224	55005049	55014670	55021651	55021797	55022169	55024300
Strand	+	+	-	-	-	-	+	+	-	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
G-401	1.81907	1.817672	1.817672	1.817672	1.828823	1.817672	1.751922	1.593395	1.526313	1.526313	1.593395
MS-1-L	1.924802	1.841059	1.924802	1.941045	1.941045	1.642375	1.58909	1.58909	1.58909	1.58909	1.638846
Hs 746T	2.204873	2.204873	2.138744	2.066825	2.066825	1.969222	1.969222	1.932551	1.682223	1.682223	1.641836
SW 13	1.783196	1.783196	1.948559	1.948559	1.957504	1.948559	1.948559	1.948559	1.948559	1.948559	1.948559
AGS	1.846831	1.846831	1.947164	1.947164	1.846831	1.837643	1.837643	1.837643	1.835276	1.835276	1.772757
NCI-H2171	1.75585	1.705756	1.75585	1.705756	1.767414	1.767414	1.997936	1.997936	1.767414	1.767414	1.705756
Hs 573.T	1.715373	1.715373	1.715373	1.809099	1.715373	1.715373	1.715373	1.809099	1.715373	1.715373	1.715373
HT1080	1.986508	1.986508	1.986508	1.865332	1.986508	1.816381	1.816381	1.816381	1.816381	1.739348	1.816381
COR-L 23/CPR	1.776189	1.776189	1.843528	1.843528	1.918754	1.843528	1.654175	1.843528	1.843528	1.843528	1.875111
HGC-27	1.761355	1.761355	1.785761	1.923706	1.923706	1.785761	1.615122	1.731732	1.52983	1.52983	1.601522
SBC-3	1.801166	1.801166	1.801166	1.93826	1.93826	1.93826	1.93826	1.93826	1.924757	1.924757	1.81675
COLO 792	1.892893	1.771931	1.892893	1.892893	1.892893	1.677516	1.484478	1.484478	1.484478	1.484478	1.677516
H-EMC-SS	1.823506	1.719494	1.719494	1.823506	1.912559	1.719494	1.719494	1.719494	1.507312	1.507312	1.502876
J82	1.626425	1.407796	1.407796	1.965283	2.17536	1.407796	1.11728	1.139138	1.139138	1.139138	1.433789
Hs 600.T	1.784564	1.784564	1.784564	1.963369	2.002754	1.963369	2.002754	1.963369	1.963369	1.963369	1.878814
Hs 781.T	1.748382	1.698541	1.748382	1.748382	2.115358	2.115358	2.115358	2.09597	1.698541	1.698541	1.490337
BPH-1	2.244272	2.244272	2.244272	2.020815	1.894345	1.894345	2.020815	2.020815	1.894345	1.894345	1.827414
COR-L23	1.759082	1.735192	1.735192	1.735192	1.892174	1.561057	1.878047	1.696891	1.696891	1.696891	1.637717
A673	1.827812	1.827812	1.865483	1.886627	1.908259	1.886627	1.908259	1.916587	1.908259	1.908259	1.916587
COLO 205	1.884661	1.682771	2.025366	2.025366	2.025366	1.985406	1.635099	1.852166	1.849826	1.849826	1.849826
RMG-I	1.737184	1.438314	1.758486	2.040656	2.040656	2.040656	2.040656	1.431506	1.312974	1.312974	1.312974
WI-26 VA4	1.783087	1.713541	1.783087	1.783087	1.8475	1.726973	1.726973	1.726973	1.726973	1.726973	1.8475
Eos-HL-60	1.760023	1.705468	1.760023	1.796386	1.858641	1.760023	1.705468	1.705468	1.705468	1.705468	1.730191
GP5D	1.899757	1.95283	1.899757	1.899757	1.95283	1.899757	1.895243	1.895243	1.895243	1.85253	1.85253
MES-SA/Dx-5	1.874252	1.800223	1.757404	1.757404	1.72794	1.711007	1.651571	1.651571	1.651571	1.651571	1.651571
SBC-2	1.959275	1.959275	2.043841	1.98024	1.959275	1.98024	1.98024	1.959275	1.901041	1.901041	1.845355
MCAS	1.439734	1.374175	1.374175	1.374175	1.374175	1.217486	1.217486	1.217486	1.229853	1.229853	1.229853
Ishikawa (Heraklio) 02 ER-	1.994195	1.994195	2.05326	1.994195	1.994195	1.666995	1.511108	1.511108	1.511108	1.511108	1.598722
MOLT-4F	1.186421	1.186421	1.186421	1.34409	1.423201	1.186421	1.186421	1.328055	1.328055	1.328055	1.328055
HARA	1.602715	1.55086	1.55086	1.602715	1.786491	1.55086	1.828838	1.839077	1.828838	1.828838	1.828838
NCI-H1944	1.711629	1.422761	1.422761	1.422761	1.711629	1.422761	1.419957	1.419957	1.419957	1.419957	1.422761
BICR 31	1.771196	1.771196	1.771196	1.784285	1.795813	1.784285	1.795813	1.820084	1.820084	1.820084	1.834161
D341 Med	1.840297	1.846435	1.86155	1.846435	1.86155	1.67146	1.67146	1.704237	1.573611	1.704237	1.86155
KHM-3S	1.37396	1.37396	2.211611	2.211611	2.211611	1.374014	1.478763	1.830353	1.478763	1.478763	1.593262
COR-L88	1.577822	1.788572	1.788572	1.788572	1.825085	1.788572	1.788572	1.881687	1.881687	1.881687	1.818932
NCI-H358	1.786631	1.652141	1.786631	1.652141	1.652141	1.548313	1.507089	1.548313	1.535903	1.535903	1.535903
1321N	1.647857	1.647857	1.647857	1.734085	1.785668	1.785668	1.734085	1.647857	1.520919	1.520919	1.413247
KYSE-510	1.717203	1.717203	1.904876	1.598059	1.598059	1.598059	1.598059	1.598059	1.582295	1.582295	1.582295
TCO-1	1.365812	1.365812	1.717426	1.717426	1.717426	1.34771	1.143509	1.151322	1.151322	1.151322	1.432718
NCI-H441	1.776251	1.776251	1.844426	1.844426	1.844426	1.692289	1.692289	1.692289	1.525735	1.525735	1.525735
SNU-398	1.528998	1.528998	1.593063	1.644383	1.656516	1.644383	1.644383	1.593063	1.546896	1.546896	1.544988
NCI-H1299	1.50363	1.455492	1.50363	1.515344	1.515344	1.50363	1.50363	1.515344	1.50363	1.50363	1.515344
ASH-3	1.3333496	1.2063767	1.3333496	1.3333496	1.3502046	1.3333496	0.968895	0.968895	0.6361593	0.6361593	0.9018576
MCF7	1.150625	1.131194	1.131194	1.131194	1.131194	1.111632	1.111632	1.111632	1.111632	1.111632	1.111632
SKOV3	0.7176477	0.7347421	0.7764232	0.7347421	0.8286913	0.7347421	0.7176477	0.7347421	0.7347421	0.7293017	0.7347421
ChaGo-K-1	0.7397704	0.7397704	1.2019286	1.2019286	1.3197395	1.3197395	0.7397704	0.7397704	0.7397704	0.7397704	1.1272925
A373 C6	1.0501332	1.0501332	1.0501332	0.869123	0.869123	0.7096039	0.7096039	0.8517779	0.8517779	0.7096039	0.8517779

ProbeSetID	SNP_A-208295	SNP_A-198999	SNP_A-198999	SNP_A-198999	SNP_A-209933	SNP_A-430318	SNP_A-424631	SNP_A-427328	SNP_A-218384	SNP_A-425218	SNP_A-223415
Affy.SNP.ID	6417094	2569063	2569064	2569066	6658895	9748516	10659347	5794037	7934783	11486260	8763615
dbSNP	rs4947984	rs2075109	rs2075110	rs11506105	rs11536635	rs7781264	rs2241055	rs6968014	rs845558	rs845559	rs7795728
Chromosome	7	7	7	7	7	7	7	7	7	7	7
Position	54985525	54993112	54993368	54994386	55003224	55005049	55014670	55021651	55021797	55022169	55024300
Strand	+	+	-	-	-	-	+	+	-	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
MES-SA	0.6483078	0.7352789	0.7352789	0.7352789	0.7512244	0.7512244	0.7512244	0.7512244	0.7512244	0.7512244	0.7540253
OAW28	0.7357038	0.727912	0.7357038	0.7357038	0.8646361	0.7357038	0.727912	0.727912	0.4894576	0.4894576	0.4894576
NCI-H460	0.6484461	0.7269926	0.8016441	0.8016441	0.8345223	0.8016441	0.4878499	0.6580405	0.4878499	0.4878499	0.6580405

ProbeSetID	SNP_A-1942958	SNP_A-2108798	SNP_A-2278627	SNP_A-1932698	SNP_A-1789168	SNP_A-2267510	SNP_A-1894491
Affy.SNP.ID	12448814	6811344	9418599	12306602	10155780	9252596	11748262
dbSNP	rs7795743	rs6593210	rs6970262	rs17518446	rs2293348	rs10269512	rs2293347
Chromosome	7	7	7	7	7	7	7
Position	55024339	55028395	55033972	55037176	55040966	55041369	55043125
Strand	-	-	+	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
KYSE-520	6.541713	6.541713	6.750218	6.750218	6.541713	6.541713	6.411673
OE21	6.425512	6.425512	6.548302	6.425512	6.392043	6.298607	6.219647
H3255	6.111655	6.111655	6.135911	6.135911	6.135911	6.135911	6.111655
NCI-H1573	5.705464	5.928978	5.937855	5.937855	6.13795	6.13795	5.928978
HSC-2	5.273699	5.472184	5.472184	5.472184	5.472184	5.472184	5.273699
HCC-827	5.01294	5.040266	5.33513	5.33513	5.33513	5.33513	5.093873
Hs 852.T	5.603882	5.608417	5.608417	5.608417	5.76554	5.603882	5.513761
KYSE-30	5.028042	5.055517	5.166432	5.166432	5.166432	5.166432	4.953463
A431	4.322911	4.322911	4.362625	4.410616	4.410616	4.410616	4.362625
BT-20	4.849844	5.006697	5.150625	5.150625	5.150625	5.006697	4.849844
Ca9-22	3.862511	3.939181	4.317663	4.317663	4.464566	4.464566	4.464566
NCI-H1568	4.168339	4.168339	4.112106	4.112106	4.044856	3.859537	3.84486
PC-9	4.307727	4.307727	4.307727	4.307727	4.456492	4.307727	4.206566
PNT1A	3.877783	3.877783	3.911622	3.911622	3.971824	3.971824	3.877783
LUDLU-1	3.88201	3.88201	3.86236	3.86236	3.936454	3.893806	3.86236
RERF-LC-MA	3.655638	3.775051	3.904667	3.904667	4.161315	4.161315	3.775051
U373 MG	3.184345	3.089149	2.932795	2.932795	2.932795	3.116349	3.116349
KYSE-450	3.971591	4.009916	4.009916	4.009916	4.203999	4.203999	4.115517
KOSC-2 cl3-43	3.223591	3.266909	3.266909	3.266909	3.223591	3.223591	3.266909
LU65	3.428992	3.428992	3.507654	3.507654	3.428992	3.398481	3.388789
PC-14	3.476991	3.476991	3.476991	3.476991	3.522971	3.476991	3.388218
NCI-H596	3.477121	3.631725	3.820114	3.820114	3.828509	3.820114	3.568589
PC-3 [JPC-3]	3.647499	3.655638	3.711979	3.711979	3.711979	3.711979	3.711979
647-V	3.070951	3.113251	3.348126	3.348126	3.348126	3.348126	3.644922
HSC-3	3.026608	3.083277	3.299672	3.299672	3.299672	3.299672	3.299672
CASKI	3.593694	3.593694	3.590495	3.593694	3.700607	3.593694	3.590495
JHH-4	3.023907	3.11823	3.11823	3.11823	3.11823	3.11823	3.11823
CAL-85-1	2.800803	3.170723	3.170723	3.170723	3.170723	3.170723	3.248024
BT-549	3.097396	3.097396	3.183837	3.183837	3.183837	3.183837	3.097396
HSC-4	2.977606	3.347547	3.347547	3.347547	3.347547	3.347547	3.359685
NUGC-4	3.101801	3.319849	3.479696	3.479696	3.479696	3.479696	3.479696
BICR10	3.18014	3.367842	3.436105	3.436105	3.492273	3.50429	3.492273
KYSE-140	3.296535	3.34572	3.52114	3.52114	3.525412	3.525412	3.52114
SNU-5	3.120892	3.176976	3.238084	3.238084	3.293975	3.293975	3.238084
SK-MEL-3	3.346054	3.346054	3.346054	3.346054	3.41406	3.346054	3.302025
KYSE-180	3.298136	3.344869	3.298136	3.298136	3.475356	3.475356	3.336038
PNT2	3.189804	3.189804	3.234785	3.321656	3.4784	3.4784	3.190368
G-361	3.073389	3.073389	3.116153	3.116153	3.116153	3.08182	2.947491
KYSE-220	3.291717	3.336992	3.336992	3.397895	3.397895	3.336992	3.283453
WM-115	3.015894	3.015894	3.077434	3.077434	3.077434	3.098836	3.077434
KYSE-50	3.03857	3.066828	3.066828	3.266165	3.266165	3.066828	3.03857
OSC-20	2.432376	3.051196	3.051196	3.051196	3.051196	3.082574	3.082574

ProbeSetID	SNP_A-1942958	SNP_A-2108798	SNP_A-2278625	SNP_A-1932698	SNP_A-1789168	SNP_A-2267510	SNP_A-1894491
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dbSNP	rs7795743	rs6593210	rs6970262	rs17518446	rs2293348	rs10269512	rs2293347
Chromosome	7	7	7	7	7	7	7
Position	55024339	55028395	55033972	55037176	55040966	55041369	55043125
Strand	-	-	+	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
SAT	2.614624	2.676061	2.676061	2.676061	2.676061	2.798889	2.798889
NCI-H1650	2.728585	2.793338	2.793338	2.793338	2.793338	3.033402	3.033402
SK-CO-1	3.013955	3.013955	3.013955	3.013955	3.013955	3.095936	3.013955
NCI-H1792	3.009861	3.009861	3.009861	3.009861	3.009861	3.234917	3.234917
NCI-H1703	2.893624	2.893624	2.986434	2.986434	3.094199	3.094199	2.986434
PCI 4B	2.971565	2.971565	3.107213	3.15257	3.178233	3.178233	3.15257
NCI-H1563	2.555684	2.846403	2.87226	3.181664	3.181664	3.181664	3.181664
NCI-H841	2.753814	2.79542	2.882085	2.882085	2.882085	2.922523	2.922523
LOU-NH91	2.897802	2.98114	2.99139	2.99139	2.99139	2.99139	2.98114
HuH-7	2.379725	3.270729	2.731956	2.731956	2.379725	2.416123	2.731956
LC-1 sq	2.446536	2.565947	2.565947	2.565947	2.565947	2.565947	2.565947
NCI-H522	2.78193	2.801644	2.802037	2.814618	2.814618	2.915157	2.915157
201T	2.609267	2.764925	2.847092	2.915693	3.02501	3.2021	3.2021
G-292 Clone A141B1	2.868771	2.989528	2.989528	2.989528	3.022632	3.022632	3.022632
RPMI-7951	2.878353	2.878353	2.878353	2.878353	3.127878	3.203298	3.203298
BEN	2.876586	2.918863	2.970854	2.970854	2.970854	2.970854	2.918863
NCI-H2228	2.613528	2.674109	2.674109	2.925508	3.175354	2.674109	2.627571
HCC-44	2.455113	2.663089	2.882818	2.882818	2.663089	2.663089	2.455113
NCI-H1734	3.045608	3.045608	3.058822	3.058822	2.75287	2.696374	2.693619
YKG-1	2.431245	2.590012	2.590012	2.590012	2.590012	2.769647	2.769647
MOG-G-CCM	2.8508	2.954609	2.971719	3.053239	3.053239	3.12277	3.053239
SK-BR-3	2.757023	2.822852	2.855212	2.855212	2.855212	2.855212	2.822852
COLO 858	2.916039	2.916039	2.916039	2.916039	2.996389	2.996389	2.916039
SCC-4	2.185984	2.435504	2.435504	2.435504	2.435504	2.594535	2.697983
TE7	2.498442	2.703754	2.703754	2.703754	2.703754	3.168338	3.168338
NCI-H1755	2.623436	2.623436	2.623436	2.778252	2.778252	3.076361	2.778252
CaR-1	2.724468	2.967637	2.967637	2.967637	2.967637	2.967637	2.967637
OVISE	2.530334	2.714789	2.714789	2.714789	2.714789	2.714789	2.714789
Calu-3	2.741489	2.741489	2.923078	2.923078	2.923078	2.923078	2.72247
NCI-H1623	2.667981	2.766131	2.667981	2.766131	2.598267	2.766131	2.57433
BICR 78	2.756535	2.782249	2.852669	2.950242	2.950242	3.069826	2.950242
LS 123	2.324575	2.604931	2.604931	2.689975	2.689975	2.689975	2.566137
AN3CA	2.673527	2.673527	2.673527	2.673527	2.673527	2.673527	2.673527
639-V	1.990462	2.53712	2.530298	2.530298	2.530298	2.530298	2.530298
WM 266-4	2.480336	2.785745	2.785745	2.785745	2.480336	2.785745	2.480336
SKN-3	2.250244	2.532985	2.744946	2.744946	2.250244	2.642794	2.642794
NCI-H2085	2.79704	2.850295	2.850295	2.903862	2.903862	2.903862	2.850295
RKN	2.063101	2.461623	2.461623	2.461623	2.461623	2.461623	2.782106
MKN45	2.86907	2.86907	2.86907	2.86907	2.86907	2.902026	2.902026
KHOS/NP	2.589447	2.651613	2.748688	2.771481	2.748688	2.772573	2.651613
SNU-16	2.766308	2.780158	2.780158	2.848673	2.977904	2.977904	2.766308
HCE7	2.292985	2.394863	2.448525	2.448525	2.448525	2.551707	2.630124
OCUG-1	2.314269	2.952639	2.952639	2.952639	2.952639	2.952639	2.952639
NL20-TA	2.601934	2.66247	2.668004	2.681307	2.758765	2.78248	2.78248
KYSE-70	2.746283	2.806601	2.806601	2.806601	2.983142	2.983142	2.826907
MOG-G-UVW	2.598888	2.734518	2.734518	2.734518	2.734518	2.734518	2.734518
NCI-H2342	2.847511	2.847511	2.832199	2.832199	2.832199	2.832199	2.832199

ProbeSetID	SNP_A-1942958	SNP_A-2108798	SNP_A-2278625	SNP_A-1932698	SNP_A-1789168	SNP_A-2267510	SNP_A-1894491
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dbSNP	rs7795743	rs6593210	rs6970262	rs17518446	rs2293348	rs10269512	rs2293347
Chromosome	7	7	7	7	7	7	7
Position	55024339	55028395	55033972	55037176	55040966	55041369	55043125
Strand	-	-	+	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
ABC-1	2.723537	2.939772	2.96053	2.96053	3.062139	3.062139	3.062139
NCI-H650	2.58833	2.58833	2.58833	2.58833	2.58833	2.611007	2.611007
FTC 133	2.548702	2.569538	2.939084	2.943852	2.990019	2.990019	2.943852
MDST8	2.636425	2.636425	2.741869	2.874178	2.9639	2.9639	2.894767
KYSE-150	2.762855	2.830133	2.879597	2.879597	3.252738	3.252738	3.03435
SW 48	2.67219	2.67219	2.759752	2.759752	2.851531	2.868931	2.868931
NCI-H2023	2.564334	2.564334	2.564334	2.766441	2.85108	2.916501	2.916501
COLO-680N	2.640357	2.663319	2.663319	2.663319	2.663319	2.750802	2.663319
KP-4	2.048398	2.328633	2.328633	2.328633	2.328633	2.739983	2.739983
SW-1710	2.437476	2.730691	2.730691	2.730691	2.705348	2.705348	2.705348
KATO III	1.878288	1.890556	1.946576	2.307987	2.751491	2.769072	2.769072
RERF-GC-1B	2.825715	2.825715	2.866517	2.866517	2.956567	2.956567	2.825715
CAL-29	2.141685	2.706624	2.706624	2.706624	2.706624	2.706624	2.706624
Sci-1	2.54205	2.544978	2.544978	2.544978	2.544978	2.544978	2.544978
NCI-H810	2.472446	2.538342	2.653592	2.653592	2.653592	2.653592	2.59087
T.Tn	2.523736	2.667878	2.667878	2.667878	2.779827	2.667878	2.523736
DOK	2.655651	2.655651	2.6613	2.664442	2.781798	2.793481	2.781798
COLO 741	2.609221	2.680281	2.680281	2.773798	2.807114	3.018834	3.018834
NCI-H1651	2.354866	2.575299	2.805717	2.805717	2.575299	2.692359	2.692359
RWPE-1	2.536462	2.768929	2.865427	2.865427	2.972986	2.972986	2.768929
KYSE-270	2.448476	2.496037	2.542469	2.542469	2.542469	2.542469	2.542469
NCI-H196	2.270952	2.303925	2.630657	2.748477	2.748477	2.748477	2.308259
SK-LU-1	2.471319	2.531522	2.531522	2.720189	2.720189	2.531522	2.528556
NCI-H1693 (ship16)	2.479835	2.600704	2.600704	2.692768	2.692768	2.854966	2.854966
NCI-H2135	2.630657	2.630657	2.609828	2.609828	2.609828	2.585234	2.503596
VMRC-RCW	1.884856	2.27149	2.27149	2.35285	2.35285	2.488398	2.488398
RERF-LC-Sq1	2.480897	2.759641	2.590541	2.590541	2.480897	2.480897	2.31313
A375.S2	2.531246	2.531246	2.531246	2.498988	2.498988	2.498988	2.432239
SW 1116	2.529397	2.529397	2.529397	2.714676	2.714676	2.729671	2.714676
CAL-148	1.642784	2.092339	2.092339	2.092339	2.092339	2.156523	2.289268
22RV1	2.588024	2.727004	2.790992	2.848651	2.886184	2.896206	2.896206
IM-95	2.74111	2.74111	2.74111	2.761736	2.761736	2.67307	2.620052
HT 115	2.547736	2.547736	2.693744	2.700315	3.018601	3.018601	2.700315
NCI-H727	2.339427	2.547115	2.547115	2.626979	2.626979	2.676438	2.676438
T.T	2.718233	2.763563	2.718233	2.718233	2.718233	2.640253	2.516519
OE33	2.559775	2.644722	2.672932	2.672932	2.672932	2.672932	2.644722
PLC/PRF/5	2.42185	2.407918	2.70834	2.711645	2.70834	2.70834	2.70834
SAS	2.345589	2.432885	2.432885	2.432885	2.432885	2.711081	2.850704
MKN74	2.623959	2.652753	2.690032	2.706985	2.852733	2.852733	2.706985
HMCB	2.776699	2.776699	2.610774	2.610774	2.514927	2.610774	2.514927
A2058	2.016357	2.32992	2.016357	2.016357	2.016357	2.142441	2.142441
HCC-366	2.205961	2.270019	2.270019	2.270019	2.270019	2.270019	2.270019
P31/FUJ	2.434213	2.465395	2.465395	2.465395	2.465395	2.800507	2.800507
SW 1417	2.448328	2.448328	2.448328	2.448328	2.328401	2.468635	2.328401
KMRC-1	2.233142	2.365791	2.356371	2.356371	2.336617	2.336617	2.336617
OAW42	2.346319	2.346319	2.540041	2.540041	2.761082	2.761082	2.688964
PE/CA-PJ15	2.35957	2.436756	2.436756	2.436756	2.35957	2.35957	2.295291

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dbSNP	rs7795743	rs6593210	rs6970262	rs17518446	rs2293348	rs10269512	rs2293347
Chromosome	7	7	7	7	7	7	7
Position	55024339	55028395	55033972	55037176	55040966	55041369	55043125
Strand	-	-	+	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
NUGC-3	2.560321	2.581828	2.581828	2.712795	2.791442	2.935665	2.935665
DMS 273	2.353233	2.441353	2.559299	2.559299	2.646371	2.68338	2.68338
MKN1	2.503256	2.510723	2.510723	2.510723	2.510723	2.619563	2.619563
RERF-LC-Ad2	2.522424	2.645783	2.645783	2.778418	2.778418	2.883594	2.883594
TASK1	1.985566	2.051985	2.051985	2.051985	2.051985	2.110627	2.110627
NCI-H23	2.414686	2.426184	2.414686	2.414686	2.382569	2.414686	2.414686
KP-1N	2.062621	2.258487	2.258487	2.080368	2.080368	2.258487	2.43662
NCI-H711	2.329148	2.403418	2.583997	2.583997	2.583997	2.619819	2.619819
NCI-H661	2.382443	2.587741	2.541799	2.541799	2.382443	2.4056	2.382443
EBC-1	2.465322	2.609022	2.655375	2.655375	2.655375	2.609022	2.465322
JHH-6	2.054832	2.135456	2.135456	2.135456	2.135456	2.135456	2.33311
NB69	2.335551	2.396158	2.589271	2.589271	2.589271	2.396158	2.448982
SR	2.462951	2.50236	2.50236	2.616104	2.50236	2.550419	2.50236
T84	2.566326	2.566326	2.613213	2.708385	2.708385	2.708385	2.732359
HO-1-N-1	1.712729	1.8454	1.8454	1.8454	1.8454	1.8454	1.978816
BALL-1	2.368865	2.602836	2.602836	2.602836	2.602836	2.602836	2.602836
Nthy-ori 3-1	2.493947	2.530838	2.530838	2.530838	2.530838	2.558741	2.558741
FTC 238	2.489348	2.508995	2.51256	2.51256	2.55635	2.610015	2.672418
HuCCT1	2.31344	2.437847	2.437847	2.437847	2.31344	2.31344	2.437847
HOS	2.318935	2.318935	2.349989	2.395869	2.585245	2.685439	2.585245
NCI-H2170	2.446388	2.446388	2.446388	2.446388	2.446388	2.517785	2.517785
Hs 936.T	2.484876	2.484876	2.518533	2.532458	2.567737	2.682047	2.567737
NCI-H1793	2.511014	2.519979	2.519979	2.691372	2.691372	2.556612	2.556612
MRK-nu-1	2.286447	2.413086	2.413086	2.413086	2.413086	2.413086	2.475091
KHOS-312H	2.239629	2.277249	2.390015	2.45728	2.45728	2.597364	2.597364
LCLC-103H	2.222316	2.265022	2.327538	2.327538	2.327538	2.327538	2.327538
H69V	2.431481	2.43786	2.431481	2.430786	2.430786	2.41046	2.310255
GCT	2.236108	2.236108	2.236108	2.269506	2.269506	2.320782	2.320782
COLO 857	2.333804	2.354369	2.375079	2.375079	2.413311	2.413311	2.375079
RERF-LC-MS	2.365385	2.514021	2.655375	2.723537	2.689657	2.689657	2.689657
MKN7	2.403268	2.573762	2.585563	2.585563	2.573762	2.248656	2.195222
A-427	1.762107	2.207949	2.207949	1.908824	1.908824	2.007866	2.059711
T47D	2.602215	2.66364	2.66364	2.679381	2.697972	2.697972	2.697972
BT-483	2.488228	2.5445	2.488228	2.5445	2.5445	2.634654	2.634654
SKN	1.780613	1.822156	1.822156	1.822156	1.822156	2.038009	2.038009
HCC202	2.346588	2.353884	2.445795	2.649817	2.703087	2.7065	2.703087
OCUM-1	2.44066	2.501923	2.538066	2.538066	2.538066	2.538066	2.538066
NCI-H2122	2.22956	2.22956	2.22956	2.22956	2.22956	2.232357	2.22956
HT 29	2.408332	2.408332	2.406803	2.3897	2.3897	2.341542	2.341542
huH-1	2.066148	2.243304	2.243304	2.243304	2.243304	2.243304	2.358245
HCC-15	2.304055	2.53157	2.53157	2.53157	2.506042	2.506042	2.304055
KHOS-240S	2.201928	2.226694	2.270544	2.360932	2.360932	2.430102	2.430102
JIMT-1	2.19723	2.219697	2.219697	2.219697	2.219697	2.219697	2.219697
KON	1.85467	2.509611	2.509611	2.230133	2.230133	2.230133	2.230133
COLO 853	2.298154	2.401611	2.487923	2.515252	2.52548	2.52548	2.487923
U-2 OS	2.261862	2.314087	2.365385	2.365385	2.365385	2.314087	2.248656
ME-180	1.900086	2.093513	2.093513	2.093513	2.093513	2.132771	2.254884

ProbeSetID	SNP_A-1942958	SNP_A-2108798	SNP_A-2278625	SNP_A-1932698	SNP_A-1789168	SNP_A-2267510	SNP_A-1894491
Affy_SNP.ID	12448814	6811344	9418599	12306602	10155780	9252596	11748262
dbSNP	rs7795743	rs6593210	rs6970262	rs17518446	rs2293348	rs10269512	rs2293347
Chromosome	7	7	7	7	7	7	7
Position	55024339	55028395	55033972	55037176	55040966	55041369	55043125
Strand	-	-	+	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
FU97	2.052679	2.159773	2.254738	2.254738	2.052679	2.052679	2.254738
Calu-1	2.170006	2.170006	2.170006	2.170006	2.170006	2.284199	2.284199
NCI-H2452	1.940146	2.212134	2.212134	2.254633	2.254633	2.254633	2.212134
UMC-11	2.126688	2.096765	2.126688	2.096765	1.960387	1.960387	1.960387
COLO 684	2.212563	2.378156	2.378156	2.378156	2.41071	2.41071	2.41071
SVts-8	0.9501698	0.9501698	0.9501698	1.0117819	1.0117819	1.0117819	1.0919367
HEC-1	2.320859	2.322098	2.322098	2.322098	2.322098	2.322098	2.389813
A-375	1.832082	1.915903	1.915903	1.915903	1.915903	1.915903	2.216273
OUMS-23	1.685491	1.931901	1.931901	1.931901	1.931901	2.110794	2.110794
EJ138	2.164656	2.211356	2.268455	2.313764	2.405425	2.405425	2.382873
HO-1-u-1	1.689687	1.920771	1.920771	1.920771	1.920771	1.967545	1.967545
HT55	2.243901	2.243901	2.243901	2.243901	2.272868	2.329199	2.272868
NCI-H2347	2.001399	2.388061	2.388061	2.388061	2.388061	2.388061	2.115455
REFL-LC-KJ	2.205061	2.262099	2.306975	2.306975	2.262099	2.262099	2.262099
COR-L279	2.014118	2.014118	2.014118	2.136998	2.136998	2.136998	2.136998
SW 1573	1.883685	1.891665	1.891665	1.891665	1.891665	2.043755	2.221568
SW 900	2.034048	2.281348	2.281348	2.281348	2.281348	2.281348	2.281348
HT 1376	2.059429	2.118046	2.159418	2.241633	2.441143	2.441143	2.441143
HUP-T3	1.539662	2.121769	2.121769	1.933909	1.933909	1.933909	2.121769
STC 1	1.850844	1.970534	2.017892	2.273407	2.273407	2.273407	2.051163
NCI-H2286	2.040044	2.456837	2.456837	2.456837	2.456837	2.373318	2.245054
VMRC-LCD	2.085289	2.127006	2.181707	2.378068	2.378068	2.378068	2.378068
H69AR	2.260438	2.260438	2.260438	2.338349	2.338349	2.385449	2.246644
VMRC-LCP	1.831011	2.033007	2.033007	1.831011	1.831011	2.033007	2.031366
FLYA13	1.931325	2.023249	2.023249	2.023249	2.023249	2.023249	2.182086
HUP-T4	1.972064	2.183373	2.183373	2.183373	2.183373	2.183373	2.462853
COR-L 105	2.132468	2.251725	2.320575	2.320575	2.350232	2.320575	2.251725
KMRC-20	2.16531	2.333379	2.350449	2.350449	2.393617	2.393617	2.393617
NCI-H2081	2.092801	2.092801	2.211008	2.397692	2.397692	2.463577	2.463577
VM-CUB1	2.165746	2.198927	2.035944	2.198927	2.354739	2.354739	2.532578
BU25 TK	2.282905	2.29009	2.29009	2.29009	2.282905	2.282905	2.217036
RCM-1	1.842065	1.848869	1.848869	1.848869	1.848869	2.161261	2.509091
NCI-H2172	2.230693	2.293714	2.293714	2.333971	2.333971	2.333971	2.293714
SK-NEP-1	1.941163	2.244683	2.301574	2.301574	2.301574	2.378979	2.378979
D1.1	2.279843	2.279843	2.260794	2.279843	2.260794	2.260794	2.19475
NCI-H2196	2.140394	2.244378	2.244378	2.193374	2.193374	2.413574	2.431134
NCI-H1435	1.898427	1.917121	2.009822	2.092577	2.092577	2.092577	2.092577
GRANTA-519	1.804641	1.804641	1.864773	1.649764	1.649764	1.813981	1.944738
SK-MES	2.306703	2.336886	2.370236	2.363401	2.363401	2.363401	2.363401
NCI-H1437	2.248458	2.250628	2.248458	2.250628	2.250628	2.263218	2.26468
BFTC-905	2.133832	2.243861	2.133832	2.133832	1.919496	1.919496	2.021316
DMS 153	2.258751	2.258751	2.258751	2.108197	2.108197	2.108197	2.108197
Detroit 562	2.108572	2.108572	2.209197	2.284552	2.28548	2.28548	2.28548
RT4	1.959495	2.047064	2.047064	2.069854	2.069854	2.069854	2.047064
273T	2.118821	2.118821	2.118821	2.118821	2.238925	2.280484	2.280484
SW 1463	2.148696	2.173568	2.213447	2.213447	2.330885	2.354164	2.354164
H69	2.156017	2.156017	2.328916	2.328916	2.156017	2.156017	2.156017

ProbeSetID	SNP_A-1942958	SNP_A-2108798	SNP_A-2278627	SNP_A-1932698	SNP_A-1789168	SNP_A-2267510	SNP_A-1894491
Affy_SNP.ID	12448814	6811344	9418599	12306602	10155780	9252596	11748262
dbSNP	rs7795743	rs6593210	rs6970262	rs17518446	rs2293348	rs10269512	rs2293347
Chromosome	7	7	7	7	7	7	7
Position	55024339	55028395	55033972	55037176	55040966	55041369	55043125
Strand	-	-	+	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
Hs 195.T	1.799016	1.956948	2.155443	2.236919	2.236919	2.236919	2.112477
NCI-H1781	2.171801	2.247319	2.247319	2.163688	2.163688	2.126868	2.026067
WDR	2.162829	2.172046	2.223745	2.223745	2.223745	2.223745	2.162829
SK-MEL-1	2.062084	1.906158	1.906158	1.906158	1.906158	1.906158	1.906158
SUIT-2	1.838055	2.085219	2.085219	2.085219	2.085219	2.085219	2.085219
WPE1-NB26	1.761136	2.073752	2.203354	2.203354	2.203354	2.203354	2.073752
HCC-78	1.867009	2.201821	2.201821	2.201821	2.201821	2.201821	2.184347
A549	2.035032	2.035032	2.035032	2.085975	2.161206	2.263679	2.263679
LK-2	1.978147	2.127061	2.127586	2.127586	2.127061	2.127061	2.127061
NCI-H2330	2.156509	2.224546	2.224546	2.224546	2.311498	2.311498	1.897919
COLO 685	2.019726	2.019726	2.019726	2.072571	2.239868	2.251765	2.251765
BICR22	2.184063	2.289111	2.289111	2.210753	2.210753	2.303224	2.308816
MEWO	2.036371	2.036371	2.020271	2.020271	1.94025	2.020271	2.036371
IHH-4	2.100181	2.182411	2.185944	2.185944	2.185944	2.185944	2.376181
NCI-N87	2.247822	2.247822	2.247822	2.344745	2.436657	2.453313	2.453313
KYSE-410	2.134727	2.148257	2.151736	2.151736	2.148257	2.151736	2.151736
DMS 92	1.636081	1.656093	1.656093	1.702063	1.702063	1.859567	2.315006
RO82-W-1	2.004411	2.010139	2.010139	2.010139	2.004411	2.004411	1.691068
NCI-H82	1.966903	1.966903	1.966903	1.991649	1.991649	2.046809	2.046809
C32	1.797686	1.884015	1.884015	1.929522	1.929522	1.963939	1.963939
Hs 633T	2.104526	2.124078	2.104526	2.104526	2.104526	2.104526	2.07167
NCI-H2405	1.937671	1.937877	2.129959	2.161384	2.161384	2.269822	2.269822
Hs 895.T	2.092898	2.200825	2.200825	2.225307	2.225307	2.243079	2.225307
MG 63	2.179905	2.184537	2.184537	2.184537	2.226801	2.226801	2.238885
CAL-120	1.892219	1.994527	2.080522	2.080522	1.892219	1.892219	2.080522
H33HJ-JA1	2.006183	2.006183	2.006183	2.006183	1.885757	1.943267	1.943267
JSC-1	1.990434	2.008499	2.078136	2.086395	2.165337	2.241646	2.280458
Hs 751.T	1.794977	1.893087	1.893087	1.893087	1.893087	1.794977	1.794977
HMVII	1.72518	1.72518	1.72518	1.775175	1.775175	1.836803	1.836803
fR2	1.664549	1.854655	1.854655	1.854655	1.854655	1.854655	1.854655
COLO 201	1.962785	1.962785	2.076465	2.076465	2.092619	2.092619	2.092619
NCI-H1915	1.828272	1.90729	1.90729	2.014764	2.014764	2.045176	2.163266
IMR-90	1.459358	1.615237	1.615237	1.615237	1.615237	2.043201	2.080901
RT112/84	1.749311	1.808945	1.808945	1.808945	1.808945	1.808945	1.808945
Hs 894(E).Lu	2.077378	2.077378	2.077378	2.077378	2.177572	2.177572	2.043826
LNCaP clone FGC	2.051674	2.185957	2.185957	2.216045	2.256178	2.318858	2.256178
NCI-H1581	2.002797	2.002797	2.054393	2.054393	2.099498	2.129283	2.129283
HCT 116	2.071572	2.071572	2.160074	2.160074	2.160074	2.160074	2.160074
Hs 228.T	1.944797	2.134038	2.214572	2.214572	2.134038	2.134038	2.003403
JURKAT EC-1	2.000563	2.000563	2.000563	1.89412	1.89412	1.934647	1.934647
SK-N-BE(2)	2.096877	2.120275	2.120275	2.125418	2.153501	2.153501	2.125418
LS180	2.029767	2.029767	2.093401	2.062282	2.020786	2.062282	2.050625
CAL-51	1.772086	1.819224	2.092144	2.092144	2.092144	2.092144	2.181165
SKG-IIIb	1.21404	1.295495	1.295495	1.295495	1.295495	1.797639	1.82036
P116 T	1.870482	2.025981	2.025981	2.02621	2.059174	2.059174	2.106306
Hs 417.Lu	1.952229	1.952229	1.939734	1.952229	1.952229	1.963369	1.939734
WIL2 NS	2.081925	2.081925	2.081925	2.086241	2.152639	2.209197	2.209197

ProbeSetID	SNP_A-1942958	SNP_A-2108798	SNP_A-2278627	SNP_A-1932698	SNP_A-1789168	SNP_A-2267510	SNP_A-1894491
Affy_SNP.ID	12448814	6811344	9418599	12306602	10155780	9252596	11748262
dbSNP	rs7795743	rs6593210	rs6970262	rs17518446	rs2293348	rs10269512	rs2293347
Chromosome	7	7	7	7	7	7	7
Position	55024339	55028395	55033972	55037176	55040966	55041369	55043125
Strand	-	-	+	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
COR-L95	2.037368	2.037368	2.037368	2.091655	2.091655	2.091655	2.037368
SK-N-FI	1.756132	1.774692	2.083187	2.090536	2.175944	2.175944	2.175944
WILCL	1.935119	1.935119	1.970666	2.204806	2.204806	2.204806	2.204806
P32/ISH	1.986001	2.003403	2.003403	2.021746	2.021746	2.021746	2.399916
EPLC-272H	1.610255	1.898846	1.898846	1.784502	1.784502	1.898846	2.011691
A2780 ADR	1.744349	1.744349	1.857885	2.0519	2.0909	2.0909	2.0519
C8166	1.97582	2.057591	2.057591	2.063778	2.283376	2.283376	2.283376
Hs 908.T	1.973244	1.973244	1.973244	1.988754	1.988754	2.344873	2.337746
NCI-H322	1.846024	1.846024	1.846024	1.984087	1.846024	1.822739	1.846024
NCI-H1417	1.652987	1.597926	1.597926	1.652987	1.652987	1.742315	1.742315
HBE4-E6/E7	1.805521	1.891021	1.891021	1.891021	1.891021	2.031751	2.100293
NCI-H1522	1.67921	1.7155	1.7155	1.7155	2.170428	2.170428	2.170428
LU99A	2.057138	2.099262	2.099262	2.168129	2.057138	2.057138	1.882213
HCT-8	1.997763	1.997763	1.949998	1.949998	1.949998	1.949998	2.033863
SBC-5	1.961557	2.052538	2.052538	2.052538	2.10376	2.10376	2.10376
Ishikawa	1.972341	1.974802	1.972341	1.974802	1.974802	1.974802	1.974802
U266B1	2.049136	2.049136	2.014377	2.014377	2.014377	2.014377	1.966465
PA-TU-8988T	1.651734	1.942208	1.942208	1.942208	1.942208	1.942208	1.942208
HRT-18	1.676177	1.738448	1.738448	1.738448	1.738448	1.738448	1.738448
C170	1.932418	1.965516	2.000058	2.000058	2.000058	2.132192	2.132192
J GAMMA 1	2.044622	2.044622	2.044622	2.044622	1.996853	1.964027	1.996853
LS174T	1.974656	1.974656	1.94712	1.94712	1.94712	1.974656	1.94712
NCI-H526	1.966202	2.021918	2.087039	2.089208	2.113777	2.113777	2.113777
HLE	1.617495	1.741637	1.814781	1.814781	1.741637	1.741637	1.814781
23132/87	1.61471	1.61471	1.867145	2.208673	2.208673	2.208673	2.208673
Hs 834.T	1.838878	1.991128	2.056473	2.056473	2.33311	2.33311	2.088522
Hs 891.T	1.810424	1.810424	1.921157	2.072458	2.072458	2.107099	2.072458
Hs 839.T	1.166692	1.443568	1.443568	1.325869	1.325869	1.443568	1.590819
SNU-1	2.031652	2.031652	2.057676	2.241009	2.057676	2.057676	2.278926
EB1	2.052226	2.164002	2.052226	2.164002	2.164002	2.164002	2.164002
NCI-H2009	1.78136	1.835337	1.835337	1.922239	1.922239	1.960197	1.960197
NCI-H1048	1.822202	1.822202	1.822202	1.994339	1.994339	2.135966	2.135966
A2780	1.937641	1.947913	1.972341	2.080214	2.080214	2.080214	1.947913
NCI-H1666	2.208673	2.099262	2.099262	2.208673	2.099262	2.168129	2.099262
G-402	1.718269	1.770852	1.770852	1.770852	1.770852	1.770852	2.045247
JHH-7	1.661971	1.661971	1.661971	1.871221	1.871221	2.021989	2.201579
OUS-11	1.688964	2.095105	2.095105	2.095105	2.095105	2.268784	2.268784
AZ-521	1.814335	1.814335	1.814335	1.953739	2.374965	2.374965	2.374965
K562/ADM	1.33344	1.33344	1.33344	1.33344	1.33344	2.15439	2.15439
NCI-H889	1.828073	1.910581	1.910581	1.972603	1.972603	1.972603	1.910581
OE19	1.797794	1.755159	1.755159	1.901713	2.072008	2.152995	2.19363
HT 728.T	2.055115	2.075397	2.075397	2.075397	2.142001	2.258276	2.258276
GA-10	1.98587	1.98587	1.967487	1.967487	1.967487	1.967487	1.98587
Hs 737.T	1.949191	1.949191	1.96947	1.96947	2.034404	2.034404	2.040343
NCI-H838	1.758925	1.882333	1.882333	1.91724	1.91724	1.91724	1.972938
NCI-H1155	1.929477	1.934957	1.929477	2.004397	2.128731	2.128731	2.128731
A2780cis	1.763768	1.782278	1.916349	2.204214	1.916349	2.204214	2.204214

ProbeSetID	SNP_A-1942958	SNP_A-2108798	SNP_A-2278625	SNP_A-1932695	SNP_A-1789168	SNP_A-2267510	SNP_A-1894491
Affy_SNP.ID	12448814	6811344	9418599	12306602	10155780	9252596	11748262
dbSNP	rs7795743	rs6593210	rs6970262	rs17518446	rs2293348	rs10269512	rs2293347
Chromosome	7	7	7	7	7	7	7
Position	55024339	55028395	55033972	55037176	55040966	55041369	55043125
Strand	-	-	+	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
G-401	1.593395	1.982084	1.982084	1.982084	1.831271	1.831271	1.982084
MS-1-L	1.638846	2.20244	2.20244	1.638846	1.573126	1.638846	1.779663
Hs 746T	1.682223	1.641836	1.641836	1.797345	1.797345	1.932551	1.797345
SW 13	1.948559	1.948559	1.957504	1.95828	1.903399	1.903399	1.903399
AGS	1.835276	1.837643	1.837643	2.154677	2.163116	2.192213	2.192213
NCI-H2171	1.578423	1.767414	1.89759	1.997936	2.104651	2.104651	2.096849
Hs 573.T	1.683012	1.809099	2.076535	2.076535	2.076535	2.083523	2.076535
HT1080	1.917626	1.917626	1.917626	1.917626	1.917626	1.992908	1.992908
COR-L 23/CPR	1.875111	1.875111	1.875111	1.98545	1.98545	2.075425	2.075425
HGC-27	1.601522	2.029767	1.982519	1.982519	1.601522	1.601522	1.63932
SBC-3	1.81675	1.924757	1.924757	1.924757	2.094979	2.094979	2.094979
COLO 792	1.677516	1.693749	1.693749	1.693749	1.693749	1.697789	1.69995
H-EMC-SS	1.502876	1.502876	1.502876	1.502876	1.502876	1.443603	2.027625
J82	1.433789	1.495497	1.495497	1.495497	1.495497	1.570666	1.616737
Hs 600.T	1.878814	1.963369	2.233222	2.067516	1.878814	1.830629	1.830629
Hs 781.T	1.699102	1.855626	1.855626	1.855626	1.699102	1.791955	1.699102
BPH-1	1.827414	1.880394	1.894345	2.029111	2.040087	2.128179	2.128179
COR-L23	1.637717	1.696891	1.696891	1.878047	1.878047	1.878047	1.696891
A673	1.916587	1.916587	1.916587	2.006701	2.21275	2.21275	2.21275
COLO 205	1.849826	1.849826	1.849826	1.849826	1.849826	1.849826	1.849826
RMG-I	1.312974	1.431506	1.431506	1.419534	1.419534	1.431506	1.745547
WI-26 VA4	1.8475	1.8475	1.817949	1.8475	1.892818	2.100488	2.11997
Eos-HL-60	1.740422	1.865891	1.740422	1.730191	1.640415	1.713764	1.730191
GP5D	1.824748	1.824748	1.85253	1.844487	1.824748	1.844487	1.85253
MES-SA/Dx-5	1.658839	1.614825	1.614825	1.658839	1.688964	1.8738	1.958807
SBC-2	1.845355	1.845355	1.845355	1.845355	1.845355	1.845355	1.831715
MCAS	1.229853	1.687759	1.687759	1.687759	1.467004	1.467004	1.687759
Ishikawa (Heraklio) 02 ER-	1.598722	1.621215	1.621215	1.746209	1.746209	1.767023	1.897202
MOLT-4F	1.328055	1.328055	1.328055	1.328055	1.328055	1.910596	1.960899
HARA	1.828838	1.839077	1.839077	1.828838	1.828838	1.828838	1.676806
NCI-H1944	1.557056	1.557056	1.557056	1.765161	1.765161	1.820729	2.029696
BICR 31	1.834161	1.834161	1.820084	1.820084	1.800455	1.800455	1.800455
D341 Med	1.86155	1.86155	1.707305	1.720146	1.720146	1.720146	1.720146
KHM-3S	1.593262	1.593262	1.593262	1.758423	1.758423	1.830353	1.830353
COR-L88	1.881687	1.818932	1.849674	1.881687	1.904488	1.904488	1.854564
NCI-H358	1.535903	1.618022	1.618022	1.810733	1.810733	1.900534	2.015568
1321N	1.413247	1.520919	1.520919	1.520919	1.520919	1.520919	1.426987
KYSE-510	1.582295	1.582295	1.582295	1.923602	1.923602	1.923602	1.923602
TCO-1	1.432718	1.432718	1.432718	1.432718	1.432718	1.503767	1.529881
NCI-H441	1.525735	1.692289	1.692289	1.692289	1.692289	1.525735	1.746209
SNU-398	1.544988	1.546896	1.546896	1.546896	1.544988	1.544988	1.544988
NCI-H1299	1.515344	1.610634	1.704797	1.752456	1.610634	1.610634	1.752456
ASH-3	0.9018576	0.9018576	0.9018576	0.9018576	0.9018576	0.9293574	1.0402108
MCF7	1.169925	1.227063	1.264536	1.598175	1.598175	1.598175	1.598175
SKOV3	0.7545363	0.7545363	0.7545363	0.9342876	0.8522504	0.9069193	0.9342876
ChaGo-K-1	1.1272925	0.9341206	0.9341206	0.9341206	0.9341206	1.1272925	1.220304
A373 C6	0.869123	0.869123	0.9306744	0.9306744	0.9306744	0.9306744	1.2244407

ProbeSetID	SNP_A-1942958	SNP_A-2108798	SNP_A-2278627	SNP_A-1932698	SNP_A-1789168	SNP_A-2267510	SNP_A-1894491
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Position	55024339	55028395	55033972	55037176	55040966	55041369	55043125
Strand	-	-	+	+	+	+	+
Cytoband	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2	p11.2
MES-SA	0.6719725	0.6719725	0.7540253	0.8882739	0.8882739	0.7540253	0.8882739
OAW28	0.4894576	0.4894576	0.4894576	0.8394229	0.8394229	0.8394229	0.8818619
NCI-H460	0.6580405	0.4669172	0.4669172	0.6580405	0.6580405	0.6580405	0.7038671

