

Supplementary Table 7: List of antibodies used in RPPA and normalized linear scores associated with Cre control and TRIM24-driven tumors.

Antibodies	Cre-1	Cre-2	Trim24COE-897.1	Trim24COE-897.2	Trim24COE-64.1	Trim24COE-64.2
N-Cadherin	0.95770565	0.96599156	1.013776543	1.000375075	1.009022537	1.039834126
B-Raf	0.93865789	0.95576823	1.004370039	1.005287521	1.018259421	1.022198541
NF-kB-p65_pS536	0.95314089	0.92805396	1.001754007	1.004232252	1.137117894	1.154919515
b-Catenin_pT41_S45	0.87332292	0.84090276	1.002564931	1.003448897	1.167114294	1.128178054
Elk1_pS383	0.95821458	0.93415014	1.059989743	1.091141023	0.996342409	0.965474792
MIF	0.94842029	0.9369278	1.054374118	1.05714489	0.993507045	0.968190064
Bak	0.94585477	0.96625489	1.048399927	1.112049507	0.951182948	0.972248109
Sox2	1.02089994	0.98114902	1.095423358	1.150353426	1	0.895960296
Merlin	0.92583504	0.91450189	1.173486123	1.191485695	0.970345282	0.990370813
43377	0.73937169	0.76362871	1.015100212	1.01611125	1.007722736	1
MDM2_pS166	0.74069458	0.72945429	1.000222744	1.005711457	1.060342506	1.003464052
Jagged1	0.78419102	0.80685208	1.004836182	1.00125486	1.049099815	1.056763221
YB1_pS102	0.80492401	0.8103776	1.005341674	1.000766553	1.103368337	1.083126009
PAICS	0.77533462	0.80223282	1.09604395	1.074497406	0.971477958	0.989286111
MMP2	0.70854228	0.69974975	1.049075424	1.052845499	0.998964175	0.962964071
EGFR_pY1173	0.72817686	0.73298452	1.00369186	1.002360279	1.131120488	1.102966984
4E-BP1_pS65	0.74190259	0.76565059	1.00539377	1.000716228	1.159419462	1.146934794
GCN5L2	0.60572109	0.59799206	1.128832609	1.111152632	1.005645494	0.956565738
IR-b	0.61533501	0.6179095	1.080004618	1.067711457	1.021049954	0.941813733
PDGFR-b	0.62428942	0.61669428	1.071547421	1.163867705	0.961924744	0.998434699
Raptor	0.63577728	0.61992393	1.018305676	1.010383143	1.004575074	1.017248347
eIF4G	0.61667181	0.62843111	1.024446798	1.0365794	0.998544691	0.994387118
AMPKa	0.5230509	0.49322317	1.019786421	1.071598536	1.009133727	0.960633925
eEF2	0.44266536	0.4406616	1.050583591	1.037856409	0.993179174	0.968504049
PKM2	0.43553005	0.4287731	1.003444839	1.002598902	1.080579535	0.997941694
PAK1	0.47139788	0.47791086	1.000170356	1.005762064	1.180133951	1.097836812
b-Actin	0.69406779	0.69469409	1.003592632	1.002456134	1.223329232	1.195718159
RSK	0.64772628	0.63442144	1.004425854	1.001651238	1.315556286	1.228071322
Mcl-1	0.78616194	0.80480197	1.002966441	1.003061036	1.236430512	1.184764446
A-Raf	0.75248921	0.75179688	1.004822744	1.001267841	1.284141458	1.21815923
Histone-H3	0.77820968	0.765203	1.002197122	1.003804202	1.627911794	1.615273309
p38_pT180_Y182	1.00895339	1.1069253	1.07129813	1.044458375	0.759116035	0.828368356
PAK4	1.02278883	1.04992974	1.057996429	1.051361107	0.760871935	0.809729321
IGFRb	1.02244761	1.01506334	1.058709344	1.075740122	0.832603272	0.868937342
PEA-15_pS116	1.07283672	1.11588341	1.009879291	1.020347367	0.869910747	0.877175558
Glutaminase	1.016186	1.02083732	1.17965329	1.173482699	0.826838791	0.849115622
Gys_pS641	0.98081555	1.05345319	1.191643591	1.245571264	0.675831924	0.633371381
Caveolin-1	1.04543624	0.99386503	1.338080589	1.248747682	0.578615008	0.553732926
c-Myc	0.92739218	0.98529259	1.369447929	1.429798911	0.99236179	0.953582509
DJ1	0.93786434	0.88057058	1.392180806	1.305276179	1.037174172	0.926372455
HSP70	0.73440859	0.73078934	1.523382853	1.529354536	0.987792555	0.973662518
Cox2	0.6274945	0.65146765	1.593415081	1.513981154	0.978844178	0.982231887
Gys	0.90865274	0.92948856	1.636258644	1.6137447	0.993871481	0.967841064
PAR	1.00381898	1.03224122	1.706421078	1.679261818	0.786306659	0.802611745
p16INK4a	0.65017043	0.6951588	1.789476326	1.873252634	0.966336294	0.99421
MCT4	0.49305181	0.47591022	1.864229649	1.877121069	0.950604808	1.009275181
Connexin-43	0.38754564	0.39145443	2.256522525	2.304875909	0.97829135	0.9827613
Ets-1	1.06001323	1.02765388	1.013962239	1.014873886	1.003775365	0.936077714
Cyclin-D1	1.00742668	1.0181704	1.032596757	1.037494557	0.964708668	0.948587385
ARID1A	1.00069189	0.98020102	1.019351782	1.018705226	0.965248697	0.973071597
Beclin	1.07356965	1.05105947	1.009174632	1.02892593	0.966350279	0.948819656
TTF1	1.07907206	1.0817636	1.003884499	1.019499773	0.946344312	0.943975806
XRCC1	1.46898485	1.46704647	1.000480532	1.003395618	0.924688655	0.942840458
GATA6	1.30911959	1.34667823	1.005685757	1.000434167	0.939036855	0.932042162
Rab25	1.73131391	1.81516832	1.003371215	1.002670023	0.622065086	0.573538217
S6_pS240_S244	0.60277173	0.6194196	1.005211327	1.000892468	1.786186132	1.710506087
PMS2	2.03949966	2.07899107	1.001336015	1.004636034	0.763075146	0.811253703
Progesterone-Receptor	2.09149289	2.32927119	1.005670622	1.000448788	0.864227164	0.897852552
YAP_pS127	0.69084171	0.67689099	5.575401155	5.468664352	1.026926175	0.936186399
PDK1	1.00919165	1.02728696	4.621940608	4.76723522	0.583590944	0.542776479
PD-L1	1.01998813	1.01733129	3.640996363	3.54180767	0.44119037	0.449639369

FoxO3a-(p_S318-S321)	0.91478852	0.89605532	0.996012062	1.009778991	1.099797219	1.044327508
Rictor_pT1135	0.87250192	0.86802619	0.990627141	1.014980844	1.05536394	1.050868467
Creb	0.87601138	0.85330058	0.998349243	1.007521266	1.072356788	1.082102767
INPP4b	0.98801732	0.98079465	0.985246513	1.016599763	1.000362815	1.034422785
p27_pT198	0.8423934	0.98254956	0.982984675	1.042845812	0.998566104	1.005085513
c-IAP2	0.85227779	0.84478427	0.988407739	1.017124793	1.180105952	1.118059596
Notch3	0.83934347	0.88074735	0.996067705	1.009725239	1.202167067	1.191879707
Stat3_pY705	0.7611933	0.75973186	0.990165764	1.015426535	1.032566642	1.039285727
WIPI1	0.69840255	0.73851097	0.968788679	1.05432796	1.053199169	1.050481179
Bcl-xL	0.78541886	0.80370202	0.984318452	1.027766055	1.037949432	1.027317403
JNK_pT183_Y185	0.68259562	0.67447767	0.991847978	1.01380151	1.141672115	1.087434338
Bad_pS112	0.75170368	0.81027075	0.993058762	1.012631889	1.117049534	1.124961173
TSC1	0.62967664	0.62293449	0.99513263	1.010628525	1.150485832	1.039479565
mTOR	0.62906152	0.62550366	0.999667191	1.006248124	1.116781313	1.037175603
p44-42-MAPK	0.58848151	0.60367781	0.998981717	1.006910294	1.190240952	1.131067172
Chk1_pS296	0.55245693	0.55850188	0.989838063	1.015743096	1.078185952	1.085044015
4E-BP1	0.50527012	0.48886308	0.978794025	1.026411675	1.094387802	1.049345415
Atg7	0.6621546	0.6555573	0.996379501	1.009424043	1.244692341	1.194660116
Rictor	0.76579999	0.77938872	0.999022681	1.006870723	1.22899673	1.176347567
Caspase-3	0.83510657	0.8408703	0.943604696	1.06040469	1.396419268	1.322360208
RPA32_pS4_S8	0.81102035	0.79844949	0.98955253	1.016018921	1.290037092	1.260828435
Bim	0.65088587	0.61081407	0.98783159	1.017681356	1.446783804	1.403170232
PKCa	0.61846729	0.6204974	0.986877818	1.018602703	1.434816877	1.353448059
Smad3	0.78080847	0.75145026	0.988851127	1.01669648	1.536316932	1.427104832
NAPSIN-A	0.77835101	0.74756803	0.9910535	1.014568979	1.482065764	1.394009714
c-Kit	0.79662859	0.77416759	0.974104017	1.030942239	1.654111329	1.520108515
TIGAR	0.51647754	0.5194255	0.972439259	1.032550401	1.408777107	1.350346391
p38-MAPK	0.53914503	0.55317422	0.973110309	1.031902164	1.366224166	1.252218744
Fibronectin	0.33641277	0.30531015	0.87807914	1.325712692	1.142273011	1.612263353
Akt	0.48303607	0.50465493	0.999139259	1.006758108	1.59743719	1.516142293
Stat3	0.32952512	0.33577567	0.978254349	1.026933004	1.470586091	1.414420881
ERCC5	1.1051744	1.09613648	0.997520801	1.008321543	0.768017171	0.872931087
Slfn11	1.05894589	1.06583021	0.866868135	1.033982449	0.980780917	0.962225527
PDHK1	1.04013935	1.02721158	0.994473762	1.003644182	0.984287918	0.981322642
XPF	1.01751115	1.02843576	0.996329525	1.008463722	0.997688748	0.964185477
SDHA	1.2493162	1.29207533	0.980329469	1.024928431	0.877168056	0.870078072
B7-H4	1.44559706	1.51408947	0.995581999	1.010194433	0.913418095	0.918655646
ER	1.52021354	1.5356687	0.999978339	1.005947553	0.921971017	0.911230859
eEF2K	1.61075322	1.57563835	0.97847859	1.007500729	0.895982897	0.93884425
CD4	1.68622748	1.51462776	0.889807824	1.011170652	0.976637065	0.873625883
HER2_pY1248	1.43232438	1.43233738	0.971146283	1.03055569	0.898440444	0.916400909
CD20	1.40601202	1.3578484	0.995153306	1.010608553	0.909150023	0.925246367
CD134	1.70792592	1.72257286	0.970795339	1.0063258	0.891799848	0.93998801
p53	1.6560036	1.7179942	0.991324424	1.014307266	0.938788543	0.882855009
WIPI2	1.75703578	1.83282433	0.987730476	1.017127997	0.864481267	0.929472382
IGFBP2	1.80281664	1.87271596	0.576046608	1.017572018	0.934195161	0.92904014
HSP27_pS82	1.72735068	1.73147702	0.994732856	1.011014709	0.754341591	0.776645504
MSI2	1.74282752	1.63334885	0.96890813	1.035961484	0.78419669	0.754949871
XBP-1	1.78842145	1.60328167	0.992393714	1.013274328	0.608953563	0.616414305
IRF-1	1.9373607	1.84295318	0.973742519	1.031291447	0.609821939	0.606045845
Jak2	0.69623381	0.68189884	0.969136405	1.035740969	1.850043496	1.520802849
S6_pS235_S236	0.43150118	0.45657145	0.987265034	1.018228651	2.214268734	2.087396262
MEK1	2.1722262	2.31307969	0.952778346	1.023048181	0.881653336	0.923709254
Wee1	2.39507779	2.45482866	0.998035322	1.007824515	0.808276256	0.85851111
cdc25C	2.3083811	2.40512976	0.997820096	1.008032424	0.861114843	0.908834425
LDHA	2.39032055	2.31766609	0.982534903	1.010369771	0.839468691	0.93605132
Cyclin-B1	3.03042353	3.05846575	0.909923328	1.066376539	0.904442679	0.881530344
ACC_pS79	2.87526877	3.09970187	0.974692996	1.030373283	0.902950159	0.907646344
PLK1	3.22267075	3.23460335	0.94992282	1.05430136	0.768919456	0.872641601
TRIM25	2.47867311	3.23886427	0.975716804	1.004656506	0.863081462	0.941613019
AR	3.22636431	3.44988141	0.978135676	1.027047642	0.710262755	0.184126953
ACC1	2.78707071	3.9232403	0.99268569	1.011328954	0.924269643	0.935117583
PARP	0.91295788	0.94417033	1.00938308	0.996862541	1.102181842	1.068976847

Smad1	0.88383979	0.92490696	1.023059319	0.983651246	1.070330482	1.077408908
IGF1R_pY1135_Y1136	0.87181429	0.87908299	1.013522189	0.992864147	1.042733976	1.064873886
Heregulin	0.8152416	0.83834345	1.016445824	0.990039906	1.032827856	0.996696536
Stathmin-1	0.81236588	0.82729125	1.009117295	0.99711929	1.034629652	1.005454378
HER3	0.85047242	0.84900981	1.012040387	0.994295572	1.056445599	1.032556184
Bax	0.83196745	0.84097896	1.056352372	0.967364161	1.021167506	1.007599506
PKC-delta_pS664	0.8576418	0.88654681	1.016072004	0.992994983	1.006768468	1.000088161
GSK-3a-b_pS21_S9	0.93452198	0.94466179	1.045347525	0.982223769	1.006062327	1.016710646
p27-Kip-1	0.95719588	0.94664962	1.00946849	0.996780035	1.185374391	1.151806234
Notch1	0.93806647	0.93075214	1.006709058	0.999445655	1.07989667	1.147507273
PRAS40_pT246	0.95548834	0.90359065	1.017713339	0.988815481	1.127249245	1.089536854
LC3A-B	0.86218011	0.89329774	1.030566669	0.976399119	1.236047628	1.119786657
D-a-Tubulin	0.73832928	0.74374322	1.00998343	0.996282601	1.032285862	1.029152648
Paxillin	0.75868597	0.78356299	1.017670832	0.988856544	1.067515793	1.017100855
C-Raf	0.80554494	0.80746598	1.006623691	0.99952812	1.052653887	1.019291358
HER3_pY1289	0.72321021	0.72756075	1.009840471	0.996420699	1.1470054	1.077255872
PI3K-p110-a	0.70049792	0.71051547	1.01283046	0.993532359	1.128303371	1.085744646
EGFR	0.6281988	0.75124003	1.044143528	0.999684365	0.988313176	1.168826949
VASP	0.58705097	0.5725404	1.015191268	0.991251811	1.010931431	0.986113952
14-3-3-zeta	0.64104989	0.64931598	1.006774319	0.999382613	1.087846632	1.032783008
B-Raf_pS445	0.58298146	0.58125215	1.017244845	0.989268048	1.173747582	1.099597722
VEGFR-2	0.38303018	0.411461	1.007791646	0.998399871	1.089040985	1.020855637
P-Met_pY1234_Y1235	0.66840454	0.66599255	1.006135204	1	1.257933351	1.170684416
p70-S6K1	0.64266595	0.67370687	1.007947069	0.998249733	1.321219391	1.137181945
MEK1_pS217_S221	0.73936578	0.74160832	1.025659968	0.98113901	1.370625972	1.289030515
PEA-15	0.63107855	0.67331155	1.055522417	0.95229178	1.302403469	1.293579084
Lck	0.76591797	0.76171113	1.021069239	0.985573671	1.230574385	1.164130281
mTOR_pS2448	0.88758077	0.86165335	1.030857353	0.976118318	1.408052696	1.304668666
Src_pY527	0.83063801	0.79768743	1.019546507	0.987044635	1.45075901	1.314146669
HES1	0.82170207	0.84740104	1.037587546	0.969616928	1.303043835	1.356930765
PKC-b-II_pS660	0.75774193	0.75972566	1.009968927	0.996296611	1.587102521	1.459610881
Akt_pT308	0.97985065	0.97378977	1.002836779	0.991154629	1.48776601	1.525666823
NDRG1_pT346	0.44375355	0.45372687	1.010400248	0.995879953	1.361829125	1.32308577
53BP1	0.50820329	0.54359348	1.00670421	0.999450339	1.279580172	1.199904339
AMPKa_pT172	0.46809512	0.42102011	1.056181289	0.951655307	1.231349355	1.206760597
eIF4E	0.22779392	0.21689893	1.010743185	0.995548675	1.176238745	1.149649689
CDK1	0.43290513	0.47023242	1.021332115	0.985319731	1.528717572	1.527679642
PKA-a	0.43940353	0.46132892	1.013349829	0.993030648	1.700699273	1.570745086
YAP	1.15647243	1.17140605	1.01185006	0.99447943	0.690007364	0.702134728
MSH6	1.12205509	1.11771275	1.010053293	0.996215113	0.652079348	0.666789391
PLC-gamma2_pY759	1.06584587	1.0651898	1.000512936	0.968056824	0.998268854	1.023048952
Tuberin_pT1462	1.08322259	1.08530311	1.021236395	0.9854122197	0.95482691	0.939779936
Glutamate-D1-2	1.08083284	1.0843439	1.013429101	0.99295407	0.912080895	0.949331799
Aurora-B	1.21754628	1.2138738	1.028993273	0.977919026	0.827016391	0.822801035
Bid	1.44297588	1.52104345	1.009240354	0.997000415	0.960585644	0.933489166
SOD2	1.35568744	1.34334497	1.002407618	0.992953837	0.961651354	0.95300523
Atg3	1.46782406	1.38233375	1.259718136	0.713554939	0.730822286	0.735007757
G6PD	1.68461864	1.68312461	1.023825537	0.982911077	0.868530428	0.272988334
PTEN	0.69246563	0.68669996	1.008776518	0.997448482	1.933228981	1.887037303
PDK1_pS241	0.65096201	0.62488709	1.010385413	0.995894284	1.749732821	1.451804865
Caspase-7-cleaved	0.78213752	0.75645176	1.007914053	0.998281626	2.992213468	2.948743874
SLC1A5	2.08267382	2.15468981	1.012438421	0.993911071	0.826674123	0.869362929
DUSP4	2.07813915	2.1901166	1.014485049	0.991934021	0.772335203	0.818569158
FAK_pY397	1.97356026	1.97105416	1.015614741	0.990842735	0.894339356	0.895497381
RIP	2.47208517	2.55730423	1.022530654	0.984161939	0.868499999	0.905884974
Myosin-IIa_pS1943	2.35315314	2.29383514	1.03591129	0.971236197	0.939285966	0.945412377
TFAM	2.28365585	2.28492934	1.012937337	0.977721101	0.962318241	0.967074643
Myosin-Heavy-Chain-11	2.49446961	2.42168402	1.00633978	0.999802379	0.14547361	0.130406315
TFRC	2.87521756	2.91972729	1.01415684	0.992251072	0.843663481	0.853234588
FASN	3.7487033	3.5803249	1.008802581	0.997423305	0.727719767	0.740005739
Axl	5.85303227	6.56876227	1.035414296	0.968184937	0.974049945	0.945937837
Rab11	0.95616478	0.97359004	0.990614611	0.993613346	1.007739026	1.021239506
COG3	0.98699194	0.98134479	0.997257479	0.983545408	1.018971577	1.042228133

TAZ	1.0146941	1.02221303	0.802434841	0.783993784	1.453749147	1.379186863
Akt_pS473	1	1.03576278	0.715535417	0.716052287	1.410149815	1.426473848
Granzyme-B	1.02178699	1.01567252	0.851115549	0.819017858	1.352741278	1.267926546
TUFM	1.0066442	1.02963602	0.74006413	0.717169716	1.300140216	1.2590373
ER-a_pS118	1.0178195	1.01933103	0.835375613	0.930202334	1.4257499	1.365625596
U-Histone-H2B	1.05196614	0.98784366	0.797113927	0.898686068	1.385722131	1.394328432
c-Abl	1.04326834	1.08023288	0.962836349	0.96281468	1.019583975	1.013283829
Rad50	1.06648839	1.06375539	0.94582001	0.924384698	0.997662257	1.020446122
Claudin-7	1.07301632	1.10046716	0.962944257	0.975380119	0.92250077	1.027182092
JNK2	1.09923924	1.10987698	0.96794635	0.921180212	0.981037712	0.98013126
SHP-2_pY542	1.09447287	1.12661828	0.89323635	0.900769877	0.971242525	1.004146864
Src-(phospho-Y416)	0.98401374	1.06367317	0.969659796	0.975886008	1.012504827	1.019946486
14-3-3-beta	0.97442371	1.0244141	0.924668734	0.940167139	1.050308777	1.00644215
c-Jun_pS73	0.97179907	1.09505358	0.961279114	0.941101232	1.047864458	1.089759026
GCLM	1.01833059	1.04629899	0.869703473	0.873363873	1.043127339	1.064634129
Tyro3	1.02049929	1.0260504	0.889861765	0.882504	1.041079899	1.038708278
PI3K-p85	1.01304438	1.02373427	0.923358237	0.933342392	1.121253269	1.056914619
Mnk1	0.99721043	1.01383574	0.969551826	0.950821381	1.224256833	1.207992681
AMPK-a2_pS345	1.02282209	1.01348357	0.93342689	0.951176076	1.218417039	1.21290479
DM-K9-Histone-H3	0.94812436	0.99943171	0.999301505	0.963267423	1.239230856	1.216169322
RBM15	1.0094201	1.0270763	0.848069913	0.83059644	1.166649557	1.260223694
ATM	1.02327645	0.99567235	0.7651204	0.969115074	1.181806766	1.169028815
b-Catenin	1.00768728	1.02867417	0.715585636	0.754566961	1.137175395	1.083885124
ULK1_pS757	1.07168377	1.09251484	0.766320368	0.760577691	0.992757383	1.030753818
Collagen-VI	1.26495527	1.2556548	0.998267253	0.98072355	0.947394598	0.964911063
LRP6_pS1490	1.2476854	1.26909208	0.979744173	0.980415414	0.962848903	0.965211024
FoxM1	1.24129734	1.24959562	0.99206759	0.994611553	0.936760002	0.951391492
eIF4E_pS209	1.25538307	1.28665702	0.951305435	0.95639551	0.974866028	0.986041543
UBAC1	1.12347852	1.15070104	1	0.993209211	0.993245075	0.952756632
IRS1	1.12929758	1.15265787	0.968997964	0.959254541	0.966515402	0.985810525
ATM_pS1981	1.15980325	1.18171595	0.995732634	0.983105855	0.91628955	0.962591957
MAPK_pT202_Y204	1.16994866	1.20125015	0.930493244	0.937910969	0.974032228	0.986840027
ZAP-70	1.17233396	1.17444384	0.918193205	0.931840668	0.998032953	0.963855851
ATR_pS428	1.21239917	1.24141851	0.899000764	0.885565724	0.995351325	0.966423898
Myt1	1.15929089	1.21788061	0.8643983	0.865939282	0.973147709	0.987687082
FAK	1.2352298	1.11826196	0.879991236	0.960254794	0.971782567	0.984836808
Chk2_pT68	1.1358238	1.10849629	0.837819021	0.919138909	1.002139852	0.959922899
p70-S6K_pT389	1.23696259	1.28400106	0.835313257	0.848737763	0.986705796	0.974703247
C-Raf_pS338	1.26112337	1.28760164	0.789070019	0.826842989	0.976914386	0.984079942
Cdc2_pY15	1.271923	1.28851853	0.890532941	0.870978196	0.992063222	0.969572735
FoxO3a	1.51934487	1.49581407	0.979795378	0.986833221	0.981895068	0.958963474
p21	1.53591374	1.58250624	0.996124099	0.97419196	1.002796297	0.959294258
PREX1	1.54521088	1.6151031	0.98278879	0.98716597	0.932360122	0.958639553
DM-Histone-H3	1.53645157	1.57426037	0.87995061	0.849946021	0.954781386	1.005275501
Tuberin	1.47940305	1.4972051	0.892988818	0.87386753	0.956843884	1.00330036
Shc_pY317	1.33645901	1.39632395	0.996374375	0.975604842	0.916864647	0.969893978
Wee1_pS642	1.32693066	1.37369501	0.954115325	0.952997683	0.969361285	0.991313133
Stat5a	1.37398749	1.3676816	0.902350378	0.895854891	0.995494091	0.96628718
STING	1.32793823	1.44233911	0.903387593	0.893838122	0.951136032	1.008766457
Rb_pS807_S811	1.46358008	1.53505457	0.661937832	0.673930401	0.982978035	0.978273119
Pcd4	1.44835008	1.48096733	0.63813397	0.654814582	0.950313462	1.009554187
p90RSK_pT573	1.38105957	1.41370876	0.746715923	0.719633541	0.96808216	0.99253808
E-Cadherin	1.34270239	1.38357239	0.549320653	0.567493241	0.942302296	1.01722604
Rad51	1.71138261	1.74605694	0.96588917	0.949404736	0.931297854	0.995399027
BRD4	1.72023411	1.76501665	0.944952936	0.901068866	0.984839312	0.976490676
P-Cadherin	1.66347625	1.77241588	0.989815616	0.983258715	0.990768229	0.962443153
Hexokinase-II	1.67699128	1.75294174	0.883349804	0.882129187	1.006660252	0.955593961
Gab2	1.0379509	1	0.985523569	0.964756375	1.967419315	1.913087567
MERIT40_pS29	2.03358923	2.17180395	0.995793402	0.984097513	0.879089365	0.961626607
Cox-IV	2.34988899	2.85726102	0.980097619	0.976174993	0.972476833	0.969338952