

Downregulated probe set in ALDH- DOX vs. CTRL

Probe	Symbol	GenBank	Gene	UniGene	Fold Change	DOX ALDH-
204114_at	NID2	NM_007361	22795	Hs.369840	4.010527511	9.977249254
230036_at	SAMD9L	BE669858	219285	Hs.489118	2.43456334	9.01219676
231644_at		AW016812			-2.28442119	3.760793278
226603_at	SAMD9L	BE966604	219285	Hs.489118	2.253728547	8.839771326
242477_at	TTC39B	BF056282	158219	Hs.563630	2.242497192	7.368197941
1559722_at		BC042892			-2.221829239	5.229921317
217332_at	RP11-159J2.1	AL133018	647288	Hs.567920	2.174029493	6.139308902
235643_at	SAMD9L	BE886225	219285	Hs.489118	2.135736034	8.625327442
211548_s_at	HPGD	J05594	3248	Hs.596913	2.12649599	9.413755752
244610_x_at		AA572726			-2.115147586	4.848868296
230607_at		T86874			-2.099198155	5.01923339
236826_at	TTC39B	H83092	158219	Hs.563630	2.095892213	6.064166248
1565701_at		AL832624			-2.046879451	4.971579139
230332_at	ZCCHC7	AA872187	84186	Hs.654700	-2.043889025	5.360124356
215206_at		AK025143			-2.017442549	5.214683376
214295_at	KIAA0485	AW129056	57235	Hs.604754	-1.990997056	5.707951367
1556865_at		AF143885			-1.984807195	5.805319923
233401_at		BF723605			-1.967534559	4.463644502
228367_at	ALPK2	BE551416	115701	Hs.628152	1.963883128	8.521473284
232565_at		AK025052			-1.9460451	5.904248272
232882_at		AA079839			-1.940352935	5.955747586
238558_at		AI445833			-1.931886747	6.998535735
1553155_x_at	ATP6V0D2	NM_152565	245972	Hs.436360	1.921061049	8.695275418
230850_at		AI652647			-1.919355579	4.681656603
232333_at		AU147805			-1.918927244	5.337175905
215898_at	TTL5	AK021879	23093	Hs.709609	-1.916386887	4.618970142
235705_at		BF676361			-1.87895106	7.100564992
239946_at		AA776723			-1.87398213	5.530788182
203914_x_at	HPGD	NM_000860	3248	Hs.596913	1.847250176	9.124719435
1556352_at		AI692624			-1.842119819	5.638534852
243993_at		AA436887			-1.840232533	4.852075649
242188_at		AI743332			-1.837797297	4.794329111
242471_at		AI916641			-1.827734629	5.6611017
1569727_at		BC028245			1.81512361	7.18984308
241960_at	CSMD1	AA705177	64478	Hs.571466	1.807335235	7.526109148
216628_at		AL117447			-1.806715435	4.570919
215889_at	SKIL	X15217	6498	Hs.581632	-1.792590443	4.300606165
243874_at	LPP	AI079544	4026	Hs.444362	-1.7892799	5.953401452
244414_at		AI148006			-1.784667992	5.612596379
232264_at		AK022204			-1.781948978	7.101680259
1553153_at	ATP6V0D2	NM_152565	245972	Hs.436360	1.781658898	8.643414057
235811_at		AW590853			-1.761472083	5.217772631
241391_at		AA654772			-1.751722268	4.495276984
236987_at	ALPK2	AI741514	115701	Hs.628152	1.75078243	7.744788588
1564378_a_at		AK025101			-1.73803763	5.390712969
239965_at	LOC151878	AW009761	151878	Hs.680377	-1.733473828	5.863726509
1559723_s_at		BC042892			-1.730407817	4.740115765
241245_at	SFRS4	AV647470	6429	Hs.469970	-1.7243532	5.909951144

1562330_s_at	CSMD1	AB067477	64478 Hs.571466	1.722298982	7.416253855
236010_at		AI373107		-1.716358291	4.38815632
224235_at		AF119853		-1.707169413	5.131655531
236134_at	DCAF7	AA769995	10238 Hs.410596	-1.705251892	5.144583198
243271_at		AI064690		1.69933647	6.51387498
81737_at	NPIPL3	AI424872	23117 Hs.552700, Hs.6	-1.694763082	5.280099879
235912_at		BE552155		-1.691760243	5.026105519
237310_at		AI743607		-1.690836941	6.370005181
215802_at		AK000144		-1.682369067	4.715546557
1565689_at		BG400570		-1.680952426	5.211456217
242405_at		BF358386		-1.680614987	5.797690185
232307_at		AK021554		-1.67155057	5.119570994
227838_at		AW070250		1.667526065	7.092993757
238774_at		AW960454		-1.656047306	5.641757396
1559491_at		AL390180		-1.637091221	5.986215604
236355_s_at		AI076172		-1.636515011	5.452618512
206548_at		NM_024880		-1.621858313	4.752670264
202772_at	HMGCL	NM_000191	3155 Hs.533444, Hs.6	1.613732346	10.46353821
1560109_s_at	NUB1	AA193477	51667 Hs.647082	-1.61217186	4.427439129
237456_at		AI655806		-1.601058632	5.552052246
232000_at	TTC39B	AW001030	158219 Hs.563630	1.594870215	6.592156687
241214_at		AI939471		-1.594027526	6.427490364
1553151_at	ATP6V0D2	AY079172	245972 Hs.436360	1.593822177	6.127799328
242732_at		BG010493		-1.591835851	5.392188124
239512_at	SFRS4	R05895	6429 Hs.469970	-1.582467727	6.288944841
229281_at	NPAS3	N51682	64067 Hs.657892	1.577868541	7.043269326
215078_at	SOD2	AL050388	6648 Hs.487046	-1.570939987	8.567688082
203913_s_at	HPGD	AL574184	3248 Hs.596913	1.567976068	6.925410339
235757_at		AA814006		-1.567429339	5.477107509
233674_at		AK026286		-1.544754952	6.659942388
239264_at		AW973078		-1.534596082	5.022928547
240263_at		N74924		-1.527998042	5.414833397
232615_at		AA632758		-1.526607253	7.56380803
222087_at	PVT1	AW451806	5820 Hs.133107, Hs.6	-1.520964032	6.506095299
209200_at	MEF2C	AL536517	4208 Hs.653394	1.511102888	10.07206684
234605_at	CDC14B	AK024886	8555 Hs.40582	-1.506428939	5.25551146
1559156_at		BC036508		-1.493012385	7.32083969
242114_at	BOLA2	BF088991	552900 Hs.444600, Hs.6	-1.485496393	5.842246534
239414_at		BF942260		-1.472552963	7.416295721
243088_at		W84667		-1.462678259	6.463625368
236976_at	FANCA	AI569792	2175 Hs.719210	-1.462081352	5.850309365
1568846_at		BC017718		-1.460480063	8.251496935
235786_at		AI806781		-1.458479649	4.855168606
1562144_at		AF075103		-1.456709094	4.707724893
1556672_a_at	RBM6	AI190489	10180 Hs.439480	-1.447891544	7.135064691
242859_at		BE156563		-1.446764951	6.658800272
1558714_at		BC043430		-1.446140631	5.239233791
242886_at		AW007763		-1.445700255	6.777535898
242143_at		BE674964		-1.433048473	5.196505471
1557718_at	PPP2R5C	AL834350	5527 Hs.368264, Hs.7	-1.426546004	5.463246136

241457_at		AI821935		-1.425353889	5.156465629
238883_at		AW975051		-1.421705845	5.662421595
235680_at		AI914925		-1.421080784	5.485447387
242038_at	LRRC8B	BG037106	23507 Hs.482017, Hs.6	-1.415413841	5.505180003
239227_at		AW182675		-1.413753058	5.98710379
227266_s_at	FYB	BF679849	2533 Hs.370503	1.408725618	6.327739611
1558410_s_at		AW974642		-1.40857607	5.886075734
1569208_a_at		BC020895		-1.408207411	5.257810203
1564886_at		AL359595		-1.405432263	6.67497196
206408_at	LRRTM2	NM_015564	26045 Hs.656653	-1.400483571	4.79290933
239908_at		AA496799		-1.400434197	7.224845963
243003_at		AV702197		-1.400193546	7.451847543
1560680_at		AL833513		-1.391417609	6.415653397
229858_at		AU146893		-1.391347821	6.616081109
242476_at		AI436356		-1.389218591	5.897007174
240798_at		BE467916		-1.384782965	4.989657664
231205_at		BF055351		-1.377684726	5.525099202
243474_at		W87425		-1.374184785	4.875604885
235847_at		BF111312		-1.37376709	5.635878127
234095_at		AU155112		-1.371839766	6.22693389
228623_at		AI224133		-1.366709103	7.92092966
238619_at		AA417078		-1.365610656	6.878220062
228573_at	ANTXR2	BE673665	118429 Hs.162963	1.365158879	9.950818451
210109_at	C7orf54	AF191492	27099 Hs.657377	-1.364271879	5.85157939
236216_at		AA598661		-1.363306066	6.151186082
241932_at		AI073803		-1.355742401	5.365774836
239348_at		AI285970		-1.352580885	5.746394193
243149_at		AI467945		-1.350407784	4.716783739
209199_s_at	MEF2C	N22468	4208 Hs.653394	1.349690435	9.634230601
1567223_at	HMGA2	U29113	8091 Hs.505924	-1.34327983	5.283316032
242868_at		T70087		-1.342577424	5.356776997
231552_at		AW451785		-1.341249019	6.1933929
240499_at		AA482221		-1.339816138	4.765993075
202720_at	TES	NM_015641	26136 Hs.592286, Hs.6	1.335902451	7.010780215
1564798_at		AK000811		-1.332835723	4.971195426
223557_s_at	TMEFF2	AB017269	23671 Hs.144513	1.331031212	9.443686909
241372_at	ZC3H6	R34135	376940 Hs.190477	1.319913899	6.21684905
1558385_at		AL832806		-1.318738931	5.92696848
244682_at	CAMSAP1	H11471	157922 Hs.522493	-1.315917022	5.537793623
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1558569_at	UNQ6228	AL832308	100131541 Hs.661972	-1.311239402	6.732864364
206638_at	HTR2B	NM_000867	3357 Hs.421649	-1.310845368	6.300984803
239567_at		AW974998		-1.307590707	5.166559629
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239519_at		AA927670		-1.305373616	6.949290327
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236947_at		AI686664		-1.287292079	5.387679882
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242983_at		AI806626		-1.284876621	6.055331801
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234159_at		AK025182		-1.280392834	6.1933929
242126_at		T53962		-1.279693914	5.073080785
239274_at		AV729557		-1.276573674	5.289077341
244473_at		R37637		-1.274649646	6.19136422
242240_at		AI805700		-1.27052649	7.981482171
1554213_at	ARHGEF10	BC036809	9639 Hs.98594	-1.269774342	6.235483101
1555074_a_at	KCNH5	BC043409	27133 Hs.27043	-1.268741856	5.152148854
230590_at		BE675486		-1.256709358	6.36043427
220969_s_at		NM_030892		-1.255823081	5.873399357
242068_at		AA608834		-1.255091279	6.334370986
243670_at		BE670161		-1.254393993	6.298453529
219210_s_at	RAB8B	NM_016530	51762 Hs.389733	1.252531426	8.896902106
1557539_at		BC008052		-1.249058254	5.257753161
243827_at		AL038125		-1.246508175	5.090613928
228885_at	MAMDC2	AI862120	256691 Hs.547172	1.244099341	7.938658868
222180_at		AU147889		-1.2424408	4.878445785
231199_at		AA701676		-1.239247755	4.97295984
217482_at		AK021987		-1.239191413	6.746678136
241681_at		AW296451		-1.237660534	8.523250339
241797_at		AI904095		-1.233064386	5.574602342
242492_at	CLNS1A	AA412065	1207 Hs.430733, Hs.5!	-1.230846072	6.81827937
227443_at	C9orf150	AI972386	286343 Hs.445356	1.230363171	9.490543805
232002_at		AW843302		-1.227981951	7.165781121
213606_s_at	ARHGDI A	AI571798	396 Hs.159161	-1.227016296	7.92480366
239957_at		AW510793		-1.226766046	5.674927094
1562059_at		AW020871		-1.223453938	4.989517584
212070_at	GPR56	AL554008	9289 Hs.513633	1.223395523	9.650393353
207746_at	POLQ	NM_014125	10721 Hs.241517	-1.22320161	5.035558454
231223_at	CSMD1	R41565	64478 Hs.571466	1.222024021	10.21594423
238563_at		AV762916		-1.220644296	5.629190794
215318_at	CG012	AL049782	116829 Hs.687692	-1.218691805	5.088582525
232835_at		AA533080		-1.216728418	6.538249517
1558599_at		AA532745		-1.216018231	4.930059765
229013_at	LOC145783	BF111029	145783 Hs.620525	1.210304432	6.928918182
234224_at		AL137541		-1.209278201	5.029854733
239238_at		AI208857		-1.209129626	6.872820177
240787_at		AW449433		-1.209108521	5.429919799
238172_at		AA192765		-1.205867076	6.452511679
239448_at		AI475033		-1.204860943	5.958505685
1569664_at		BC035915		-1.204625094	5.889268693
241762_at	FBXO32	BF244402	114907 Hs.403933	-1.201540977	5.216605613
214215_s_at	LARP4B	AW514174	23185 Hs.681734, Hs.7	-1.201398131	5.011255105
241320_at		AI821449		-1.19843653	6.438203212
1557597_at		BC036212		-1.197701235	5.270676936
241301_at		AA694187		-1.194646051	5.847261909
225345_s_at		AU155376		-1.191962436	7.13975899
239469_at		BF513404		-1.191351279	6.084684925
1552829_at		NM_018505		-1.186675757	5.871886366

239923_at		AI056872		-1.185689302	5.263571335
232465_at		AK021749		-1.184736202	5.814342576
215979_s_at	SLC7A1	AK022999	6541 Hs.14846	-1.179773588	7.40288594
240302_at		AW450681		-1.176980078	5.624145396
232134_at		AW139789		-1.173025017	8.52800424
1570108_at		BC037848		-1.167714757	6.111813249
230099_at		AI139993		-1.16735959	5.646997613
213556_at	LOC390940	BE673445	390940 Hs.22049	1.165040076	8.365901634
232711_at		AU146197		-1.163456902	7.171100511
239606_at		AA669135		-1.161466094	6.667950381
242343_x_at		H57111		-1.161073721	6.334415364
232592_at		AU146731		-1.160449332	6.553674955
231343_at		AW300131		1.157345144	9.058293683
232778_at		AK026036		-1.157341022	6.426497012
244457_at		BF724206		-1.157010166	5.497985021
231950_at	ZNF658	AW874634	26149 Hs.522147	1.154698581	6.332787832
1566079_at	RPS16P5	AL833001	647190 Hs.675022	-1.154158212	5.740014468
235454_at		AI436561		-1.154153809	5.997101564
238438_at	CNOT6L	R67226	246175 Hs.592519	-1.148244828	4.950103549
222453_at	CYBRD1	AL136693	79901 Hs.715531	-1.146416232	8.898566249
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220941_s_at	C21orf91	NM_017447	54149 Hs.293811	1.144823601	9.391385285
240247_at		AI653240		-1.141014777	7.761099423
222282_at		AV761453		-1.140469185	7.165076325
205933_at	SETBP1	NM_015559	26040 Hs.435458	1.137791562	8.388530219
244347_at		BE825318		-1.130701943	6.342499118
240008_at		AI955765		-1.12656662	5.436576671
1560926_at		AF085924		-1.126257237	7.358212315
239757_at	ZFAND6	AA741493	54469 Hs.719106	-1.125568584	5.935947012
240257_at	SYNJ2	AI971212	8871 Hs.434494	-1.124140779	6.128558185
241893_at		BE927766		-1.123412459	5.081078414
233315_at		AK024947		-1.123186393	5.934117093
243768_at		AA026388		-1.122398537	7.346260473
201775_s_at	KIAA0494	AA676790	9813 Hs.719205	-1.120432801	8.679352994
244356_at		AL079909		-1.11896753	7.54442516
237377_at		AA069425		-1.11873928	4.94904123
236346_at		BF115793		-1.118574924	7.480749926
233921_s_at		AK022078		-1.118204719	7.089373219
236595_at		AA776458		-1.117894154	6.201153958
1569477_at		BC025999		-1.116530721	6.40072714
1567224_at	HMGA2	U29113	8091 Hs.505924	-1.115914833	4.921529594
220410_s_at	CAMSAP1	NM_018627	157922 Hs.522493	-1.115500301	6.264168824
238706_at	PAPD4	BG168850	167153 Hs.418198	-1.114722037	5.152322974
240248_at		AA778783		-1.11255464	6.328362658
243589_at	KIAA1267	AI823453	284058 Hs.648744	-1.11165161	6.383715285
228484_s_at	FOXO1	AI472322	2308 Hs.370666	-1.111237419	6.837802011
227478_at	SETBP1	BF739885	26040 Hs.435458	1.110884956	8.985904131
242449_at		BG054682		-1.106983063	5.823983663
1555854_at		AA594609		-1.10647736	8.971570236
244358_at		AW372457		-1.105973871	7.742020115

244633_at		AA404996		-1.105410517	5.918017712
239892_at		AW593666		-1.104384516	6.780422129
1556240_at		AI339498		-1.103723408	5.218451371
233800_at		AA805082		-1.102331772	5.4661987
239597_at		AA993566		-1.10086361	5.862675207
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