		Differen	nce from median of	controls							
Well Name	Well#	1st	2nd	3rd	Average	Standard deviation	SSMD	GeneSymbol	Geneld	GeneAccession	GINumber
E01	49	-1.266	-1.134	-1.060	-1.153	0.104	-6.248	Dynamin-2			
F01 E12	61 60	-1.246 -1.211	-1.127 -1.111	-1.033 -1.053	-1.135 -1.125	0.107 0.080	-6.012 -7.940	Dynamin-2 Dynamin-2			
F12	72	-1.221	-1.100	-1.040	-1.120	0.092	-6.854	Dynamin-2			
G12	84	-1.024	-0.691	-0.905	-0.873	0.169	-2.914	Clathrin heavy chain			
H12 G01	96 73	-1.044 -0.932	-0.586 -0.604	-0.889 -0.887	-0.840 -0.807	0.233 0.178	-2.035 -2.559	Clathrin heavy chain Clathrin heavy chain			
H01	73 85	-0.952 -0.952	-0.553	-0.856	-0.787	0.208	-2.539	Clathrin heavy chain			
F09	69	-0.608	-0.474	-0.503	-0.528	0.070	-4.239	PPP2R2A	5520	NM_002717	32307111
D07	43	-0.382	-0.246	-0.348	-0.325	0.070	-2.606	PTPRH	5794	NM_002842	67190343
E08 E06	56 54	-0.297 -0.356	-0.334 -0.236	-0.291 -0.313	-0.307 -0.302	0.023 0.061	-7.494 -2.793	INPP4A PR48	3631 28227	NM_001566 NM_199326	4504704 41349490
D09	45	-0.269	-0.269	-0.313	-0.278	0.016	-2.793 -9.780	FLJ23751	92370	NM_152282	22748648
H07	91	-0.309	-0.141	-0.258	-0.236	0.087	-1.538	SET	6418	NM_003011	142361523
F05	65	-0.201	-0.195	-0.229	-0.208	0.018	-6.554	DKFZP566K0524	26095	NM_001042365	109138576
D05 F10	41 70	-0.214 -0.196	-0.158 -0.144	-0.247 -0.200	-0.206 -0.180	0.045 0.031	-2.575 -3.253	PTPN2 PFKFB3	5771 5209	NM_002828 NM_004566	18104978 42476167
E04	52	-0.173	-0.154	-0.194	-0.174	0.020	-5.002	EYA2	2139	NM_172110	26667229
G02	74	-0.166	-0.176	-0.158	-0.167	0.009	-10.001	PTPN11	5781	NM_002834	33356176
G09	81	-0.220	-0.104	-0.166	-0.163	0.058	-1.581	FLJ22405	64419	NM_022485	117938273
F07 B07	67 19	-0.171 -0.143	-0.158 -0.156	-0.157 -0.173	-0.162 -0.157	0.008 0.015	-11.419 -5.907	G6PC MTMR4	2538 9110	NM_000151 NM_004687	119393889 31377795
F04	64	-0.167	-0.155	-0.142	-0.155	0.013	-6.943	PPP1CC	5501	NM_002710	4506006
H08	92	-0.134	-0.128	-0.175	-0.146	0.026	-3.220	NUDT3	11165	NM_006703	37622350
B04	16 50	-0.109	-0.158	-0.153	-0.140	0.027	-2.950 2.417	PTPN5	84867	NM_001039970	90652860
E02 C03	50 27	-0.119 -0.127	-0.151 -0.129	-0.111 -0.120	-0.127 -0.125	0.021 0.005	-3.417 -14.473	PHPT1 NUDT14	29085 256281	NM_014172 NM_177533	142373598 37594466
C11	35	-0.126	-0.065	-0.163	-0.118	0.049	-1.349	PPP2R4	5524	NM_178003	30065648
B02	14	-0.126	-0.113	-0.086	-0.108	0.020	-3.030	SGPP1	81537	NM_030791	40254975
F06	66 75	-0.108 -0.084	-0.101 0.111	-0.103 -0.105	-0.104	0.003	-17.617 2.061	NT5C3 PPFIA1	51251	NM_016489 NM 003626	70608210
G03 H04	75 88	-0.084 -0.121	-0.111 -0.076	-0.105 -0.100	-0.100 -0.099	0.014 0.022	-3.961 -2.512	DUSP3	8500 1845	NM_004090	29171751 37655179
A08	8	-0.116	-0.106	-0.075	-0.099	0.021	-2.615	CDC25A	993	NM_201567	42490759
F03	63	-0.097	-0.115	-0.083	-0.098	0.016	-3.451	DUSP21	63904	NM_022076	33942082
H11	95 30	-0.128	-0.067 -0.144	-0.096 -0.067	-0.097 -0.094	0.031	-1.794 1.227	CILP CTDP1	8483	NM_003613	51944961
D03 C06	39 30	-0.072 -0.081	-0.144 -0.102	-0.067	-0.092	0.043 0.010	-1.237 -4.989	PTPRD	9150 5789	NM_048368 NM_130393	67188549 104486447
C09	33	-0.091	-0.076	-0.092	-0.086	0.009	-5.295	PSPH	5723	NM_004577	46249387
G08	80	-0.116	-0.085	-0.058	-0.086	0.029	-1.674	FLJ25449	151649	NM_144714	109627669
E10	58	-0.053	-0.117	-0.081	-0.084	0.032	-1.492	PGAM2	5224	NM_000290	50593009
H09 G07	93 79	-0.100 -0.092	-0.110 -0.085	-0.041 -0.069	-0.083 -0.082	0.037 0.012	-1.259 -3.804	NT5C1A PPP2R3A	84618 5523	NM_032526 NM_002718	14210537 32967594
F11	71	-0.084	-0.077	-0.073	-0.078	0.006	-7.842	PTPN18	26469	NM_014369	18375654
C05	29	-0.104	-0.076	-0.053	-0.078	0.025	-1.725	DUSP5	1847	NM_004419	62865889
F08	68	-0.144	-0.036	-0.053	-0.077	0.058	-0.750	ACP2 PTPN6	53 5777	NM_001610	4557009
H06 C04	90 28	-0.068 -0.040	-0.078 -0.083	-0.062 -0.076	-0.069 -0.067	0.008 0.023	-4.764 -1.619	EYA4	2070	NM_080548 NM_172103	94557302 98991762
D06	42	-0.069	-0.082	-0.040	-0.064	0.022	-1.661	PIB5PA	27124	NM_001002837	50726959
D08	44	-0.013	-0.091	-0.085	-0.063	0.043	-0.819	PPM1B	5495	NM_001033556	75813617
D04	40 10	-0.047	-0.073	-0.069	-0.063	0.014	-2.506	NUDT15	55270	NM_018283	8922791
A10 G06	10 78	-0.126 -0.059	-0.037 -0.034	-0.024 -0.088	-0.062 -0.060	0.056 0.027	-0.630 -1.260	PFKFB1 PPP3R1	5207 5534	NM_002625 NM_000945	118582285 45238847
D02	38	-0.052	-0.048	-0.071	-0.057	0.012	-2.661	PTPRT	11122	NM_007050	148539859
A04	4	-0.072	-0.031	-0.055	-0.053	0.021	-1.428	PPP1R3C	5507	NM_005398	21314622
F02 H02	62 86	-0.120 -0.047	0.004 -0.081	-0.027 -0.013	-0.048 -0.047	0.064 0.034	-0.419 -0.787	PPM1A DUSP14	5494 11072	NM_021003 NM_007026	29557948 148596940
G04	76	-0.066	-0.052	-0.013	-0.047	0.034	-1.142	PPP1R14B	26472	XM_370630	41200335
H05	89	-0.052	-0.075	-0.003	-0.043	0.037	-0.664	ILKAP	80895	 NM_030768	29171685
A01	1	-0.055	-0.052	-0.013	-0.040	0.023	-0.967	Non-targeted control	.===		
E05 B03	53 15	-0.067 -0.053	-0.030 -0.044	0.006 0.009	-0.030 -0.029	0.037 0.033	-0.465 -0.492	DLG1 PPP1R3A	1739 5506	NM_001098424 NM_002711	148539627 34304342
D01	37	-0.044	-0.006	-0.037	-0.029	0.020	-0.432	Non-targeted control	3300	14141_002711	34304342
D11	47	-0.030	-0.044	-0.004	-0.026	0.020	-0.736	NUDT1	4521	NM_002452	40288273
B09	21	-0.093	-0.001	0.018	-0.025	0.059	-0.241	G6PC3	92579	NM_138387	24308433
E07 C10	55 34	-0.025 -0.017	-0.035 -0.054	0.000 0.010	-0.020 -0.020	0.018 0.032	-0.636 -0.351	MK-STYX CDC25C	51657 995	NM_016086 NM_022809	32481212 125625348
G05	34 77	-0.017 -0.017	0.022	-0.060	-0.020	0.032	-0.351 -0.254	PTPRA	5786	NM_080841	125987586
B08	20	-0.015	-0.017	-0.019	-0.017	0.002	-5.752	PPP1R1B	84152	NM_181505	142371393
H03	87	-0.062	0.008	0.016	-0.013	0.043	-0.168	SSH1	54434	NM_018984	40254883
C08 B10	32 22	-0.011 -0.048	0.003 -0.020	-0.016 0.047	-0.008 -0.007	0.010 0.049	-0.458 -0.079	MTM1 FLJ32332	4534 132160	NM_000252 NM_144641	4557895 21389470
C01	25	-0.012	0.002	-0.010	-0.007	0.007	-0.518	Non-targeted control	132100	14141_144041	21303470
A12	12	-0.003	-0.045	0.036	-0.004	0.040	-0.056	Non-targeted control			
A03	3	-0.029	-0.023	0.046	-0.002	0.042	-0.025	DUSP10	11221	NM_144729	21536332
D12 B11	48 23	0.011 -0.027	-0.002 0.010	-0.005 0.029	0.001 0.004	0.009 0.029	0.093 0.076	Non-targeted control PTPRO	5800	NM_002848	13677212
H10	94	-0.055	-0.007	0.029	0.004	0.066	0.070	DUSP19	142679	NM_080876	31377625
A02	2	-0.042	0.009	0.047	0.005	0.045	0.064	DOT1L	84444	 NM_032482	163965372
D10	46	0.008	-0.014	0.030	0.008	0.022	0.213	PPP4C	5531	NM_002720	4506026
B05 E11	17 59	-0.033 -0.005	0.015 0.018	0.048 0.022	0.010 0.012	0.041 0.015	0.139 0.455	DUSP11 EPM2A	8446 7957	NM_003584 NM_001018041	4503414 66346727
B01	13	0.003	0.018	0.022	0.012	0.015	0.455	Non-targeted control	1331	14141_001010041	003 4 0/2/
G10	82	-0.046	0.055	0.045	0.018	0.055	0.183	INPP5E	56623	NM_019892	38327538
E03	51	-0.025	0.006	0.075	0.019	0.051	0.205	PTPN21	11099	NM_007039	150418004
C02 C12	26 36	-0.021 0.052	0.036 0.012	0.048 0.005	0.021 0.023	0.037 0.025	0.324 0.518	ALPL Non-targeted control	249	NM_000478	116734716
A05	5	-0.005	0.012	0.005	0.023	0.025	0.320	CTDSP1	58190	NM_021198	10864008
C07	31	0.039	0.042	-0.003	0.026	0.025	0.593	DUSP4	1846	NM_001394	58331238
A11	11	0.004	0.040	0.096	0.047	0.047	0.566	ALPPL2	251	NM_031313	157266295
B12 A09	24 9	0.049 0.017	0.023 0.034	0.076 0.109	0.050 0.053	0.026 0.049	1.061 0.611	Non-targeted control PPP3CA	5530	NM_000944	142369518
G11	83	0.017	0.049	0.109	0.066	0.049	1.474	KIAA0310	9919	NM_014866	124378038
A06	6	0.086	0.022	0.140	0.083	0.059	0.793	PTPRJ	5795	NM_001098503	148728159
E09	57	0.049	0.086	0.117	0.084	0.034	1.404	PTPN13	5783	NM_006264	5453991

0.098 NM_152422 B06 18 0.057 0.107 0.132 0.038 1.458 PTPDC1 138639 30089951 0.099 A07 0.172 56953 NM_020201 74229030 0.067 0.059 0.063 0.888 NT5M 7

		Differer	nce from median of	controls							
Well Name	Well#	1st	2nd	3rd	Average	Standard deviation	SSMD	GeneSymbol	Geneld	GeneAccession	GINumber
F01 E01	61 40	-1.238 -1.233	-1.115 -1.118	-1.152 -1.114	-1.168	0.063 0.068	-10.424 -9.633	Dynamin-2			
E12	49 60	-1.233 -1.198	-1.118 -1.076	-1.114 -1.142	-1.155 -1.139	0.068	-9.633 -10.536	Dynamin-2 Dynamin-2			
F12	72	-1.213	-1.060	-1.124	-1.132	0.077	-8.292	Dynamin-2			
G12	84	-1.008	-0.751	-0.998	-0.919	0.146	-3.563	Clathrin heavy chain			
H12	96 95	-1.020	-0.714	-0.956	-0.897	0.162	-3.128	Clathrin heavy chain			
H01 G01	85 73	-0.960 -0.893	-0.622 -0.652	-0.907 -0.900	-0.830 -0.815	0.181 0.141	-2.580 -3.257	Clathrin heavy chain Clathrin heavy chain			
F09	69	-0.365	-0.372	-0.442	-0.393	0.042	-5.247	THTPA	79178	NM_024328	54607163
G10	82	-0.330	-0.188	-0.389	-0.302	0.103	-1.652	PPP3R2	5535	NM_147180	34303920
B05	17	-0.288	-0.184	-0.366	-0.279	0.091	-1.726	PPP1R7	5510	NM_002712	4506012
F06	66	-0.256	-0.167	-0.197	-0.207	0.045	-2.594	DKFZP761G058	152926	NM_152542	31542532
C07 G04	31 76	-0.209 -0.235	-0.124 -0.154	-0.283 -0.209	-0.205 -0.199	0.079 0.041	-1.456 -2.722	PPP1R2 SSH3	5504 54961	NM_006241 NM_017857	34147499 24586674
G02	74	-0.226	-0.148	-0.186	-0.187	0.039	-2.689	PPP2R5D	5528	NM_006245	31083266
C08	32	-0.143	-0.203	-0.195	-0.180	0.033	-3.132	WBP11	51729	NM_016312	18375679
G09	81	-0.302	-0.020	-0.190	-0.171	0.142	-0.680	PPP1R12B	4660	NM_032103	14195596
C05 D03	29 39	-0.164 -0.187	-0.131 -0.092	-0.209 -0.179	-0.168 -0.153	0.039 0.052	-2.404 -1.646	PPP1R1C BPNT1	151242 10380	NM_001080545 NM_006085	122937385 116812594
B03	15	-0.187	-0.083	-0.179	-0.147	0.056	-1.495	PTPN14	5784	NM_005401	34328898
H06	90	-0.177	-0.067	-0.176	-0.140	0.063	-1.255	NT5E	4907	NM_002526	4505466
H08	92	-0.162	-0.108	-0.150	-0.140	0.029	-2.766	MTMR6	9107	NM_004685	134142347
A06	6	-0.188	-0.089	-0.135	-0.137	0.049	-1.573	NT5C	30833	NM_014595	7657032
D09 A09	45 9	-0.136 -0.192	-0.050 -0.064	-0.212 -0.132	-0.132 -0.129	0.081 0.064	-0.922 -1.143	PPP2R1B PTPN23	5519 25930	NM_181699 NM_015466	124269711 110681717
B04	16	-0.123	-0.102	-0.132	-0.121	0.019	-3.632	PPP2R2B	5521	NM_181676	32307116
F04	64	-0.143	-0.099	-0.100	-0.114	0.025	-2.577	DUSP15	128853	NM_177991	29826316
C06	30	-0.098	-0.061	-0.168	-0.109	0.054	-1.133	PTP4A3	11156	NM_007079	14589853
B07	19	-0.213	-0.041	-0.068	-0.107	0.093	-0.653	KIAA0377	9677	NM_014659	7662083
D10 G07	46 79	-0.102 -0.035	-0.113 -0.123	-0.102 -0.154	-0.105 -0.104	0.006 0.062	-9.176 -0.948	FBP2 PTPN4	8789 5775	NM_003837 NM_002830	22907027 18104987
H05	89	-0.127	-0.052	-0.126	-0.102	0.043	-1.329	DUSP18	150290	NM_152511	51093844
C02	26	-0.061	-0.133	-0.107	-0.100	0.036	-1.567	PPP5C	5536	 NM_006247	20127496
D04	40	-0.119	-0.073	-0.102	-0.098	0.023	-2.395	ACP5	54	NM_001611	161377449
C04	28	-0.147	-0.058	-0.084	-0.096 -0.096	0.046	-1.188	PPAP2C	8612 4659	NM_003712	29171743 4505316
E08 F05	56 65	-0.230 -0.106	-0.018 -0.087	-0.039 -0.092	-0.095	0.117 0.010	-0.463 -5.456	PPP1R12A PPP1R13B	23368	NM_002480 NM_015316	121114286
E02	50	-0.100	-0.062	-0.120	-0.094	0.029	-1.795	PPFIA4	8497	NM_015053	55769553
F02	62	-0.117	-0.039	-0.109	-0.088	0.043	-1.161	NUDT4	11163	NM_019094	98985815
A08	8	-0.092	-0.089	-0.046	-0.075	0.026	-1.662	PPP2R1A	5518	NM_014225	32455242
C09 F07	33 67	-0.071 -0.090	-0.007 -0.060	-0.137 -0.060	-0.072 -0.070	0.065 0.017	-0.625 -2.321	PPP1R1A MTMR7	5502 9108	NM_006741 NM_004686	155030237 142379887
E03	51	-0.093	-0.026	-0.079	-0.066	0.017	-2.321	IGBP1	3476	NM_001551	48375183
D05	41	-0.151	-0.040	-0.006	-0.065	0.076	-0.487	INPP5A	3632	NM_005539	109702905
C11	35	-0.140	0.015	-0.063	-0.063	0.078	-0.458	PTPLB	201562	NM_198402	158819030
D08	44	-0.047	-0.064	-0.067	-0.059	0.011	-3.152	PPM1F	9647	NM_014634	29826333
F03 H11	63 95	-0.051 -0.059	-0.038 -0.031	-0.076 -0.074	-0.055 -0.054	0.019 0.022	-1.647 -1.407	PTPN1 Empty	5770	NM_002827	18104977
D06	42	-0.049	-0.017	-0.088	-0.051	0.035	-0.821	PPM1E	22843	NM_014906	30089947
E06	54	-0.050	-0.065	-0.034	-0.049	0.015	-1.816	PPP1R8	5511	NM_014110	20336237
E04	52	-0.096	-0.009	-0.030	-0.045	0.045	-0.559	DUSP13	51207	NM_016364	56237017
F08	68	-0.067	-0.039	-0.025	-0.044	0.021	-1.154	CTDSP2 SGPP2	10106	NM_005730	93004101
C10 A01	34 1	-0.015 -0.054	-0.056 -0.035	-0.051 -0.031	-0.041 -0.040	0.022 0.012	-1.023 -1.897	Non-targeted control	130367	NM_152386	23510288
G03	75	-0.019	-0.046	-0.053	-0.039	0.018	-1.219	ACPP	55	NM_001099	161377405
A11	11	-0.058	-0.031	-0.022	-0.037	0.019	-1.107	PPAP2B	8613	NM_003713	29171739
F10	70 53	-0.019	-0.015	-0.077	-0.037	0.035	-0.603	DUSP22	56940	NM_020185	34147625
E05 D11	53 47	-0.050 -0.045	-0.007 -0.025	-0.052 -0.034	-0.036 -0.035	0.026 0.010	-0.801 -2.010	SBF1 DUSP7	6305 1849	NM_002972 NM 001947	37574611 85662716
D01	37	-0.027	-0.033	-0.036	-0.032	0.004	-4.233	Non-targeted control	1043	14141_001547	03002710
E09	57	-0.123	0.034	0.001	-0.030	0.083	-0.202	C9ORF67	84814	NM_032728	34147435
H04	88	-0.040	0.008	-0.054	-0.029	0.032	-0.500	PTPRG	5793	NM_002841	18860897
H03 H09	87 93	-0.004 0.011	-0.004 -0.025	-0.072 -0.056	-0.027 -0.023	0.039 0.033	-0.387 -0.392	PPFIA3 PPM1G	8541 5496	NM_003660 NM_002707	41872688 29826283
B11	23	-0.036	-0.002	-0.025	-0.023	0.033	-0.532	PPP6C	5537	NM_002721	20127429
C01	25	-0.006	-0.020	-0.033	-0.020	0.013	-0.845	Non-targeted control		_	
A07	7	0.008	-0.031	-0.035	-0.019	0.024	-0.456	PPM2C	54704	NM_018444	8923959
B09	21 77	-0.015	-0.004	-0.034	-0.017	0.016	-0.635	TENS1 LPPR4	64759	NM_022748	134152712
G05 G06	77 78	-0.008 0.013	0.005 -0.047	-0.049 -0.016	-0.017 -0.017	0.028 0.030	-0.345 -0.315	PTPN22	9890 26191	NM_014839 NM_012411	45827807 15619017
D07	43	-0.009	-0.023	-0.014	-0.015	0.007	-1.208	FBP1	2203	NM_000507	160298191
G11	83	-0.037	0.002	0.009	-0.009	0.025	-0.198	CDC14B	8555	NM_001077181	116008457
B01	13	-0.015	-0.004	-0.008	-0.009	0.006	-0.885	Non-targeted control	250		0.470.40.45
H10 A03	94 3	-0.042 0.000	-0.008 -0.015	0.036 0.010	-0.005 -0.001	0.039 0.013	-0.072 -0.063	ALPP G6PC2	250 57818	NM_001632 NM_001081686	94721245 126273545
H02	86	-0.016	-0.013	0.010	0.000	0.013	-0.005	NUDT5	11164	NM_014142	37594463
F11	71	0.040	-0.017	-0.023	0.000	0.035	0.000	PTPRR	5801	 NM_002849	19743915
C03	27	0.009	0.000	-0.006	0.001	0.007	0.089	TPTE2	93492	NM_130785	40549428
G08	80	0.045	-0.008	-0.026	0.004	0.037	0.055	PPFIBP2	8495	NM_003621	57163846
E11 E10	59 58	-0.050 -0.044	0.058 0.042	0.023 0.068	0.011 0.022	0.055 0.059	0.109 0.210	CDC25B NUDT12	994 83594	NM_004358 NM_031438	47078250 37574615
D12	48	0.006	0.042	0.008	0.025	0.039	0.798	Non-targeted control	55554	051430	5,5,7013
C12	36	0.067	0.004	0.008	0.026	0.036	0.416	Non-targeted control			
A04	4	-0.029	0.075	0.034	0.027	0.052	0.290	PFKFB2	5208	NM_001018053	64762444
A12	12	0.060	0.005	0.019	0.028	0.028	0.558	Non-targeted control	E120F	NINA 046364	151201212
A02 B06	2 18	0.029 0.036	0.028 0.041	0.048 0.052	0.035 0.043	0.011 0.008	1.747 3.099	ACP6 SSH2	51205 85464	NM_016361 NM_033389	151301212 37674209
B10	22	0.049	0.061	0.045	0.052	0.008	3.452	PON2	5445	NM_001018161	66529395
B08	20	0.010	0.045	0.103	0.053	0.047	0.635	LCK	3932	 NM_001042771	112789547
B12	24	0.041	0.025	0.097	0.054	0.038	0.810	Non-targeted control	4000	NINA COSTA	4700470
A10 E07	10 55	0.021 0.020	0.044 0.078	0.102 0.118	0.055 0.072	0.042 0.049	0.749 0.828	CDKN3 NT5C1B	1033 93034	NM_005192 NM_033253	17981703 50593107
H07	91	0.020	0.054	0.109	0.082	0.027	1.689	PPM1L	151742	NM_139245	63003904
D02	38	0.081	0.064	0.117	0.087	0.027	1.809	ALPI	248	NM_001631	157266291

0.093 B02 14 0.078 0.113 0.088 0.018 2.907 IMPA2 3613 NM_014214 7657235 0.120 A05 0.106 0.081 0.174 0.048 PFKFB4 5210 NM_004567 19923257 1.407 5

		Differen	ce from median of	controls							
Well Name	Well#	1st	2nd	3rd	Average	Standard deviation	SSMD	GeneSymbol	Geneld	GeneAccession	GINumber
E01	49	-1.174	-1.149	-1.219	-1.181	0.036	-18.749	Dynamin-2			
F01	61 60	-1.155	-1.105 1.073	-1.230	-1.163	0.063	-10.428	Dynamin-2			
E12 F12	60 72	-1.188 -1.130	-1.072 -1.103	-1.213 -1.239	-1.158 -1.157	0.075 0.072	-8.661 -9.083	Dynamin-2 Dynamin-2			
H12	96	-0.998	-0.730	-1.018	-0.915	0.160	-3.217	Clathrin heavy chain			
G12	84	-0.985	-0.664	-1.068	-0.906	0.214	-2.391	Clathrin heavy chain			
G01	73	-0.874	-0.671	-0.957	-0.834	0.147	-3.198	Clathrin heavy chain			
H01	85	-0.833	-0.652	-0.976	-0.820	0.162	-2.850	Clathrin heavy chain	2625	NINA 005544	64005476
G10 E10	82 58	-0.302 -0.337	-0.360 -0.205	-0.510 -0.271	-0.391 -0.271	0.107 0.066	-2.059 -2.315	INPP5D PPP2CA	3635 5515	NM_005541 NM 002715	64085176 57222566
G04	76	-0.212	-0.234	-0.271	-0.271	0.082	-1.849	PTPRE	5791	NM 006504	40805849
F03	63	-0.337	-0.143	-0.194	-0.224	0.101	-1.258	PPP1R15B	84919	NM_032833	41282220
H06	90	-0.141	-0.172	-0.260	-0.191	0.062	-1.744	PPP2R5B	5526	NM_006244	30795206
H07	91	-0.219	-0.101	-0.215	-0.178	0.067	-1.499	RNGTT	8732	NM_003800	134142827
E09	57 25	-0.206	-0.073	-0.241	-0.174	0.089	-1.105	INPPL1	3636	NM_001567	4755141
C11 C06	35 30	-0.246 -0.180	-0.008 -0.089	-0.232 -0.202	-0.162 -0.157	0.133 0.060	-0.685 -1.475	TIMM50 PPAP2A	92609 8611	NM_001001563 NM_003711	48526508 29171735
G09	81	-0.138	-0.125	-0.181	-0.148	0.030	-2.819	SACM1L	22908	NM_014016	7662337
F06	66	-0.117	-0.115	-0.194	-0.142	0.045	-1.781	PALD	27143	 NM_014431	51100971
F11	71	-0.184	-0.043	-0.191	-0.139	0.083	-0.945	PPP3CB	5532	NM_021132	11036639
G07	79	-0.101	-0.098	-0.217	-0.139	0.068	-1.154	PSPHL	8781	XM_926720	89024914
H04	88	-0.100 0.136	-0.091 0.111	-0.214	-0.135	0.069	-1.110 -4.754	NUDT2 PPP1CA	318	NM_001161	22265329 45827797
H08 F07	92 67	-0.136 -0.076	-0.111 -0.147	-0.139 -0.162	-0.129 -0.128	0.015 0.046	-4.754 -1.563	PGAM1	5499 5223	NM_206873 NM_002629	31543395
C05	29	-0.096	-0.120	-0.144	-0.120	0.024	-2.823	DUSP9	1852	NM 001395	4503420
E05	53	-0.153	-0.094	-0.108	-0.118	0.031	-2.154	PPP2R5A	5525	_ NM_006243	30795205
G08	80	-0.089	-0.097	-0.161	-0.116	0.040	-1.646	DUSP6	1848	NM_001946	42764682
H02	86	-0.096	-0.102	-0.144	-0.114	0.026	-2.469	PTPRK	5796	NM_002844	18860901
E07 B06	55 18	-0.126 -0.106	-0.063 -0.075	-0.139 -0.139	-0.109 -0.107	0.041 0.032	-1.514 -1.878	DUSP12 DLG7	11266 9787	NM_007240 NM_014750	6005955 21361644
C08	32	-0.137	-0.017	-0.139	-0.107	0.032	-0.790	INPP4B	8821	NM_001101669	156104896
F02	62	-0.116	-0.092	-0.082	-0.097	0.017	-3.142	PTPRB	5787	NM_002837	157952212
H11	95	-0.081	-0.059	-0.148	-0.096	0.047	-1.165	PPP4R1	9989	NM_005134	108936951
H05	89	-0.059	-0.076	-0.144	-0.093	0.045	-1.169	SKIP	51763	NM_016532	18765708
D11	47	-0.137	0.015	-0.155	-0.092	0.093	-0.560	DUSP1	1843	NM_004417	7108342
C09 B09	33 21	-0.102 -0.157	-0.072 -0.021	-0.101 -0.092	-0.092 -0.090	0.017 0.068	-3.018 -0.748	MTMR3 PPP1CB	8897 5500	NM_153051 NM_002709	148276975 4506004
E11	59	-0.137	-0.021	-0.032	-0.030	0.063	-0.748	TA-PP2C	160760	NM_139283	21281678
G03	75	-0.051	-0.075	-0.128	-0.085	0.039	-1.212	PTPRN	5798	NM_002846	18860905
G06	78	-0.055	-0.082	-0.114	-0.084	0.030	-1.604	PTPRS	5802	NM_130853	104487294
H03	87	-0.116	-0.030	-0.097	-0.081	0.045	-1.011	PPAPDC1	196051	NM_001030059	73611919
F04	64	-0.188	0.012	-0.066	-0.081	0.101	-0.452	INPP5B	3633	NM_005540	113722124
H09 G05	93 77	-0.036 -0.052	-0.082 -0.048	-0.100 -0.118	-0.073 -0.073	0.033 0.039	-1.240 -1.050	BPGM DUSP16	669 80824	NM_001724 NM_030640	40353767 38372910
D06	42	-0.060	-0.046	-0.110	-0.072	0.034	-1.197	PPP2R5E	5529	NM_006246	31083295
H10	94	-0.052	-0.051	-0.103	-0.069	0.030	-1.314	PPFIA2	8499	_ NM_003625	29171754
B07	19	-0.021	-0.058	-0.125	-0.068	0.053	-0.725	PTPN20A	653129	NM_001042397	109138574
F05	65	-0.077	-0.052	-0.055	-0.061	0.014	-2.510	PTP4A1	7803	NM_003463	62865860
G02 E08	74 56	-0.029 -0.061	-0.047 0.008	-0.107 -0.090	-0.061 -0.048	0.041 0.050	-0.837 -0.535	MTMR8 PPP3CC	55613 5533	NM_017677 NM_005605	21361758 34147659
E02	50	-0.069	-0.033	-0.033	-0.045	0.021	-1.224	ACPT	93650	NM_033068	18375509
A06	6	-0.042	-0.027	-0.066	-0.045	0.020	-1.294	PTPRU	10076	NM_133178	110735403
D01	37	-0.058	0.000	-0.076	-0.044	0.040	-0.631	Non-targeted control			
F10	70	-0.093	0.004	-0.036	-0.041	0.049	-0.482	PTPRM	5797	NM_002845	157419149
D05 B08	41 20	-0.039 -0.059	-0.031 0.007	-0.052 -0.059	-0.040 -0.037	0.010 0.038	-2.183 -0.548	PPP2CZ SYNJ1	333926 8867	NM_005167 NM 003895	65506327 44921605
B03	20 15	-0.006	-0.053	-0.03 9 -0.045	-0.037	0.038	-0.348 -0.788	PON1	5444	NM 000446	31880793
C01	25	-0.028	-0.017	-0.045	-0.030	0.014	-1.193	Non-targeted control			0200700
B05	17	0.023	-0.058	-0.052	-0.029	0.045	-0.360	PPP2R2D	55844	NM_001003656	51093852
F09	69	-0.089	0.033	-0.008	-0.021	0.062	-0.195	SYNJ2	8871	NM_003898	52851404
A07 G11	7 83	0.028 0.016	-0.051 -0.005	-0.039 -0.072	-0.021 -0.020	0.043 0.046	-0.271 -0.251	DUSP2 PTEN	1844 5728	NM_004418 NM_000314	12707563 110224474
C12	36	-0.019	-0.030	-0.072	-0.020	0.013	-0.251	Non-targeted control	3728	14141_000314	110224474
A08	8	-0.035	-0.017	0.001	-0.017	0.018	-0.535	PPP1R3D	5509	NM_006242	37622904
C07	31	-0.003	-0.011	-0.034	-0.016	0.016	-0.553	PDP2	57546	NM_020786	41349496
A01	1	-0.025	-0.015	0.004	-0.012	0.015	-0.452	Non-targeted control			
E06 E04	54 52	-0.044 -0.013	0.036 0.030	-0.016 -0.030	-0.008 -0.004	0.041 0.031	-0.113 -0.074	PTPMT1 PTPLA	114971 9200	NM_175732 NM_014241	148224883 82659104
B11	23	0.013	-0.007	-0.030	-0.004	0.031	-0.074	IMPA1	3612	NM_005536	8393607
C02	26	-0.019	0.001	0.016	-0.001	0.017	-0.018	PPM1D	8493	NM_003620	29558519
B01	13	0.022	0.020	-0.031	0.004	0.030	0.068	Non-targeted control		_	
B02	14	0.077	-0.009	-0.045	0.008	0.063	0.070	STYX	6815	NM_145251	34222188
D09	45	-0.020	0.002	0.045	0.009	0.033	0.155	MTMR2	8898	NM_016156	20357517
E03 A11	51 11	0.015 0.086	0.023 -0.006	-0.003 -0.035	0.012 0.015	0.013 0.063	0.509 0.136	PTPRZ1 ENPP7	5803 339221	NM_002851 NM_178543	91208427 31341740
C03	27	0.031	0.010	0.007	0.015	0.003	0.682	PTPRC	5788	NM_080923	115385974
B04	16	0.044	0.001	0.024	0.023	0.021	0.613	MGC10067	134510	NM_145049	142347678
D08	44	0.060	0.025	-0.003	0.027	0.032	0.483	OCRL	4952	NM_000276	21396493
A03	3	0.010	0.052	0.022	0.028	0.022	0.724	MINPP1	9562	NM_004897	164420775
A05 D02	5 38	-0.003 -0.002	0.038	0.051 0.063	0.029	0.028	0.569	PTPN12 EYA1	5782 2138	NM_002835	18375651 26667218
D02 A10	38 10	-0.002 0.083	0.025 -0.004	0.063 0.019	0.029 0.033	0.033 0.045	0.498 0.411	EYA1 PGAM4	2138 441531	NM_172059 NM_001029891	2666/218 84370365
D12	48	0.019	0.035	0.049	0.033	0.045	1.277	Non-targeted control	. 41331	001023031	5 157 0505
B12	24	0.062	0.036	0.027	0.042	0.018	1.273	Non-targeted control			
A12	12	0.071	0.000	0.068	0.046	0.040	0.648	Non-targeted control			
F08	68	0.030	0.097	0.022	0.050	0.041	0.682	NUDT9	53343	NM_024047	37594462
C10 D07	34 43	-0.037 0.050	0.116 0.055	0.105 0.078	0.061 0.061	0.086 0.015	0.403 2.356	ACP1 PPP2CB	52 5516	NM_001040649 NM_004156	96304456 57634540
D07	43 40	0.058	0.053	0.078	0.061	0.013	1.620	MTMR1	8776	NM_003828	88999580
A09	9	0.095	0.062	0.050	0.069	0.023	1.689	PTP4A2	8073	NM_080392	88758598
A02	2	0.104	0.065	0.044	0.071	0.031	1.295	PPP1R16B	26051	NM_015568	34335276
D10	46	0.063	0.070	0.088	0.074	0.013	3.225	PPEF1	5475	NM_152226	23312377

B10	22	0.098	0.083	0.051	0.077	0.024	1.811	DUSP8	1850	NM_004420	153281157
A04	4	0.106	0.062	0.102	0.090	0.024	2.087	PTPN9	5780	NM_002833	18375663
C04	28	0.045	0.087	0.140	0.091	0.048	1.074	PPAPDC2	403313	NM_203453	66773039
D03	39	0.097	0.108	0.111	0.105	0.007	7.940	NT5C2	22978	NM_012229	20149601