

Improved Prediction of Gene Expression through Integrating Cell Signalling Models with Machine Learning: supplemental document

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supplementary table 1. The accuracy of the first round model.

Gene	R2	Gene	R2	Gene	R2	Gene	R2
DDR1	0.085	RRAGA	-0.021	SDHB	0.048	HES1	0.078
RHOA	0.018	SMC4	0.318	GTF2A2	0.076	PSMG1	0.199
RNPS1	0.044	NIPSNAP1	0.139	CCNB2	0.412	DDB2	0.373
RPS6	-0.038	MYBL2	0.276	LIG1	0.403	CCNA2	0.437
CSRP1	-0.044	ELAVL1	0.027	EBP	0.038	EML3	0.028
XBP1	0.543	KIAA0100	0.025	WRB	0.155	PRAF2	0.123
SKP1	0.007	TP53	0.331	SYNE2	-0.04	MRPL19	0.097
CAPN1	0.141	PSME2	0.242	OXCT1	0.34	MNAT1	-0.062
CTSD	0.288	LBR	0.062	NNT	0.132	RAB4A	-0.073
MAT2A	0.122	SCCPDH	0.125	GAA	0.133	BCL2	0.061
STMN1	0.341	SMARCD2	0.079	SLC37A4	0.03	RFC2	-0.026
CIRBP	0.17	NET1	0.367	HPRT1	0.104	SLC25A13	-0.009
PSME1	0.254	HDAC2	0.038	FAH	0.1	MRPL12	0.134
PGAM1	0.014	HSPB1	0.073	POP4	0.07	CEBPA	0.026
STAT1	-0.01	CDC25B	0.319	CDC20	0.262	SLC5A6	-0.011
GNAS	-0.023	ATMIN	0.044	NUP88	0.035	AURKA	0.359
HIF1A	-0.012	PSRC1	0.29	MSH6	0.088	CCNH	0.002
PAICS	0.08	UBE2A	0.094	PIN1	-0.002	CDC45	0.033
HADH	0.257	COASY	-0.078	ETFB	0.01	ENOSF1	0.074
SMARCC1	-0.025	BIRC5	0.532	CRYZ	0.168	CDK2	0.343
PIP4K2B	0.014	ARHGAP1	0.246	UBE2C	0.489	ELOVL6	0.492
PFKL	0.009	CPNE3	0.034	LYPLA1	-0.001	CREB1	-0.102
PGRMC1	0.048	NUP62	0.087	TIMELESS	0.502	PPIC	0.068
MBNL1	0.081	PCM1	0.02	PAN2	0.006	BRCA1	0.162
PCNA	0.242	NUP93	0.055	RB1	0.068	CCDC85B	0.101
ILK	0.051	PLK1	0.162	TBP	-0.011	CCP110	0.013
SYPL1	0.045	CDK4	0.446	HAT1	-0.081	CDC25A	0.03
NUDCD3	0.051	HSD17B10	0.134	S100A4	0.195	NCK1	-0.1
TOP2A	0.543	CSK	-0.063	B4GAT1	0.278	TMEM5	0.013
EBNA1BP2	0.135	TCEAL4	0.364	ABCB6	0.153	MELK	0.188

SCP2	0.114	USP1	0.204	RFC5	0.159	CCNF	0.239
TMEM109	0.231	MPC2	0.287	CDK1	0.46	TCFL5	0.014
TRAP1	0.026	DECR1	0.19	TLE1	0.052	ICAM3	0.067
SOX4	0.143	ERBB3	0.234	PAFAH1B3	0.114	ATF5	0.036
SCRN1	0.087	PRSS23	0.465	DCK	0.167	CCNE2	-0.011
NFKBIA	0.215	IKBKAP	-0.035	DYNLT3	-0.036	LSM6	0.2
G3BP1	-0.001	GALE	0.509	BAMBI	0.354	ITGAE	0.069
RPA1	0.117	PRKCD	0.007	SLC35A1	0.4	SGCB	-0.036
MCM3	0.378	S100A13	0.445	ITGB1BP1	-0.009	PLS1	0.019
SMC1A	-0.025	NRIP1	0.244	TIMM17B	0.025	PCMT1	0.192
PWP1	-0.04	EAPP	0.024	UGDH	0.094	INPP4B	0.07
ICMT	0.208	POLR2K	0.108	EZH2	0.141	CHEK1	0.047
RUVBL1	0.205	NRAS	0.029	MYCBP	0.212	DAG1	0.111
GHR	0.168	EED	0.073	ASAH1	0.034	ZDHHC6	0.02
SCYL3	-0.013	GATA3	0.353	SMARCA4	0.074	TIMM9	0.123
POLG2	-0.073	CETN3	0.002	ATP5S	0.142	GMNN	0.216
NOLC1	0.075	TGFB3	0.254	PARP2	0.01	CDCA4	0.256
POLE2	0.344	HIST1H2BK	0.22	AKR7A2	0.238	NENF	0.114
RAD51C	0.116	ANXA7	0.03	SPDEF	0.543	DNAJC15	-0.015
KIF14	0.079	PUF60	-0.009	CCNB1	0.188	HEBP1	0.193
SLC35A3	0.003	CFLAR	0.018	ERBB2	-0.015	CD320	0.031
AKT1	0.253	RPL39L	-0.03	VDAC1	-0.095	CHMP4A	0.086
MUC1	0.199	CHEK2	0.187	GAPDH	-0.086	HEATR1	-0.11
CASP2	-0.117	HMG20B	0.097	CNDP2	0.082	CISD1	0.238
PARP1	0.176	TPM1	0.162	ADI1	0.161	SUV39H1	0.178
FDFT1	0.095	BDH1	0.482	TMEM50A	0.198	VPS28	-0.053
CCND1	0.042	LSM5	0.112	HACD3	0.031	KIF20A	0.346
OXA1L	-0.092	TXNDC9	-0.092	YKT6	0.437	METRNL	0.029
CDC42	-0.018	HN1L	0.178	BZW2	-0.026	LAGE3	0.285
ADH5	-0.053	ZMIZ1	-0.068	EVL	0.036	PXMP2	0.274
CYCS	-0.017	PDS5A	-0.013	MIF	0.156	AMDHD2	0.155
COPS7A	0.046	XPO7	-0.001	IARS2	0.285	FBXO11	-0.1
GLOD4	0.09	COG4	0.014	KLHDC2	0.205	MTERF3	0.217
DLD	-0.068	TMEM97	0.071	STUB1	0.136	PSMD10	-0.023
IFRD2	0.034	TFDP1	0.22	ENOPH1	-0.054	TSEN2	0.089
TSPAN6	0.012	KAT6B	0.005	MRPS2	-0.078	PRR7	0.105
CDKN1B	-0.009	ADO	-0.03	NUP85	0.17	GTPBP8	-0.077
PRPF4	-0.031	JADE2	0.179	FIS1	0.012	DENND2D	-0.043
CBR1	-0.016	PCCB	0.077	NUSAP1	0.25	ARPP19	-0.1
NR2F6	0.212	KIAA1033	0.062	MRPS16	0.068	ATG3	-0.042
TSPAN4	0.38	RRP1B	0.047	NPDC1	-0.004	WDR61	0.091
PSIP1	0.035	THAP11	-0.008	DERA	0.11	CANT1	-0.021
AURKB	0.284	C2CD5	0.26	UBR7	0.102	FAM63A	0.095
RPA3	0.252	POLR2I	-0.047	ISOC1	0.188	ATF1	0.028
HDGFRP3	0.061	RPIA	0.289	CCDC92	0.09	TSTA3	0.092
RRS1	0.036	PLCB3	0.161	SCAND1	0.202		

supplementary table 2. The accuracy of the seven models is illustrated as follows: cn is the common neighbour model, lpi is local path index model, ent is the relative-entropy model, loc is the locally linear embedding model, lgin is the laplacian eigenmaps embedding model, n2v is the node2vec model, and depwlc is the deepwalk model.

Gene	baseline	cn	ib	ent	locallylinear	lap _{egn}	n2v	deepwalk	stacking
DDR1	0.45	0.4532	0.4516	0.4519	0.4517	0.4527	0.4515	0.4518	0.4538
RHOA	0.4643	0.4636	0.4633	0.4637	0.4652	0.4636	0.4648	0.4622	0.4662
RNPS1	0.4	0.3988	0.4021	0.4043	0.4001	0.4007	0.4006	0.4015	0.4036
RPS6	0.4167	0.4145	0.4146	0.4146	0.4128	0.4139	0.4167	0.4162	0.4167
CSRP1	0.1946	0.1933	0.193	0.1923	0.1939	0.1935	0.1935	0.1969	0.1964
XBP1	0.7518	0.7516	0.7511	0.7518	0.7517	0.7515	0.7502	0.7512	0.7526
SKP1	0.4822	0.4826	0.4829	0.483	0.4838	0.4834	0.4812	0.4824	0.4846
CAPN1	0.4592	0.4585	0.4595	0.4593	0.4601	0.458	0.4582	0.4581	0.4627
CTSD	0.6112	0.6114	0.6114	0.6122	0.6104	0.6114	0.6114	0.6107	0.613
MAT2A	0.4682	0.4695	0.4717	0.4698	0.4718	0.471	0.4701	0.4718	0.4725
STMN1	0.676	0.6759	0.677	0.677	0.6773	0.6776	0.6762	0.6771	0.678
CIRBP	0.5163	0.5138	0.5141	0.5127	0.5119	0.5135	0.5132	0.5121	0.5155
PSME1	0.6093	0.6103	0.6111	0.6116	0.611	0.6107	0.6115	0.6111	0.6124
PGAM1	0.2458	0.2486	0.247	0.2484	0.248	0.2479	0.2488	0.2454	0.2508
STAT1	0.2715	0.2731	0.2729	0.2743	0.2673	0.2689	0.2713	0.2674	0.2737
GNAS	0.2093	0.2114	0.2093	0.2094	0.2093	0.2084	0.2111	0.21	0.2122
HIF1A	0.3849	0.3884	0.3889	0.3895	0.3876	0.3894	0.3869	0.3864	0.3901
PAICS	0.5619	0.5634	0.5626	0.5641	0.5621	0.5646	0.563	0.5632	0.5651
HADH	0.5623	0.562	0.5628	0.5633	0.5626	0.5631	0.5619	0.5637	0.5639
SMARCC1	0.4122	0.4144	0.4134	0.4149	0.4149	0.4169	0.4141	0.417	0.4169
PIP4K2B	0.3865	0.3904	0.3913	0.3925	0.3894	0.3922	0.3899	0.3882	0.3924
PFKL	0.2845	0.2866	0.2883	0.2868	0.287	0.2868	0.286	0.2887	0.2886
PGRMC1	0.5239	0.5243	0.5252	0.5244	0.5254	0.5243	0.5255	0.5242	0.527
MBNL1	0.4416	0.4401	0.4441	0.4425	0.4436	0.444	0.4391	0.4436	0.444
PCNA	0.4139	0.4114	0.4113	0.412	0.4121	0.413	0.4138	0.4112	0.4145
ILK	0.4552	0.4547	0.4528	0.4515	0.4539	0.453	0.4545	0.454	0.4554
SYPL1	0.4451	0.4451	0.4476	0.4477	0.4479	0.4455	0.4448	0.4482	0.4484
NUDCD3	0.4956	0.4928	0.4936	0.4912	0.4935	0.4949	0.4932	0.4926	0.4951
TOP2A	0.7958	0.7948	0.7956	0.7948	0.7966	0.7949	0.7946	0.795	0.7964
EBNA1BP2	0.5359	0.5367	0.5368	0.5363	0.5377	0.5364	0.5366	0.537	0.5389
SCP2	0.4834	0.4861	0.4881	0.4871	0.4883	0.4875	0.485	0.4877	0.4884
TMEM109	0.5836	0.5846	0.5828	0.5815	0.5837	0.5808	0.5835	0.5833	0.5849
TRAP1	0.4347	0.4346	0.4326	0.4349	0.4366	0.4342	0.4353	0.4359	0.4366
SOX4	0.5297	0.5311	0.5313	0.5318	0.535	0.5318	0.5327	0.5314	0.5339
SCRN1	0.4654	0.4625	0.4625	0.4626	0.4639	0.4628	0.4648	0.4647	0.4656
NFKBIA	0.4707	0.472	0.4689	0.4711	0.4702	0.468	0.4729	0.4701	0.4729
G3BP1	0.2792	0.2768	0.2787	0.2787	0.2739	0.2814	0.2769	0.2746	0.28
RPA1	0.5569	0.5566	0.555	0.5536	0.5565	0.5555	0.5564	0.5575	0.5581
MCM3	0.6614	0.6621	0.6619	0.6608	0.6622	0.6621	0.6627	0.6607	0.6631

SMC1A	0.1879	0.1896	0.1931	0.1926	0.1921	0.1915	0.1895	0.1942	0.1938
PWP1	0.3422	0.3393	0.3403	0.3429	0.343	0.3405	0.3389	0.3406	0.343
ICMT	0.5028	0.5027	0.5034	0.5023	0.5022	0.5015	0.5018	0.5031	0.5042
RUVBL1	0.5557	0.5571	0.5564	0.5574	0.555	0.5567	0.5551	0.5575	0.5581
RRAGA	0.4428	0.4445	0.4458	0.4461	0.4457	0.4469	0.445	0.4425	0.4471
SMC4	0.6742	0.6743	0.6758	0.6752	0.6747	0.6747	0.6742	0.6751	0.6763
NIPSNAP1	0.5525	0.5531	0.556	0.5541	0.5543	0.5523	0.5531	0.554	0.556
MYBL2	0.5776	0.5772	0.5784	0.5797	0.581	0.58	0.579	0.579	0.5808
ELAVL1	0.4433	0.4446	0.4426	0.4433	0.442	0.4431	0.4431	0.4407	0.4452
KIAA0100	0.4004	0.3996	0.3992	0.4009	0.4034	0.4	0.4011	0.401	0.4031
TP53	0.5886	0.59	0.5923	0.5911	0.5927	0.5922	0.5884	0.5924	0.5927
PSME2	0.5581	0.5575	0.5563	0.5562	0.5588	0.5583	0.5604	0.5575	0.5596
LBR	0.5088	0.5101	0.5081	0.509	0.5099	0.51	0.5102	0.5101	0.5113
SCCPDH	0.483	0.4821	0.4826	0.4841	0.4857	0.4835	0.4824	0.4848	0.4859
SMARCD2	0.4616	0.4631	0.4627	0.4627	0.4652	0.4633	0.4601	0.4614	0.4645
NET1	0.6266	0.6274	0.6264	0.6256	0.6262	0.6267	0.6273	0.6265	0.6278
HDAC2	0.5316	0.532	0.5327	0.533	0.5337	0.5335	0.5332	0.5354	0.5352
HSPB1	0.4742	0.4707	0.4694	0.4706	0.4726	0.471	0.4732	0.4725	0.4737
CDC25B	0.5267	0.5273	0.5265	0.5279	0.5293	0.5283	0.5265	0.5283	0.5291
ATMIN	0.483	0.483	0.4849	0.4824	0.4838	0.4817	0.4838	0.4817	0.4854
PSRC1	0.5967	0.5962	0.5966	0.5955	0.5982	0.5972	0.5968	0.5985	0.5986
UBE2A	0.4857	0.4837	0.4835	0.4842	0.4838	0.4822	0.484	0.4824	0.4862
COASY	0.4036	0.4032	0.4045	0.4052	0.4025	0.4042	0.4034	0.403	0.4055
BIRC5	0.7816	0.7814	0.7809	0.7826	0.7823	0.7825	0.7811	0.7821	0.7829
ARHGAP1	0.6153	0.6143	0.6141	0.6144	0.6143	0.6153	0.6142	0.6145	0.6161
CPNE3	0.4638	0.4657	0.4655	0.465	0.4661	0.4667	0.467	0.4646	0.4684
NUP62	0.536	0.5364	0.5373	0.5368	0.5392	0.5359	0.5345	0.5388	0.5387
PCM1	0.4049	0.4026	0.4068	0.4051	0.4067	0.4069	0.4042	0.404	0.4074
NUP93	0.4355	0.4336	0.4275	0.4284	0.4315	0.431	0.4346	0.4317	0.4347
PLK1	0.3966	0.3959	0.3959	0.3954	0.3962	0.3973	0.3969	0.397	0.398
CDK4	0.6708	0.67	0.67	0.6702	0.669	0.6698	0.6699	0.6695	0.6711
HSD17B10	0.5422	0.5426	0.5407	0.5419	0.5427	0.54	0.54	0.5409	0.5434
CSK	0.4011	0.4026	0.4022	0.4015	0.4019	0.4039	0.4017	0.4006	0.4044
TCEAL4	0.651	0.6514	0.6515	0.6506	0.6517	0.6514	0.6515	0.6518	0.6525
USP1	0.585	0.5839	0.5833	0.5844	0.5841	0.5843	0.5838	0.586	0.5858
MPC2	0.6136	0.6123	0.6156	0.6146	0.6151	0.6138	0.613	0.6162	0.6159
DECR1	0.5276	0.5292	0.5283	0.5287	0.5281	0.5278	0.5288	0.5267	0.5303
ERBB3	0.4996	0.5002	0.4997	0.5008	0.4995	0.5005	0.4997	0.5009	0.502
PRSS23	0.6763	0.6775	0.676	0.6768	0.6794	0.6782	0.6767	0.6787	0.6791
IKBKAP	0.3325	0.3346	0.332	0.3339	0.3352	0.3342	0.3336	0.3314	0.3361
GALE	0.7267	0.7261	0.7262	0.7258	0.7271	0.7267	0.7272	0.7259	0.7279
PRKCD	0.4411	0.4405	0.4401	0.4411	0.4393	0.4408	0.4404	0.4399	0.4424
S100A13	0.7024	0.7026	0.703	0.7043	0.7043	0.7036	0.7034	0.7044	0.7046
NRIP1	0.4643	0.462	0.4636	0.4653	0.4653	0.4646	0.4617	0.4643	0.4664
EAPP	0.2905	0.2906	0.2923	0.293	0.292	0.291	0.2927	0.2893	0.2941
POLR2K	0.5375	0.5371	0.536	0.5369	0.5364	0.5385	0.5356	0.5363	0.5387
NRAS	0.5165	0.5161	0.5188	0.5178	0.5191	0.5176	0.5177	0.5167	0.5198

SDHB	0.5161	0.5161	0.5192	0.5176	0.5175	0.5187	0.5157	0.5176	0.5196
GTF2A2	0.4871	0.487	0.4874	0.4872	0.4893	0.4891	0.4876	0.4868	0.4905
CCNB2	0.6618	0.6644	0.6629	0.6641	0.6632	0.6644	0.6632	0.6629	0.6651
LIG1	0.671	0.6715	0.672	0.6715	0.6721	0.6715	0.671	0.6708	0.6729
EBP	0.4182	0.4159	0.4161	0.4171	0.4161	0.4177	0.4165	0.4152	0.4191
WRB	0.5282	0.5294	0.53	0.5293	0.5307	0.5304	0.5288	0.5322	0.5312
SYNE2	0.3888	0.3883	0.3903	0.3901	0.3889	0.39	0.3891	0.3884	0.3914
OXCT1	0.6251	0.6266	0.6249	0.6246	0.6263	0.6256	0.6256	0.6258	0.6268
NNT	0.4925	0.4908	0.4909	0.4911	0.4928	0.4919	0.4921	0.4916	0.4939
GAA	0.5445	0.5456	0.545	0.5449	0.5441	0.546	0.5453	0.5447	0.5471
SLC37A4	0.4011	0.4023	0.4009	0.398	0.3999	0.4019	0.4023	0.403	0.403
HPRT1	0.4113	0.4112	0.4107	0.4131	0.4136	0.4127	0.4106	0.4121	0.4139
FAH	0.4888	0.4882	0.4877	0.4887	0.4891	0.4887	0.4896	0.4903	0.4914
POP4	0.5105	0.5122	0.5111	0.5096	0.512	0.511	0.5113	0.513	0.5132
CDC20	0.5349	0.5357	0.535	0.536	0.5377	0.5368	0.5363	0.5365	0.5377
NUP88	0.4804	0.479	0.4772	0.4769	0.4804	0.4791	0.4812	0.4818	0.4813
MSH6	0.4968	0.496	0.497	0.4976	0.4971	0.4982	0.4971	0.4958	0.4988
PIN1	0.3268	0.3296	0.3306	0.3283	0.3301	0.3305	0.3303	0.332	0.3318
ETFB	0.4607	0.4609	0.4634	0.4638	0.46	0.4645	0.4605	0.4636	0.4642
CRYZ	0.5211	0.5216	0.5237	0.5246	0.5226	0.523	0.5225	0.5231	0.5244
UBE2C	0.751	0.751	0.7524	0.7524	0.7538	0.7514	0.752	0.7512	0.7533
LYPLA1	0.2788	0.2769	0.2799	0.2776	0.2818	0.2806	0.2765	0.2791	0.2812
TIMELESS	0.7269	0.7257	0.7244	0.7242	0.7258	0.7247	0.7264	0.7233	0.7265
PAN2	0.2264	0.2277	0.2297	0.2256	0.2268	0.2329	0.2289	0.2283	0.231
RB1	0.4562	0.4552	0.4565	0.4575	0.4571	0.4567	0.4554	0.4571	0.4586
TBP	0.4453	0.4462	0.4446	0.4447	0.4469	0.4442	0.4464	0.4468	0.4476
HAT1	0.1991	0.2	0.202	0.2022	0.2002	0.1971	0.2022	0.1991	0.2031
S100A4	0.5006	0.5005	0.4998	0.5018	0.503	0.5022	0.5011	0.5016	0.5029
B4GAT1	0.61	0.6096	0.6135	0.6116	0.6095	0.6121	0.6098	0.6121	0.6124
ABCB6	0.4977	0.4975	0.4979	0.5002	0.5012	0.5018	0.4973	0.5012	0.5017
RFC5	0.4189	0.4213	0.422	0.4226	0.4229	0.4224	0.4202	0.4223	0.4232
CDK1	0.6971	0.6971	0.697	0.697	0.6971	0.6974	0.6963	0.6977	0.6984
TLE1	0.3466	0.3472	0.3487	0.3493	0.3509	0.3484	0.3492	0.3495	0.3511
PAFAH1B3	0.5619	0.5607	0.5606	0.5609	0.5606	0.5611	0.5609	0.5623	0.5629
DCK	0.4165	0.4176	0.4187	0.4189	0.4158	0.4184	0.4186	0.4174	0.4197
DYNLT3	0.3763	0.3784	0.377	0.3725	0.3763	0.3748	0.378	0.3787	0.3787
BAMBI	0.5989	0.6003	0.5989	0.5996	0.6024	0.5994	0.6007	0.601	0.6018
SLC35A1	0.673	0.6743	0.6753	0.6758	0.6755	0.6755	0.6746	0.6754	0.6762
ITGB1BP1	0.1526	0.1527	0.1585	0.1554	0.1534	0.1529	0.1557	0.1513	0.1565
TIMM17B	0.3699	0.371	0.3745	0.3745	0.3728	0.3751	0.3734	0.3708	0.3749
UGDH	0.2253	0.2263	0.2262	0.2245	0.2289	0.2296	0.2231	0.2264	0.229
EZH2	0.4902	0.4907	0.49	0.4895	0.4881	0.489	0.492	0.4913	0.4919
MYCBP	0.5828	0.5823	0.5831	0.5836	0.5837	0.5825	0.5819	0.5837	0.5847
HES1	0.4417	0.4432	0.4437	0.4411	0.4455	0.4416	0.4412	0.4443	0.4452
PSMG1	0.518	0.5205	0.5197	0.522	0.5214	0.5222	0.5188	0.5222	0.5224
DDB2	0.6351	0.6347	0.6324	0.6343	0.6361	0.6325	0.6351	0.6339	0.6359
CCNA2	0.7343	0.7343	0.7339	0.7332	0.7354	0.7332	0.7344	0.7348	0.7352

EML3	0.4451	0.4439	0.4479	0.4454	0.4462	0.4449	0.4438	0.4436	0.4475
PRAF2	0.5112	0.5123	0.5137	0.5136	0.5143	0.5128	0.5109	0.5124	0.5147
MRPL19	0.4781	0.4782	0.4777	0.4797	0.4789	0.4797	0.4786	0.4782	0.4802
MNAT1	0.3133	0.3149	0.3136	0.315	0.3153	0.3144	0.3151	0.3168	0.3165
RAB4A	0.3676	0.3711	0.369	0.3697	0.3682	0.3692	0.3719	0.3706	0.3721
BCL2	0.4264	0.4259	0.4286	0.4265	0.4266	0.4287	0.4263	0.4298	0.4292
RFC2	0.3452	0.3429	0.3434	0.3417	0.3466	0.3439	0.3448	0.3447	0.3463
SLC25A13	0.2891	0.2882	0.2884	0.2908	0.2885	0.2854	0.2877	0.2896	0.2906
MRPL12	0.5619	0.5623	0.5618	0.5602	0.563	0.5621	0.5636	0.5616	0.5641
CEBPA	0.4241	0.4251	0.4274	0.4287	0.429	0.4288	0.4269	0.4286	0.4297
SLC5A6	0.2991	0.2976	0.3005	0.2998	0.2971	0.2978	0.2993	0.3002	0.3013
AURKA	0.6638	0.6641	0.6651	0.6663	0.6653	0.6635	0.6644	0.6629	0.6659
CCNH	0.4035	0.4004	0.4032	0.4044	0.4041	0.4058	0.404	0.4082	0.4066
CDC45	0.2986	0.3022	0.3001	0.2987	0.3009	0.3	0.3016	0.2985	0.3028
ENOSF1	0.4783	0.4804	0.4807	0.4811	0.4821	0.4805	0.4801	0.4831	0.4823
CDK2	0.6066	0.6065	0.6067	0.6067	0.6052	0.6082	0.6067	0.6067	0.6081
ELOVL6	0.7138	0.714	0.7129	0.7128	0.7138	0.7112	0.7124	0.7132	0.7147
CREB1	0.1637	0.1672	0.1681	0.17	0.166	0.1699	0.1685	0.1676	0.1701
PPIC	0.414	0.4145	0.4123	0.4137	0.4131	0.4134	0.4127	0.4148	0.4157
BRCA1	0.4585	0.4601	0.4592	0.4598	0.4596	0.4608	0.4612	0.4584	0.4615
CCDC85B	0.3908	0.3912	0.3921	0.3931	0.3931	0.3937	0.3917	0.3938	0.3947
CCP110	0.3567	0.359	0.3604	0.3616	0.3584	0.3595	0.3587	0.3589	0.3613
CDC25A	0.4217	0.4217	0.4224	0.4253	0.4233	0.4229	0.4207	0.4221	0.425
NCK1	0.2086	0.2097	0.2069	0.206	0.2083	0.2055	0.2082	0.2073	0.2106
TMEM5	0.3733	0.3737	0.3726	0.3752	0.3738	0.3729	0.3746	0.3743	0.376
MELK	0.2727	0.2695	0.2689	0.2689	0.2722	0.2651	0.272	0.2705	0.272
CCNF	0.5626	0.5636	0.5615	0.5628	0.5643	0.5634	0.5645	0.5634	0.5649
TCFL5	0.3565	0.3545	0.3564	0.3558	0.3563	0.3562	0.357	0.359	0.3587
ICAM3	0.5135	0.5141	0.5145	0.5152	0.5134	0.5141	0.5147	0.5124	0.5162
ATF5	0.3238	0.3225	0.3216	0.3221	0.3224	0.3241	0.323	0.3241	0.325
CCNE2	0.2928	0.2921	0.289	0.2929	0.2927	0.2941	0.2949	0.291	0.2951
LSM6	0.5487	0.5495	0.5494	0.55	0.5491	0.5488	0.5487	0.5511	0.5508
ITGAE	0.347	0.3461	0.3455	0.3485	0.349	0.3495	0.3439	0.3444	0.3497
SGCB	0.239	0.2411	0.2433	0.2426	0.2402	0.2403	0.2412	0.246	0.2439
PLS1	0.335	0.3363	0.3343	0.3339	0.335	0.3324	0.3339	0.336	0.3369
PCMT1	0.5858	0.5852	0.5861	0.5871	0.5862	0.588	0.5852	0.5858	0.5876
INPP4B	0.4622	0.4615	0.4618	0.46	0.4647	0.4614	0.4621	0.4602	0.4643
CHEK1	0.3956	0.3983	0.3976	0.3972	0.3973	0.3988	0.3988	0.3945	0.3996
DAG1	0.4767	0.4777	0.4762	0.4756	0.4763	0.4768	0.4774	0.4761	0.4786
GHR	0.4814	0.4833	0.4806	0.481	0.4834	0.4821	0.4829	0.4795	0.4842
SCYL3	0.275	0.2756	0.2803	0.2792	0.2786	0.28	0.2755	0.2776	0.2806
POLG2	0.1879	0.1835	0.1868	0.1847	0.191	0.1868	0.1877	0.1879	0.1898
NOLC1	0.4819	0.4836	0.4845	0.4869	0.4837	0.4855	0.4838	0.4818	0.4867
POLE2	0.6244	0.626	0.6266	0.6266	0.6259	0.6261	0.6253	0.6265	0.6274
RAD51C	0.3238	0.3246	0.3263	0.3261	0.3233	0.3277	0.3261	0.3262	0.3273
KIF14	0.2685	0.2662	0.2674	0.2669	0.2653	0.2651	0.2688	0.2657	0.2687
SLC35A3	0.4422	0.4409	0.4397	0.4391	0.4408	0.4383	0.4402	0.4399	0.4419

AKT1	0.5895	0.5901	0.591	0.589	0.5899	0.5883	0.5903	0.5898	0.5915
MUC1	0.5587	0.5559	0.558	0.5583	0.5586	0.5584	0.5576	0.5582	0.5602
CASP2	0.2185	0.2206	0.2189	0.2216	0.2199	0.2207	0.2199	0.226	0.223
PARP1	0.5533	0.5515	0.5505	0.5524	0.5514	0.5495	0.5517	0.5516	0.5532
FDFT1	0.4287	0.4301	0.4282	0.4296	0.4325	0.427	0.4306	0.4278	0.4319
CCND1	0.3906	0.3914	0.3908	0.3928	0.3928	0.3923	0.394	0.3919	0.3946
OXA1L	0.1624	0.1622	0.1576	0.1586	0.163	0.1625	0.162	0.1648	0.1636
CDC42	0.3164	0.3202	0.3193	0.3186	0.3186	0.321	0.3227	0.3204	0.3222
ADH5	0.2575	0.2561	0.2576	0.26	0.256	0.2607	0.2555	0.2586	0.2605
CYCS	0.2857	0.286	0.2861	0.2862	0.2837	0.288	0.2847	0.2858	0.2879
COPS7A	0.3624	0.3647	0.3635	0.3635	0.3618	0.3641	0.3647	0.3631	0.366
GLOD4	0.5245	0.5264	0.5279	0.5279	0.5286	0.5289	0.5281	0.5277	0.5292
DLI	0.241	0.241	0.2402	0.2393	0.2384	0.2417	0.239	0.2414	0.2426
IFRD2	0.2813	0.2864	0.285	0.2842	0.2851	0.2846	0.2838	0.2841	0.2867
TSPAN6	0.3665	0.3705	0.3707	0.3731	0.3701	0.3707	0.37	0.3714	0.373
CDKN1B	0.2094	0.2047	0.2051	0.2026	0.202	0.2058	0.2044	0.2055	0.2072
PRPF4	0.2239	0.223	0.2223	0.2202	0.2212	0.222	0.225	0.223	0.2248
CBR1	0.4013	0.4015	0.4038	0.405	0.403	0.4033	0.4029	0.4059	0.4051
NR2F6	0.4343	0.432	0.4323	0.4315	0.4323	0.4335	0.4324	0.4328	0.4345
TSPAN4	0.5372	0.5391	0.5377	0.5397	0.5372	0.538	0.5401	0.5389	0.5402
PSIP1	0.237	0.2374	0.2364	0.2333	0.2372	0.2359	0.2378	0.2368	0.2393
AURKB	0.5254	0.5261	0.5261	0.5257	0.526	0.526	0.5266	0.5251	0.5277
RPA3	0.5603	0.5592	0.5614	0.5607	0.5601	0.5605	0.5601	0.5607	0.562
HDGFRP3	0.3709	0.3711	0.371	0.3701	0.3723	0.3717	0.3714	0.3687	0.3733
RRS1	0.2586	0.2616	0.2624	0.2587	0.2596	0.2588	0.2606	0.2596	0.2626
EED	0.4031	0.405	0.4047	0.4038	0.4039	0.405	0.4049	0.4043	0.4063
GATA3	0.6088	0.6087	0.6071	0.6079	0.6086	0.6086	0.6069	0.6101	0.6098
CETN3	0.4343	0.4348	0.4377	0.4389	0.4373	0.4391	0.4351	0.4366	0.4389
TGFB3	0.5191	0.5196	0.5181	0.519	0.518	0.519	0.5192	0.5196	0.5206
HIST1H2BK	0.4033	0.4024	0.4035	0.4026	0.4016	0.4035	0.4024	0.4002	0.4047
ANXA7	0.5195	0.5181	0.5193	0.5198	0.5177	0.5182	0.5202	0.5181	0.5211
PUF60	0.261	0.2588	0.2574	0.2573	0.2587	0.2558	0.2583	0.2586	0.261
CFLAR	0.3329	0.3354	0.3395	0.3417	0.3359	0.3382	0.3335	0.3349	0.3391
RPL39L	0.4126	0.415	0.4149	0.4139	0.4139	0.4152	0.4128	0.4137	0.4162
CHEK2	0.4908	0.4903	0.4917	0.4915	0.4937	0.4904	0.4905	0.4893	0.493
HMG20B	0.3731	0.373	0.3752	0.3761	0.3731	0.3764	0.3741	0.3738	0.3767
TPM1	0.5297	0.5299	0.5308	0.5307	0.5301	0.5294	0.5294	0.5307	0.5316
BDH1	0.7226	0.7229	0.7214	0.7219	0.7207	0.7208	0.7226	0.7221	0.7233
LSM5	0.3134	0.3122	0.3122	0.3116	0.3138	0.3125	0.3136	0.312	0.3146
TXNDC9	0.2033	0.2011	0.2026	0.2002	0.2005	0.201	0.2026	0.2007	0.2041
HN1L	0.5504	0.5505	0.5472	0.5479	0.5499	0.5477	0.5491	0.5506	0.5506
ZMIZ1	0.366	0.365	0.3686	0.3661	0.3655	0.3678	0.3675	0.3689	0.3688
PDS5A	0.4265	0.4274	0.4296	0.4308	0.4324	0.4276	0.4297	0.4285	0.431
XPO7	0.3139	0.3155	0.3139	0.3134	0.3165	0.3162	0.3142	0.3167	0.3175
COG4	0.4244	0.4251	0.4221	0.4229	0.4258	0.4225	0.4235	0.424	0.4261
TMEM97	0.4558	0.455	0.456	0.4537	0.4609	0.4533	0.4566	0.4584	0.4578
TFDP1	0.5556	0.5546	0.5547	0.5561	0.5567	0.5547	0.5562	0.5581	0.5576

KAT6B	0.3658	0.3681	0.3699	0.3669	0.3701	0.3676	0.3672	0.3635	0.3699
ADO	0.2842	0.2865	0.2875	0.2843	0.2823	0.2853	0.2851	0.2861	0.2876
JADE2	0.4849	0.4846	0.4833	0.4849	0.4843	0.4843	0.4855	0.4835	0.4866
PCCB	0.2735	0.2781	0.2745	0.2748	0.2738	0.2737	0.2773	0.2783	0.2776
KIAA1033	0.1856	0.1873	0.1903	0.1892	0.1857	0.1909	0.187	0.1927	0.1907
RRP1B	0.4777	0.4796	0.4814	0.4794	0.4806	0.4789	0.4785	0.4795	0.4818
THAP11	0.3153	0.3166	0.3177	0.3182	0.3197	0.3178	0.316	0.3186	0.3198
C2CD5	0.4357	0.4349	0.4368	0.4365	0.437	0.4355	0.4365	0.4374	0.4382
POLR2I	0.2334	0.2326	0.232	0.2334	0.2313	0.2338	0.2309	0.233	0.2346
RPIA	0.5189	0.5185	0.5213	0.5224	0.5194	0.5219	0.5184	0.5189	0.5219
PLCB3	0.4166	0.419	0.4201	0.4171	0.4185	0.4189	0.419	0.4185	0.4204
ASAH1	0.4465	0.4457	0.4453	0.4464	0.4487	0.4461	0.447	0.4455	0.4485
SMARCA4	0.5055	0.505	0.503	0.5021	0.5034	0.5032	0.5071	0.5017	0.506
ATP5S	0.4624	0.4602	0.4576	0.4576	0.4579	0.458	0.4588	0.4572	0.4609
PARP2	0.3808	0.3812	0.3824	0.3822	0.3837	0.3805	0.3819	0.3816	0.3838
AKR7A2	0.4009	0.3979	0.3997	0.3982	0.3989	0.3994	0.3985	0.4003	0.4013
SPDEF	0.7371	0.7376	0.7384	0.7386	0.7392	0.738	0.7381	0.7369	0.739
CCNB1	0.3837	0.386	0.384	0.3805	0.3843	0.3828	0.383	0.3825	0.3855
ERBB2	0.3681	0.3698	0.3679	0.3678	0.366	0.3679	0.3669	0.3687	0.3705
VDAC1	0.14	0.1411	0.1391	0.14	0.1419	0.1452	0.1408	0.1395	0.1437
GAPDH	0.2432	0.2428	0.2388	0.2422	0.2413	0.2428	0.2429	0.2396	0.244
CNDP2	0.3714	0.3729	0.3733	0.3742	0.3735	0.3734	0.3733	0.3717	0.3755
ADI1	0.3571	0.3573	0.3555	0.3543	0.3557	0.3538	0.3554	0.3547	0.3582
TMEM50A	0.532	0.5311	0.5318	0.5294	0.5331	0.5297	0.5319	0.5293	0.5332
HACD3	0.2863	0.2859	0.2865	0.2875	0.2876	0.2887	0.2877	0.2863	0.2897
YKT6	0.733	0.7339	0.7338	0.7321	0.734	0.7347	0.7333	0.7337	0.7351
BZW2	0.3537	0.3557	0.3565	0.356	0.3545	0.3575	0.3532	0.3566	0.3577
EVL	0.4192	0.4176	0.4194	0.4186	0.418	0.4179	0.4194	0.4196	0.4206
MIF	0.4915	0.4931	0.4957	0.4951	0.4928	0.4965	0.492	0.495	0.4956
IARS2	0.4712	0.4698	0.4701	0.4704	0.4728	0.4701	0.4704	0.4691	0.4721
KLHDC2	0.5697	0.5711	0.5693	0.5686	0.5677	0.5677	0.5717	0.5694	0.5713
STUB1	0.3342	0.3356	0.3338	0.3358	0.3376	0.3374	0.3347	0.3351	0.3377
ENOPH1	0.2858	0.2885	0.2917	0.2897	0.2867	0.2885	0.2875	0.2911	0.2906
MRPS2	0.2632	0.2639	0.2603	0.2603	0.258	0.2616	0.2634	0.2629	0.2642
NUP85	0.424	0.4248	0.4216	0.4252	0.4233	0.4274	0.4219	0.4246	0.4258
FIS1	0.2984	0.2943	0.2992	0.2989	0.2984	0.2992	0.3009	0.299	0.3008
NUSAP1	0.4675	0.4699	0.4706	0.4733	0.4698	0.471	0.4693	0.469	0.4723
MRPS16	0.3476	0.347	0.3484	0.3494	0.3435	0.3501	0.3469	0.3443	0.3495
NPDC1	0.2553	0.258	0.2593	0.2561	0.2577	0.256	0.2571	0.2608	0.2603
DERA	0.3533	0.3531	0.3556	0.3563	0.3533	0.3516	0.357	0.356	0.357
UBR7	0.4379	0.4389	0.438	0.4394	0.4374	0.4371	0.4375	0.4363	0.44
ISOC1	0.3885	0.3892	0.3882	0.39	0.3887	0.3867	0.3895	0.3898	0.3908
CCDC92	0.3958	0.3974	0.3968	0.3966	0.3972	0.3971	0.3966	0.396	0.3988
SCAND1	0.4478	0.4465	0.4466	0.4467	0.4461	0.4467	0.448	0.4467	0.4489
ZDHHC6	0.3468	0.3465	0.3439	0.3469	0.3475	0.347	0.3467	0.3441	0.3484
TIMM9	0.3882	0.3871	0.3832	0.3826	0.3856	0.3834	0.3869	0.382	0.3875
GMNN	0.547	0.5464	0.5503	0.5491	0.549	0.5488	0.5476	0.5474	0.5498

CDCA4	0.5363	0.5355	0.5365	0.5371	0.5375	0.5353	0.5358	0.534	0.538
NENF	0.4988	0.4968	0.4976	0.4989	0.4942	0.4986	0.4975	0.4956	0.4995
DNAJC15	0.3155	0.3166	0.317	0.3154	0.322	0.3152	0.3158	0.3155	0.3196
HEBP1	0.4344	0.4352	0.4367	0.4352	0.4371	0.4357	0.4368	0.4367	0.4376
CD320	0.3378	0.3387	0.3382	0.3371	0.3388	0.3377	0.3384	0.3396	0.3402
CHMP4A	0.3617	0.3616	0.3607	0.3628	0.3639	0.3616	0.365	0.3642	0.3649
HEATR1	0.2344	0.2352	0.2364	0.2343	0.2365	0.2355	0.2367	0.2337	0.2383
CISD1	0.5626	0.561	0.5621	0.563	0.5633	0.5613	0.5611	0.5625	0.5637
SUV39H1	0.4118	0.4106	0.4075	0.4102	0.4122	0.4097	0.4105	0.4097	0.4127
VPS28	0.337	0.3366	0.3326	0.3348	0.3386	0.3352	0.3362	0.3385	0.3379
KIF20A	0.5461	0.5452	0.5433	0.544	0.5458	0.5455	0.5441	0.5448	0.5467
METRNL	0.2418	0.2464	0.2441	0.2433	0.2415	0.2451	0.2421	0.2389	0.2461
LAGE3	0.5423	0.5429	0.5419	0.5408	0.5416	0.5424	0.5431	0.5409	0.5439
PXMP2	0.5113	0.5101	0.5115	0.5107	0.5098	0.5127	0.5116	0.5107	0.5126
AMDHD2	0.4398	0.4391	0.4399	0.4387	0.439	0.4383	0.4398	0.4418	0.4414
FBXO11	0.1828	0.1869	0.1878	0.1869	0.1877	0.1866	0.1882	0.1937	0.1895
MTERF3	0.4058	0.4035	0.4056	0.4037	0.4048	0.4042	0.4032	0.4029	0.406
PSMD10	0.3725	0.3724	0.3742	0.3708	0.3703	0.3707	0.3733	0.3739	0.3745
TSEN2	0.4133	0.4154	0.4179	0.4155	0.4165	0.4146	0.4152	0.4151	0.4173
PRR7	0.4656	0.4683	0.4693	0.4715	0.4686	0.4688	0.4658	0.4695	0.4703
GTPBP8	0.2136	0.2129	0.2159	0.2158	0.2135	0.2171	0.2134	0.2161	0.2171
DENND2D	0.3045	0.3033	0.3013	0.3023	0.3004	0.3019	0.3025	0.3008	0.3045
ARPP19	0.2112	0.2096	0.2116	0.2105	0.2101	0.2066	0.2083	0.2126	0.2128
ATG3	0.2906	0.2902	0.2919	0.2908	0.2909	0.2893	0.289	0.2883	0.293
WDR61	0.4329	0.432	0.4339	0.4348	0.4332	0.4323	0.4319	0.4373	0.4353
CANT1	0.2319	0.2347	0.231	0.2312	0.232	0.2324	0.2312	0.2325	0.2346
FAM63A	0.4159	0.4157	0.4125	0.4156	0.4148	0.4178	0.417	0.4208	0.4183
ATF1	0.4881	0.4888	0.4901	0.4905	0.4884	0.4902	0.4886	0.4901	0.4911
TSTA3	0.4897	0.4897	0.4908	0.491	0.4905	0.4925	0.4891	0.4917	0.4924

supplementary table 3. The top 5 genes participating in improving the predictions for each gene, the last column demonstrates the importance of the first gene.

index	gene 1	gene 2	gene 3	gene 4	gene 5	gene importance
DDR1	SCRN1	NNT	ERBB3	NIPSNAP1	BCL2	0.211
RHOA	SKP1	ELAVL1	SCP2	PAICS	NIPSNAP1	0.29
RNPS1	RRP1B	GLOD4	TMEM109	NUP62	SYPL1	0.194
RPS6	S100A13	SLC35A1	CPNE3	PSME1	NRAS	0.133
CSRP1	GALE	HSD17B10	GAPDH	STUB1	CHMP4A	0.072
XBP1	SPDEF	NET1	NRAS	HN1L	PRSS23	0.297
SKP1	NRAS	RHOA	LAGE3	ANXA7	TMEM50A	0.247
CAPN1	MUC1	THAP11	HIST1H2BK	S100A4	HES1	0.544
CTSD	AKT1	PRSS23	NPDC1	EML3	PRR7	0.279
MAT2A	CDC25A	AURKB	MSH6	TSEN2	NR2F6	0.234
STMN1	BIRC5	TIMELESS	EZH2	TOP2A	ANXA7	0.259
CIRBP	GAA	ARHGAP1	DDB2	TCEAL4	PRAF2	0.249
PSME1	DECR1	NIPSNAP1	PSME2	SKP1	NRAS	0.335
PGAM1	PAICS	HSD17B10	MYCBP	TFDP1	MIF	0.158
STAT1	UBE2A	NNT	ARHGAP1	SYNE2	RPA1	0.151
GNAS	GAA	SMARCA4	RPA1	NIPSNAP1	ENOSF1	0.082
HIF1A	CCNH	TCEAL4	RB1	TIMM17B	CTSD	0.196
PAICS	HDAC2	PSMG1	ICAM3	LAGE3	BDH1	0.336
HADH	SCP2	CISD1	CHEK2	OXCT1	S100A13	0.124
SMARCC1	MBNL1	SYNE2	HN1L	NOLC1	CNDP2	0.18
PIP4K2B	DAG1	NNT	B4GAT1	NIPSNAP1	GAA	0.201
PFKL	GAA	LIG1	ELAVL1	PARP1	MIF	0.124
PGRMC1	RPA1	SDHB	HN1L	UBE2A	ANXA7	0.308
MBNL1	SMC4	SMARCC1	LBR	GLOD4	HN1L	0.223
PCNA	CCNA2	PPIC	UBE2C	HPRT1	UGDH	0.288
ILK	TSTA3	NUDCD3	PSME2	S100A4	TPM1	0.101
SYPL1	PIP4K2B	NNT	PDS5A	RPS6	HDAC2	0.22
NUDCD3	ENOSF1	TSTA3	PIP4K2B	RAB4A	SKP1	0.165
TOP2A	UBE2C	CDK1	YKT6	KIF20A	SMC4	0.352
EBNA1BP2	SDHB	NOLC1	NUP93	RPIA	TPM1	0.293
SCP2	NNT	RHOA	CRYZ	NRAS	DECR1	0.24
TMEM109	TIMELESS	PAFAH1B3	MCM3	CDK4	RFC2	0.362
TRAP1	MRPL12	TSTA3	SMARCA4	RUVBL1	PAICS	0.114
SOX4	TLE1	NFKBIA	GATA3	NET1	BAMBI	0.147
SCRN1	DDR1	PSRC1	TPM1	RB1	CHEK1	0.245
NFKBIA	SOX4	DAG1	GAA	GATA3	HSD17B10	0.288
G3BP1	TMEM97	POLR2K	CPNE3	CDCA4	USP1	0.116
RPA1	PGRMC1	TIMELESS	SMARCA4	ATMIN	SYNE2	0.204
MCM3	TIMELESS	TMEM109	SMC4	CDK2	NUP85	0.578
SMC1A	XPO7	TIMELESS	OXCT1	MAT2A	GAPDH	0.067
PWP1	MRPL19	EZH2	TBP	NOLC1	EBNA1BP2	0.112
ICMT	TIMELESS	TMEM109	RPA1	PARP1	AKT1	0.171

RUVBL1	MYBL2	TFDP1	PAFAH1B3	SDHB	TSEN2	0.32
RRAGA	PRAF2	SCRN1	CCDC92	GALE	TPM1	0.094
SMC4	AURKA	CCNA2	YKT6	TOP2A	NUP62	0.413
NIPSNAP1	PSME1	BDH1	PIP4K2B	CDK4	RPA1	0.249
MYBL2	TMEM109	RUVBL1	TFDP1	BIRC5	TIMELESS	0.137
ELAVL1	PAICS	RHOA	HACD3	SCRN1	RB1	0.355
KIAA0100	DAG1	HSD17B10	ENOSF1	HN1L	GAA	0.084
TP53	ERBB3	CDK4	DDB2	NR2F6	JADE2	0.309
PSME2	PSME1	HSD17B10	SDHB	MYCBP	ILK	0.18
LBR	SMC4	AURKA	EZH2	HDAC2	PCMT1	0.106
SCCPDH	INPP4B	GAA	SDHB	ANXA7	DECR1	0.265
SMARCD2	TSTA3	COPS7A	COG4	SMARCA4	AKT1	0.184
NET1	XBP1	TP53	GATA3	BAMBI	GAA	0.357
HDAC2	PAICS	NRAS	POP4	GALE	PSMD10	0.277
HSPB1	S100A13	SMARCD2	MIF	RPS6	ILK	0.067
CDC25B	OXCT1	YKT6	TSPAN4	SLC35A1	MSH6	0.222
ATMIN	DAG1	NUP62	PCM1	RRP1B	PCMT1	0.051
PSRC1	BIRC5	AURKA	AURKB	SCRN1	YKT6	0.299
UBE2A	CPNE3	ANXA7	EBNA1BP2	PGRMC1	EAPP	0.26
COASY	MYCBP	TSTA3	SKP1	GAPDH	ICAM3	0.079
BIRC5	CCNA2	POP4	YKT6	NUSAP1	AURKA	0.51
ARHGAP1	AKT1	PLCB3	LIG1	TIMELESS	TMEM109	0.183
CPNE3	UBE2A	SKP1	PCMT1	ATF1	RPS6	0.252
NUP62	SMC4	JADE2	PAICS	HN1L	LIG1	0.345
PCM1	PDS5A	HN1L	NUP62	XPO7	ENOSF1	0.108
NUP93	EBNA1BP2	FAH	CCNH	GATA3	NOLC1	0.32
PLK1	BRCA1	AURKA	BIRC5	CDC20	KIF14	0.163
CDK4	ERBB3	BDH1	TMEM109	TP53	PSME1	0.274
HSD17B10	SDHB	SMARCA4	DAG1	NIPSNAP1	GALE	0.351
CSK	AKT1	CEBPA	COG4	MRPL12	PRKCD	0.168
TCEAL4	DECR1	NR2F6	ERBB3	CDK4	PSME1	0.184
USP1	CDCA4	SMC4	YKT6	ATF1	NUP62	0.313
MPC2	GTF2A2	MYCBP	ELOVL6	RPA3	ITGAE	0.313
DECR1	PSME1	TCEAL4	DDB2	TMEM50A	DYNLT3	0.279
ERBB3	NNT	PIP4K2B	GATA3	SOX4	KLHDC2	0.152
PRSS23	SPDEF	CTSD	XBP1	NRIP1	EZH2	0.33
IKBKAP	TIMELESS	RPA1	EZH2	AKT1	TSEN2	0.18
GALE	DDB2	CDC25B	ABCB6	TSPAN4	MRPL12	0.138
PRKCD	PIP4K2B	CSK	GAA	BAMBI	SYPL1	0.236
S100A13	ICMT	PSME1	TSPAN4	SLC35A1	DECR1	0.187
NRIP1	XBP1	C2CD5	SOX4	GATA3	PRSS23	0.266
EAPP	SLC35A1	ATF1	UBE2A	TSPAN6	PSME1	0.157
POLR2K	NRAS	SKP1	MYCBP	CRYZ	CCNH	0.318
NRAS	HDAC2	POLR2K	SKP1	CRYZ	LYPLA1	0.309
SDHB	ANXA7	HSD17B10	PGRMC1	PAICS	EBNA1BP2	0.316
GTF2A2	POP4	MPC2	CRYZ	SLC35A1	DNAJC15	0.357
CCNB2	CCNA2	BIRC5	TOP2A	AURKA	KIF20A	0.441

LIG1	ARHGAP1	TIMELESS	S100A13	GAA	RPA1	0.273
EBP	PAICS	FDFT1	LAGE3	TMEM97	ICAM3	0.262
WRB	PAICS	CISD1	PCMT1	MYCBP	NIPSNAP1	0.185
SYNE2	RPA1	SMARCC1	TIMELESS	MSH6	PDS5A	0.077
OXCT1	PARP1	TIMELESS	NENF	GHR	ELOVL6	0.278
NNT	SCP2	RPA1	TOP2A	SYPL1	PIP4K2B	0.277
GAA	PRAF2	KIAA0100	NPDC1	LIG1	PIP4K2B	0.307
SLC37A4	NUDCD3	CDK4	ARHGAP1	PRR7	WRB	0.147
HPRT1	DCK	NRAS	MYCBP	WRB	PAICS	0.123
FAH	PAICS	NUP93	LAGE3	HSD17B10	GAA	0.285
POP4	GTF2A2	HDAC2	NUP88	ANXA7	BIRC5	0.354
CDC20	AURKA	BIRC5	PLK1	YKT6	EBNA1BP2	0.357
NUP88	YKT6	EZH2	MRPL12	POP4	PARP2	0.166
MSH6	CDC25A	SMC4	TIMELESS	NUP62	PDS5A	0.176
PIN1	HSD17B10	GALE	PAFAH1B3	GHR	HMG20B	0.186
ETFB	PAFAH1B3	TSTA3	COG4	PIP4K2B	BDH1	0.288
CRYZ	NRAS	SLC35A1	ANXA7	POLR2K	DNAJC15	0.305
UBE2C	TOP2A	YKT6	CDK1	AURKA	CCNA2	0.467
LYPLA1	NRAS	SKP1	POLR2K	RPA3	ICAM3	0.154
TIMELESS	MCM3	RPA1	LIG1	BIRC5	TMEM109	0.423
PAN2	TP53	ERBB3	CDKN1B	PRPF4	POLG2	0.075
RB1	TGFB3	SCRN1	CISD1	CCNH	HADH	0.154
TBP	HDAC2	AURKA	ANXA7	ATF1	EAPP	0.098
HAT1	RPA3	POLR2K	DNAJC15	CYCS	TFDP1	0.11
S100A4	SLC35A3	ILK	MPC2	CCDC92	KLHDC2	0.102
B4GAT1	BDH1	ICAM3	ICMT	PRSS23	SPDEF	0.346
ABCB6	GALE	HSD17B10	ANXA7	PARP1	GAA	0.29
RFC5	BRCA1	BIRC5	POLE2	CDK1	PSMG1	0.154
CDK1	TOP2A	UBE2C	BIRC5	NRAS	CCNA2	0.508
TLE1	SOX4	ATMIN	ERBB2	GAA	GATA3	0.178
PAFAH1B3	ETFB	TMEM109	BDH1	BIRC5	LAGE3	0.292
DCK	HPRT1	STMN1	POLE2	CDK1	PSMG1	0.135
DYNLT3	DECR1	PSME1	POLR2K	CPNE3	B4GAT1	0.107
BAMBI	NET1	SOX4	KLHDC2	PRKCD	SLC35A1	0.304
SLC35A1	KLHDC2	S100A13	ELOVL6	ATF1	EAPP	0.17
ITGB1BP1	SDHB	ANXA7	TSTA3	SMARCD2	PCNA	0.073
TIMM17B	PSME2	RPL39L	ATG3	PRR7	HIF1A	0.169
UGDH	PCNA	SCCPDH	NET1	INPP4B	CCND1	0.042
EZH2	STMN1	CDC25A	NUP88	ZDHHC6	SMC4	0.203
MYCBP	SDHB	POLR2K	LAGE3	SKP1	MRPL12	0.181
HES1	KLHDC2	CCDC92	ZMIZ1	STMN1	ASAHI	0.141
PSMG1	PAICS	ANXA7	HDAC2	USP1	CDCA4	0.356
DDB2	GAA	DECR1	S100A13	CIRBP	DENND2D	0.284
CCNA2	BIRC5	SMC4	PAICS	AURKA	TOP2A	0.499
EML3	TMEM109	AKT1	CSK	FIS1	CTSD	0.197
PRAF2	GAA	NIPSNAP1	SKP1	ASAHI	PSME1	0.346
MRPL19	MRPL12	EBNA1BP2	SDHB	TMEM109	TIMELESS	0.123

MNAT1	POLR2K	RPA3	PCMT1	ATF1	NRAS	0.093
RAB4A	MYCBP	TMEM5	PSME1	NUDCD3	GAPDH	0.064
BCL2	SCRN1	DDR1	EVL	OXCT1	ICMT	0.141
RFC2	PAFAH1B3	TMEM109	MYBL2	NUP93	BIRC5	0.224
SLC25A13	SMARCA4	SMC4	OXCT1	MBNL1	PARP1	0.099
MRPL12	PAICS	GALE	SMC4	MYCBP	CCND1	0.342
CEBPA	MRPL12	CSK	GALE	DAG1	NUP62	0.102
SLC5A6	SMARCA4	GAPDH	CSK	MELK	SMARCD2	0.169
AURKA	SMC4	CCNA2	YKT6	CDC20	TOP2A	0.383
CCNH	CISD1	POLR2K	RB1	NUP93	ATG3	0.152
CDC45	CDK2	DERA	CDC25A	RUVBL1	POLE2	0.136
ENOSF1	SMARCA4	SDHB	PIP4K2B	GAA	PAICS	0.277
CDK2	MYBL2	CDK1	NUSAP1	ICMT	MCM3	0.142
ELOVL6	SPDEF	FDF1	CCNB2	MRPL19	HADH	0.428
CREB1	HDAC2	MTERF3	ITGAE	HES1	SCRN1	0.048
PPIC	MYCBP	PRAF2	PSMD10	RHOA	LIG1	0.159
BRCA1	SUV39H1	TIMELESS	MCM3	CDC25A	MYBL2	0.103
CCDC85B	TSTA3	TRAP1	CCND1	PRR7	ENOSF1	0.176
CCP110	MBNL1	SYNE2	TIMELESS	TMEM109	USP1	0.122
CDC25A	TIMELESS	BRCA1	TMEM109	SUV39H1	EED	0.256
NCK1	TBP	NRAS	ATF1	PSMD10	CDKN1B	0.121
TMEM5	PRR7	EBNA1BP2	RPS6	TSTA3	ENOSF1	0.079
MELK	USP1	SLC5A6	BIRC5	SMC4	CCNA2	0.088
CCNF	YKT6	CDCA4	B4GAT1	MSH6	PCMT1	0.456
TCFL5	GAA	PIP4K2B	TSPAN4	SKP1	CISD1	0.119
ICAM3	BDH1	PAICS	LAGE3	B4GAT1	MRPS16	0.186
ATF5	AURKA	PSMG1	MYBL2	MAT2A	TIMELESS	0.075
CCNE2	CDK2	CDC25A	CCNH	BRCA1	CDC45	0.121
LSM6	GMNN	S100A13	ATF1	RPA3	POLE2	0.198
ITGAE	GTF2A2	GMNN	CRYZ	ADI1	S100A4	0.176
SGCB	SKP1	POP4	RHOA	BIRC5	PFKL	0.101
PLS1	NNT	VPS28	CCDC92	TPM1	SLC35A3	0.143
PCMT1	GHR	POLR2K	LBR	CRYZ	SPDEF	0.093
INPP4B	SCCPDH	ERBB2	LBR	DENND2D	NET1	0.289
CHEK1	AURKB	OXCT1	BRCA1	HADH	SCRN1	0.137
DAG1	PIP4K2B	HN1L	KIAA0100	HSD17B10	SKP1	0.208
GHR	OXCT1	PCMT1	MPC2	DAG1	CEBPA	0.206
SCYL3	HIF1A	XPO7	PRR7	ANXA7	HEATR1	0.088
POLG2	CETN3	SLC35A1	CCDC92	CSK	PAN2	0.136
NOLC1	EBNA1BP2	TMEM109	TFDP1	NUP93	RRP1B	0.312
POLE2	CCNA2	GMNN	ATF1	CHEK2	POP4	0.42
RAD51C	PSME2	WRB	MYCBP	STMN1	OXCT1	0.106
KIF14	SMC4	PLK1	KIF20A	GHR	ITGAE	0.143
SLC35A3	RPL39L	OXCT1	SLC35A1	RB1	RPS6	0.167
AKT1	SMARCA4	TMEM109	RPA1	TIMELESS	AMDHD2	0.128
MUC1	CAPN1	THAP11	GALE	B4GAT1	GHR	0.583
CASP2	PGRMC1	PARP1	OXCT1	PAICS	TBP	0.069

PARP1	TIMELESS	ASAH1	ELAVL1	OXCT1	ARHGAP1	0.342
FDFT1	ELOVL6	EBP	PCMT1	BZW2	TMEM97	0.31
CCND1	MRPL12	SKP1	TPM1	PAICS	PFKL	0.22
OXA1L	PIP4K2B	SOX4	GAA	BDH1	NFKBIA	0.091
CDC42	MIF	PSME1	NENF	AKT1	POP4	0.153
ADH5	PSMD10	ICAM3	SKP1	RPA3	LYPLA1	0.107
CYCS	POLR2K	GTF2A2	TFDP1	DNAJC15	DYNLT3	0.112
COPS7A	SMARCD2	KLHDC2	CDK2	RPS6	FIS1	0.203
GLOD4	HN1L	SYPL1	MBNL1	HDAC2	XBP1	0.104
DLG	SDHB	MRPL19	ANXA7	IARS2	CPNE3	0.063
IFRD2	PAICS	POLR2I	NUP62	SDHB	RUVBL1	0.087
TSPAN6	SCP2	GTF2A2	EAPP	PSME1	SLC35A1	0.207
CDKN1B	ATF1	BAMBI	DDB2	TBP	LYPLA1	0.046
PRPF4	SDHB	TBP	STUB1	PARP2	LSM5	0.086
CBR1	HSD17B10	SDHB	POP4	COG4	PAICS	0.148
NR2F6	ERBB3	AURKB	ELOVL6	SPDEF	B4GAT1	0.099
TSPAN4	S100A13	LAGE3	ETFB	GAA	ENOSF1	0.353
PSIP1	TIMELESS	STMN1	PDS5A	MCM3	HADH	0.153
AURKB	CHEK1	CISD1	MYBL2	PSRC1	POLE2	0.299
RPA3	MYCBP	POLR2K	NRAS	MPC2	ICAM3	0.223
HDGFRP3	HDAC2	PAICS	NNT	SCCPDH	MBNL1	0.129
RRS1	RRP1B	PAICS	NUP62	PSMG1	AURKA	0.11
EED	USP1	CDC25A	YKT6	HDAC2	POP4	0.218
GATA3	NET1	XBP1	ERBB3	NUP93	SOX4	0.241
CETN3	TMEM50A	POLG2	DNAJC15	S100A13	HADH	0.103
TGFB3	OXCT1	RB1	BCL2	DDR1	SPDEF	0.13
HIST1H2BK	S100A13	HMG20B	GHR	CCDC92	DYNLT3	0.236
ANXA7	SDHB	SKP1	NRAS	CPNE3	TBP	0.37
PUF60	NUP88	RRP1B	COG4	BIRC5	HSD17B10	0.074
CFLAR	NRAS	RB1	CHEK1	AURKB	SCP2	0.177
RPL39L	SLC35A3	SKP1	PRR7	MYCBP	RPS6	0.152
CHEK2	CDCA4	ICAM3	YKT6	POLE2	TIMELESS	0.228
HMG20B	AKT1	CANT1	RPA1	AMDHD2	ICMT	0.076
TPM1	SMARCA4	EBNA1BP2	ENOSF1	TCEAL4	SCRN1	0.256
BDH1	MCM3	TIMELESS	B4GAT1	ICAM3	ERBB3	0.225
LSM5	POLR2K	CYCS	ATG3	PCMT1	MRPL19	0.117
TXNDC9	CRYZ	MPC2	GTPBP8	EAPP	DERA	0.098
HN1L	PGRMC1	PAICS	KIAA0100	NUP62	DAG1	0.223
ZMIZ1	ARHGAP1	SCRN1	VPS28	KAT6B	GLOD4	0.16
PDS5A	PCM1	NNT	SYNE2	SMC4	SCRN1	0.208
XPO7	PCM1	HEATR1	ATG3	HN1L	MBNL1	0.142
COG4	HSD17B10	CSK	ETFB	LIG1	CEBPA	0.212
TMEM97	CDCA4	EBP	UBE2C	FDFT1	MCM3	0.174
TFDP1	MYBL2	SDHB	NOLC1	PAICS	CDK2	0.313
KAT6B	SYNE2	EVL	ZMIZ1	BCL2	SLC37A4	0.096
ADO	SMARCA4	GAA	PCM1	SUV39H1	ATMIN	0.123
JADE2	NUP62	AURKA	FAM63A	TP53	PIP4K2B	0.357

PCCB	NIPSNAP1	HMG20B	PSME1	ADI1	STUB1	0.101
KIAA1033	ATMIN	RPS6	SLC35A1	ADI1	TLE1	0.049
RRP1B	SMC4	SMARCA4	NOLC1	EBNA1BP2	RPIA	0.231
THAP11	RUVBL1	GHR	MUC1	PCMT1	LBR	0.079
C2CD5	SPDEF	SLC35A1	NRIP1	GATA3	TOP2A	0.273
POLR2I	MIF	PAFAH1B3	ETFB	SDHB	IFRD2	0.068
RPIA	TMEM109	PAICS	EBNA1BP2	RUVBL1	ENOSF1	0.161
PLCB3	SMARCA4	CCDC92	ARHGAP1	TGFB3	AMDHD2	0.198
ASAH1	PRAF2	GAA	OXCT1	STAT1	PARP1	0.144
SMARCA4	RPA1	HSD17B10	NUP62	ENOSF1	RRP1B	0.322
ATP5S	SKP1	PSME1	RPA3	DECR1	ATF1	0.146
PARP2	USP1	NUP88	SMC4	AURKA	YKT6	0.157
AKR7A2	S100A13	CDK2	PRAF2	DECR1	MYCBP	0.176
SPDEF	ELOVL6	SLC35A1	XBP1	ARHGAP1	TCEAL4	0.204
CCNB1	CCNA2	AURKA	MAT2A	AURKB	PSRC1	0.197
ERBB2	GAA	TLE1	DDB2	ERBB3	GAPDH	0.217
VDAC1	EBNA1BP2	GAPDH	SDHB	MYCBP	CISD1	0.06
GAPDH	DAG1	TRAP1	COASY	CSK	ATG3	0.06
CNDP2	GAA	HN1L	TIMELESS	SMARCC1	ARHGAP1	0.16
ADI1	MYCBP	STMN1	ANXA7	BDH1	TFDP1	0.095
TMEM50A	SKP1	DECR1	TCEAL4	KLHDC2	CDC25B	0.316
HACD3	CISD1	AURKB	HADH	RB1	TFDP1	0.147
YKT6	BIRC5	UBE2C	AURKA	CCNF	TIMELESS	0.433
BZW2	SKP1	PRR7	TSTA3	PSME1	FDFT1	0.152
EVL	PAFAH1B3	AMDHD2	GAA	NNT	BCL2	0.152
MIF	PSME1	SKP1	BIRC5	LAGE3	MRPL12	0.278
IARS2	TIMELESS	ICMT	PCMT1	CDK4	WRB	0.18
KLHDC2	SLC35A1	ERBB3	HES1	CCDC92	BAMBI	0.372
STUB1	HSD17B10	AKT1	CHMP4A	PAFAH1B3	HMG20B	0.118
ENOPH1	SDHB	ATG3	PSMG1	PAICS	SMC4	0.101
MRPS2	MRPL12	TMEM109	CSK	RRP1B	NUP88	0.058
NUP85	TIMELESS	MCM3	TMEM109	PARP2	EZH2	0.141
FIS1	TSPAN6	HMG20B	EML3	KLHDC2	COPS7A	0.111
NUSAP1	TOP2A	CDK1	DERA	CDK2	CDC45	0.219
MRPS16	ICAM3	LAGE3	CDK4	PSME1	ATP5S	0.224
NPDC1	GAA	TSTA3	CTSD	EVL	EML3	0.212
DERA	GMNN	CDK2	NUSAP1	CDC45	MPC2	0.104
UBR7	PCMT1	SMC4	USP1	CDCA4	CCP110	0.109
ISOC1	MPC2	GHR	TIMELESS	ELOVL6	IARS2	0.202
CCDC92	GAA	RRAGA	PLCB3	HES1	GHR	0.163
SCAND1	DDB2	TCEAL4	HSPB1	PSME1	CIRBP	0.169
ZDHHC6	PAICS	EZH2	TFDP1	PCMT1	PARP2	0.186
TIMM9	ATF1	POLE2	NUP62	XBP1	TSEN2	0.122
GMNN	POLE2	BIRC5	GTF2A2	RPA3	AURKB	0.348
CDCA4	USP1	CHEK2	CCNF	LAGE3	ICAM3	0.331
NENF	PSME1	TSTA3	TSEN2	CDC42	OXCT1	0.292
DNAJC15	CRYZ	GTF2A2	RPA3	CETN3	POLR2K	0.2

HEBP1	SPDEF	GHR	CIRBP	ARHGAP1	TCEAL4	0.111
CD320	EBNA1BP2	SDHB	SMARCA4	HSD17B10	PGRMC1	0.084
CHMP4A	TPM1	SMARCA4	PCMT1	TCEAL4	CISD1	0.14
HEATR1	XPO7	SMARCC1	SCYL3	NOLC1	SOX4	0.171
CISD1	CCNH	AURKB	BDH1	RB1	SLC35A3	0.288
SUV39H1	AURKA	CDC25A	YKT6	BRCA1	ADO	0.197
VPS28	ENOSF1	SLC35A3	SMARCC1	ZMIZ1	SYNE2	0.114
KIF20A	CCNA2	AURKA	BIRC5	SMC4	TOP2A	0.244
METRNL	AKT1	PRR7	MYCBP	MELK	NUDCD3	0.092
LAGE3	ICAM3	PAFAH1B3	MYCBP	MRPS16	S100A13	0.242
PXMP2	PSME1	S100A13	MYCBP	OXCT1	TIMELESS	0.102
AMDHD2	GAA	PAFAH1B3	GALE	HSPB1	CCDC92	0.2
FBXO11	CASP2	CHEK1	HADH	RHOA	MCM3	0.051
MTERF3	POLE2	TSEN2	CCNA2	SLC35A1	POLR2K	0.141
PSMD10	SKP1	NRAS	HDAC2	RHOA	ADH5	0.218
TSEN2	TMEM109	MRPL19	CDC25A	MAT2A	MRPL12	0.119
PRR7	TSTA3	NUDCD3	ENOSF1	SDHB	CISD1	0.151
GTPBP8	CRYZ	TIMM9	SLC35A1	CYCS	TXNDC9	0.065
DENND2D	DDB2	INPP4B	DECR1	SCRN1	SYPL1	0.152
ARPP19	NRAS	CEBPA	RPS6	ATF1	CBR1	0.109
ATG3	CCNH	EBNA1BP2	EZH2	POLR2K	ENOPH1	0.093
WDR61	MYCBP	ATP5S	RPA3	POP4	CRYZ	0.27
CANT1	KIF20A	HMG20B	TLE1	MRPS2	NUP62	0.088
FAM63A	NNT	DDB2	GATA3	DDR1	TCEAL4	0.217
ATF1	NRAS	SLC35A1	POLE2	TBP	USP1	0.191
TSTA3	ETFB	GAA	MYCBP	TRAP1	COG4	0.153

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