

Table S1: The genes that significantly related to XYZ phenotype ($p < 0.05$) at Day 14

gene	p	gene	p	gene	p	gene	p	gene	p
MIR147B	0.004631	MARK2	0.007058	GOLGA6A	0.014295	LOC554223	0.015127	CLIP2	0.009233
FAM229A	0.008245	TSNARE1	0.001362	SLC10A5	0.017705	NOP10	0.005657	XBP1	0.005855
MED16	0.013121	VMA21	0.001847	EP400NL	0.007039	BATF3	0.011038	MIR671	0.017784
MIR4326	0.020243	AK2	0.019519	APOE	0.018332	MAML3	0.022283	ZNF180	0.001379
MIR3615	0.011032	EPHB4	0.013341	SKA2	0.003234	GOLGA2P5	0.018673	ZNF230	0.019404
LOC100505658	0.008709	EPHB6	0.009424	FAM159A	0.019032	MAP7D1	0.020876	TMEM106C	0.009529
DCDC5	0.013125	TMEM218	0.006845	IL4R	0.013165	PARVA	0.015672	NUP37	0.014656
TMEM44-AS1	0.01706	ALKBH3	0.000224	ILF2	0.014656	WDR12	0.01956	C1orf50	0.009431
MIR4635	0.014911	PHACTR1	0.000462	INHBB	0.001036	ECHDC1	0.000995	THOC6	0.007396
EGFLAM-AS4	0.0153	CCZ1B	0.016652	INPP5A	0.004129	LMO3	0.020593	ZXDC	0.004855
PRICKLE2-AS3	0.011201	LRWD1	0.00192	CD82	0.006009	UNC45A	0.021492	CCNJL	0.017803
SERF2	0.002389	GPR97	0.013736	KCNA5	0.012903	APOBR	0.001563	AIMP2	0.019991
GDF11	0.015721	ZSCAN23	0.011623	B3GNT8	0.005683	SLC25A40	0.015345	CALCB	0.012864
SPRY2	0.007861	FGF11	0.002819	APITD1	0.013514	PCDHGA11	0.003925	MAFK	0.005413
ABCC4	0.015363	FGFR4	0.000672	ACAT1	0.006963	PSPN	0.005017	TREML2	0.00762
SPEG	0.016244	COPZ1	0.014402	KLRB1	0.013426	SUSD2	0.007824	ZNF672	0.009208
HMG20B	0.022647	ZFP30	0.006282	C2orf82	0.001386	PANX2	0.015518	WDR76	0.020323
ANAPC10	0.005043	LMTK2	0.023037	CDHR4	0.008757	TMEM9B	0.021714	THOC7	0.00898
ADARB1	0.019061	PPP6R1	0.013295	PROB1	0.002045	PSMA6	0.018636	PUS1	0.007775
CD52	0.009301	SBNO2	0.014799	RRRC26	0.002252	C15orf39	0.012121	VWA7	0.01505
HAX1	0.006178	KDM4B	0.009431	ARL5C	0.006253	TOMM22	0.002075	B9D2	0.001766
TACC3	0.021022	KDM6B	0.001929	LAMC1	0.004907	SLC12A9	0.008921	PABPN1	0.017566
SLC30A9	0.004902	METAP1	0.002854	RPSA	0.011444	CTNNBIP1	0.011245	NETO1	0.017831

SEMA4B	0.002347	JMJD6	0.015296	LGALS1	0.000776	INTS12	0.00243	ZNF205-AS1	0.015283
PROCR	0.008165	PHLPP1	0.003522	ARHGDIG	0.021795	CAMK1D	0.0125	CAPZA1	0.022783
SH2B2	0.003367	BICD2	0.002973	OR2A20P	0.002584	SMURF1	0.019966	DYSF	0.018578
MYL12A	0.019162	ICOSLG	0.001659	ARL3	0.018774	PSMD10	0.004772	FZD7	0.023267
RFPL2	0.010585	DPY19L1	0.01976	MIRLET7B	0.001959	SMAGP	0.018452	ADPGK	0.006703
KDM5B	0.013121	ZNF629	0.021046	TBC1D3C	0.006296	CLK4	0.017819	SF3B5	0.007665
NOXA1	0.017598	ARHGEF18	0.004343	PLEKHG7	0.014346	COG6	0.014296	APH1B	0.008023
CPLX1	0.001277	COMM3D	0.002145	NUDT4P1	0.011463	WDFY4	0.022377	RIOK1	0.005116
HCP5	0.006543	ZFPM2	0.010986	SMIM4	0.004406	PRX	0.006774	MAD1L1	0.000932
EHD1	0.021701	ETHE1	0.012489	NUTM2G	0.003776	GRHL3	0.00571	STARD3NL	0.012646
TMED2	0.020624	RIMBP2	0.020788	MYH3	0.021661	PTPRK	0.018731	TMEM133	0.005172
UQCR11	0.010242	TNPO3	0.008814	MYL4	0.004979	PVRL1	0.0015	TAGLN2	0.01976
CLP1	0.006339	ZKSCAN5	0.012266	MYL6	0.023126	CXCL16	0.016495	ENKD1	0.000632
KDELR2	0.001537	EID1	0.010741	HNRRNPM	0.019728	RAB6A	0.003097	ELMOD3	0.006059
SMA5	0.005312	MTCH2	0.01507	NDUFA9	0.005121	RABIF	0.010209	PHF6	0.016656
WWP2	0.01928	FTH1P3	0.015157	NDUFS3	0.022163	RAN	0.005683	MIEN1	0.015184
ADAM29	0.006007	EPHX4	0.013609	NDUFS6	0.017621	RARA	0.015627	UTF1	0.009879
ADAMTS6	0.014751	EHBPI1L1	0.004458	RERE	0.010814	RXFP1	0.020589	C14orf142	0.000549
PADI2	0.020004	GABRA2	0.008003	NFATC1	0.016799	RELB	0.017184	KBTBD8	0.002484
PAC SIN2	0.006733	ALX3	0.019149	NFKB1	0.005641	RNASE2	0.008554	TMEM185A	0.002342
C1orf85	0.021984	TBC1D22A	0.018292	NOTCH1	0.02292	RNASE6	0.015478	TNRC18	0.023057
MGAT4B	0.000168	PRDX5	0.01713	PNP	0.001364	BCL6	0.018636	HINT2	0.020065
HENMT1	0.011009	ANAPC13	0.020255	ROR1	0.003804	RPE	0.007212	MCEE	0.017173
ATG4A	0.010982	SPEF1	0.004187	COA5	0.006956	RPL10	0.003264	CAPS2	0.015014
ZNF501	0.014479	NIPSNAP3A	2.88E-05	CHAC2	0.012513	RPL12	0.011961	HDGFRP2	0.014388
WDR92	0.017238	NPHP4	0.003371	P2RX5	0.021003	RPL15	0.008736	CBX2	0.007129

MRPL54	0.012908	MYCBP	0.022824	PA2G4	0.003694	RPL18A	0.023057	SFT2D3	0.010018
ARAP1	0.005683	GBAS	0.009366	PAX5	0.012856	RPL30	0.01295	TMEM25	0.012438
CLCN4	0.014449	BLOC1S1	0.013781	PBX2	0.015778	RPLP0	0.007212	GEMIN2	0.005363
CHCHD1	0.001189	SLC17A5	0.007915	ATP5C1	0.013426	RPS3	0.016757	RAB2B	0.01538
CPXM2	0.011169	CNNM1	0.014752	FIS1	0.000213	RPS5	0.011763	LSM10	0.017566
CLTA	0.018752	PGBD2	0.019542	TXNDC12	0.0003	RPS6	0.017184	COX14	0.014741
GTSF1	0.004125	SNORD50A	0.015979	APH1A	0.000842	RPS11	0.005434	TMEM60	0.002269
ANKRD9	0.007258	AMPD2	0.011952	SS18L2	0.015778	RPS14	0.003097	SNORD14C	0.022023
ADSSL1	0.012665	FAM184B	0.013978	NUSAP1	0.01332	RPS20	0.005434	SCIN	0.018753
NTAN1	0.002175	NFU1	0.013558	TMEM216	0.015387	RPS27	0.017898	DNAH17	0.004158
PAQR4	0.0142	ERLEC1	0.007909	COA4	0.014236	BDNF	0.022225	AOC3	0.007194
SLC38A10	0.016031	CSDC2	0.021434	PHF21A	0.009095	SAFB	0.002024	DYNLL1	0.013871
TBC1D16	0.004897	GLO1	0.012283	TMA7	0.006831	BGN	0.020155	IRS2	0.005495
ALDH16A1	0.010888	CECR2	0.018271	ATRAID	0.017039	UBE2O	0.010593	EIF3I	0.009941
ZNF57	0.005116	GNB3	0.008725	TRMT112	0.022512	ANKEF1	0.023251	VAMP8	0.000471
PLK3	0.007839	LOC283038	0.021043	UFC1	0.014515	MRPS14	0.001396	ACTN1	0.009269
KLHDC7A	0.002283	CCDC88B	0.010242	CHMP5	0.014515	C19orf33	0.019512	TNFSF13	0.01697
COL6A3	0.018191	TTC9C	0.004763	CUTA	0.004813	BLVRB	0.005	CBS	0.016029
AASDH	0.011424	HARBI1	0.00327	ATP5G1	0.013571	SH3GL1	0.012697	TNFRSF10C	0.021583
COX6B1	0.011111	SUGT1P3	0.015212	LIPT1	0.009205	PLA2G2F	0.016278	PEX11B	0.009865
COX7B	0.015528	PRTG	0.021222	ATP6V1H	0.004288	CSRNP1	0.017065	BANF1	0.004288
C7orf31	0.010866	HID1	0.004194	FAM96B	0.003277	LOC646938	0.000525	ST3GAL5	0.005705
HDX	0.022505	UBALD2	0.00421	PTRH2	0.015222	ST3GAL2	0.01909	EIF2B2	0.005748
FUND1	0.007404	HMSD	0.020175	ASB1	0.022519	TTC23	0.006601	BUD31	0.016442
SPIN4	0.020635	EPHA10	0.01561	VPS29	0.020423	MRPS24	0.023265	NAV2	0.016948
UBE2F	0.008837	LOC285074	0.015233	SELT	0.001027	MRPS15	0.016985	MVB12B	0.001728

CSF2RB	0.01881	RABL3	0.015212	RTEL1	0.022451	MRPL32	0.013338	UBE3D	0.012379
CSNK1G2	0.012993	C5orf51	0.019705	TM7SF3	0.010418	PINK1	0.021246	CCDC97	0.021317
CSPG4	0.008714	FFAR2	0.014751	PENK	0.00297	JMJD4	0.013874	TMEM263	0.022077
PRSS36	0.017935	GRK6	0.008554	PEX7	0.018873	KIAA0895L	0.01419	NFS1	0.006172
CCDC42	0.000246	GRB10	0.006327	PFKFB3	0.002683	PAX8-AS1	0.01114	SART1	0.015994
RUNDIC1	0.013736	MAGEH1	0.002379	PFKL	0.019162	SLC11A1	0.019044	RQCD1	0.002274
NLRP4	0.006589	MRPL15	0.011813	ATP5J	0.00081	FUNDC2	0.020698	SYNGR1	0.018538
C22orf42	0.020743	STXBP6	0.023023	PGM5	0.022043	UAP1	0.019256	CD3D	0.00911
MSL3P1	0.005238	GSN	0.009999	PIK3CD	0.004668	SSR2	0.000541	ZNF765	0.016998
CYP1A1	0.014111	GTF2E1	0.012124	STX18	0.014125	STAT5A	0.021911	COX7A2L	0.016599
CCDC167	0.020206	UBN1	0.003694	PMM2	0.015482	STAT5B	0.006635	TMSB10	0.001352
CLEC2L	0.022283	TIMM22	0.000164	ATP5O	0.014468	SNORA54	0.002557	IL1RL1	0.02292
CRYGN	0.002613	NENF	0.018054	RIPK4	0.008022	SURF2	0.020517	PTTG1	0.014043
WBSCR27	0.010799	OSGIN1	0.015398	POLR2I	0.002347	TBCE	0.004497	STRADA	0.017461
ADRBK1	0.011186	FTSJ2	0.003251	POLR2J	0.013692	TCEB2	0.004029	IMP4	0.021668
CYP2F1	0.019925	ANXA1	0.020939	POLR2K	0.005077	PPAN-P2RY11	0.015468	ATP6V1F	0.006037
TTC16	0.016291	HBG2	0.008554	NLE1	0.021977	MIR572	0.015996	NT5C1B	0.002168
CD55	0.015878	HIP1	0.007769	RNF186	0.021105	MIR619	0.021003	LOC93432	0.005099
ZFP1	0.015567	HMBS	0.004033	GNB1L	0.018179	ZEB1	0.020939	SDR42E1	0.0056
ZNF433	0.013041	HMGB2	0.02178	FBXL12	0.012823	DYNLT1	0.022783	CRIPTR	0.009431
DDT	0.002807	NR4A1	0.017527	TMEM70	0.00546	TEF	0.012306	PGS1	0.008859
NLRP6	0.008593	PRMT1	0.00474	PPP1R1A	0.02044	TLE3	0.007167	AATK	0.010357
ARID3A	0.013704	HSD17B3	0.013787	SOBP	0.006578	TLN1	0.021518	ZNF592	0.014468
IMMP1L	8.83E-05	HSPA1B	0.023057	ATP5SL	0.006082	GOLGA6L7P	0.000734	CROCC	0.000842
OR56B4	0.007385	HSPA8	0.018868	SHQ1	0.010392	RGPD6	0.01581	TBKBP1	0.015493
EIF2S1	0.00167	HTR2A	0.017178	TRIM62	0.019392	UBA52	0.001823	RNF40	0.01538

SNAI3-AS1	0.021195	C17orf67	0.018394	NUDT15	0.012646	UBB	0.001626	ARHGEF11	0.023265
APLF	0.003043	CCDC158	0.015409	CWF19L1	0.012823	VAV1	0.016235	GAB2	0.015578
TIGD1	0.016878	IFITM4P	0.022318	SLC39A9	0.0066	VCP	0.005948	RBM8A	0.00251
KCTD6	0.016974	SMCO2	0.019715	LGR4	0.012015	VWF	0.022512	CDC34	0.011215

Table S2: The genes that significantly related to XYZ genome ($p < 0.05$) at Day 30

Gene	ρ	Gene	ρ	Gene	ρ
SNORD12A	0.022687	TIMM9	0.008581	SMAP1	0.018805
C10orf131	0.021325	RNU4-2	0.012015	RPL36A	0.015737
TMEM221	0.000959	GLUD1	0.021323	MRPS12	0.013176
SMIM6	0.010393	GNG3	0.009899	ACSM3	0.009518
CLUHP3	0.006241	C14orf177	0.017721	CELA2A	0.004952
REXO1L2P	0.002949	FAM171A2	0.004115	ATXN2	0.020163
MIR4313	0.001961	IAH1	0.022706	SCP2	0.014205
MIR3685	0.015154	ZNF252P-AS1	0.018079	VIPAS39	0.02253
FGD5-AS1	0.007342	ILDR1	0.02258	SDHB	0.001482
LINC00888	0.01673	GRIN1	0.008271	RBKS	0.01839
INMT-FAM188B	0.007612	ZC3H7A	0.017048	SMOC2	0.008825
BLOC1S5-TXND5	0.017909	LINC00652	0.007612	CEP170P1	0.011829
MIA-RAB4B	0.000942	DROSHA	0.00647	CLEC7A	0.010494
FAM13A	0.021015	SNX12	0.017593	LOC646938	0.018382
CDK2	0.020831	HDGF	0.01641	ARMCX5	0.018626
RBM7	0.004009	HLA-J	0.002983	C8orf33	0.019355
TESK2	0.009258	HMGN2	0.004091	SLC9A5	0.022161
VAV3	0.019544	FOXA3	0.000585	SLC18A2	0.014318
GPNMB	0.009336	HNRNPC	0.00885	SLC20A2	0.019433
VTI1B	0.007879	HOXC13	0.011786	SMARCE1	0.005464
SLC19A2	0.012598	HPN	0.004866	SPOCK1	0.005391
CENPA	0.00075	TFAP2E	0.02012	SREBF2	0.018261
TRIM3	0.006108	CCDC36	0.012603	SCARNA15	0.01867
TXNIP	0.013855	KY	0.013519	SNORA81	0.005417
IGF2BP1	0.010483	ZKSCAN2	0.017445	AURKA	0.009741
MTX2	0.014553	XKR7	0.003245	SULT1A2	0.009496
EBP	0.011857	ACTBL2	0.007648	KLF9	0.018765
TCFL5	0.007714	IFNGR1	0.02001	TAF7	0.01294
IQGAP2	0.011051	FAM71F2	0.006358	SNORD86	0.00425
YWHAQ	0.009479	SOX2-OT	0.020383	SNORD89	0.005537
DSTN	0.016814	IGFBP7	0.011762	DYNLT3	0.018125
RCC1	0.018901	ILF2	0.015194	TEC	0.014156
PRDM5	0.020594	IMPG1	0.008902	TFDP1	0.014823
ZWINT	0.010342	KCNJ12	0.014996	C1QBP	0.003782
HHLA2	0.015943	KIF3C	0.011667	TMOD1	0.008758
ADCY6	0.015846	KRT10	0.002683	KRTAP19-8	0.006519
RNF139	0.022642	LRRC75A	0.004114	TUFM	0.009624
GTF3C6	0.000856	LINC00478	0.001915	OSGIN2	0.014715
CYP4F8	0.023154	TMPRSS11F	0.022089	UFD1L	0.012791
LYST	0.023158	OR52H1	0.019914	UGCG	0.013415
DDX42	0.008224	CTXN2	0.003287	KDM6A	0.001656

PDAP1	0.021275	LINC00938	0.011762	ZNF131	0.022581
CHRNA3	0.000812	LMX1B	0.018139	DDX54	0.010974
PIK3IP1	0.023082	MBNL1-AS1	0.006468	C2orf54	0.00866
KLHL32	0.01381	RGS7BP	0.010452	SEMA6D	0.000613
OSBPL9	0.010763	MIR7-1	0.003802	PIF1	0.021751
FBXO32	0.002145	EPCAM	0.022718	RPF1	0.016467
SLC26A7	0.010703	SMAD7	0.020945	EPC1	0.010338
UHRF2	0.009055	MAGOAH	0.012708	ZSCAN16	0.003219
MED12L	0.006014	CD46	0.019743	PTP4A2	0.011814
CATSPER1	0.021739	SMIM22	0.014961	ANKRD13C	0.015552
CLC	0.020078	HIST2H2BF	0.01513	CAMK2B	0.004038
DCUN1D3	0.0083	MYBPC1	0.001916	COIL	0.015754
WFIKKN2	0.019259	NDUFS1	0.016183	CAB39L	0.00715
ATP6V1G3	0.00378	NPTX1	0.01587	ISCA1	0.021111
SYT2	0.011438	SERPINE1	0.012512	GDF5	0.02091
SPATA17	0.002983	COPS7A	0.02001	RILP	0.012914
TTC32	0.016996	PCDH7	0.012815	DYNLRB2	0.020596
ADORA2A	0.017107	FAHD2A	0.00177	HIST1H4E	0.008729
PIP5KL1	0.000395	GLOD4	0.020693	KRTAP4-12	0.018167
UBE2F	0.016629	MEMO1	0.003593	CASP8	0.002175
ASB15	0.021525	GAL	0.022089	BFSP2	0.005738
CSNK2A2	0.005943	LEF1	0.006541	POLR3GL	0.016004
TMEM161B	0.010957	ZNF117	0.012299	C9orf64	0.022993
RAET1L	0.005324	ANGPT4	0.004556	NTPCR	0.006676
CYP19A1	0.012436	BIN2	0.006334	USP38	0.012178
HSFY2	0.011317	SFMBT1	0.017341	MAEL	0.014383
DBI	0.006995	CHMP5	0.021752	PPFIBP1	0.006954
DENND2C	0.010981	TMBIM4	0.012754	ITGA8	0.00668
UBXN2A	0.012576	CES1P1	0.00951	SLC25A12	0.021775
C10orf128	0.002704	PFDN4	0.007893	EIF3D	0.022065
E2F3	0.010763	PIGH	0.00735	EIF3G	0.020213
EMR1	0.014771	PKP1	0.022089	B3GALNT1	0.010623
R3HCC1	0.008417	PLXNA2	0.002124	URI1	0.018386
ETS2	0.007678	SETD4	0.002569	GGACT	0.022175
F8	0.002852	POLE3	0.009697	DCAF5	0.021098
LOC220729	0.011719	ERRFI1	0.005341	KSR1	0.021775
43167	0.015644	INO80	0.005555	KAT2B	0.022781
ALKBH3	0.004708	PCSK4	0.020211	SYNJ2	0.002245
SDK1	0.004344	ZNF586	0.013214	NAE1	0.023244
TMED4	0.020349	ANKRD49	0.011358	CPNE3	0.010416
FGF2	0.021267	TRIM68	0.022856	GYG2	0.00064
CASC3	0.012884	ZDHHC4	0.022211	CCND3	0.00301
CPEB3	0.022237	NADSYN1	0.016229	WDR34	0.01807
WDR47	0.023296	LGI2	0.019251	CCNH	0.014771

SNX13	0.001982	YOD1	0.010486	ZNF551	0.004921
RFTN1	0.002839	GNG12	0.010136	MCU	0.003364
GANAB	0.012134	KCMF1	0.022052	CCT6A	0.020611
TSPYL4	0.012312	C8orf4	0.00378	TCEAL8	0.020508
ARHGEF18	0.019216	MCCC1	0.017066	PRPF4	0.019998
TDRD7	0.015453	PTGER2	0.020417	C12orf29	0.015627
DDAH2	0.010486	MIR491	0.013621	GGTLC1	0.015302
GABARAPL1	0.020167	ERMN	0.019519	C1orf105	0.018411
ABL1	0.016571	PTMA	0.023158	PXYLP1	0.01512
ACKR1	0.013079	WDR48	0.02131	REEP6	0.017695
SNX32	0.014883	PTPN1	0.022042	BAG5	0.022088
MCM9	0.01547	SPTBN4	0.004799	CD70	0.005004
DNAJB5	0.007966	PTPN12	0.008411	DOCK4	0.009675
EGFL6	0.015708	RAP2C	0.008575	PTDSS1	0.01356
CHMP2B	0.01983	RAB6A	0.00527	NUP153	0.006177
FAM32A	0.00185	ACTA2	0.01484	NR_024132	0.00808
DECR2	0.01391	RBBP7	0.009845	NR_024557	0.007757
FAM127B	0.020555	ZNF350	0.015562	3-Sep	0.003284

Table S3: Pathway enrichment results for chest pain at Day 14

GO	Description	LogP	Enrichment	Z-score	genes
hsa03010	Ribosome	-19	9	15	FAU RPSA RPL4 RPL6 RPL10 RPL12 RPL15 RPL17 RPL18A RPL24 RPL27 RPL30 RPL29 RPL31 RPL35A RPLP0 RPS3 RPS5 RPS6 RPS9 RPS11 RPS14 RPS15 RPS20 RPS27 UBA52 MRPL15 MRPS14 MRPL14 MRPS15
hsa05171	Coronavirus disease - COVID-19	-14	5.9	11	FAU CXCL10 RPSA NFKB1 RPL4 RPL6 RPL10 RPL12 RPL15 RPL17 RPL18A RPL24 RPL27 RPL30 RPL29 RPL31 RPL35A RPLP0 RPS3 RPS5 RPS6 RPS9 RPS11 RPS14 RPS15 RPS20 RPS27 UBA52 VWF
hsa00190	Oxidative phosphorylation	-12	7.4	11	ATP5F1B ATP5F1C ATP5MC1 ATP5PF ATP6V1E1 ATP5PO COX6A1 COX6B1 COX7A2 ATP6V1F ATP5MF ATP5MG UQCR11
hsa05012	Parkinson disease	-11	5	9.6	ATP5F1B ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6A1 COX6B1 COX7A2 EIF2S1 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 PSMA1 PSMA4 PSMA6 PSMD8 TXN UBA52 UBB UQCRB COX7A2L UQCR11 SLC39A8 PINK1 MCU
hsa05016	Huntington disease	-10	4.3	8.6	ATP5F1B ATP5F1C ATP5MC1 ATP5PF ATP5PO BDNF CLTA COX6A1 COX6B1 COX7A2 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 POLR2I POLR2J POLR2K POLR2L PSMA1 PSMA4 PSMA6 PSMD8 UQCRB ULK1 COX7A2L UQCR11
hsa05014	Amyotrophic lateral sclerosis	-8.9	3.8	7.8	ATP5F1B ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6A1 COX6B1 COX7A2 EIF2S1 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 MAP2K3 PSMA1 PSMA4 PSMA6 PSMD8 UQCRB VCP ULK1 COX7A2L UQCR11 PINK1 MCU MAP1LC3B2 POM121C
hsa05208	Chemical carcinogenesis - reactive oxygen species	-8.6	4.7	8.1	ATP5F1B ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6A1 COX6B1 COX7A2 CYP2F1 MGST1 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 NFKB1 UQCRB COX7A2L GSTO1 UQCR11 PRKD2
hsa05022	Pathways of	-8.5	3.3	7.4	ATP2A3 ATP5F1B ATP5F1C ATP5MC1 ATP5PF ATP5PO BDNF COX6A1 COX6B1 COX7A2 CYP2F1 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 NFKB1 UQCRB COX7A2L GSTO1 UQCR11 PRKD2

	neurodegeneration - multiple diseases				X7A2 EIF2S1 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 NFKB1 MAP2K3 PSMA1 PSMA4 PSMA6 PSMD8 UBA52 UBB UQCRB VCP ULK1 COX7A2L UQCR11 PINK1 MCU MAP1LC3B2
hsa05010	Alzheimer disease	-8.4	3.6	7.5	ATP2A3 ATP5F1B ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6A1 COX6B1 COX7A2 EIF2S1 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 NFKB1 PSMA1 PSMA4 PSMD8 UQCRB ULK1 COX7A2L UQCR11 APH1A SLC39A8 APH1B MCU ATP5F1B ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6A1 COX6B1 COX7A2 EIF2S1 L
hsa05020	Prion disease	-8.3	4.2	7.7	AMC1 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 PSMA1 PSMA4 PSMA6 PSMD8 UQCRB COX7A2L UQCR11 MCU ATP5F1B ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6A1 COX6B1 COX7A2 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 MAP2K3 RPS6 UQCRB COX7A2L ATP5MF ATP5MG UQCR11 COA5
hsa04714	Thermogenesis	-8.3	4.5	7.8	ATP2A3 ATP5F1B ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6A1 COX6B1 COX7A2 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 MAP2K3 RPS6 UQCRB COX7A2L ATP5MF ATP5MG UQCR11 COA5
hsa05415	Diabetic cardiomyopathy	-7.1	4.4	7.2	NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 NFKB1 UQCRB COX7A2L UQCR11
hsa04120	Ubiquitin mediated proteolysis	-5.7	4.7	6.4	CDC34 SKP1 ELOB UBA52 UBB UBE2E1 UBE2E2 UBE2H WWP2 ANAPC13 PRPF19 ANAPC2 UBE2O UBE2F
hsa04932	Non-alcoholic fatty liver disease	-5.2	4.3	6	COX6A1 COX6B1 COX7A2 EIF2S1 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 NFKB1 UQCRB COX7A2L UQCR11
hsa04144	Endocytosis	-4	3	4.7	GRK2 CLTA FGFR4 GRK6 WIPF1 HGS CYTH1 ACAP1 ARPC2 EHD1 DNM3 EPS15L1 VPS37B VPS25 RAB11FIP4 MVB12B
hsa03020	RNA polymerase	-3.2	7	5.1	POLR2I POLR2J POLR2K POLR2L POLR1D
hsa03040	Spliceosome	-2.9	3.2	4	DHX8 HNRNPC HNRNPM SNRPD2 SART1 DDX23 SF3A1 LSM5 PRPF19 SF3B5
hsa04146	Peroxisome	-2.8	4	4.1	PRDX1 ABCD4 PXMP2 PEX11B PRDX5 HACL1 GSTK1
hsa04810	Regulation of actin cytoskeleton	-2.6	2.6	3.5	ACTN1 FGFR4 GSN ITGB4 LIMK2 MYH9 VAV1 LPAR2 ARPC2 SCIN FGD3 MYL12B

hsa04260	Cardiac muscle contraction	-2.6	3.8	3.9	ATP2A3 COX6A1 COX6B1 COX7A2 UQCRB COX7A2L UQCR11
hsa05017	Spinocerebellar ataxia	-2.5	3	3.5	ATP2A3 PSMA1 PSMA4 PSMA6 PSMD8 ULK1 CIC ATXN10 MCU
hsa04137	Mitophagy - animal	-2.4	3.9	3.7	UBA52 UBB ULK1 FIS1 PINK1 MAP1LC3B2
hsa05132	Salmonella infection	-2.2	2.3	3	DYNC1I2 FLNA NFKB1 MAP2K3 RPS3 SKP1 DYNLT1 TXN CYTH1 ARPC2 MYL12B ARL8A
hsa05166	Human T-cell leukemia virus 1 infection	-2.1	2.3	3	HLA-DQA1 MAD2L1 NFATC1 NFKB1 NFKB2 RAN STAT5B MAD1L1 PTTG1 ANAPC13 ANAPC2
hsa04066	HIF-1 signaling pathway	-2.1	3	3.1	MKNK2 LDHA NFKB1 PFKFB3 PFKL RPS6 ELOB
hsa04623	Cytosolic DNA-sensing pathway	-2	3.8	3.2	CXCL10 NFKB1 POLR2K POLR2L POLR1D
hsa05203	Viral carcinogenesis	-1.9	2.3	2.8	ACTN1 GSN H2BC5 NFKB1 NFKB2 STAT5B H4C4 MAD1L1 HDAC5 SCIN
	Retrograde				
hsa04723	endocannabinoid signaling	-1.9	2.6	2.8	GNB3 NDUFA4 NDUFA8 NDUFA9 NDUFB1 NDUFC1 NDUFS6 GNB5
hsa03050	Proteasome	-1.8	4.1	3.1	PSMA1 PSMA4 PSMA6 PSMD8
hsa00620	Pyruvate metabolism	-1.8	4	3.1	ACAT1 DLAT GLO1 LDHA
hsa00650	Butanoate metabolism	-1.7	5.1	3.2	ACAT1 BDH1 HADH
hsa04721	Synaptic vesicle cycle	-1.6	3	2.6	ATP6V1E1 CLTA RAB3A ATP6V1F DNM3
hsa03018	RNA degradation	-1.6	3	2.6	CNOT2 CNOT3 PFKL CNOT9 LSM5
hsa03013	Nucleocytoplasmic transport	-1.6	2.6	2.5	RAN SUMO2 TNPO3 THOC6 THOC7 POM121C
hsa00480	Glutathione metabolism	-1.5	3.3	2.5	MGST1 GSTO1 TXNDC12 GSTK1
hsa00760	Nicotinate and	-1.4	4.1	2.7	PNP NADK NT5C1B

	nicotinamide metabolism				
hsa05131	Shigellosis	-1.4	1.9	2.1	ACTN1 NFKB1 SKP1 UBA52 UBB CYTH1 ARPC2 TECPR1 MYL12B MAP1LC3B2
hsa04658	Th1 and Th2 cell differentiation	-1.3	2.6	2.2	HLA-DQA1 JAG2 NFATC1 NFKB1 STAT5B

Table S4: Pathway enrichment results for chest distress at Day 14

GO	Description	LogP	Enrichment	Z-score	genes
hsa03010	Ribosome	-23	10	17	FAU RPSA RPL3 RPL4 RPL6 RPL7 RPL7A RPL10 RPL12 RPL15 RPL17 RPL24 RPL30 RPL31 RPL34 RPL35A RPLP0 RPS3 RPS3A RPS9 RPS10 RPS14 RPS15A RPS18 RPS20 RPS24 RPS27 UBA52 RPL23 RPL35 RPL13A MRPS14 MRPS15 MRPL11
hsa05171	Coronavirus disease - COVID-19	-19	7	14	FAU RPSA PIK3CD PIK3R2 RPL3 RPL4 RPL6 RPL7 RPL7A RPL10 RPL12 RPL15 RPL17 RPL24 RPL30 RPL31 RPL34 RPL35A RPLP0 RPS3 RPS3A RPS9 RPS10 RPS14 RPS15A RPS18 RPS20 RPS24 RPS27 MAP3K7 UBA52 IKBKG RPL23 RPL35 RPL13A
hsa05012	Parkinson disease	-8.3	4.2	7.7	ATP5F1C ATP5MC3 ATP5PF ATP5PO COX7A2 COX7B COX7C EIF2S1 NDUFA4 NDUFA6 NDUFS3 PRKACB PSMA6 UBA52 UBB UBC XBP1 COX7A2L PPIF UQC R11 TXN2 SLC39A9 MCU
hsa05010	Alzheimer disease	-8.1	3.5	7.3	APOE ATP2A3 ATP5F1C ATP5MC3 ATP5PF ATP5PO COX7A2 COX7B COX7C EIF2S1 NDUFA4 NDUFA6 NDUFA8 NDUFS3 PIK3CD PIK3R2 PSMA6 XBP1 FZD1 FZD8 IKBK G IRS2 COX7A2L ATG13 PPIF APC2 UQCR11 SLC39A9 MCU
hsa05022	Pathways of neurodegeneration - multiple diseases	-7.7	3.1	6.9	ATP2A3 ATP5F1C ATP5MC3 ATP5PF ATP5PO COX7A2 COX7B COX7C EIF2S1 HIP1 NDUFA4 NDUFA6 NDUFA8 NDUFS3 MAP2K3 PSMA6 SPTBN2 UBA52 UBB UBC XBP1 FZD1 FZD8 DNAH17 COX7A2L ATG13 PPIF APC2 UQCR11 TOMM40L MCU MAP1LC 3C
hsa05208	Chemical carcinogenesis -	-7.7	4.4	7.5	ACP1 ATP5F1C ATP5MC3 ATP5PF ATP5PO COX7A2 COX7B COX7C NDUFA4 NDUF A6 NDUFA8 NDUFS3 PIK3CD PIK3R2 PRKCD IKBKG COX7A2L GSTO1 PPIF UQCR11

	reactive oxygen species				
hsa05020	Prion disease	-7.5	3.9	7.1	PRKD2 ATP5F1C ATP5MC3 ATP5PF ATP5PO COX7A2 COX7B COX7C EIF2S1 HSPA8 NDUFA4 NDUFA6 NDUFA8 NDUFS3 NOTCH1 PIK3CD PIK3R2 PRKACB PRKCD PSMA6 COX7A2L PPIF UQCR11 MCU
hsa04714	Thermogenesis	-6.7	4	6.8	ATP5F1C ATP5MC3 ATP5PF ATP5PO COX7A2 COX7B COX7C NDUFA4 NDUFA6 NDUFA8 NDUFS3 PRKACB PRKAG1 MAP2K3 COX7A2L ATP5MF UQCR11 COA4 SIRT6 COX14
hsa00190	Oxidative phosphorylation	-6.6	5.2	7.2	ATP5F1C ATP5MC3 ATP5PF ATP5PO COX7A2 COX7B COX7C NDUFA4 NDUFA6 NDUFA8 NDUFS3 PPA1 COX7A2L ATP5MF UQCR11
hsa04932	Non-alcoholic fatty liver disease	-6.6	4.8	7	COX7A2 COX7B COX7C EIF2S1 MAP3K11 NDUFA4 NDUFA6 NDUFA8 NDUFS3 PIK3CD PIK3R2 PRKAG1 XBP1 IRS2 COX7A2L UQCR11
hsa04120	Ubiquitin mediated proteolysis	-6.3	4.9	6.9	CDC34 ELOC ELOB UBA52 UBB UBC UBE2E2 CUL5 ANAPC10 WWP2 ANAPC13 ANAPC2 KLHL9 UBE2O UBE2F
hsa05415	Diabetic cardiomyopathy	-6.3	4.1	6.6	ATP2A3 ATP5F1C ATP5MC3 ATP5PF ATP5PO COX7A2 COX7B COX7C NDUFA4 NDUFA6 NDUFA8 NDUFS3 PIK3CD PIK3R2 PRKCD COX7A2L PPIF UQCR11
hsa05014	Amyotrophic lateral sclerosis	-5.3	2.9	5.5	ATP5F1C ATP5MC3 ATP5PF ATP5PO COX7A2 COX7B COX7C EIF2S1 NDUFA4 NDUFA6 NDUFS3 MAP2K3 PSMA6 SRSF3 XBP1 DNAH17 COX7A2L ATG13 UQCRC11 TOMM40L MCU MAP1LC3C
hsa05016	Huntington disease	-4.9	3	5.3	ATP5F1C ATP5MC3 ATP5PF ATP5PO COX7A2 COX7B COX7C HIP1 NDUFA4 NDUFA6 NDUFS3 POLR2D POLR2K PSMA6 DNAH17 COX7A2L ATG13 PPIF UQCR11
hsa04658	Th1 and Th2 cell differentiation	-4.5	5	5.8	CD3D JAG2 JAK3 NFATC1 NOTCH1 NOTCH3 STAT5A STAT5B IKBKG MAML3
hsa04140	Autophagy - animal	-4.2	3.9	5.2	EIF2S1 PIK3CD PIK3R2 PRKACB PRKCD MAP3K7 IRS2 VAMP8 ATG4A ATG13 DEPTOR MAP1LC3C
hsa04137	Mitophagy - animal	-3.8	5.1	5.2	TFE3 UBA52 UBB UBC TFEB FIS1 FUNDC1 MAP1LC3C

hsa05017	Spinocerebellar ataxia	-3.5	3.6	4.6	ATP2A3 PIK3CD PIK3R2 PSMA6 SPTBN2 XBP1 ATG13 PPIF CIC ATXN10 MCU
hsa05166	Human T-cell leukemia virus 1 infection	-3.4	2.9	4.3	CD3D JAK3 NFATC1 PIK3CD PIK3R2 PRKACB SPI1 STAT5A STAT5B TLN1 IKBKG ANAPC10 ANAPC13 ANAPC2
hsa03040	Spliceosome	-3.4	3.5	4.4	HNRNPC HSPA8 MAGOH SRSF3 BUD31 RBM8A CRNL1 PPIL1 MAGOHB SF3B5 PH F5A
hsa04660	T cell receptor signaling pathway	-3.4	4	4.6	CD3D CD8B NCK1 NFATC1 PIK3CD PIK3R2 MAP3K7 VAV1 IKBKG
hsa05221	Acute myeloid leukemia	-3.2	4.8	4.7	PIK3CD PIK3R2 RARA SPI1 STAT5A STAT5B IKBKG
hsa03020	RNA polymerase	-3.1	6.8	5	POLR2D POLR2K POLR3B POLR3GL POLR1F
hsa05135	Yersinia infection	-3.1	3.4	4.1	CD8B NFATC1 PIK3CD PIK3R2 MAP2K3 MAP3K7 VAV1 WAS IKBKG ARPC1A
hsa05162	Measles	-3	3.3	4.1	CD3D EIF2S1 HSPA8 JAK3 PIK3CD PIK3R2 STAT5A STAT5B MAP3K7 IKBKG
hsa04062	Chemokine signaling pathway	-3	2.9	3.9	GRK5 GRK6 JAK3 PIK3CD PIK3R2 PRKACB PRKCD CCL23 STAT5B VAV1 WAS IKB KG
hsa04666	Fc gamma R-mediated phagocytosis	-2.9	3.8	4.1	LIMK2 PIK3CD PIK3R2 PRKCD VAV1 WAS GAB2 ARPC1A
hsa04144	Endocytosis	-2.9	2.6	3.7	CAPZA1 FGFR4 GRK5 GRK6 HSPA8 SH3GL1 DNAJC6 IQSEC1 ARPC1A EHD1 CHMP5 VPS37B MVB12B ARAP1
hsa04330	Notch signaling pathway	-2.8	4.7	4.2	JAG2 NOTCH1 NOTCH3 TLE3 HEY1 MAML3
hsa04213	Longevity regulating pathway - multiple species	-2.7	4.5	4.1	HSPA8 PIK3CD PIK3R2 PRKACB PRKAG1 IRS2
hsa05170	Human immunodeficiency virus 1 infection	-2.6	2.6	3.5	CD3D LIMK2 NFATC1 PIK3CD PIK3R2 MAP2K3 MAP3K7 ELOC ELOB CUL5 IKBKG AP1S2

hsa05161	Hepatitis B	-2.5	2.9	3.5	JAK3 NFATC1 PCNA PIK3CD PIK3R2 MAP2K3 STAT5A STAT5B MAP3K7 IKBKG
hsa05131	Shigellosis	-2.5	2.4	3.4	ACTN1 PIK3CD PIK3R2 PRKCD MAP3K7 TLN1 UBA52 UBB UBC IKBKG ARPC1A TE CPR1 MAP1LC3C
hsa05200	Pathways in cancer	-2.5	1.9	3.2	CSF2RB FGFR4 GLI1 JAG2 JAK3 NOTCH1 NOTCH3 PIK3CD PIK3R2 PRKACB RARA SPI1 STAT5A STAT5B ELOC ELOB FZD1 FZD8 IKBKG GSTO1 APC2 HEY1
hsa05224	Breast cancer	-2.3	2.8	3.3	JAG2 NOTCH1 NOTCH3 PIK3CD PIK3R2 FZD1 FZD8 APC2 HEY1
hsa05220	Chronic myeloid leukemia	-2.2	3.7	3.4	PIK3CD PIK3R2 STAT5A STAT5B IKBKG GAB2
hsa05340	Primary immunodeficiency	-2	4.9	3.5	CD3D CD8B JAK3 IKBKG
hsa04659	Th17 cell differentiation	-2	3	3.1	CD3D JAK3 NFATC1 RARA STAT5A STAT5B IKBKG
hsa03013	Nucleocytoplasmic transport	-2	3	3.1	EEF1A1 KPNA2 MAGOH RBM8A TNPO3 MAGOHB THOC7
hsa04066	HIF-1 signaling pathway	-2	3	3.1	ANGPT1 PFKFB3 PIK3CD PIK3R2 ELOC ELOB MKNK1
hsa04310	Wnt signaling pathway	-2	2.5	2.9	NFATC1 PRKACB MAP3K7 TLE3 FZD1 FZD8 APC2 LGR4 NOTUM
hsa04260	Cardiac muscle contraction	-1.9	3.2	3	ATP2A3 COX7A2 COX7B COX7C COX7A2L UQCR11
hsa04530	Tight junction PD-L1 expression and	-1.9	2.5	2.8	ACTN1 CD1E MYL6 PCNA PRKACB PRKAG1 WAS ARPC1A F11R
hsa05235	PD-1 checkpoint pathway in cancer	-1.9	3.1	3	CD3D NFATC1 PIK3CD PIK3R2 MAP2K3 IKBKG
hsa04211	Longevity regulating pathway	-1.9	3.1	3	PIK3CD PIK3R2 PRKACB PRKAG1 IRS2 ATG13
hsa05203	Viral carcinogenesis	-1.9	2.3	2.7	ACTN1 GTF2E1 JAK3 PIK3CD PIK3R2 PRKACB STAT5A STAT5B H2BC21 IKBKG

	Growth hormone				
hsa04935	synthesis, secretion and action	-1.8	2.7	2.8	PIK3CD PIK3R2 PRKACB MAP2K3 STAT5A STAT5B IRS2
hsa04152	AMPK signaling pathway	-1.8	2.7	2.8	PFKFB3 PIK3CD PIK3R2 PRKAG1 MAP3K7 IRS2 EEF2K
hsa04664	Fc epsilon RI signaling pathway	-1.8	3.4	2.9	PIK3CD PIK3R2 MAP2K3 VAV1 GAB2
hsa04919	Thyroid hormone signaling pathway	-1.8	2.7	2.7	ATP2A3 NOTCH1 NOTCH3 PIK3CD PIK3R2 PRKACB MED16
hsa05211	Renal cell carcinoma	-1.8	3.4	2.9	PIK3CD PIK3R2 ELOC ELOB TFE3
hsa04930	Type II diabetes mellitus	-1.8	4	3	PIK3CD PIK3R2 PRKCD IRS2
hsa04520	Adherens junction	-1.7	3.3	2.8	ACP1 ACTN1 NECTIN1 MAP3K7 WAS
hsa05417	Lipid and atherosclerosis	-1.7	2.2	2.5	EIF2S1 HSPA8 NFATC1 PIK3CD PIK3R2 MAP2K3 MAP3K7 VAV1 XBP1 IKBKG
hsa01522	Endocrine resistance	-1.7	2.8	2.7	JAG2 NOTCH1 NOTCH3 PIK3CD PIK3R2 PRKACB
hsa05223	Non-small cell lung cancer	-1.7	3.2	2.8	JAK3 PIK3CD PIK3R2 STAT5A STAT5B
hsa04380	Osteoclast differentiation	-1.7	2.5	2.6	NFATC1 PIK3CD PIK3R2 SPI1 MAP3K7 IKBKG GAB2
	AGE-RAGE signaling pathway				
hsa04933	pathway in diabetic complications	-1.7	2.8	2.6	NFATC1 PIK3CD PIK3R2 PRKCD STAT5A STAT5B
hsa00270	Cysteine and methionine metabolism	-1.6	3.7	2.8	GCLC MDH1 MAT2B KYAT3
hsa04914	Progesterone-mediated	-1.6	2.7	2.6	PIK3CD PIK3R2 PRKACB ANAPC10 ANAPC13 ANAPC2

	oocyte maturation				
hsa04625	C-type lectin receptor signaling pathway	-1.6	2.7	2.5	BCL3 NFATC1 PIK3CD PIK3R2 PRKCD IKBKG
hsa04910	Insulin signaling pathway	-1.5	2.4	2.4	PIK3CD PIK3R2 PRKACB PRKAG1 MKNK1 IRS2 SH2B2
hsa04915	Estrogen signaling pathway	-1.5	2.3	2.4	HSPA8 KRT12 PIK3CD PIK3R2 PRKACB PRKCD RARA
hsa04662	B cell receptor signaling pathway	-1.5	2.8	2.5	NFATC1 PIK3CD PIK3R2 VAV1 IKBKG
hsa04923	Regulation of lipolysis in adipocytes	-1.5	3.3	2.6	PIK3CD PIK3R2 PRKACB IRS2
hsa04141	Protein processing in endoplasmic reticulum	-1.5	2.2	2.3	EIF2S1 HSPA8 SSR2 XBP1 UBXN8 BAG2 TRAM1 ERLEC1
hsa04668	TNF signaling pathway	-1.5	2.5	2.3	BCL3 PIK3CD PIK3R2 MAP2K3 MAP3K7 IKBKG
hsa04550	Signaling pathways regulating pluripotency of stem cells	-1.4	2.3	2.3	FGFR4 JAK3 PIK3CD PIK3R2 FZD1 FZD8 APC2
hsa04012	ErbB signaling pathway	-1.4	2.7	2.4	NCK1 PIK3CD PIK3R2 STAT5A STAT5B
hsa04670	Leukocyte				
	transendothelial migration				
	Nicotinate and				
hsa00760	nicotinamide metabolism	-1.4	4	2.6	PNP SIRT6 NADK
hsa04015	Rap1 signaling	-1.4	2	2.1	ANGPT1 FGFR4 PIK3CD PIK3R2 MAP2K3 SIPA1 TLN1 VAV1 PRKD2

	pathway				
hsa04621	NOD-like receptor signaling pathway	-1.3	2	2	PRKCD MAP3K7 IKBKG SUGT1 TXN2 NLRX1 MCU MAP1LC3C
hsa04623	Cytosolic DNA-sensing pathway	-1.3	2.9	2.3	POLR2K IKBKG POLR3B POLR3GL
hsa05217	Basal cell carcinoma	-1.3	2.9	2.3	GLI1 FZD1 FZD8 APC2
hsa04810	Regulation of actin cytoskeleton	-1.3	1.9	2	ACTN1 FGFR4 ITGB4 LIMK2 PIK3CD PIK3R2 VAV1 APC2 ARPC1A
hsa01240	Biosynthesis of cofactors	-1.3	2.1	2.1	AK2 GCLC MAT2B LIPT1 COQ3 NADK IDO2

Table S5: Pathway enrichment results for palpitations at Day 14

GO	Description	LogP	Enrichment	Z-score	genes
hsa05168	Herpes simplex virus 1 infection	-2.9	2.7	3.8	LTA NFKBIA ZNF354A ZNF17 ZNF180 CARD9 ZNF419 ZNF442 ZNF527 ZNF160 ZNF563 ZNF565 ZNF850
hsa05321	Inflammatory bowel disease	-2.4	6.3	4.3	IL12RB1 TGFB2 IL23A IL23R
hsa03022	Basal transcription factors	-2	6.9	3.9	GTF2B TAF11 GTF2IRD1
hsa05134	Legionellosis	-1.7	5.4	3.3	CLK1 NFKBIA SAR1B
hsa04659	Th17 cell differentiation	-1.7	3.8	2.9	IL12RB1 NFKBIA IL23A IL23R
	Neuroactive ligand-receptor interaction				
hsa04080	ligand-receptor interaction	-1.6	2.3	2.5	ADRB3 CHRNB2 GNRH2 HRH1 PLG PRSS1 NPFFR2 PATE2
hsa04060	Cytokine-cytokine	-1.6	2.4	2.5	IL12RB1 LTA TGFB2 ACKR4 IL23A IL23R GDF6

	receptor interaction				
hsa05164	Influenza A	-1.6	3	2.6	DNAJB1 NFKBIA PLG PRSS1 CALCOCO2

Table S6: Pathway enrichment results for purple or dark lips at Day 14

GO	Description	LogP	Enrichment	Z-score	genes
hsa00140	Steroid hormone biosynthesis	-3	8.8	5.3	CYP1A1 AKR1C1 AKR1C2 HSD17B3
hsa05217	Basal cell carcinoma	-2.9	8.6	5.2	BMP2 GADD45B AXIN2 FZD7
hsa05202	Transcriptional misregulation in cancer	-1.8	3.5	3	CEBPA MLLT3 GADD45B ZEB1 BCL11B
hsa04550	Signaling pathways regulating pluripotency of stem cells	-1.7	3.8	2.9	INHBB REST AXIN2 FZD7
hsa05224	Breast cancer	-1.6	3.7	2.8	GADD45B AXIN2 FZD7 SHC2
hsa05226	Gastric cancer	-1.6	3.6	2.8	GADD45B AXIN2 FZD7 SHC2
hsa05225	Hepatocellular carcinoma	-1.4	3.2	2.5	GADD45B AXIN2 FZD7 SHC2

Table S7: Pathway enrichment results for purple or dark tongue at Day 14

GO	Description	LogP	Enrichment	Z-score	genes
hsa03013	Nucleocytoplasmic transport	-3.6	5.6	5.2	KPNB1 RAN UPF1 SUMO3 TNPO3 THOC6 KPNA7
hsa04974	Protein digestion and absorption	-2.1	4.2	3.5	ATP1B1 COL3A1 MEP1B COL14A1 COL27A1
hsa04144	Endocytosis	-2	2.7	3	FGFR4 HSPA1B PIP5K1A IST1 CHMP4A MVB12B WIPF2 ACTR3C
hsa04630	JAK-STAT signaling pathway	-1.9	3.2	3	CCND2 IFNA5 IL5RA IL15 PIK3CD IL20
hsa04120	Ubiquitin mediated proteolysis	-1.6	3	2.6	CDC34 UBA52 UBOX5 UBE2S UBE2O
hsa04550	Signaling pathways regulating pluripotency of stem cells	-1.6	3	2.6	BMPR1B FGFR4 INHBB PIK3CD WNT4
hsa05132	Salmonella infection	-1.6	2.4	2.4	DYNC1LI2 PIK3CD TXN RIPK1 ACTR1B RHOJ ACTR3C
hsa04623	Cytosolic DNA-sensing pathway	-1.4	4.1	2.7	IFNA5 POLR2L RIPK1
hsa03010	Ribosome	-1.4	2.7	2.3	RPL3L MRPL12 UBA52 MRPL20 MRPS11

hsa04217	Necroptosis	-1.4	2.7	2.3	PARP1 IFNA5 RIPK1 CHMP4A SPATA2L
hsa04668	TNF signaling pathway	-1.4	3.1	2.4	IL15 PIK3CD VEGFC RIPK1

Table S8: Pathway enrichment results for XYZ phenotype at Day 14

GO	Description	LogP	Enrichment	Z-score	genes
hsa03010	Ribosome	-10	7	10	RPSA RPL10 RPL12 RPL15 RPL18A RPL30 RPLP0 RPS3 RPS5 RPS6 RPS11 RPS14 RPS20 RPS27 UBA52 MRPL15 MRPS14 MRPS15 MRPL32
hsa05171	Coronavirus disease - COVID-19	-6.9	4.5	7.1	RPSA NFKB1 PIK3CD RPL10 RPL12 RPL15 RPL18A RPL30 RPLP0 RPS3 RPS5 RPS6 RPS11 RPS14 RPS20 RPS27 UBA52 VWF
hsa00190	Oxidative phosphorylation	-6.2	5.6	7.1	ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6B1 COX7B NDUFA9 NDUFS3 NDUFS6 COX7A2L ATP6V1F UQCR11 ATP6V1H
hsa05012	Parkinson disease	-6	3.9	6.3	ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6B1 COX7B EIF2S1 NDUFA9 NDUFS3 NDUFS6 PSMA6 UBA52 UBB XBP1 COX7A2L UQCR11 SLC39A9 PINK1
hsa05010	Alzheimer disease	-6	3.3	6.1	APOE ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6B1 COX7B EIF2S1 NDUFA9 NDUFS3 NDUFS6 NFKB1 PIK3CD PSMA6 XBP1 FZD7 IRS2 COX7A2L UQCR11 APH1A SLC39A9 APH1B
hsa05020	Prion disease	-5.8	3.8	6.2	ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6B1 COX7B EIF2S1 HSPA1B HSPA8 LAMC1 NDUFA9 NDUFS3 NDUFS6 NOTCH1 PIK3CD PSMA6 COX7A2L UQCR11
hsa05016	Huntington disease	-5.7	3.6	6.1	ATP5F1C ATP5MC1 ATP5PF ATP5PO BDNF CLTA COX6B1 COX7B HIP1 NDUFA9 NDUFS3 NDUFS6 POLR2I POLR2J POLR2K PSMA6 DNAH17 COX7A2L UQCR11
hsa05208	Chemical carcinogenesis - reactive oxygen species	-5.7	4.2	6.3	ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6B1 COX7B CYP1A1 CYP2F1 NDUFA9 NDUFS3 NDUFS6 NFKB1 PIK3CD COX7A2L UQCR11 EPHX4
hsa05022	Pathways of neurodegeneration -	-5	2.8	5.2	ATP5F1C ATP5MC1 ATP5PF ATP5PO BDNF COX6B1 COX7B EIF2S1 HI

	multiple diseases				
hsa04714	Thermogenesis	-4.9	3.7	5.6	P1 NDUFA9 NDUFS3 NDUFS6 NFKB1 PSMA6 UBA52 UBB VCP XBP1 F ZD7 DNAH17 COX7A2L UQCR11 PINK1 ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6B1 COX7B NDUFA9 NDUFS 3 NDUFS6 RPS6 COX7A2L UQCR11 COA4 COX14 COA5 COX6B1 COX7B EIF2S1 NDUFA9 NDUFS3 NDUFS6 NFKB1 PIK3CD XB P1 IRS2 COX7A2L UQCR11
hsa04932	Non-alcoholic fatty liver disease	-4.8	4.5	5.8	ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6B1 COX7B NDUFA9 NDUFS 3 NDUFS6 NFKB1 PIK3CD COX7A2L UQCR11
hsa05415	Diabetic cardiomyopathy	-4.2	3.7	5.1	ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6B1 COX7B NDUFA9 NDUFS 3 NDUFS6 NFKB1 PIK3CD COX7A2L UQCR11 ATP5F1C ATP5MC1 ATP5PF ATP5PO COX6B1 COX7B EIF2S1 NDUFA9
hsa05014	Amyotrophic lateral sclerosis	-4.1	2.9	4.7	NDUFS3 NDUFS6 PSMA6 VCP XBP1 DNAH17 COX7A2L UQCR11 PINK 1 NUP37
hsa04144	Endocytosis	-3.9	3.2	4.7	GRK2 CAPZA1 CLTA FGFR4 GRK6 HSPA1B HSPA8 SH3GL1 EHD1 CH MP5 VPS29 SMURF1 MVB12B ARAP1
hsa05166	Human T-cell leukemia virus 1 infection	-3.9	3.4	4.7	CD3D NFATC1 NFKB1 PIK3CD RAN RELB STAT5A STAT5B TLN1 MA D1L1 PTTG1 ANAPC10 ANAPC13
hsa04120	Ubiquitin mediated proteolysis	-3.7	4.1	4.9	CDC34 ELOB UBA52 UBB ANAPC10 WWP2 ANAPC13 SMURF1 UBE2O UBE2F
hsa04658	Th1 and Th2 cell differentiation	-3.7	5	5.1	CD3D IL4R NFATC1 NFKB1 NOTCH1 STAT5A STAT5B MAML3
hsa04659	Th17 cell differentiation	-2.6	3.8	3.8	CD3D IL4R NFATC1 NFKB1 RARA STAT5A STAT5B
hsa05162	Measles	-2.5	3.3	3.7	CD3D EIF2S1 HSPA1B HSPA8 NFKB1 PIK3CD STAT5A STAT5B
hsa05134	Legionellosis	-2.5	5.1	4.1	HSPA1B HSPA8 NFKB1 VCP CLK4
hsa04330	Notch signaling pathway	-2.5	4.9	4	NOTCH1 TLE3 APH1A MAML3 APH1B
hsa05221	Acute myeloid leukemia	-2.2	4.3	3.6	NFKB1 PIK3CD RARA STAT5A STAT5B
hsa04137	Mitophagy - animal	-2.1	4	3.4	UBA52 UBB FIS1 PINK1 FUNDC1
hsa04510	Focal adhesion	-2.1	2.6	3	ACTN1 COL6A3 LAMC1 PIK3CD TLN1 VAV1 VWF MYL12A PARVA
hsa05203	Viral carcinogenesis	-2	2.6	3	ACTN1 GSN GTF2E1 NFKB1 PIK3CD STAT5A STAT5B MAD1L1 SCIN

hsa05220	Chronic myeloid leukemia	-2	3.8	3.3	NFKB1 PIK3CD STAT5A STAT5B GAB2
hsa03013	Nucleocytoplasmic transport	-2	3.2	3.1	RAN RBM8A TNPO3 NUP37 THOC6 THOC7
hsa04066	HIF-1 signaling pathway	-1.9	3.2	3	NFKB1 PDKFB3 PFKL PIK3CD RPS6 ELOB
hsa05417	Lipid and atherosclerosis	-1.9	2.4	2.8	CYP1A1 EIF2S1 HSPA1B HSPA8 NFATC1 NFKB1 PIK3CD VAV1 XBP1
hsa03040	Spliceosome	-1.9	2.8	2.8	HSPA1B HSPA8 HNRNPM BUD31 SART1 RBM8A SF3B5
hsa04260	Cardiac muscle contraction	-1.8	3.3	2.9	COX6B1 COX7B MYL4 COX7A2L UQCR11
hsa04150	mTOR signaling pathway	-1.7	2.6	2.7	GRB10 PIK3CD RPS6 FZD7 ATP6V1F ATP6V1H STRADA
hsa04062	Chemokine signaling pathway	-1.7	2.4	2.6	GRK2 GNB3 GRK6 NFKB1 PIK3CD STAT5B VAV1 CXCL16
hsa00051	Fructose and mannose metabolism	-1.7	5.3	3.3	PFKFB3 PFKL PMM2
hsa05235	PD-L1 expression and PD-1 checkpoint pathway in cancer	-1.7	3.3	2.8	CD3D NFATC1 NFKB1 PIK3CD BATF3
hsa03020	RNA polymerase	-1.7	5.1	3.2	POLR2I POLR2J POLR2K
hsa04151	PI3K-Akt signaling pathway	-1.7	2	2.4	BDNF COL6A3 FGFR4 GNB3 NR4A1 IL4R LAMC1 NFKB1 PIK3CD RPS6 VWF PHLPP1
hsa04213	Longevity regulating pathway - multiple species	-1.7	3.7	2.9	HSPA1B HSPA8 PIK3CD IRS2
hsa04666	Fc gamma R-mediated phagocytosis	-1.6	3	2.6	GSN PIK3CD VAV1 GAB2 SCIN
hsa04141	Protein processing in endoplasmic reticulum	-1.5	2.4	2.4	EIF2S1 HSPA1B HSPA8 SSR2 VCP XBP1 ERLEC1
hsa04933	AGE-RAGE signaling pathway in diabetic complications	-1.5	2.9	2.5	NFATC1 NFKB1 PIK3CD STAT5A STAT5B
hsa05207	Chemical carcinogenesis - receptor activation	-1.5	2.2	2.3	BCL6 CYP1A1 NFKB1 PIK3CD STAT5A STAT5B EPHX4 MIRLET7B
hsa04917	Prolactin signaling pathway	-1.5	3.3	2.6	NFKB1 PIK3CD STAT5A STAT5B
hsa04660	T cell receptor signaling pathway	-1.5	2.8	2.4	CD3D NFATC1 NFKB1 PIK3CD VAV1
hsa04625	C-type lectin receptor signaling	-1.5	2.8	2.4	PLK3 NFATC1 NFKB1 PIK3CD RELB

	pathway				
hsa04721	Synaptic vesicle cycle	-1.3	3	2.3	CLTA ATP6V1F CPLX1 ATP6V1H
hsa01200	Carbon metabolism	-1.3	2.5	2.2	ACAT1 PFKL RPE ADPGK MCEE
hsa01240	Biosynthesis of cofactors	-1.3	2.3	2.1	AK2 HMBS PMM2 NFS1 LIPT1 ADSS1

Table S9: Pathway enrichment results for chest pain at Day 30

GO	Description	LogP	Enrichment	Z-score	genes
hsa04360	Axon guidance	-2.5	5.2	4.1	CFL1 EFNA4 RHOD SEMA6D PARD6G
hsa04080	Neuroactive ligand-receptor interaction	-1.9	3.2	3	AVP CHRNA1 CHRNA3 GRIN1 GABRR3 UTS2B
hsa04015	Rap1 signaling pathway	-1.6	3.6	2.8	EFNA4 GRIN1 DOCK4 PARD6G
hsa04142	Lysosome	-1.5	4.3	2.8	HYAL2 ABCB9 PLA2G15

Table S10: Pathway enrichment results for chest distress at Day 30

GO	Description	LogP	Enrichment	Z-score	genes
hsa04014	Ras signaling pathway	-3	4	4.3	ETS2 GNG3 GRIN1 KITLG PRKACG REL KSR1 RASSF5
hsa05203	Viral carcinogenesis	-2.7	4	4	CDC20 DDB1 HPN JAK1 NFKBIA PRKACG REL
hsa04610	Complement and coagulation cascades	-2.2	5.5	3.8	C4A C8B F8 SERPINE1
hsa04144	Endocytosis	-2.2	3.2	3.3	ARRB1 CXCR4 CYTH3 SH3KBP1 VPS26B CHMP4B PIP5KL1
hsa04062	Chemokine signaling pathway	-2.2	3.6	3.4	ARRB1 FOXO3 GNG3 NFKBIA PRKACG CXCR4
hsa04140	Autophagy - animal	-2.1	4.1	3.5	LAMP1 PRKACG IRS2 MLST8 EIF2AK4
hsa05030	Cocaine addiction	-2.1	7.1	4	GRIN1 PRKACG PPP1R1B
hsa03040	Spliceosome	-2	4	3.3	SF3A1 SF3B2 DDX42 WBP11 CTNNBL1
hsa04713	Circadian entrainment	-2	4.8	3.5	GNG3 GRIN1 PER1 PRKACG
hsa05166	Human T-cell leukemia virus 1 infection	-1.9	3.1	3	CDC20 E2F3 ETS2 JAK1 NFKBIA PRKACG
hsa05163	Human cytomegalovirus infection	-1.9	3.1	2.9	E2F3 GNG3 JAK1 NFKBIA PRKACG CXCR4
hsa03013	Nucleocytoplasmic transport	-1.8	4.3	3.2	SUMO3 CASC3 XPO7 POM121C
hsa04213	Longevity regulating pathway - multiple species	-1.8	5.6	3.4	FOXO3 PRKACG IRS2
hsa04724	Glutamatergic synapse	-1.8	4.1	3.1	GNG3 GRIN1 PRKACG SLC38A3
hsa04726	Serotonergic synapse	-1.8	4	3	APP GNG3 HTR2B PRKACG

hsa05031	Amphetamine addiction	-1.7	5.1	3.1	GRIN1 PRKACG PPP1R1B
hsa05223	Non-small cell lung cancer	-1.6	4.8	3	E2F3 FOXO3 RASSF5
hsa04926	Relaxin signaling pathway	-1.6	3.6	2.8	ARRB1 GNG3 NFKBIA PRKACG
hsa04728	Dopaminergic synapse	-1.6	3.5	2.7	ARRB1 GNG3 PRKACG PPP1R1B
hsa05162	Measles	-1.5	3.3	2.6	CSNK2A2 JAK1 NFKBIA EIF2AK4
hsa04371	Apelin signaling pathway	-1.5	3.3	2.6	GNG3 SERPINE1 PRKACG PRKCE
hsa04151	PI3K-Akt signaling pathway	-1.5	2.3	2.3	AREG FOXO3 GNG3 JAK1 KITLG PRL MLST8
hsa04211	Longevity regulating pathway	-1.4	3.9	2.6	FOXO3 PRKACG IRS2
hsa05235	PD-L1 expression and PD-1 checkpoint pathway in cancer	-1.4	3.9	2.6	CSNK2A2 JAK1 NFKBIA
hsa04727	GABAergic synapse	-1.4	3.9	2.6	GNG3 PRKACG SLC38A3
hsa04010	MAPK signaling pathway	-1.4	2.4	2.2	AREG ARRB1 KITLG PRKACG TAOK3 CACNA2D4
hsa05032	Morphine addiction	-1.4	3.8	2.5	ARRB1 GNG3 PRKACG
hsa04218	Cellular senescence	-1.3	3	2.3	E2F3 FOXO3 SERPINE1 RASSF5
hsa05160	Hepatitis C	-1.3	3	2.3	E2F3 JAK1 NFKBIA EIF2AK4

Table S11: Pathway enrichment results for palpitations at Day 30

GO	Description	LogP	Enrichment	Z-score	genes
hsa05152	Tuberculosis	-7.6	6.7	8.3	CAMK2A CAMK2G CASP8 FCGR3B IFNGR2 IL10RB LAMP2 PP P3R1 MAPK13 RAF1 TLR4 TLR6 TLR9 TIRAP
hsa05235	PD-L1 expression and PD-1 checkpoint pathway in cancer	-5	7.8	6.9	IFNGR2 PPP3R1 MAPK13 RAF1 TLR4 CD274 TLR9 TIRAP
hsa04218	Cellular senescence	-4.1	5	5.4	CCND3 CDKN2A HLA-C CXCL8 NBN PPP3R1 MAPK13 RAF1 M CU
hsa05132	Salmonella infection	-3.8	3.8	4.8	CASP8 CXCL8 PAK1 MAPK13 RAF1 TLR4 TLR6 RIPK3 TLR9 A CTR3B TIRAP

hsa05142	Chagas disease	-3.7	5.9	5.4	CASP8 IFNGR2 CXCL8 MAPK13 TLR4 TLR6 TLR9
hsa04620	Toll-like receptor signaling pathway	-3.7	5.8	5.3	CASP8 CXCL8 MAPK13 TLR4 TLR6 TLR9 TIRAP
hsa05417	Lipid and atherosclerosis	-3.7	4	4.8	CAMK2A CAMK2G CASP8 ICAM1 CXCL8 PPP3R1 MAPK13 TLR4 TLR6 TIRAP
hsa05163	Human cytomegalovirus infection	-3.5	3.8	4.6	CASP8 CDKN2A GNG3 HLA-C CXCL8 IL10RB PPP3R1 MAPK13 RAF1 SP1
hsa04613	Neutrophil extracellular trap formation	-3.4	4.1	4.6	FCGR3B FPR2 MAPK13 RAF1 SEPLIG TLR4 H4C5 H4-16 H2BC18
hsa05200	Pathways in cancer	-3.3	2.6	4	CAMK2A CAMK2G CASP8 CCND3 CDKN2A FGF7 GNG3 IFNR2 CXCL8 IL15RA JAK3 RAF1 SP1 TGFA RALBP1 DAPK2
hsa04217	Necroptosis	-3.3	4.4	4.6	CAMK2A CAMK2G CASP8 IFNGR2 JAK3 TLR4 RIPK3 CHMP1B
hsa04650	Natural killer cell mediated cytotoxicity	-3.1	4.6	4.5	FCGR3B HLA-C ICAM1 IFNGR2 PAK1 PPP3R1 RAF1
hsa05219	Bladder cancer	-2.9	8.4	5.2	CDKN2A CXCL8 RAF1 DAPK2
hsa04140	Autophagy - animal	-2.9	4.3	4.2	LAMP2 RAF1 DAPK2 GABARAPL1 WIPI1 SUPT20H VMP1
hsa04360	Axon guidance	-2.9	3.8	4.1	CAMK2A CAMK2G PAK1 PPP3R1 RAF1 SEMA7A PLXNC1 LRR C4
hsa04621	NOD-like receptor signaling pathway	-2.9	3.8	4.1	CASP8 DEFA3 CXCL8 MAPK13 TLR4 RIPK3 GABARAPL1 MCU
hsa05214	Glioma	-2.7	5.8	4.5	CAMK2A CAMK2G CDKN2A RAF1 TGFA
hsa05167	Kaposi sarcoma-associated herpesvirus infection	-2.7	3.6	3.9	CASP8 GNG3 HLA-C ICAM1 CXCL8 PPP3R1 MAPK13 RAF1
hsa05144	Malaria	-2.6	6.9	4.5	ICAM1 CXCL8 TLR4 TLR9
hsa05203	Viral carcinogenesis	-2.6	3.4	3.7	CASP8 CCND3 CDKN2A HLA-C JAK3 H4C5 H4-16 H2BC18
hsa05161	Hepatitis B	-2.5	3.7	3.8	CASP8 CXCL8 JAK3 MAPK13 RAF1 TLR4 TIRAP
hsa04012	ErbB signaling pathway	-2.5	5.1	4.1	CAMK2A CAMK2G PAK1 RAF1 TGFA
hsa05170	Human immunodeficiency virus 1 infection	-2.5	3.3	3.6	CASP8 GNG3 HLA-C PAK1 PPP3R1 MAPK13 RAF1 TLR4
hsa05164	Influenza A	-2.4	3.5	3.6	CASP8 CCND3 ICAM1 IFNGR2 CXCL8 RAF1 TLR4

hsa05150	Staphylococcus aureus infection	-2.3	4.5	3.7	DEFA3 FCGR3B FPR2 ICAM1 SELPLG
hsa04666	Fc gamma R-mediated phagocytosis	-2.3	4.5	3.7	FCGR3B INPP5D PAK1 RAF1 ACTR3B
hsa05202	Transcriptional misregulation in cancer	-2.1	3.2	3.2	BCL6 DEFA3 EYA1 CXCL8 CDK14 SP1 KDM6A
hsa04625	C-type lectin receptor signaling pathway	-2.1	4.2	3.5	CASP8 PAK1 PPP3R1 MAPK13 RAF1
hsa04720	Long-term potentiation	-2.1	5.2	3.7	CAMK2A CAMK2G PPP3R1 RAF1
hsa04514	Cell adhesion molecules	-2.1	3.5	3.3	HLA-C ICAM1 SELPLG MPZL1 CD274 LRRC4
hsa04664	Fc epsilon RI signaling pathway	-2.1	5.1	3.7	ALOX5AP INPP5D MAPK13 RAF1
hsa05130	Pathogenic Escherichia coli infection	-2.1	3.1	3.2	CASP8 CXCL8 PAK1 MAPK13 TLR4 ACTR3B TIRAP
hsa05223	Non-small cell lung cancer	-2	4.8	3.5	CDKN2A JAK3 RAF1 TGFA
hsa05145	Toxoplasmosis	-2	3.9	3.3	CASP8 IFNGR2 IL10RB MAPK13 TLR4
hsa04725	Cholinergic synapse	-2	3.8	3.3	CAMK2A CAMK2G GNG3 KCNJ2 KCNQ2
hsa04115	p53 signaling pathway	-2	4.7	3.5	CASP8 CCND3 CDKN2A COP1
hsa00260	Glycine, serine and threonine metabolism	-2	6.5	3.8	ALAS1 AOC2 AOC3
hsa04630	JAK-STAT signaling pathway	-1.9	3.2	3	CCND3 IFNGR2 IL10RB IL15RA JAK3 RAF1
hsa05133	Pertussis	-1.9	4.6	3.4	CXCL8 MAPK13 TLR4 TIRAP
hsa05212	Pancreatic cancer	-1.9	4.6	3.4	CDKN2A RAF1 TGFA RALBP1
hsa05140	Leishmaniasis	-1.9	4.5	3.3	FCGR3B IFNGR2 MAPK13 TLR4
hsa04310	Wnt signaling pathway	-1.9	3.1	3	CAMK2A CAMK2G CCND3 PPP3R1 DAAM2 SOST
hsa05166	Human T-cell leukemia virus 1 infection	-1.8	2.7	2.8	CCND3 CDKN2A HLA-C ICAM1 IL15RA JAK3 PPP3R1
hsa04926	Relaxin signaling pathway	-1.8	3.4	2.9	ACTA2 GNG3 MAPK13 RAF1 RXFP2
hsa04068	FoxO signaling pathway	-1.7	3.3	2.9	BCL6 FOXO4 MAPK13 RAF1 GABARAPL1
hsa04014	Ras signaling pathway	-1.7	2.6	2.7	FGF7 GNG3 FOXO4 PAK1 RAF1 TGFA RALBP1
hsa04658	Th1 and Th2 cell differentiation	-1.7	3.8	2.9	IFNGR2 JAK3 PPP3R1 MAPK13
hsa04151	PI3K-Akt signaling pathway	-1.6	2.2	2.5	CCND3 FGF7 GNG3 IBSP ITGB6 JAK3 RAF1 TGFA TLR4
hsa04912	GnRH signaling pathway	-1.6	3.7	2.8	CAMK2A CAMK2G MAPK13 RAF1
hsa05162	Measles	-1.6	3.1	2.7	CASP8 CCND3 JAK3 TLR4 TLR9

hsa04936	Alcoholic liver disease	-1.6	3	2.6	CASP8 CXCL8 MAPK13 TLR4 TIRAP
hsa01522	Endocrine resistance	-1.6	3.5	2.7	CDKN2A MAPK13 RAF1 SP1
hsa05134	Legionellosis	-1.5	4.6	2.9	CASP8 CXCL8 TLR4
hsa04933	AGE-RAGE signaling pathway in diabetic complications	-1.5	3.5	2.7	ICAM1 CXCL8 MAPK13 THBD
hsa05169	Epstein-Barr virus infection	-1.5	2.6	2.4	CASP8 CCND3 HLA-C ICAM1 JAK3 MAPK13
hsa04370	VEGF signaling pathway	-1.5	4.4	2.8	PPP3R1 MAPK13 RAF1
hsa04145	Phagosome	-1.5	2.8	2.5	FCGR3B HLA-C LAMP2 TLR4 TLR6
hsa05416	Viral myocarditis	-1.5	4.3	2.8	CASP8 HLA-C ICAM1
hsa05205	Proteoglycans in cancer	-1.5	2.5	2.4	CAMK2A CAMK2G PAK1 MAPK13 RAF1 TLR4
hsa04660	T cell receptor signaling pathway	-1.5	3.3	2.6	PAK1 PPP3R1 MAPK13 RAF1
hsa04064	NF-kappa B signaling pathway	-1.5	3.3	2.6	ICAM1 CXCL8 TLR4 TIRAP
hsa04921	Oxytocin signaling pathway	-1.5	2.8	2.4	CAMK2A CAMK2G KCNJ2 PPP3R1 RAF1
hsa04922	Glucagon signaling pathway	-1.4	3.2	2.5	CAMK2A CAMK2G PPP3R1 PPP4R3C
hsa04659	Th17 cell differentiation	-1.4	3.2	2.5	IFNGR2 JAK3 PPP3R1 MAPK13
hsa04066	HIF-1 signaling pathway	-1.4	3.2	2.5	CAMK2A CAMK2G IFNGR2 TLR4
hsa04668	TNF signaling pathway	-1.4	3.1	2.4	CASP8 ICAM1 MAPK13 RIPK3
hsa04810	Regulation of actin cytoskeleton	-1.4	2.4	2.2	FGF7 ITGB6 PAK1 RAF1 ACTR3B FGD3
hsa05031	Amphetamine addiction	-1.3	3.8	2.5	CAMK2A CAMK2G PPP3R1
hsa05211	Renal cell carcinoma	-1.3	3.8	2.5	PAK1 RAF1 TGFA
hsa05120	Epithelial cell signaling in Helicobacter pylori infection	-1.3	3.7	2.5	CXCL8 PAK1 MAPK13
hsa04622	RIG-I-like receptor signaling pathway	-1.3	3.7	2.5	CASP8 CXCL8 MAPK13
hsa04722	Neurotrophin signaling pathway	-1.3	2.9	2.3	CAMK2A CAMK2G MAPK13 RAF1

Table S12: Pathway enrichment results for purple or dark lips at Day 30

GO	Description	LogP	Enrichment	Z-score	genes
hsa05014	Amyotrophic lateral sclerosis	-16	3.1	10	ATP5F1A ATP5F1B ATP5F1E ATP5PB ATP5MC3 ATP5PO CASP3 COX6C COX7A 2 COX7B COX7C EIF2S1 HNRNPA1 HNRNPA2B1 KIF5B NCBP1 NDUFA4 NDUFA 9 NDUFAB1 NDUFB2 NDUFB4 NDUFB5 NDUFS3 NDUFS4 NUP88 PIK3C3 PPP3C B PPP3CC PSMA1 PSMA2 PSMA3 PSMA5 PSMA6 PSMA7 PSMC6 PSMD1 PSMD4 PSMD9 RAB5A SDHA SDHB SRSF7 SOD1 VDAC1 TUBA1A BECN1 COX7A2L EIF 2AK3 MTR3 NUP58 RB1CC1 PSMD6 NUP153 ALYREF DCTN6 UQCR11 TARDB P CHMP2B TBK1 NUP54 CYCS ATG2B WDR41 NUP107 DERL1 SEH1L DNAL1 TO MM40L C9orf72 SLC25A5 ATP5F1A ATP5F1B ATP5F1E ATP5PB ATP5MC3 ATP5PO CASP3 COX6 C COX7A2 COX7B COX7C CREB1 ATF2 CSNK2A2 EIF2S1 FYN HSPA8 KIF5B ND UFA4 NDUFA9 NDUFAB1 NDUFB2 NDUFB4 NDUFB5 NDUFS3 NDUFS4 PIK3CA PPP3CB PPP3CC PRKACB MAPK3 PRNP PSMA1 PSMA2 PSMA3 PSMA5 PSMA6 P SMA7 PSMC6 PSMD1 PSMD4 PSMD9 SDHA SDHB SOD1 VDAC1 VDAC2 VDAC3 TUBA1A COX7A2L EIF2AK3 PSMD6 UQCR11 CYCS SLC25A5 FAS ATP5F1A ATP5F1B ATP5F1E ATP5PB ATP5MC3 ATP5PO CALM2 CAMK2D CASP3 CASP7 COX6C COX7A2 COX7B COX7C CSNK2A2 CTNNB1 EIF 2S1 KIF5B KRAS ATXN3 NDUFA4 NDUFA9 NDUFAB1 NDUFB2 NDUFB4 NDUF B5 NDUFS3 NDUFS4 NRAS PIK3C3 PPP3CB PPP3CC PRKCB MAPK3 PRNP PSMA 1 PSMA2 PSMA3 PSMA5 PSMA6 PSMA7 PSMC6 PSMD1 PSMD4 PSMD9 RAB5A R PS27A SDHA SDHB SOD1 VDAC1 VDAC2 VDAC3 TUBA1A AXIN2 BECN1 COX7 A2L EIF2AK3 RB1CC1 PSMD6 DCTN6 UQCR11 FRAT2 TARDBP CHMP2B TBK1 CYCS ATG2B WDR41 DERL1 DNAL1 TOMM40L WNT3A CSNK1A1L C9orf72
hsa05020	Prion disease	-14	3.2	9.6	
hsa05022	Pathways of neurodegeneration - multiple diseases	-14	2.6	9.1	
hsa05010	Alzheimer disease	-14	2.8	9.2	SLC25A5 FAS ATP5F1A ATP5F1B ATP5F1E ATP5PB ATP5MC3 ATP5PO CALM2

					CASP3 CASP7 COX6C COX7A2 COX7B COX7C CSNK2A2 CTNNB1 EIF2S1 KIF5B KRAS NDUFA4 NDUFA9 NDUFAB1 NDUFB2 NDUFB4 NDUFB5 NDUFS3 NDUF S4 SLC11A2 NRAS PIK3C3 PIK3CA PPP3CB PPP3CC MAPK3 PSMA1 PSMA2 PSM A3 PSMA5 PSMA6 PSMA7 PSMC6 PSMD1 PSMD4 PSMD9 SDHA SDHB ADAM17 VDAC1 VDAC2 VDAC3 TUBA1A AXIN2 BECN1 NAE1 COX7A2L EIF2AK3 RB1C C1 PSMD6 UQCR11 FRAT2 CYCS ATG2B SLC39A9 SLC39A10 WNT3A CSNK1A1 L
hsa05016	Huntington disease	-14	3.1	9.3	SLC25A5 ATP5F1A ATP5F1B ATP5F1E ATP5PB ATP5MC3 ATP5PO CASP3 COX6 C COX7A2 COX7B COX7C CREB1 HDAC2 KIF5B NDUFA4 NDUFA9 NDUFAB1 ND UFB2 NDUFB4 NDUFB5 NDUFS3 NDUFS4 NRF1 PIK3C3 POLR2B POLR2D POL R2H POLR2K PSMA1 PSMA2 PSMA3 PSMA5 PSMA6 PSMA7 PSMC6 PSMD1 PSM D4 PSMD9 REST SDHA SDHB SOD1 TFAM VDAC1 VDAC2 VDAC3 TUBA1A BEC N1 COX7A2L TBPL1 RB1CC1 PSMD6 DCTN6 UQCR11 CYCS ATG2B DNAL1 DHX15 HNRNPA1 HNRNPC HSPA8 MAGOH NCBP1 SNU13 PLRG1 SRSF1 SRSF7 TRA2B SNRPF SRSF9 SNRNP40 DHX38 RBM8A ALYREF SMNDC1 BCAS2 SRSF1 0 SRSF8 SNRNP27 NCBP2 SF3B3 SF3B1 LSM5 RBMX CRNKL1 CDC40 SF3B6 LS M8 PRPF38B PRPF40A RBM22 RBM25 PHF5A RBM17 ISY1-RAB43
hsa03040	Spliceosome	-14	4.2	9.9	SLC25A5 ATP5F1A ATP5F1B ATP5F1E ATP5PB ATP5MC3 ATP5PO CALM2 CAM K2D CASP3 COX6C COX7A2 COX7B COX7C EIF2S1 GNAI3 KIF5B NDUFA4 NDU FA9 NDUFAB1 NDUFB2 NDUFB4 NDUFB5 NDUFS3 NDUFS4 NFE2L2 SLC11A2 PR KACB PSMA1 PSMA2 PSMA3 PSMA5 PSMA6 PSMA7 PSMC6 PSMD1 PSMD4 PS MD9 RPS27A SDHA SDHB SOD1 VDAC1 VDAC2 VDAC3 TUBA1A COX7A2L EIF 2AK3 PSMD6 UQCR11 CYCS SLC39A9 SLC39A10
hsa05012	Parkinson disease	-13	3.2	9.3	FAU RPL5 RPL6 RPL7 RPL15 RPL17 RPL24 RPL27 RPL30 RPL31 RPL34 RPL35A R PL39 RPS3A RPS6 RPS7 RPS10 RPS15A RPS17 RPS18 RPS20 RPS21 RPS23 RPS27 RPS27A RPS29 RPL23 RPL35 RPS27L RSL24D1 MRPL30 MRPL16 MRPL20 MRPS1
hsa03010	Ribosome	-11	3.7	8.7	

hsa04120	Ubiquitin mediated proteolysis	-9.8	3.6	8.1	4 MRPS11 RPL36A-HNRNPH2 BIRC2 UBE2K MDM2 NEDD4 RPS27A SKP1 ELOC UBE2D2 UBE2E1 UBE2I UBE2N UBE3A CUL4B CUL3 CUL2 UBA3 UBE4A RNF7 UBE3C UBA2 SAE1 WWP1 FBXW11 ANAPC13 HERC4 ANAPC4 UBR5 UBA6 UBE2W FBXW7 KLHL9 UBE2F CTSL EIF2S1 HMGB1 IGBP1 KRAS NRAS PIK3C3 PIK3CA PPP2CA PPP2CB PRKA A1 PRKACB PRKCQ MAPK3 RHEB MAP3K7 UVRAG BECN1 EIF2AK3 RB1CC1 R RAS2 MRAS TBK1 SH3GLB1 ATG2B WDR41 ATG3 RAB33B ATG4C ATG4A C9orf72 ATP5F1A ATP5F1B ATP5F1E ATP5PB ATP5MC3 ATP5PO COX6C COX7A2 COX7B COX7C COX15 NDUFA4 NDUFA9 NDUFAB1 NDUFB2 NDUFB4 NDUFB5 NDUFS3 NDUFS4 SDHA SDHB COX7A2L ATP6V1G1 ATP5MG UQCR11 PPA2 ATP6V1D CYCS ATP6V1E2 ATP6V1C2 FAU IFNA13 IL6ST OAS2 PIK3CA PRKCB MAPK3 RPL5 RPL6 RPL7 RPL15 RPL17 RPL24 RPL27 RPL30 RPL31 RPL34 RPL35A RPL39 RPS3A RPS6 RPS7 RPS10 RPS15A RPS17 RPS18 RPS20 RPS21 RPS23 RPS27 RPS27A RPS29 ADAM17 MAP3K7 RPL23 RPL35 TBK1 RPS27L IRAK4 RSL24D1 RPL36A-HNRNPH2 ATP5F1A ATP5F1B ATP5F1E ATP5PB ATP5MC3 ATP5PO BMP8B COX6C COX7A2 COX7B COX7C COX15 CREB1 ATF2 KRAS NDUFA4 NDUFA9 NDUFAB1 NDUFB2 NDUFS4 NRAS PRKAA1 PRKACB PRKAG1 RHEB RPS6 SDHA SDHB SMARCE1 COX7A2L ATP5MG UQCR11 PRKAG2 COA1 COX14 COX20 NDUFAF8 BIRC2 ARF6 RHOG RHOH CASP3 CASP7 CDC42 CTNNB1 DYN1I2 HSP90AA1 KIF5B M6PR PIK3C3 PIK3CA MAPK3 PTPRC RAB5A RALA SKP1 MAP3K7 DYNLT3 TUBA1A RIPK2 TNFRSF10B CYTH1 ABI1 ARPC2 MYL12A EXOC5 DCTN6 LY96 IRAK4 SNX9 CYCS ARL8B EXOC2 ELMO2 RILP MYL12B TIRAP SNX33 EEF1A1 TNPO1 MAGOH NCBP1 NUP88 RAN SUMO3 SUMO2 UBE2I SUMO1 NUP
hsa04140	Autophagy - animal	-9.3	3.5	7.8	
hsa00190	Oxidative phosphorylation	-9.2	3.6	7.8	
hsa05171	Coronavirus disease - COVID-19	-8.9	2.8	7.3	
hsa04714	Thermogenesis	-8.4	2.8	7	
hsa05132	Salmonella infection	-8	2.7	6.7	
hsa03013	Nucleocytoplasmic	-7.4	3.6	6.9	

	transport				
hsa05208	Chemical carcinogenesis - reactive oxygen species	-7.4	2.7	6.5	58 RBM8A NUP153 ALYREF IPO8 IPO7 NCBP2 XPO7 NMD3 PHAX NUP54 NUP107 THOC7 SEH1L ABL1 ABL2 ACP1 SLC25A5 ATP5F1A ATP5F1B ATP5F1E ATP5PB ATP5MC3 ATP5PO COX6C COX7A2 COX7B COX7C KRAS NDUFA4 NDUFA9 NDUFAB1 NDUB2 NDUFB4 NDUFB5 NDUFS3 NDUFS4 NFE2L2 NRAS PIK3CA MAPK3 PTPN1 SDHA SDHB SOD1 VDAC1 VDAC2 VDAC3 COX7A2L UQCR11 PRKD3 SLC25A5 GTF2B ATXN3 OPA1 PIK3C3 PIK3CA PRKCB PSMA1 PSMA2 PSMA3 PSMA5 PSMA6 PSMA7 PSMC6 PSMD1 PSMD4 PSMD9 VDAC1 VDAC2 VDAC3 BCN1 TBPL1 RB1CC1 PSMD6 PUM2 ATXN10 CYCS ATG2B MAGOH NCBP1 PPP1CB PPP1CC PPP2CA PPP2CB PPP2R2A PPP2R5C RNGTT RB8A ALYREF HBS1L NUDT21 CPSF6 NCBP2 TARDBP DAZAP1 PPP2R3B PCF11 PPP2R3C PAPOLG BCL2L2-PABPN1
hsa05017	Spinocerebellar ataxia	-7.3	3.2	6.6	SMA5 PSMA6 PSMA7 PSMC6 PSMD1 PSMD4 PSMD9 VDAC1 VDAC2 VDAC3 BCN1 TBPL1 RB1CC1 PSMD6 PUM2 ATXN10 CYCS ATG2B MAGOH NCBP1 PPP1CB PPP1CC PPP2CA PPP2CB PPP2R2A PPP2R5C RNGTT RB8A ALYREF HBS1L NUDT21 CPSF6 NCBP2 TARDBP DAZAP1 PPP2R3B PCF11 PPP2R3C PAPOLG BCL2L2-PABPN1
hsa03015	mRNA surveillance pathway	-7	3.7	6.7	M8A ALYREF HBS1L NUDT21 CPSF6 NCBP2 TARDBP DAZAP1 PPP2R3B PCF11 PPP2R3C PAPOLG BCL2L2-PABPN1
hsa03050	Proteasome	-6.4	4.9	6.8	PSMA1 PSMA2 PSMA3 PSMA5 PSMA6 PSMA7 PSMC6 PSMD1 PSMD4 PSMD9 PSME2 PSMD6 PSME4 POMP AMFR EIF2S1 DNAJA1 HSPA8 HSP90AA1 ATXN3 NFE2L2 SKP1 SSR3 SEC62 UBAM2D2 UFD1 MBTPS1 EIF2AK3 BAG2 SEC24D DNAJA2 SEC24B ERP29 UBXN4 TRAM1 SEC61G ERLEC1 YOD1 NGLY1 SAR1A DNAJC1 DERL1 STT3B FAS CASP3 CASP7 CDC42 COX6C COX7A2 COX7B COX7C EIF2S1 NDUFA4 NDUFA9 NDUFAB1 NDUFB2 NDUFB4 NDUFB5 NDUFS3 NDUFS4 PIK3CA PRKAA1 PRKAG1 SDHA SDHB COX7A2L EIF2AK3 UQCR11 PRKAG2 CYCS SLC25A5 ATP5F1A ATP5F1B ATP5F1E ATP5PB ATP5MC3 ATP5PO CAMK2D COX6C COX7A2 COX7B COX7C NDUFA4 NDUFA9 NDUFAB1 NDUFB2 NDUFB4 NDUFB5 NDUFS3 NDUFS4 PIK3CA PPP1CB PPP1CC PRKCB SDHA SDHB VDAC1 VDAC2 VDAC3 COX7A2L TBC1D4 UQCR11
hsa04141	Protein processing in endoplasmic reticulum	-6.1	2.7	5.8	ARF4 ARF6 CAPZA1 CAPZA2 CDC42 HSPA8 KIF5B MDM2 NEDD4 RAB4A RAB5A SNX1 SNX4 USP8 CYTH1 VPS26A ARPC2 STAM2 WWP1 RAB11FIP2 WASHC4
hsa04932	Non-alcoholic fatty liver disease	-6	2.8	5.8	UFA9 NDUFAB1 NDUFB2 NDUFB4 NDUFB5 NDUFS3 NDUFS4 PIK3CA PRKAA1 PRKAG1 SDHA SDHB COX7A2L EIF2AK3 UQCR11 PRKAG2 CYCS SLC25A5 ATP5F1A ATP5F1B ATP5F1E ATP5PB ATP5MC3 ATP5PO CAMK2D COX6C COX7A2 COX7B COX7C NDUFA4 NDUFA9 NDUFAB1 NDUFB2 NDUFB4 NDUFB5 NDUFS3 NDUFS4 PIK3CA PPP1CB PPP1CC PRKCB SDHA SDHB VDAC1 VDAC2 VDAC3 COX7A2L TBC1D4 UQCR11
hsa05415	Diabetic cardiomyopathy	-6	2.5	5.7	ARF4 ARF6 CAPZA1 CAPZA2 CDC42 HSPA8 KIF5B MDM2 NEDD4 RAB4A RAB5A SNX1 SNX4 USP8 CYTH1 VPS26A ARPC2 STAM2 WWP1 RAB11FIP2 WASHC4
hsa04144	Endocytosis	-5.6	2.3	5.3	

hsa04660	T cell receptor signaling pathway	-5.3	3.1	5.5	ACAP2 CHMP2B SNX5 CHMP2A CHMP4A SNX12 SH3GLB1 CHMP5 VTA1 CHMP3 CHMP1B SNX6 ARAP2 VPS37A AGAP6 CD3G CD8A CDC42 MAP3K8 DLG1 FYN ITK KRAS NCK1 NRAS PIK3CA PPP3CB PPP3CC PRKCQ MAPK3 PTPRC MAP3K7 BCL10 VAV3 MALT1
hsa01200	Carbon metabolism	-5.1	2.9	5.4	ADH5 DLAT DLD FH GLUD1 GOT1 GOT2 IDH3A MDH1 PGK1 PGK2 PRPS1 PRPS2 PSPH SDHA SDHB SHMT1 SUCLG2 SUCLA2 PSAT1 HKDC1 SLC25A5 CALM2 E2F4 ETS1 KRAS MDM2 NBN NRAS PIK3CA PPP1CB PPP1CC PP3CB PPP3CC MAPK3 RB1 RHEB VDAC1 VDAC2 VDAC3 RAS2 MRAS FBXW11 TRPM7 HIPK1 LIN9
hsa04218	Cellular senescence	-4.9	2.6	5.1	PP3CB PPP3CC MAPK3 RB1 RHEB VDAC1 VDAC2 VDAC3 RAS2 MRAS FBXW11 TRPM7 HIPK1 LIN9 ASAHI FYN GNAI3 KRAS NRAS PIK3CA PPP2CA PPP2CB PPP2R2A PPP2R5C PRKCB MAPK3 ROCK1 NSMAF SGPL1 SPTLC1 GNA13 PPP2R3B PPP2R3C SGPP1 CERS6
hsa00020	Citrate cycle (TCA cycle)	-4.2	4.8	5.4	DLAT DLD FH IDH3A MDH1 SDHA SDHB SUCLG2 SUCLA2 FAS CALM2 CAMK2D CASP3 CASP7 CDC42 CYP2A7 EIF2S1 HSPA8 HSP90AA1 INFE2L2 NRAS PIK3CA PPP3CB PPP3CC MAPK3 RAP1B MAP3K7 TNFRSF10B EIF2AK3 VAV3 LY96 TBK1 IRAK4 CYCS MIB1 TIRAP
hsa05417	Lipid and atherosclerosis	-4.2	2.2	4.4	FNA13 KRAS NFE2L2 NRAS PIK3CA PPP3CB PPP3CC MAPK3 RAP1B MAP3K7 TNFRSF10B EIF2AK3 VAV3 LY96 TBK1 IRAK4 CYCS MIB1 TIRAP BTG1 EXOSC10 TOB1 EXOSC8 EXOSC7 MTREX LSM5 CNOT7 LSM8 XRN1 CNOT6 TENT4B PNPT1 DCP2 CNOT6L
hsa03018	RNA degradation	-4	3.1	4.7	CALM2 IMPA1 IMPA2 INPP1 INPP4A ITPK1 MTM1 OCRL PIK3C3 PIK3CA PIP4K2A PRKCB MTMR1 MTMR6 SACM1L BPNT2 PIKFYVE
hsa04070	Phosphatidylinositol signaling system	-4	2.8	4.6	IGBP1 PIK3C3 PPP2CA PPP2CB BECN1 ATG2B ATG3 ATG4C ATG4A CSNK2A2 DKC1 FBL SNU13 RAN RPP38 WDR43 REXO2 GNL3 NOB1 NMD3 FCF1 UTP18 SBDS XRN1 LSG1 NOL6 SPATA5
hsa04136	Autophagy - other	-4	4.5	5.1	IMPA1 IMPA2 INPP1 INPP4A ITPK1 MTM1 OCRL PIK3C3 PIK3CA PIP4K2A MTMR1 MTMR6 SACM1L PIKFYVE
hsa03008	Ribosome biogenesis in eukaryotes	-3.9	2.7	4.5	FAS B2M CALM2 CASP3 CCR3 CREB1 ATF2 CTNNB1 GNAI3 IFNA13 IL10RA ITGB1 EXOSC10 TOB1 EXOSC8 EXOSC7 MTREX LSM5 CNOT7 LSM8 XRN1 CNOT6 TENT4B PNPT1 DCP2 CNOT6L
hsa00562	Inositol phosphate metabolism	-3.9	3.1	4.6	ACAP2 CHMP2B SNX5 CHMP2A CHMP4A SNX12 SH3GLB1 CHMP5 VTA1 CHMP3 CHMP1B SNX6 ARAP2 VPS37A AGAP6 CD3G CD8A CDC42 MAP3K8 DLG1 FYN ITK KRAS NCK1 NRAS PIK3CA PPP3CB PPP3CC PRKCQ MAPK3 PTPRC MAP3K7 BCL10 VAV3 MALT1
hsa05163	Human cytomegalovirus	-3.8	2.1	4.2	ASAH1 FYN GNAI3 KRAS NRAS PIK3CA PPP2CA PPP2CB PPP2R2A PPP2R5C PRKCB MAPK3 ROCK1 NSMAF SGPL1 SPTLC1 GNA13 PPP2R3B PPP2R3C SGPP1 CERS6 BTG1 EXOSC10 TOB1 EXOSC8 EXOSC7 MTREX LSM5 CNOT7 LSM8 XRN1 CNOT6 TENT4B PNPT1 DCP2 CNOT6L

	infection				
hsa05205	Proteoglycans in cancer	-3.7	2.1	4.1	GAV KRAS MDM2 NRAS PIK3CA PPP3CB PPP3CC PRKACB PRKCB MAPK3 PTG ER2 RB1 RHEB ROCK1 GNA13 GNB5 TBK1 CYCS FAS CAMK2D CASP3 CD44 CDC42 CTNNB1 CTSL EIF4B IGF2 ITGAV KRAS MD M2 PPP1R12A NRAS PIK3CA PPP1CB PPP1CC PRKACB PRKCB MAPK3 ROCK1 RPS6 TFAP4 VAV3 RRAS2 MRAS WNT3A ARNTL CALM2 CAMK2D CREB1 ATF2 GNAI3 KIF5B PPP1CB PPP1CC PPP2CA P PP2CB PPP2R2A PPP2R5C PPP3CB PPP3CC PRKACB PRKCB GNB5 PPP2R3B PPP 2R3C
hsa04728	Dopaminergic synapse	-3.7	2.4	4.3	PP2CB PPP2R2A PPP2R5C PPP3CB PPP3CC PRKACB PRKCB GNB5 PPP2R3B PPP 2R3C
hsa04910	Insulin signaling pathway	-3.5	2.4	4.1	CALM2 EIF4E KRAS NRAS PIK3CA PPP1CB PPP1CC PRKAA1 PRKACB PRKAG1 MAPK3 PTPN1 RHEB RPS6 SHC1 FLOT1 RHOQ PRKAG2 HKDC1 SOCS4 ARF6 CD44 CDC42 MDM2 PIK3C3 PIK3CA PRKCQ MAPK3 ROCK1 RPS27A SKP1
hsa05131	Shigellosis	-3.5	2	3.9	MAP3K7 UBE2D2 UBE2N VDAC1 BECN1 RIPK2 BCL10 CYTH1 ARPC2 MYL12A MALT1 FBXW11 TBK1 CYCS ELMO2 HKDC1 SHARPIN TIFA MYL12B SLC25A5 BIRC2 FAS CAMK2D GLUD1 H2AZ1 HMGB1 HSP90AA1 IFNA13 IFNG
hsa04217	Necroptosis	-3.5	2.2	4	R1 VDAC1 VDAC2 VDAC3 TNFRSF10B CHMP2B CHMP2A CHMP4A CHMP5 CH MP3 TRPM7 CHMP1B SHARPIN CASP3 CDC42 CCR3 CREB1 ATF2 DDX3X DLG1 GTF2B GTF2E2 GTF2H3 HDAC2
hsa05203	Viral carcinogenesis	-3.4	2.1	3.9	IL6ST KRAS MDM2 NRAS PIK3CA PRKACB MAPK3 RASA2 RB1 UBE3A VDAC3 H4C13 KAT2B TBPL1 YWHAQ
hsa04137	Mitophagy - animal	-3.4	2.9	4.2	CSNK2A2 KRAS NRAS RPS27A BECN1 TAX1BP1 USP8 EIF2AK3 RRAS2 MRAS B CL2L13 TBK1 TOMM7
hsa04919	Thyroid hormone signaling pathway	-3.3	2.4	4	ATP1B3 CTNNB1 HDAC2 ITGAV KRAS MDM2 NRAS PFKFB2 PIK3CA PRKACB PRKCB MAPK3 RHEB KAT2B TBC1D4 MED13 MED4 MED30 ABL1 FAS ARF6 CASP3 CASP7 CDC42 FYN MYO5A NCK1 MAPK3 ROCK1 MAP3
hsa05130	Pathogenic Escherichia coli infection	-3.3	2	3.8	K7 TMBIM6 TUBA1A TNFRSF10B CYTH1 SEC24D ABI1 ARPC2 SEC24B GNA13 TMED10 IRAK4 CYCS TIRAP

hsa04014	Ras signaling pathway	-3.2	1.9	3.7	ABL1 ABL2 ARF6 CALM2 CDC42 EFNA1 ETS1 IGF2 KRAS NF1 NRAS PIK3CA PRKACB PRKCB MAPK3 RAB5A RALA RAP1B RASA2 SHC1 KSR1 GNB5 RRAS2 M RAS RASA3 TBK1 EXOC2 PLA2G12B
hsa04720	Long-term potentiation	-3.1	2.9	4	CALM2 CAMK2D KRAS NRAS PPP1CB PPP1CC PPP3CB PPP3CC PRKACB PRKCB MAPK3 RAP1B BIRC2 FAS B2M CASP3 EIF2S1 IFNA13 IFNGR1 OAS2 PIK3CA PPP1CB PPP1CC R HEB SRSF1 SRSF7 MAP3K7 ZNF12 ZNF23 ZNF124 ZNF140 ZNF223 SRSF9 EIF2B2 ZNF235 EIF2AK3 ZNF432 ALYREF ZNF443 ZNF267 SRSF8 TBK1 IRAK4 CYCS ZFP14 ZNF77 ZNF333 ZNF551 ZNF468 ZNF486 ZNF764 ZNF792 ZNF540 ZNF584 ZNF718 ZNF780A ZFP82 ZNF680 ZNF761 ZNF674 ZNF605
hsa05168	Herpes simplex virus 1 infection	-3.1	1.6	3.4	POLR2B POLR2D POLR2H POLR2K POLR1D POLR3B POLR2M POLR1B CALM2 CAMK2D PPP1CB PPP1CC PPP2CA PPP2CB PPP2R5C PPP3CB PPP3CC PRKACB MAPK3 SKP1 YWHAQ CPEB3 FBXW11 ANAPC13 ANAPC4 CPEB2 FAS CASP3 CDC42 CREB1 CTNNB1 DLG1 HDAC2 IFNA13 ITGA1 ITGA4 ITGAV KRAS MDM2 NRAS PIK3CA PPP2CA PPP2CB PPP2R2A PPP2R5C PRKACB MAPK3 RB1 RHEB UBE3A AXIN2 TBPL1 ATP6V1G1 PPP2R3B TBK1 ATP6V1D PPP2R3C WNT3A ATP6V1E2 CSNK1A1L ATP6V1C2 EIF4B EIF4E KRAS NRAS PIK3CA PRKAA1 PRKCB MAPK3 RHEB RPS6 ATP6V1 T RICKTOR
hsa03020	RNA polymerase	-3	3.8	4.2	
hsa04114	Oocyte meiosis	-2.9	2.2	3.6	
hsa05165	Human papillomavirus infection	-2.8	1.7	3.3	
hsa04150	mTOR signaling pathway	-2.8	2.1	3.5	
hsa03022	Basal transcription factors	-2.8	3.2	3.8	CCNH GTF2B GTF2E2 GTF2H3 TAF7 TAF12 TAF13 TBPL1 TAF5L
hsa04662	B cell receptor signaling pathway	-2.8	2.6	3.6	BTK FCGR2B KRAS NRAS PIK3CA PPP3CB PPP3CC PRKCB MAPK3 BCL10 VAV3 MALT1 DAPP1
hsa05170	Human immunodeficiency virus	-2.8	1.9	3.4	AP1G1 FAS B2M CALM2 CASP3 CD3G GNAI3 IFNA13 KRAS NRAS PIK3CA PPP3CB PPP3CC PRKCB MAPK3 SKP1 MAP3K7 ELOC CUL4B RNF7 GNB5 FBXW11 T

	1 infection				BK1 IRAK4 CYCS
hsa05169	Epstein-Barr virus infection	-2.8	1.9	3.4	FAS B2M BTK CASP3 CD3G CD44 CD58 HDAC2 IFNA13 MDM2 NEDD4 OAS2 PIK3CA PSMC6 PSMD1 PSMD4 RB1 MAP3K7 SAP30 PSMD6 TBK1 IRAK4 CYCS SAP30L
hsa05160	Hepatitis C	-2.8	2.1	3.4	FAS CASP3 CTNNB1 EIF2S1 IFNA13 EIF3E KRAS NRAS OAS2 PIK3CA PPP2CA PPP2CB PPP2R2A MAPK3 RB1 EIF2AK3 YWHAQ TBK1 CYCS RSAD2 CALM2 CAMK2D CASP3 CREB1 FCGR2B IFNA13 IFNGR1 IL10RA PIK3C3 PPP3C
hsa05152	Tuberculosis	-2.7	2	3.4	B PPP3CC MAPK3 RAB5A RFX5 RIPK2 KSR1 BCL10 MALT1 IRAK4 CYCS CLEC7A TIRAP
hsa01230	Biosynthesis of amino acids	-2.7	2.6	3.5	ACY1 GOT1 GOT2 IDH3A PGK1 PGK2 PRPS1 PRPS2 PSPH SHMT1 MAT2B PSAT1 FAS CALM2 CASP3 CCR3 CREB1 CTNNB1 IFNA13 IFNGR1 IL6ST KRAS NRAS PIK3C3 PIK3CA PPP3CB PPP3CC MAPK3 RB1 RPS27A BECN1 GNB5 TBK1 CYCS ATG3
hsa05167	Kaposi sarcoma-associated herpesvirus infection	-2.7	1.9	3.3	CDC42 ITGA1 ITGA4 ITGAE ITGAV KRAS PPP1R12A NRAS PIK3CA PIP4K2A PPRAS2 MRAS MYL12B PIKFYVE SPATA13 BIRC2 HSP90AA1 IFNA13 OAS2 PKN2 MAPK3 MAP3K7 VDAC1 VDAC2 VDAC3 RIPK2 SUGT1 CARD8 TBK1 IRAK4 TRPM7 ERBIN DHX33 SHARPIN GBP4 NEK7 NLRP6
hsa04810	Regulation of actin cytoskeleton	-2.6	1.8	3.2	P1CB PPP1CC MAPK3 ROCK1 TMSB4X ARPC2 VAV3 MYL12A GNA13 IQGAP2 RAS2 MRAS MYL12B PIKFYVE SPATA13 BIRC2 HSP90AA1 IFNA13 OAS2 PKN2 MAPK3 MAP3K7 VDAC1 VDAC2 VDAC3 RIPK2 SUGT1 CARD8 TBK1 IRAK4 TRPM7 ERBIN DHX33 SHARPIN GBP4 NEK7 NLRP6 CALM2 GNAI3 KRAS NRAS NRF1 PIK3C3 PRKAA1 PRKACB PRKAG1 MAPK3 RPS6 TFAM BECN1 GNA13 GNB5 RRAS2 MRAS PRKAG2
hsa04621	NOD-like receptor signaling pathway	-2.6	1.9	3.2	ARNTL CREB1 PRKAA1 PRKAG1 SKP1 FBXW11 PRKAG2 FAS CASP3 CREB1 ATF2 DDX3X IFNA13 KRAS NRAS PCNA PIK3CA PRKCB MAPK3 RB1 MAP3K7 VDAC3 YWHAQ TBK1 IRAK4 CYCS TIRAP
hsa04371	Apelin signaling pathway	-2.6	2.1	3.3	CASP3 CASP7 CLK1 EEF1A1 HSPA8 HBS1L BCL2L13 CYCS SAR1A CLK4 ARNTL CREB1 PRKAA1 PRKAG1 SKP1 FBXW11 PRKAG2 FAS CASP3 CREB1 ATF2 DDX3X IFNA13 KRAS NRAS PCNA PIK3CA PRKCB MAPK3 RB1 MAP3K7 VDAC3 YWHAQ TBK1 IRAK4 CYCS TIRAP
hsa05134	Legionellosis	-2.6	2.8	3.6	CASP3 CASP7 CLK1 EEF1A1 HSPA8 HBS1L BCL2L13 CYCS SAR1A CLK4 ARNTL CREB1 PRKAA1 PRKAG1 SKP1 FBXW11 PRKAG2 FAS CASP3 CREB1 ATF2 DDX3X IFNA13 KRAS NRAS PCNA PIK3CA PRKCB MAPK3 RB1 MAP3K7 VDAC3 YWHAQ TBK1 IRAK4 CYCS TIRAP
hsa04710	Circadian rhythm	-2.6	3.6	3.8	CASP3 CASP7 CLK1 EEF1A1 HSPA8 HBS1L BCL2L13 CYCS SAR1A CLK4 ARNTL CREB1 PRKAA1 PRKAG1 SKP1 FBXW11 PRKAG2 FAS CASP3 CREB1 ATF2 DDX3X IFNA13 KRAS NRAS PCNA PIK3CA PRKCB MAPK3 RB1 MAP3K7 VDAC3 YWHAQ TBK1 IRAK4 CYCS TIRAP
hsa05161	Hepatitis B	-2.6	2	3.2	CASP3 CASP7 CLK1 EEF1A1 HSPA8 HBS1L BCL2L13 CYCS SAR1A CLK4 ARNTL CREB1 PRKAA1 PRKAG1 SKP1 FBXW11 PRKAG2 FAS CASP3 CREB1 ATF2 DDX3X IFNA13 KRAS NRAS PCNA PIK3CA PRKCB MAPK3 RB1 MAP3K7 VDAC3 YWHAQ TBK1 IRAK4 CYCS TIRAP

hsa05166	Human T-cell leukemia virus 1 infection	-2.5	1.8	3.1	SLC25A5 B2M CD3G CREB1 ATF2 DLG1 ETS1 IL15 KRAS NRAS PIK3CA PPP3CB PPP3CC PRKACB MAPK3 RAN RANBP1 RB1 SPI1 VDAC1 VDAC2 VDAC3 KAT2B TBPL1 ANAPC13 ANAPC4
hsa05235	PD-L1 expression and PD-1 checkpoint pathway in cancer	-2.5	2.4	3.3	CD3G CSNK2A2 IFNGR1 KRAS NRAS PIK3CA PPP3CB PPP3CC PRKCQ MAPK3 EML4 BATF3 TIRAP
hsa04211	Longevity regulating pathway	-2.5	2.4	3.3	CREB1 ATF2 EIF4E KRAS NRAS PIK3CA PRKAA1 PRKACB PRKAG1 RHEB RB1 CC1 SESN1 PRKAG2
hsa05031	Amphetamine addiction	-2.5	2.6	3.4	CALM2 CAMK2D CREB1 ATF2 HDAC2 PPP1CB PPP1CC PPP3CB PPP3CC PRKACB PRKCB
hsa00470	D-Amino acid metabolism	-2.4	8.1	4.4	DAO GLS SRR
hsa04213	Longevity regulating pathway - multiple species	-2.3	2.6	3.2	HDAC2 HSPA8 KRAS NRAS PIK3CA PRKAA1 PRKACB PRKAG1 SOD1 PRKAG2
hsa04625	C-type lectin receptor signaling pathway	-2.3	2.2	3.1	CALM2 KRAS MDM2 NRAS PIK3CA PPP3CB PPP3CC MAPK3 KSR1 BCL10 MALT1 RRAS2 MRAS CLEC7A
hsa04261	Adrenergic signaling in cardiomyocytes	-2.3	1.9	3	ATP1B3 ATP2B1 CALM2 CAMK2D CREB1 ATF2 GNAI3 PPP1CB PPP1CC PPP2CA PPP2CB PPP2R2A PPP2R5C PRKACB MAPK3 SCN4B PPP2R3B PPP2R3C
hsa05162	Measles	-2.2	2	3	FAS CASP3 CD3G CSNK2A2 EIF2S1 FCGR2B HSPA8 IFNA13 CD46 OAS2 PIK3CA MAP3K7 EIF3H EIF2AK3 TBK1 IRAK4 CYCS
hsa04145	Phagosome	-2.2	1.9	2.9	CTSL DYNC1I2 FCGR2B ITGAV M6PR PIK3C3 RAB5A TUBA1A STX7 MARCO ATP6V1G1 SEC61G ATP6V1D CLEC7A RILP ATP6V1E2 PIKFYVE ATP6V1C2
hsa04650	Natural killer cell mediated cytotoxicity	-2.1	2	2.9	FAS CASP3 CD48 FYN IFNA13 IFNGR1 KRAS NRAS PIK3CA PPP3CB PPP3CC PRKCB MAPK3 SHC1 TNFRSF10B VAV3
hsa04152	AMPK signaling	-2.1	2	2.9	CREB1 PFKFB2 PIK3CA PPP2CA PPP2CB PPP2R2A PPP2R5C PRKAA1 PRKAG1 R

	pathway				
hsa00240	Pyrimidine metabolism	-2	2.5	2.9	AB2A RHEB MAP3K7 PPP2R3B PRKAG2 PPP2R3C
hsa05211	Renal cell carcinoma	-2	2.3	2.9	DCK DCTD DUT NME1 NT5E RRM1 UMPS RRM2B CMPK1
hsa00900	Terpenoid backbone biosynthesis	-2	3.7	3.2	CDC42 ETS1 FH KRAS NRAS PIK3CA MAPK3 RAP1B ELOC CUL2
hsa00220	Arginine biosynthesis	-2	3.7	3.2	FNTA HMGCS1 IDI1 ICMT NUS1
hsa04210	Apoptosis	-2	1.9	2.7	ACY1 GLS GLUD1 GOT1 GOT2
hsa05110	Vibrio cholerae infection	-1.9	2.6	2.9	BIRC2 FAS CASP3 CASP7 CTSC CTSL EIF2S1 IL3 KRAS NRAS PIK3CA MAPK3 T
hsa00270	Cysteine and methionine metabolism	-1.9	2.6	2.9	UBA1A TNFRSF10B EIF2AK3 CYCS
hsa04110	Cell cycle	-1.9	1.9	2.7	PRKACB SLC12A2 ATP6V1G1 KDELR2 SEC61G ATP6V1D ATP6V1E2 ATP6V1C2
hsa03430	Mismatch repair	-1.9	3.5	3.1	AHCY GOT1 GOT2 LDHA MDH1 MAT2B PSAT1 ENOPH1
hsa03060	Protein export	-1.9	3.5	3.1	PRKACB SLC12A2 ATP6V1G1 KDELR2 SEC61G ATP6V1D ATP6V1E2 ATP6V1C2
hsa04146	Peroxisome	-1.9	2.2	2.7	ORCA ANAPC13 ANAPC4
hsa04350	TGF-beta signaling pathway	-1.9	2.1	2.6	MSH6 PCNA RFCl RPA1 RPA2
hsa01240	Biosynthesis of cofactors	-1.8	1.8	2.7	SRP14 SEC62 SEC11A SEC61G SPCS1
hsa01524	Platinum drug resistance	-1.8	2.2	2.7	ABCD2 DAO PEX1 PEX6 SOD1 GNPAT PEX3 AGPS ECI2 FAR1 NUDT19
hsa04921	Oxytocin signaling pathway	-1.8	1.8	2.5	ACVR1 BMP8B E2F4 ID2 SMAD5 PPP2CA PPP2CB MAPK3 ROCK1 SKP1 TFDP1 HJV
hsa04022	cGMP-PKG signaling pathway	-1.8	1.7	2.5	ALPL COX15 CPOX DLD GCH1 NME1 SHMT1 UGP2 UMPS UROD KMO MTHFD2
hsa04010	MAPK signaling	-1.8	1.5	2.4	MAT2B PSAT1 CMPK1 PDXP COQ5
					BIRC2 FAS CASP3 ABCC2 MSH6 MDM2 PIK3CA MAPK3 TOP2B CYCS
					CALM2 CAMK2D GNAI3 KRAS PPP1R12A NRAS PPP1CB PPP1CC PPP3CB PPP3C
					C PRKAA1 PRKACB PRKAG1 PRKCB MAPK3 ROCK1 PRKAG2
					SLC25A5 ATP1B3 ATP2B1 CALM2 CREB1 ATF2 GNAI3 PPP1R12A PPP1CB PPP1CC PPP3CB PPP3C
					MAPK3 ROCK1 VDAC1 VDAC2 VDAC3 GNA13
					FAS CASP3 CDC42 MAP3K8 ATF2 DUSP5 EFNA1 HSPA8 IGF2 KRAS STMN1 MA

	pathway				
hsa05214	Glioma	-1.8	2.1	2.6	P3K4 NF1 NRAS PPM1A PPM1B PPP3CB PPP3CC PRKACB PRKCB MAPK3 RAP1B RASA2 MAP3K7 MAP3K2 RRAS2 MRAS IRAK4
hsa04066	HIF-1 signaling pathway	-1.8	1.9	2.5	CALM2 CAMK2D KRAS MDM2 NRAS PIK3CA PRKCB MAPK3 RB1 SHC1 CAMK2D EIF4E IFNGR1 LDHA PGK1 PGK2 PIK3CA PRKCB MAPK3 RPS6 ELOC CUL2 HKDC1
hsa05164	Influenza A	-1.7	1.7	2.4	SLC25A5 FAS CASP3 EIF2S1 IFNA13 IFNGR1 OAS2 PIK3CA PRKCB MAPK3 VDA C1 TNFRSF10B TBK1 IRAK4 CYCS TMPRSS4 RSAD2 BCL2L2-PABPN1
hsa00563	Glycosylphosphatidylinositol (GPI)-anchor biosynthesis	-1.7	3.1	2.8	PIGB PIGK PIGO PIGY PIGS
hsa04611	Platelet activation	-1.7	1.8	2.4	BTK FYN GNAI3 PPP1R12A PIK3CA PPP1CB PPP1CC PRKACB MAPK3 RAP1B RACK1 MYL12A GNA13 MYL12B
hsa04916	Melanogenesis	-1.6	1.9	2.4	CALM2 CAMK2D CREB1 CTNNB1 GNAI3 KRAS NRAS POMC PRKACB PRKCB MAPK3 WNT3A
hsa05135	Yersinia infection	-1.6	1.8	2.3	ARF6 RHOG CD8A CDC42 ITGA4 PIK3CA PKN2 MAPK3 ROCK1 MAP3K7 ARPC2 VAV3 TBK1 IRAK4 ELMO2
hsa04670	Leukocyte transendothelial migration	-1.6	1.8	2.3	RHOH CDC42 CTNNB1 GNAI3 ITGA4 ITK PIK3CA PRKCB RAP1B ROCK1 VAV3 MYL12A MYL12B
hsa00620	Pyruvate metabolism	-1.6	2.4	2.5	ADH5 DLAT DLD FH GLO1 LDHA MDH1
hsa03420	Nucleotide excision repair	-1.6	2.4	2.5	CCNH GTF2H3 PCNA RFCl RPA1 RPA2 CUL4B
hsa04960	Aldosterone-regulated sodium reabsorption	-1.6	2.6	2.5	ATP1B3 KRAS NR3C2 PIK3CA PRKCB MAPK3
hsa01250	Biosynthesis of nucleotide sugars	-1.6	2.6	2.5	UAP1 UGP2 FPGT NANS UXS1 HKDC1

hsa04510	Focal adhesion	-1.6	1.6	2.2	BIRC2 CDC42 CTNNB1 FYN ITGA1 ITGA4 ITGAV PPP1R12A PIK3CA PPP1CB PP1CC PRKCB MAPK3 RAP1B ROCK1 SHC1 ZYX VAV3 MYL12A MYL12B
hsa01210	2-Oxocarboxylic acid metabolism	-1.6	3.4	2.7	ACY1 GOT1 GOT2 IDH3A
hsa04064	NF-kappa B signaling pathway	-1.6	1.9	2.3	BIRC2 BTK CSNK2A2 PRKCB PRKCQ MAP3K7 UBE2I BCL10 MALT1 LY96 IRAK4 TIRAP
hsa00280	Valine, leucine and isoleucine degradation	-1.6	2.4	2.4	ACADM DLD HADH HMGCS1 HIBADH ACAD8 MCCC1
hsa04370	VEGF signaling pathway	-1.5	2.2	2.3	CDC42 KRAS NRAS PIK3CA PPP3CB PPP3CC PRKCB MAPK3 ACADM ADH5 FAS CASP3 CTNNB1 IFNA13 PRKAA1 PRKAG1 TRA2B MAP3K7 LY96 TBK1 IRAK4 PRKAG2 TIRAP
hsa04936	Alcoholic liver disease	-1.5	1.7	2.2	LY96 TBK1 IRAK4 PRKAG2 TIRAP
hsa04520	Adherens junction	-1.5	2	2.3	ACP1 CDC42 CSNK2A2 CTNNB1 FYN MAPK3 PTPN1 MAP3K7 FARP2
hsa04730	Long-term depression	-1.5	2.1	2.3	GNAI3 KRAS NRAS PPP2CA PPP2CB PRKCB MAPK3 GNA13 CTNNB1 IGF2 KRAS NFE2L2 NRAS PIK3CA PRKCB MAPK3 RB1 SHC1 SMARCE1 AXIN2 FRAT2 PBRM1 PHF10 WNT3A CSNK1A1L
hsa05225	Hepatocellular carcinoma	-1.5	1.6	2.1	ABL1 CALM2 CAMK2D CDC42 KRAS NRAS PIK3CA MAPK3 RAP1B SHC1 RIPK2 PRDM4 IRAK4
hsa04722	Neurotrophin signaling pathway	-1.5	1.8	2.1	CDC42 DLG1 NEDD4 PCNA PPP2CA PPP2CB PPP2R2A PRKAA1 PRKACB PRKA G1 ROCK1 TUBA1A ARPC2 MYL12A PRKAG2 RAB8B MYL12B
hsa04530	Tight junction	-1.5	1.6	2.1	ABL1 CAMK2D CDC42 EFNA1 FYN GNAI3 KRAS NCK1 NRAS PIK3CA PPP3CB PPP3CC MAPK3 ROCK1 RYK SEMA3B MYL12A MYL12B
hsa04360	Axon guidance	-1.5	1.6	2.1	AP1G1 AGA ASA1 CTSC CLN5 CTSL FUCA2 M6PR SLC11A2 CD164 LAPTM4A GGA3 DMXL2 SLC17A5
hsa04142	Lysosome	-1.5	1.7	2.1	BIRC2 BMP8B CTNNB1 DLG1 ID2 PPP1CB PPP1CC PPP2CA PPP2CB PPP2R2A TP53BP2 AXIN2 LIMD1 YWHAQ FBXW11 WNT3A
hsa04390	Hippo signaling pathway	-1.5	1.6	2.1	DAO DLD PSPH SHMT1 PSAT1 SRR
hsa00260	Glycine, serine and threonine metabolism	-1.5	2.4	2.3	

	threonine metabolism				
hsa04931	Insulin resistance	-1.4	1.8	2.1	CREB1 PIK3CA PPP1CB PPP1CC PRKAA1 PRKAG1 PRKCB PRKCQ PTPN1 OGT TBC1D4 PRKAG2
hsa04115	p53 signaling pathway	-1.4	2	2.2	FAS CASP3 MDM2 PPM1D TNFRSF10B SESN1 RRM2B CYCS ZMAT3
hsa04012	ErbB signaling pathway	-1.4	1.9	2.1	ABL1 ABL2 CAMK2D KRAS NCK1 NRAS PIK3CA PRKCB MAPK3 SHC1
hsa05210	Colorectal cancer	-1.4	1.9	2.1	CASP3 CTNNB1 MSH6 KRAS NRAS PIK3CA MAPK3 RALA AXIN2 CYCS
hsa04925	Aldosterone synthesis and secretion	-1.4	1.8	2.1	ATF1 ATP1B3 ATP2B1 CALM2 CAMK2D CREB1 ATF2 POMC PRKACB PRKCB PRKD3
hsa04723	Retrograde endocannabinoid signaling	-1.4	1.6	2	GABRG2 GNAI3 NDUFA4 NDUFA9 NDUFAB1 NDUFB2 NDUFB4 NDUFB5 NDUFS3 NDUFS4 PRKACB PRKCB MAPK3 GNB5 SLC17A8
hsa04668	TNF signaling pathway	-1.3	1.7	2	BIRC2 FAS CASP3 CASP7 MAP3K8 CREB1 ATF2 IL15 PIK3CA MAPK3 MAP3K7 BAG4
hsa05220	Chronic myeloid leukemia	-1.3	1.9	2	ABL1 HDAC2 KRAS MDM2 NRAS PIK3CA MAPK3 RB1 SHC1
hsa04215	Apoptosis - multiple species	-1.3	2.5	2.2	BIRC2 CASP3 CASP7 BECN1 CYCS
hsa04915	Estrogen signaling pathway	-1.3	1.6	1.9	CALM2 CREB1 ATF2 GNAI3 HSPA8 HSP90AA1 KRAS NRAS PIK3CA POMC PRKACB MAPK3 SHC1 KRT23

Table S13: Pathway enrichment results for purple or dark tongue at Day 30

GO	Description	LogP	Enrichment	Z-score	genes
hsa05168	Herpes simplex virus 1 infection	-7.2	2.4	6.2	BAK1 C5 CASP8 EIF2S1 IKBKB IRAK1 NFKB1 OAS2 OAS3 POU2F1 PPP1CB PPP1CC PTPN11 RNASEL SRSF2 SRSF6 STAT2 TAP1 ZNF8 ZNF33A ZNF43 ZNF84 ZNF90 ZNF136 ZNF175 TNFRSF14 ZNF443 ZNF544 IRAK4 ZNF571 TLR9 ZNF83 MA

					VS ZNF471 ZNF529 ZNF250 ZNF426 ZNF557 ZNF333 ZNF587 ZNF764 ZNF641 ZN F621 ZNF699 ZNF605
hsa00514	Other types of O-glycan biosynthesis	-6	6.2	7.1	GALNT3 B4GALT1 MFNG OGT PLOD3 C1GALT1C1 GALNT7 GALNT11 COLGA LT1 B3GLCT GXYLT1 SLC25A6 BAK1 CASP8 EIF2S1 IKBKB KPNA1 CIITA NFKB1 OAS2 OAS3 RNASE
hsa05164	Influenza A	-5.7	3.2	5.8	L STAT2 XPO1 TRIM25 PABPN1 TNFRSF10B TNFRSF10A CALCOCO2 KPNA6 IR AK4 MAVS ARF6 BAK1 CASP8 DYNC1LI2 FLNB IKBKB IRAK1 KPNA1 M6PR NFKB1 PFN2
hsa05132	Salmonella infection	-5	2.7	5.2	PIK3C2A TNFRSF10B TNFRSF10A WASL TLR6 RIPK3 ARHGEF26 VPS41 FHOD1 IRAK4 TLR9 VPS33A AHNAK2 TAB3
hsa03015	mRNA surveillance pathway	-4.1	3.5	5	PNN PPP1CB PPP1CC PPP2CB PPP2R2A PPP2R5B PABPN1 ACIN1 PCF11 PPP2R2 D PAPOLG UPF3A PABPC1L
hsa04371	Apelin signaling pathway	-4.1	3	4.8	ACTA2 APLNR MEF2A BORCS8-MEF2B MEF2D MYL4 PDE3B SLC8A2 HDAC4 MRAS PIK3R4 PRKAG2 GNG13 GNB4 CALML4 MEF2B
hsa05162	Measles	-3.6	2.9	4.3	BAK1 CASP8 CDK2 EIF2S1 FCGR2B IKBKB IRAK1 JAK3 NFKB1 OAS2 OAS3 ST AT2 IRAK4 TLR9 MAVS
hsa04722	Neurotrophin signaling pathway	-3.2	2.9	4.1	ABL1 CAMK2A CRK GAB1 IKBKB IRAK1 MAP3K3 NFKB1 PDPK1 PTPN11 SH2B 3 IRAK4 CALML4
hsa04621	NOD-like receptor signaling pathway	-3.2	2.4	3.9	CASP8 IKBKB NFKB1 OAS2 OAS3 RNASEL STAT2 VDAC2 RIPK3 CARD8 IRAK4 ERBIN DHX33 MAVS MCU GBP4 TAB3
hsa05160	Hepatitis C	-3	2.5	3.8	BAK1 CASP8 CDK2 EIF2S1 IKBKB NFKB1 OAS2 OAS3 PPP2CB PPP2R2A RNASE L STAT2 CLDN16 PPP2R2D MAVS
hsa05169	Epstein-Barr virus infection	-2.7	2.2	3.5	BAK1 BTK CASP8 CDK2 HDAC2 IKBKB IRAK1 JAK3 NFKB1 OAS2 OAS3 PSMD 12 STAT2 TAP1 IRAK4 MAVS SAP30L
hsa00310	Lysine degradation	-2.6	3.4	3.7	ACAT2 EZH1 NSD2 BBOX1 PLOD3 KMT5B SMYD2 COLGALT1
hsa04014	Ras signaling pathway	-2.5	2.1	3.2	ABL1 ARF6 EFNA1 FLT4 GAB1 IKBKB NFKB1 PTPN11 STK4 BRAP RASA4 PAK

hsa05130	Pathogenic Escherichia coli infection	-2.5	2.1	3.2	4 MRAS GNG13 PAK6 GNB4 PLA2G12B CALML4 ABL1 ARF6 BAK1 CASP8 IKBKB IRAK1 NCK1 NFKB1 PTPN11 TNFRSF10B TNF RSF10A WASL CLDN16 IRAK4 WIPF2 TAB3
hsa00260	Glycine, serine and threonine metabolism	-2.4	4	3.7	ALAS2 SARDH PGAM2 SHMT1 AOC3 GCAT
hsa00511	Other glycan degradation	-2.4	5.9	4.1	FUCA2 MAN2C1 MAN2B1 GBA2
hsa04142	Lysosome	-2.4	2.4	3.2	AP1G1 CTSE CTSO FUCA2 GALC M6PR MAN2B1 NAGLU LAPTM4A ATP6V0A2 LAMP3 NAGPA
hsa04261	Adrenergic signaling in cardiomyocytes	-2.3	2.3	3.1	CAMK2A CREB1 MYH7 MYL4 PPP1CB PPP1CC PPP2CB PPP2R2A PPP2R5B SLC 8A2 TNNI3 PPP2R2D CALML4
hsa05142	Chagas disease	-2.3	2.6	3.2	CASP8 IKBKB IRAK1 NFKB1 PPP2CB PPP2R2A TLR6 IRAK4 TLR9 PPP2R2D
hsa04910	Insulin signaling pathway	-2.2	2.3	3.1	CRK IKBKB PDE3B PDPK1 PHKA2 PPP1CB PPP1CC PYGB MKNK1 PRKAG2 PPP 1R3F CALML4
hsa04330	Notch signaling pathway	-2.2	3.1	3.3	CTBP1 HDAC2 MFNG TLE1 DTX4 DLL1 DTX3L
hsa05170	Human immunodeficiency virus 1 infection	-2.2	2	2.9	AP1G1 BAK1 CASP8 CRK IKBKB IRAK1 NFKB1 TAP1 TNFRSF1B CUL4B PAK4 I RAK4 GNG13 PAK6 GNB4 CALML4
hsa04140	Autophagy - animal	-2.1	2.3	3	EIF2S1 PDPK1 PPP2CB ULK1 SNAP29 MRAS WIPI2 PIK3R4 SUPT20H RRAGC M TMR14 ATG4C
hsa04130	SNARE interactions in vesicular transport	-2.1	4	3.4	STX16 SNAP23 SNAP29 GOSR1 STX6
hsa03013	Nucleocytoplasmic transport	-2.1	2.4	3	KPNA1 PNN XPO1 NUP155 ACIN1 KPNA6 NUP62 SENP2 XPO4 UPF3A
hsa04931	Insulin resistance	-2.1	2.4	3	CREB1 IKBKB NFKB1 PDPK1 PPP1CB PPP1CC PTPN11 PYGB OGT PRKAG2
hsa05220	Chronic myeloid leukemia	-2.1	2.8	3.1	ABL1 BAK1 CRK CTBP1 HDAC2 IKBKB NFKB1 PTPN11
hsa05017	Spinocerebellar ataxia	-2.1	2.2	2.9	SLC25A6 ATXN3 PSMD12 ATXN2 VDAC2 ULK1 NOP56 AFG3L2 WIPI2 PIK3R4 T WNK MCU

hsa00600	Sphingolipid metabolism	-2	3.2	3.1	GALC SGPL1 SPTLC1 GBA2 CERK ACER2
hsa03040	Spliceosome	-2	2.2	2.8	SRSF2 SRSF6 DHX16 PRPF4 DHX38 DDX46 SMNDC1 ACIN1 PRPF40B RBMX PRPF40A RBM25
hsa00512	Mucin type O-glycan biosynthesis	-2	3.7	3.2	GALNT3 GCNT1 C1GALT1C1 GALNT7 GALNT11
hsa04022	cGMP-PKG signaling pathway	-2	2.1	2.7	SLC25A6 CREB1 MEF2A BORCS8-MEF2B MEF2D MYH7 PDE3B PPP1CB PPP1CC SLC8A2 VDAC2 CALML4 MEF2B CAMK2A CREB1 PPP1CB PPP1CC PPP2CB PPP2R2A PPP2R5B GNG13 PPP2R2D GNB4 CALML4
hsa04728	Dopaminergic synapse	-1.9	2.2	2.7	CAMK2A CREB1 PPP1CB PPP1CC PPP2CB PPP2R2A PPP2R5B GNG13 PPP2R2D GNB4 CALML4
hsa01200	Carbon metabolism	-1.9	2.3	2.8	ACAT2 DLAT GOT2 PGAM2 PGK2 PRPS2 SHMT1 H6PD ADPGK PGP
hsa04210	Apoptosis	-1.8	2.1	2.6	BAK1 CASP2 CASP8 CTSO EIF2S1 IKBKB NFKB1 PDPK1 TNFRSF10B TNFRSF10A HTRA2
hsa04071	Sphingolipid signaling pathway	-1.8	2.2	2.6	NFKB1 PDPK1 PPP2CB PPP2R2A PPP2R5B NSMAF SGPL1 SPTLC1 PPP2R2D ACER2
hsa04152	AMPK signaling pathway	-1.8	2.2	2.6	CREB1 EEF2 PDPK1 PPP2CB PPP2R2A PPP2R5B ULK1 EEF2K PRKAG2 PPP2R2D
hsa04144	Endocytosis	-1.8	1.8	2.5	GRK2 AP2A1 ARF6 CAV1 EPS15 SNX1 WASL RNF41 STAM2 ARFGEF1 SPART RUFY2 RAB22A SNX6 RBSN WIPF2 WASHC2C
hsa04622	RIG-I-like receptor signaling pathway	-1.8	2.6	2.7	CASP8 IKBKB NFKB1 TRIM25 RNF125 MAVS DHX58
hsa04620	Toll-like receptor signaling pathway	-1.8	2.3	2.6	CASP8 IKBKB IRAK1 IRF5 NFKB1 TLR6 IRAK4 TLR9 TOLLIP
hsa05152	Tuberculosis	-1.7	1.9	2.4	CAMK2A CASP8 CREB1 FCGR2B IRAK1 CIITA NFKB1 NFYB TLR6 ATP6V0A2 IRAK4 TLR9 CALML4
hsa04922	Glucagon signaling pathway	-1.7	2.2	2.5	CAMK2A CREB1 PDE3B PGAM2 PHKA2 PYGB PRKAG2 CALML4 LDHAL6A
hsa00562	Inositol phosphate	-1.7	2.5	2.6	PIK3C2A PI4KB MTMR6 MINPP1 CDIPT MTMR14 ITPKC

	metabolism				
hsa05410	Hypertrophic cardiomyopathy	-1.7	2.4	2.5	ITGA4 ITGAV ITGB6 MYBPC3 MYH7 SLC8A2 TNNI3 PRKAG2
hsa04360	Axon guidance	-1.7	1.9	2.4	ABL1 CAMK2A EFNA1 NCK1 PDPK1 PTPN11 SLIT3 TRPC1 SEMA3B PAK4 SEM A4D SRGAP2 PAK6
hsa03022	Basal transcription factors	-1.6	2.9	2.6	ERCC3 GTF2E1 MNAT1 TAF1 TAF5
hsa04114	Oocyte meiosis	-1.6	2	2.3	CAMK2A CDK2 PPP1CB PPP1CC PPP2CB PPP2R5B SMC1A CDC16 ANAPC7 CALML4
hsa05165	Human papillomavirus infection	-1.6	1.6	2.2	ATP6V1A BAK1 CASP8 CDK2 CREB1 FZD2 HDAC2 IKBKB ITGA4 ITGAV ITGB6 LLGL1 MFNG NFKB1 PPP2CB PPP2R2A PPP2R5B STAT2 ATP6V0A2 PPP2R2D
hsa04141	Protein processing in endoplasmic reticulum	-1.5	1.9	2.2	BAK1 EIF2S1 ATXN3 SSR3 MBTPS1 MAN1B1 SEC61A1 NPLOC4 NGLY1 ERO1B EDEM3 UBXN2A
hsa04070	Phosphatidylinositol signaling system	-1.5	2.2	2.3	PIK3C2A PI4KB DGKD MTMR6 CDIPT MTMR14 ITPKC CALML4
hsa04136	Autophagy - other	-1.5	3.3	2.6	PPP2CB WIPI2 PIK3R4 ATG4C
hsa03420	Nucleotide excision repair	-1.5	2.8	2.5	ERCC3 MNAT1 RFC2 XPC CUL4B
hsa00280	Valine, leucine and isoleucine degradation	-1.5	2.8	2.4	ACAT2 HADHB HMGCS1 MCCC1 MCCC2
hsa05417	Lipid and atherosclerosis	-1.5	1.7	2.1	CAMK2A CASP8 EIF2S1 IKBKB IRAK1 NFKB1 PDPK1 POU2F1 VAV2 TNFRSF10B TNFRSF10A TLR6 IRAK4 CALML4
hsa04218	Cellular senescence	-1.5	1.9	2.1	SLC25A6 CDK2 FOXM1 NFKB1 PPP1CB PPP1CC RBBP4 VDAC2 MRAS MCU CALML4
hsa05135	Yersinia infection	-1.5	1.9	2.2	ARF6 CRK IKBKB IRAK1 ITGA4 NFKB1 VAV2 WASL IRAK4 WIPF2
hsa04217	Necroptosis	-1.4	1.8	2.1	SLC25A6 CAMK2A CASP8 JAK3 PYGB STAT2 VDAC2 TNFRSF10B TNFRSF10A RIPK3 H2AW
hsa05144	Malaria	-1.4	2.6	2.3	GYPC HBA1 HBA2 HBB TLR9

hsa00010	Glycolysis / Gluconeogenesis	-1.4	2.4	2.2	DLAT PGAM2 PGK2 MINPP1 ADPGK LDHAL6A
hsa04012	ErbB signaling pathway	-1.4	2.2	2.2	ABL1 CAMK2A CRK GAB1 NCK1 PAK4 PAK6
hsa04510	Focal adhesion	-1.4	1.7	2	CAV1 CRK FLNB FLT4 ITGA4 ITGAV ITGB6 PDPK1 PPP1CB PPP1CC VAV2 PAK4 PAK6
				2	TRIM37 UBB UBA7 CUL4B CDC16 HERC4 FZR1 ANAPC7 UBA6 UBE2Z
hsa04120	Ubiquitin mediated proteolysis	-1.4	1.9	2	IKBKB NCK1 NFKB1 PDCD1 PDPK1 VAV2 PAK4 PAK6
hsa04660	T cell receptor signaling pathway	-1.4	2	2.1	BTK IKBKB IRAK1 NFKB1 TRIM25 CARD10 IRAK4 TAB3
hsa04064	NF-kappa B signaling pathway	-1.4	2	2.1	CAMK2A CREB1 HDAC2 PPP1CB PPP1CC CALML4
hsa05031	Amphetamine addiction	-1.3	2.3	2.1	CRK GAB1 PTPN11 PAK4 PAK6 FLCN
hsa05211	Renal cell carcinoma	-1.3	2.3	2.1	ANK3 CAMK2A CAV1 FLNB FZD2 GAB1 ITGAV PDPK1 PPP1CB PPP1CC PTPN11 VAV2 MRAS
hsa05205	Proteoglycans in cancer	-1.3	1.7	1.9	
hsa00900	Terpenoid backbone biosynthesis	-1.3	3.6	2.4	ACAT2 FNTA HMGCS1
hsa00230	Purine metabolism	-1.3	1.9	2	ADSS2 AMPD3 GUK1 HPRT1 PDE3B PDE7A PPAT PRPS2 PAICS
hsa04110	Cell cycle	-1.3	1.9	2	ABL1 CDK2 HDAC2 ZBTB17 SMC1A CDC16 DBF4 FZR1 ANAPC7
hsa05143	African trypanosomiasis	-1.3	2.9	2.2	HBA1 HBA2 HBB TLR9
hsa00250	Alanine, aspartate and glutamate metabolism	-1.3	2.9	2.2	ADSS2 GOT2 PPAT DDO

Table S14: Pathway enrichment results for XYZ phenotype at Day 30

GO	Description	LogP	Enrichment	Z-score	genes
hsa04110	Cell cycle	-3.6	5.6	5.2	ABL1 CCND3 CCNH CDK2 E2F3 TFDP1 YWHAQ

hsa04725	Cholinergic synapse	-3	5.4	4.6	ADCY6 CAMK2B CHRNA3 GNG3 KCNJ12 GNG12
hsa05203	Viral carcinogenesis	-3	4	4.2	CASP8 CCND3 CDK2 HPN H4C5 KAT2B YWHAQ H2BC18
hsa04114	Oocyte meiosis	-2.7	4.6	4.2	ADCY6 CAMK2B CDK2 AURKA YWHAQ CPEB3
hsa04014	Ras signaling pathway	-2.6	3.5	3.8	ABL1 ETS2 FGF2 GNG3 GRIN1 KSR1 ANGPT4 GNG12
hsa05200	Pathways in cancer	-2.6	2.5	3.4	ABL1 ADCY6 CAMK2B CASP8 CCND3 CDK2 E2F3 FGF2 GNG3 IFNGR1 PTGER2 LEF1 GNG12
hsa05034	Alcoholism	-2.6	3.8	3.8	ADORA2A GNG3 GRIN1 SLC18A2 H4C5 GNG12 H2BC18
hsa04713	Circadian entrainment	-2.5	5.2	4.2	ADCY6 CAMK2B GNG3 GRIN1 GNG12
hsa05167	Kaposi sarcoma-associated herpesvirus infection	-2.5	3.6	3.7	CASP8 E2F3 FGF2 GNG3 IFNGR1 LEF1 GNG12
hsa04390	Hippo signaling pathway	-2.3	3.9	3.6	CCND3 SMAD7 SERPINE1 GDF5 YWHAQ LEF1
hsa04015	Rap1 signaling pathway	-2.3	3.4	3.4	ADCY6 ADORA2A FGF2 GRIN1 DOCK4 VAV3 ANGPT4
hsa04217	Necroptosis	-2.3	3.8	3.6	CAMK2B CASP8 GLUD1 IFNGR1 CHMP2B CHMP5
hsa04115	p53 signaling pathway	-2.2	5.5	3.9	CASP8 CCND3 CDK2 SERPINE1
hsa04727	GABAergic synapse	-1.9	4.5	3.3	ADCY6 GNG3 GABARAPL1 GNG12
hsa05162	Measles	-1.9	3.6	3.1	CASP8 CCND3 CDK2 CSNK2A2 CD46
hsa04371	Apelin signaling pathway	-1.9	3.6	3.1	ACTA2 ADCY6 GNG3 SERPINE1 GNG12
hsa03040	Spliceosome	-1.8	3.4	3	HNRNPC MAGOH PRPF4 DDX42 RNU4-2
hsa04914	Progesterone-mediated oocyte maturation	-1.7	4	3	ADCY6 CDK2 AURKA CPEB3
hsa04934	Cushing syndrome	-1.7	3.3	2.8	ADCY6 CAMK2B CDK2 E2F3 LEF1
hsa04218	Cellular senescence	-1.7	3.2	2.8	CCND3 CDK2 E2F3 SERPINE1 MCU
hsa04024	cAMP signaling pathway	-1.6	2.7	2.6	ADCY6 ADORA2A CAMK2B GRIN1 PTGER2 VAV3
hsa04066	HIF-1 signaling pathway	-1.6	3.7	2.8	CAMK2B IFNGR1 SERPINE1 ANGPT4
hsa05166	Human T-cell leukemia virus 1 infection	-1.6	2.7	2.6	ADCY6 CCND3 CDK2 E2F3 ETS2 KAT2B

hsa05163	Human cytomegalovirus infection	-1.6	2.7	2.6	ADCY6 CASP8 E2F3 GNG3 PTGER2 GNG12
hsa04151	PI3K-Akt signaling pathway	-1.6	2.3	2.4	CCND3 CDK2 FGF2 GNG3 ITGA8 YWHAQ ANGPT4 GNG12
hsa04724	Glutamatergic synapse	-1.6	3.5	2.7	ADCY6 GNG3 GRIN1 GNG12
hsa05031	Amphetamine addiction	-1.5	4.4	2.8	CAMK2B GRIN1 SLC18A2
hsa04520	Adherens junction	-1.5	4.3	2.8	CSNK2A2 PTPN1 LEF1
hsa05152	Tuberculosis	-1.5	2.8	2.4	CAMK2B CASP8 IFNGR1 KSR1 CLEC7A
hsa04926	Relaxin signaling pathway	-1.4	3.1	2.4	ACTA2 ADCY6 GNG3 GNG12
hsa04971	Gastric acid secretion	-1.4	4	2.6	ADCY6 CAMK2B SLC26A7
hsa04144	Endocytosis	-1.4	2.4	2.2	CHMP2B SNX12 CHMP5 SMAP1 PIP5KL1 SNX32
hsa04728	Dopaminergic synapse	-1.4	3.1	2.4	CAMK2B GNG3 SLC18A2 GNG12
hsa05022	Pathways of neurodegeneration - multiple diseases	-1.3	1.9	2	CAMK2B CASP8 CSNK2A2 GRIN1 NDUFS1 ATXN2 SDHB CHMP2B M CU
hsa05012	Parkinson disease	-1.3	2.3	2.1	ADORA2A CAMK2B NDUFS1 SDHB SLC18A2 MCU

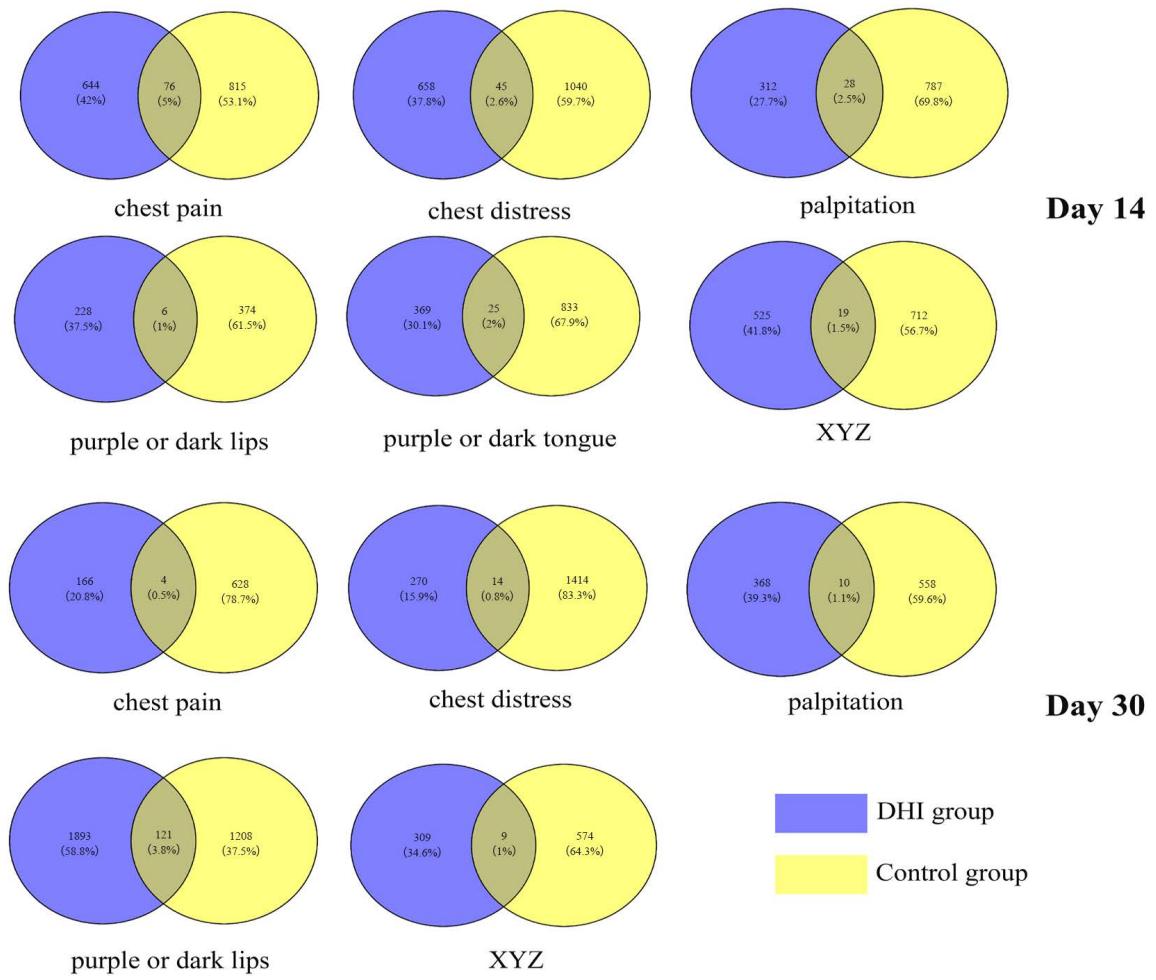


Figure S1: Comparison of the number of PARGSs between DHI and control group