$Table \ 1: The \ 500 \ most \ strongly \ \textit{COMT} - correlated \ genes \ (ranking \ on \ correlation \ p\text{-value}) \ in \ Prefrontal \ Cortex.$

1	RPL18P13	CAT	COMT	ACY1	CHCHD8	GATM	MY06	RPL18AP3	C1orf122	YBX1P2
2	CDC14B	EDNRB	PDLIM3	C6orf48	UBP1 ¹	PGCP	CADPS2	PPIB	ECH1	DTYMK
3	HSCB	TTLL4	PNKD	CSTB	MEGF10	H3F3C	POLR2F	HIP1R	RPS2P8	ZFAND3
4	CLU	SERF2	PTTG1IP	PKM2	RPL19	COPE	AASS	NKRF	RPS14	C12orf39
5	SPAG9	LIMS1	SNAP23	PABPC1	PPAP2B	ZMAT5	SCNM1	RPS5	RPL13A	BCAN
6	NKAIN4	UBE3A	CCDC28A	C3orf70	RPL12	RUSC1	ELP4	CAMSAP1L1	PDE6D	GSN
7	THSD1	MMP28	LRRC3B	ALDH2	IDH2	ADA	ABCA5	PGLS	IVD	QKI
8	GRSF1	HSD17B6	ELOVL4	PGM1	RAB11FIP5	APOE	PIR	RGC32	OLIG1	ACP6
9	CPT2	DFFA	RAB3GAP1	CHCHD5	GP\$N2	APCDD1	MLC1	SLC25A20	RPL35P5	SCAMP1
10	RPS3	ARL6IP6	ARHGAP24	HSD17B8	ZHX2	IQCK	LZTFL1	SH3BGRL2	PIK3C2A	WDR69
11	KPNA4	ACAA2	GABARAPL1	ABTB2	RARS	BGLAP	RAG1AP1	YIF1A	PRDX4	GPIAP1
12	PCCB	DENR	ALAD	FAM36A	RHBDD1	SDAD1	OPTN	PPP2CA	SCAMP2	STAT3
13	DNAJA2	S100A16	HIST1H2AC	TTYH1	PMF1	CIB1	DIRC2	WIPI1	MYCBP2	RBP1
14	SH3GL2	PJA2	ATP6V0E1	CLCN2	TPT1	NEFM	MAPRE1	GPR17	S100A13	H3F3AP
15	CBR1	WDFY2	ELOVL2	CYP2J2	MGC35154	CCNB1	GALNT10	SGPL1	GLUL	NME5
16	KIF21A	PLA2G5	SMARCA2	THRA	KRT10	SASH1	EIF3S4	$F\!AU$	CRB1	SPATA18
17	C6orf72	EEF1D	RELA	MON1B	MATR3	PSMC1	PHGDH	PTPRZ1	PREPL	GNG5
18	PFN1	NT5C	SEL1L	CHDH	TNFSF13	NSMCE1	PCDHGC3	VPS36	NME6	CTNNA1
19	MCF2	CXorf38	RPS17	SAE1	LRP4	CECR1	FER	TST	GNAI2	NUBP1
20	FTL	HIP1	HSF2	AGPAT3	ALDH1A1	ATPAF1	RAB9A	DYNLT1	AKR7A3	SNAP91
21	CLK3	LGALS3BP	C9orf23	SYT4	EIF4EBP3	RPS9	C14orf169	IGFBP7	CINP	CD302
22	ARHGEF3	PSME1	CADPS	C9orf140	KUA-UEV	DAG1	PRR23B	C1orf61	IFITM5	LDHD
23	CDC37L1	HSD17B10	DPY19L3	GRHPR	VPS72	ATG5	GABRG2	TMEM136	TOMM34	HLA-A
24	CST3	FAM83B	C5orf5	NDFIP1	GRN	GM2A	TFPT	VPS52	PHPT1	ACO2
25	TP53AP1	LOC401589	MTM1	RPL36	PARP4	LIPI	GSTM2	SYPL1	MRPL53	RPL8
26	RPS11	C11orf48	FAM102A	PXMP2	MGST1	CCPG1	KIAA1600	NEFH	SYNJ1	OACT2

Table 1: (Top 500 Positively Correlated Genes in PF Crtx Continued)

27	GNG12	PON2	C16orf14	PECR	TRIM37	LIX1L	YARS	LPHN2	SMAP1	SELENBP1
28	POPDC3	RFXANK	$\mathcal{N}\!P\!L$	TCEAL2	CASC1	CD83	BCKDHA	TSC22D4	APRIN	FAH
29	MKL2	SLC6A15	LIN28B	MEF2C	ATP6V1C1	MTCH1	ARL15	C2orf18	SLC25A1	CRYL1
30	IGSF1	SFRS15	PYCR2	PTPN4	C11orf59	ARSF	CLDN10	TGFB2	DOCK1	CDH12
31	ENO2	C15orf37	OXR1	PIP3-E	ANXA5	BCAT2	CACNA2D3	RSPO3	TMEM134	RPL30
32	PLCB1	DERA	RANBP3L	<i>NAPB</i>	ITGB5	C2orf34	PRKCSH	GPR37L1	MERTK	C2orf68
33	SLC25A12	ASRGL1	TOMM70A	PCDHGA12	THYN1	MTA3	C2orf28	IRF2BP2	GNA13	TOPBP1
34	MYH10	FADS1	EPN2	PITPNC1	PCMT1	CTSH	<i>ZBTB20</i>	C9orf46	PCTK2	ENPP5
35	CAPZA2	DNAJC5G	CCNF	ADHFE1	TMBIM4	MARCKSL1	C10orf26	SLC30A5	RPS13	AGXT2L1
36	PIP5KL1	ACOX2	CCDC63	OLIG2	ZNF775	RALYL	PNPO	AFF2	MSRB2	COL5A2
37	WASL	ZNF804B	ACSS1	SLC39A10	AP3M1	DCP2	HYI	VMA21	DBI	ZFHX4
38	EDG1	WDR16	ADORA2B	RPP40	CYP4X1	PPAP2A	PRICKLE1	FBXO8	C5orf4	HDGF
39	MDS032	RPLP0	S100B	EPHX2	FLJ37357	PPP2R5C	CRTAC1	NCKAP1	FAM26C	GNA12
40	ABHD3	TSPAN12	PXMP3	B3GNT1	OR5K3	TTC19	NDEL1	TP53BP2	FLJ23049	PANK2

Table 1: (Top 500 Positively Correlated Genes in PF Crtx Continued)

41	CUL3	FYN	AP3M2	UCRC	EPB41L3	USP54	ZNF533	RPS4X	B4GALNT2	RP1-32F7.2
42	FUCA2	RRNAD1	TAS2R8	KRT222P	TAX1BP3	NR2E1	AHI1	SLC39A12	MC3R	SS18
43	ADARB1	FGFR2	GPRASP2	MIF4GD	SLC7A6OS	LARGE	ACTL6A	ZYG11B	SLC9A9	GPR177
44	ACSBG1	RIT1	MLL2	ORC4L	PLAT	FLJ10986	GLS	HHLA3	PI4KII	DERL2
45	IL5RA	C6orf129	FGF1	PACS2	SORCS2	LGALS3	ALS2CR13	UBE2D2	NARS	ACTR6
46	PIPOX	PHLPP	KLK7	CD82	EPB41L5	LUZP4	ACOX1	GRM1	REEP5	ALX4
47	OPLAH	CYB5A	AFF3	ZNF490	C3orf63	CAMTA1	ACAA1	AK2	LRFN5	ZNF396
48	GJB6	KIAA1161	SLCO1C1	CDC14A	RPL11	LDB2	CDC2L1	PRC1	LGALS13	NDUFV2
49	RGS20	RIOK1	HSDL1	HLA-F	RPIB9	C5orf32	PELI2	MRPS36	KIAA0528	EDG7
50	MTMR4	C1orf198	VDAC3	DLL1	MAG1	TUFM	PCDHGA1	SEC13	PSAT1	IDH1
51	ZBTB24									

 $^{^{1}}$ The expression levels of genes in boldface are negatively correlated with COMT expression.

Table 2: The 500 most strongly COMT – correlated genes (ranking on correlation p-value) in Cerebellum.

1	PGCP	CAT	PCCB	FUCA2	TMEM170	RAB9A	OACT2	ARHGEF12	COMT	CYB5D2
2	TMEM106C	ACTG1	SEPP1	KUA-UEV	CBR1	VPS41	PIGT	GRHPR	PFN1	MARCKSL1
3	PYCR2	FNTA	C1orf71	MYO6	PDE6D	C3orf70	DIRC2	GRM3	PDE4B	B3GAT1
4	QKI	PPAP2B	PPIB	ABCA9	SNAP91	TYW1	CDS1	C3orf1	PTPRZ1	GPR177
5	SYPL1	GLE1L	MAP1B	C5orf4	RGS17	RPIB9	CELSR3	CPT2	FGFR2	PPAP2A
6	C8orf61	RUSC1	TTC19	HIP1	TP53AP1	ISOC1	LRRC8D	DAD1	SLC31A2	HIP1R
7	CD9	HHLA3	DFFA	ATP6V0E1	EDNRB	LYPLAL1	CAMSAP1L1	LYRM2	CRTAP	PXMP3
8	SPAG9	PPP2R5C	MOG	TMEM42	TMEM165	ZMYND17	SERPINI1	PRTFDC1	NCAM2	PRMT2

Table 2: (Top 500 Positively Correlated Genes in Crblm Continued)

9	RNF150	PRKCE	MAPRE1	ABCA8	FANCL	SYT4	SUZ12	OMG	CHCHD8	CDC14B
10	HSPA2	CLDN11	ARMC10	PRG-3	TNRC5	C10orf78	HNMT	SEC22C	CKS1B	GFOD1
11	CGI-38	$\mathcal{N}SDHL$	ALS2CR13	SUMF1	SPTLC2	FEZ1	PXK	GSN	FAM80B	COPE
12	PEPD	C12orf31	SERBP1	MYCBP2	IQCK	BRP44L	GALNT10	CPEB3	C10orf58	LAMP2
13	PRDX4	AASS	IGSF11	PIR	C14orf24	C20orf23	PAOX	NLGN3	ZYG11B	HSD17B6
14	PMF1	YBX1P2	ZNF398	TUBA1A	SCN2A	RNF13	ATG3	ZNF540	SLC44A1	KLHL4
15	ATP5S	GPR27	TMEM59	SH3BGR	RNH1	TMEM87A	BSN	C19orf42	SFT2D1	EDC4
16	LIN9	PHF16	RARS	SLC25A26	DTD1	C6orf48	ZMAT5	$G\!ATM$	ACO2	SLCO3A1
17	ABTB2	RHBDD1	SGPL1	ALG14	TTYH2	TYMS	GTDC1	NRIP3	SLC24A2	LGALS3BP
18	RAG1AP1	CREG1	SLC9A9	SPCS1	LEPREL2	MAPRE2	FLJ40142	STOM	MMS19L	C1orf122
19	GMPR2	GPR135	PTTG1IP	DSCAML1	TMBIM4	ACBD5	TTLL7	USP54	PSEN1	FBXO7
20	PSMB2	C10orf26	PLEKHA5	MLYCD	$\mathcal{N}\!C\!ST\!\mathcal{N}$	C6orf72	DHRS7	PDGFRA	UGT8	FA2H
21	SCRG1	SPIRE1	FLJ20054	CNDP1	DOCK10	APBB2	DYNC1I2	CTSL1	CXorf57	POLR2G
22	MRPL24	CNOT6	DHRS7B	D15WSU75E	RFFL	CHD9	WDFY1	MTM1	NRBP2	YIF1B
23	KNOP1	ACOT8	FBXL10	NSMCE1	ARL6IP6	POLD4	C10orf90	SNAPC4	STK36	PHKA2
24	EIF2AK1	DAZAP2	HSD17B12	GAPDH	GBA	TMED10	EIF3S2	FKSG30	LGMN	TMCO1
25	KIAA0196	CDC2L5	LIPA	ECH1	ARPP-21	IVD	FAM13A1	ASPA	DYNLT1	M6PRBP1
26	SETDB2	ERCC1	SH3GL2	PQLC3	ETFA	EDIL3	ZDHHC9	JTV1	C1orf57	DCTN6
27	COX6A1	SOX10	AK2	RYR2	RAB3GAP1	ANXA5	TCEA2	C11orf49	ENPP6	ELOVL4
28	CAPN13	RNF130	KIAA1853	C18orf10	CDKN1C	ZNF562	PFDN1	TM4SF11	NDN	CA14
29	C1orf19	TRIM45	SLC45A3	PIGK	PHYHIPL	CADM4	TNFRSF25	RIPK2	DNAPTP6	C20orf169-DBNDD2
30	SLC30A7	ENPP2	RPS6KA4	CD82	BTG3	ALAD	CYP27A1	ARMCX6	KIAA0256	ERBB3
31	FOXO3A	DERA	RPAIN	ZNF488	CDKAL1	CLCN3	KLK6	SEC11C	CHCHD5	MYO1D
32	RHBDL2	PHF11	ENOSF1	C9orf46	GREM1	TMEM98	AHCTF1	DIAPH1	LARS2	RPN1
33	USP47	ZNF174	C20orf116	EPC2	DHRS4	PCDHAC1	GALNT9	MYLK	TMEM63A	TAF5
34	NOLA2	PAQR4	FRMD6	SCCPDH	RAB11FIP5	RSAD1	GAL3ST1	PTPN13	ADC	KCNQ2

Table 2: (Top 500 Positively Correlated Genes in Crblm Continued)

35	DEFB103B	SRPX	GM2A	TMEM125	POLR2F	PSPH	BICD1	SLCO1A2	HSF2	SEMA4D
36	NDUFA11	CNTN2	MALT1	C11orf31	CHRNB1	MAP3K12	SHC4	PADI2	KCTD18	SLC25A20
37	FOXJ2	SUMF2	HRASLS3	RPS2P8	ARPP-19	DMRT2	GOLGA7	NCOA3	SNAP25	ADSSL1
38	GPR37	RPUSD4	HSD17B4	ACTL6A	AP3B1	C3orf63	ZIM2	PRRT2	CDC123	GLT25D2
39	GLTP	ZNF529	EVI2A	ARL5A	SALL1	EMG1	ARHGAP1	KIAA0672	S100B	TUFT1
40	CCNE2	DDX19A	GAB1	SNAP23	PIP3-E	CAP1	TST	ZHX1	NIN72	VPS52

Table 2: (Top 500 Positively Correlated Genes in Crblm Continued)

41	TSPAN9	TM9SF1	FOLH1	CCNB1	AP1GBP1	DTYMK	BTBD16	INTU	AFMID	LACTB2
42	TMEM136	MAG1	SERF2	NUDCD2	C10orf32	LRRC37B	CA2	MON1B	CCDC21	HOXD1
43	YIPF1	RPS6KB1	PLCL1	KIF13B	SETD6	HPN	CRTAC1	MAP3K5	LOC731950	EXOSC5
44	LITAF	COL16A1	$BV\!ES$	KIAA0240	MOBKL2B	MGC35308	$\mathcal{N}\!M\!B$	RAB33A	RASSF2	RRNAD1
45	COL4A3BP	GPAA1	PMP22	SLC5A11	GPX4	PLEKHA6	CLU	UGCG	STX2	UBE2G1
46	FSTL5	DSCR3	POLG	CXADR	ZNF521	RTKN	TAX1BP1	HLCS	HSD17B10	ZNF354A
47	ADARB1	KBTBD8	LTC4S	SLC35F1	TGFA	ZNF547	CCS	TRIM13	PHC2	CRADD
48	TALD01	TBCC	$D\!AK$	GABRB3	GJB1	BTF3	KCTD2	RIMS3	ZNF326	WDR42A
49	NUDT5	TSPAN3	AHCY	SCN3A	CAPZA2	DLL1	CD302	LZTS2	FAM55C	ELOVL6
50	CRYGS	ADIPOR1	CYB561D2	AGPAT3	PER2	ZMAT3	ABCF2	PPP2R5E	TMEM16D	HIAT1
51	THRAP2									

 $Table \ 3: \ The \ 500 \ most \ strongly \ \textit{COMT} - correlated \ genes \ (ranking \ on \ correlation \ p\text{-value}) \ in \ Temporal \ Cortex.$

1	GATM	CAT	MMP28	SNAP23	CDC14B	THSD1	PTTG1IP	DTYMK	PPAP2B	COMT
2	ALDH2	EDNRB	ECH1	ARL6IP6	SLC25A20	GLUL	MEGF10	IL17RB	GSTM2	HSCB
3	PIR	DPY19L3	PNPO	SLC39A12	HIST1H2AC	MTM1	LRP4	CSTB	CLDN10	RGC32
4	ACY1	ASRGL1	CRB1	PGM1	LRRC3B	ARHGAP24	CYP2J2	WDFY2	RAB9A	C2orf34
5	AASS	AQP4	PTPRZ1	MLC1	PON2	SASH1	ZFAND3	RIT1	PSMC1	MERTK
6	QKI	DOCK7	MYO6	RANBP3L	LIMS1	AGXT2L1	C3orf70	IQCK	HIP1	APCDD1
7	THBS4	HSD17B6	RGN	SPAG9	ITPR2	TNFSF13	SGPL1	ACAA2	NR2E1	TPD52L1
8	GPR177	EMX2	BBS2	ATP1B2	ACP6	GPR98	TTLL4	PXMP3	CBR1	MGST1
9	CHDH	RAB5C	ALDH6A1	C4 or f 19	RELA	PITPNC1	<i>TP53BP2</i>	C14orf159	PSAT1	PXMP2
10	LAMA1	LRIG1	ATP6V0E1	PPP2R5C	SLC1A2	LHX2	PCDHGC3	CHST7	COQ9	CADPS2

Table 3: (Top 500 Positively Correlated Genes in Temp Crtx Continued)

11	IDH1	CHCHD8	CA2	APOE	TSC22D4	AGPAT3	CHPT1	LGALS3	FYN	SLCO1C1
12	<i>TJP2</i>	CLU	VPS72	LARS2	POLR2F	CPT2	CCNB1	EDG1	DFFA	DDR1
13	MARCKSL1	CLCC1	ZFHX4	DBI	PDLIM3	SYPL1	IDH2	BCAN	SLC15A2	ZNF22
14	SCRG1	TMEM136	ACSBG1	PIPOX	ZHX2	FLJ32310	ACOX2	GRN	DVL3	ZMAT5
15	CYP4X1	ACTL6A	RAD1	CDKN2C	ATPAF1	PNKD	ELOVL2	SOX21	RFC2	ETFDH
16	GNG12	IL33	PRDX4	HIP1R	GPAM	FAM89A	HRSP12	RHBDD1	DYNLT1	AK3L2
17	EPB41L5	RPL18P13	CTSH	PPIB	COPE	PHLPP	MIF4GD	MAPRE1	SOX9	ETFA
18	C6orf72	KRT10	C12orf39	FBXO8	LINC00299	PPAPDC1B	ACAA1	FAM36A	PCCB	SLC9A9
19	GALNT10	APPL2	OLIG1	ABHD3	GNA12	RGS20	C4orf18	CCDC25	PCDHGA12	PIK3C2A
20	GPR37L1	PFN1	YBX1P2	PSD2	ACO2	TMEM70	LDHD	TNIK	TMEM123	NPL
21	TRAF5	NKX2-2	INPP5A	C1orf122	CST3	BCKDHA	CXorf38	HSPA2	EIF4EBP3	IRX5
22	GLI3	SLC30A9	IL10RB	SCAMP2	PDE6D	IRF2BP2	TST	SFT2D1	MSI2	ATP1A2
23	SLC7A10	IGFBP7	GNA13	KUA-UEV	SEC22C	PARP4	C10orf110	GNG5	RARS	NTRK2
24	AP3M1	HNRPU	GPSN2	TMEM38B	PDE1A	GSTK1	MPP3	SUPT16H	ZNF622	PRKD1
25	FGF2	TTYH1	TTC21B	NWD1	C9orf61	TM9SF1	$O\!AF$	BCAR3	CDKN3	GSN
26	PHKG1	CAMSAP1L1	ARL15	PHGDH	TNFAIP6	C5orf32	UBP1	ITGAV	HIBCH	GPT2
27	GNAI2	RAD9B	PGCP	ADORA2B	ABCA5	PPAP2A	C9orf123	PLA2G5	C5orf5	S100A13
28	GALNT4	SPARCL1	ADA	BBOX1	TSPAN12	RUSC1	SPON1	TNRC5	MSRB2	BEST4
29	PDGFRA	RBM4B	IGSF11	AKR7A3	PCDHB5	SERF2	LSAMP	CUL3	CSRP1	LMO7
30	LRCH1	PARP3	C5orf4	USP15	FZD8	FADS1	CRYL1	MTUS1	TRPS1	CASC1
31	DHRS4L2	TBC1D14	CHRNB1	PLA2R1	SH3GL2	SLC44A3	AMOT	C3orf63	S100A1	ADHFE1
32	SHH	KIAA1161	CLK3	ACSL6	TLK1	TCEAL4	LYN	RHOBTB3	ALDH7A1	C9orf46
33	ABTB2	SLC7A6OS	DNASE2	NUDT5	<i>ZBTB20</i>	HSF2	SORCS2	GPR17	EPHX1	NLGN3
34	RET	SFRS15	METT11D1	ACOX1	EIF2B3	POU3F2	ELP4	CD37	GRHPR	PYCR2
35	DDO	PYGM	H3F3AP4	DERA	OPHN1	PHF21B	CCDC121	UBE2D2	FNTA	NPAS3
36	DHX32	ITGB5	TUBB2B	APOC2	PARP9	CDC42EP4	APRIN	POLR3C	STOX1	LOC400455

Table 3: (Top 500 Positively Correlated Genes in Temp Crtx Continued)

37	CHSY-2	RFTN2	RNF182	WASF3	CALB1	EPHX2	KIAA0789	SH3BP4	GFM1	HSF1
38	FTL	TOMM70A	ERO1L	DOCK1	C22orf9	C20orf107	SH3BGRL3	DHRS4	LCN8	FAM102A
39	DAG1	ELOVL4	C16orf14	EFHD1	C1orf74	RAB7L1	CYFIP1	MON1B	PI4KII	CD82
40	SEL1L	DKFZP686D0972	FAH	PHF17	PER2	C19orf54	NME6	OLIG2	NEK10	SLC30A5

Table 3: (Top 500 Positively Correlated Genes in Temp Crtx Continued)

41	SLC35B2	GRSF1	FAM96A	TSEN34	C2orf18	BA16L21.2.1	CD302	ZNF385	C1orf198	ELOVL5
42	OMA1	CDC37L1	USP14	NUBP1	SLC7A11	ANKFYI	POLR3A	DHTKD1	SLC4A4	OTX1
43	SRI	ZWILCH	TH1L	KCTD18	KAL1	STAT3	PEPD	DLL1	TMEM127	CADM4
44	SGCE	APP	RERG	PCBD1	ARPC5L	TLCD1	BAAT	UBE3A	CNTFR	MRPL53
45	GPX7	SAMD5	PCMT1	TMEM80	RYR1	SH3BP5	TPP1	C20orf72	ALDH4A1	PHACTR4
46	POGZ	FLT3	CSRP2	DIRC2	F3	HHLA3	RFXANK	EPB41L3	MIXL1	HSD17B10
47	LMNB1	GAL3ST4	RHEBL1	CTNNA1	TTC19	MCF2	FAM118B	LMX1B	IVD	GBP7
48	TAF5	ZNF544	ZNF263	MRPL41	HIGD2A	GPR56	ALAD	LEPROT	TM4SF18	TP53AP1
49	STS	ASCL1	UBE2J1	HLA-A	KIAA0195	PPP1R3D	COL5A3	SELENBP1	HERPUD2	LZTFL1
50	ANKRD15	ASXL2	MCCC2	GNG7	NFE2L2	BZRAP1	NKAIN4	ZWINT	MAT1A	LRRC8A
51	AKR1C3									

 $Table \ 4: \ The \ 500 \ most \ strongly \ \textit{COMT}-correlated \ genes \ (ranking \ on \ correlation \ p-value) \ in \ Pons.$

1	KLK3	KUA-UEV	COMT	HIST1H2AC	SYPL1	FNTA	TMEM170	PRDX4	GATM	CAT
2	PSEN1	CLU	SLCO3A1	SPAG9	MEGF10	C20orf23	HSPA2	AGPS	QKI	B3GAT1
3	ADIPOR1	PIGM	PIGT	ARL6IP6	LRRC63	GSN	MOG	MTM1	SEC22C	LGALS3BP
4	THRAP6	TMBIM4	C7orf11	UGT8	PPIB	SPTLC2	POLR2F	PET112L	TP53AP1	CD9
5	PFN1	PCSK6	HIST1H4H	DTYMK	<i>TJP2</i>	SEPT10	SFT2D1	CBR1	C9orf77	BRP44L
6	RAB9A	CHCHD8	KLHL4	$BV\!ES$	STX2	ACTL6A	BTD	PGCP	FNTB	C6orf129
7	ATG3	SLC9A9	PIR	MARCKSL1	TLE4	C10orf78	SCHIP1	EDIL3	HIP1	C8orf61
8	SSR4	MYO1D	LRRC8D	FA2H	SNAP23	PRELID1	MOBKL2B	DYNC1I2	STXBP3	DYNLT1
9	AASS	SEPP1	PPP2R5C	DAD1	PTTG1IP	ABCA9	EEF2K	PXK	USH1C	SLC31A2
10	AFMID	C1orf122	KCTD3	TIMP2	ACTG1	CAMSAP1L1	STAMBP	HSCB	C2orf28	PRKCSH

Table 4: (Top 500 Positively Correlated Genes in Pons Continued)

11	CDKN1C	DAZAP2	DHRS4L2	KRT10	PAPSS1	HINT3	RHBDD1	PHF16	MYLK	ATP8A1
12	MAPRE2	ELOVL6	DIXDC1	GPR27	MYO6	CKS1B	TMEM87A	PIGG	TALDO1	TMEM116
13	ALDH3A2	DICER1	$MPD\mathcal{Z}$	PQLC3	IGSF11	NRBP2	SLC44A1	KLK6	AGPAT3	ANXA5
14	SCCPDH	ABCA8	EMILIN2	C20orf4	HRASLS3	RNH1	FBXO32	NFE2L3	BTG3	POLR2G
15	MED8	TIA1	CTNNA1	TSPAN8	DHRS4	FLJ11506	SALL1	ENPP6	KIAA1026	ENPP2
16	C9orf164	ARHGEF12	RKHD1	TMEM38B	CTTNBP2	WDFY2	FLOT2	ATP5S	C5orf4	ST18
17	SASH1	EXDL2	ATP6V0E1	DSCAML1	C22orf5	$\mathcal{N}\!L\mathcal{N}$	PDE4B	CRYL1	LASS2	MIF4GD
18	MYO1E	MLH1	PPAP2A	ELOVL1	SEC23B	EIF2B1	MTSS1	CDC14B	C10orf32	LITAF
19	CYP27A1	REEP3	GPC5	RARS	C3orf63	SGCE	FBXO7	C5orf37	BRD7	TFCP2
20	BRMS1	HMGCL	GLTP	PDE6D	WDR57	TMEM5	C3orf70	<i>PSPH</i>	MTUS1	PARP4
21	RAB11FIP5	NSMCE1	AK2	SCRG1	HIAT1	CA14	C20orf116	RFFL	APBB2	PADI2
22	PCCB	CRIPT	RRNAD1	CADM4	SERF2	ZMYM5	TMC7	CLDN11	YIF1A	COL16A1
23	ZFAND3	C1orf77	TMEM59	ERBB2IP	LIPA	CMTM5	KIAA0196	PRPF19	UBE2G1	HARS
24	EFS	HHLA3	OMA1	CYP2J2	KIAA0672	M6PRBP1	HNRPC	DNAJC5	LYRM2	STT3A
25	PGLS	CECR1	C6orf72	INTS4	BCAP31	SNX24	SIRT2	TRIP4	KIAA0892	SLC12A2
26	BBX	PLEKHG4	PMF1	PRPSAP2	CYB5A	TTLL4	SORT1	ALG13	SCNM1	H3F3AP4
27	ITGAV	TESK2	KIAA0776	FIS1	SS18	CPEB3	SRPRB	CDS1	RBM4B	ANGPTL2
28	UNC50	RP2	ZDHHC14	KLHDC1	RNF13	CANT1	GALC	SSFA2	INTU	SPP1
29	PTPRD	RNF130	BAT5	PHF11	LUZP1	SGK2	C10orf90	CNTNAP4	VAMP3	IMPDH2
30	HOXD1	NETO2	PIK3C2A	FBXO8	ARPC5L	FAM123B	ACY1	TOR1AIP1	UBP1	CYP20A1
31	GM2A	HIST2H2BE	TTLL11	PXMP3	EFCBP1	DKFZP434A0131	OACT2	C20orf39	ARMC10	MAP7
32	C11orf67	C1orf198	ANAPC5	ATG4C	ZCCHC11	B4GALNT1	DYRK4	ZMAT5	ELOVL4	LDLRAP1
33	YAF2	LAMP2	TMEM140	ELOVL2	CXADR	TTYH2	TTLL7	ANKFY1	ST3GAL3	RALBP1
34	PHF5A	C1orf135	$\mathcal{N}SMCE4A$	PRKCE	FAM13C1	CAPN3	DDX19A	SP2	NEK7	GEMIN8
35	KIAA0319	TBC1D2	ASCC1	INSM2	PDXK	RPIB9	DOCK10	RHBDL2	OPRL1	FUSIP1
36	GRM3	ADCY5	DVL3	FAM38B	POLR2I	SOX10	KIAA1217	GRHPR	SUPV3L1	FN3KRP

Table 4: (Top 500 Positively Correlated Genes in Pons Continued)

37	MTP18	C20orf46	PPIE	PPAP2B	FBXW7	HNRPK	MGC3731	PEMT	KIF13B	RTKN
38	SERPINI1	MGC3207	CAPZA2	ASPA	$\mathcal{N}VL$	DOLPP1	CRYAB	HPS5	AGPAT4	CLASP1
39	MAPRE1	ZNF277P	MAL	TMEM125	PSMC1	NUDT5	C18orf54	APOD	NKX2-2	CNOT4
40	TTC19	SFT2D3	TNRC5	EPB41L2	PIGO	HIP1R	TMEM165	GAL3ST1	KIAA0256	C6orf106

Table 4: (Top 500 Positively Correlated Genes in Pons Continued)

41	C3orf1	RHOG	KIAA0100	NAPE-PLD	APIP	FEZ1	HNRPDL	LIPT1	C14orf24	LOC732402
42	GPR37	POLR3A	POU3F2	TGFA	PLCB1	TTF2	RIPK2	C10orf39	PRRG1	TMEM42
43	RPLP0	DPP3	SGPL1	NPC1	STMN1	PAQR4	C14orf122	RPL18P13	FAM96A	NCAPD2
44	SLC35B2	LSM7	EFHD1	ACAA2	PPAP2C	CHMP2B	FBXO31	SAV1	SLC39A6	DPF2
45	STX16	EIF1	SYNJ1	FUNDC2	SPTLC1	SLCO1A2	SEC61B	ACBD5	ISOC1	DACH1
46	EI24	CCNB1	DDX28	PDE4A	SDAD1	AP3M1	C1orf19	ABTB2	MGC59937	HIATL2
47	DUSP3	KIF6	TMEM46	SLC13A3	COPE	GPR3	TMEM63A	C20orf149	PIGN	ARID1A
48	CKMT2	MGC16169	SYVN1	RELA	ATP5G2	ALS2CR2	FLJ14803	TMED4	EXOSC5	ADCY9
49	PJA2	TBC1D5	PIP5K2A	DDX5	ZFYVE26	C8orf72	SEMA4D	RIOK1	SNAP91	KRAS
50	RDX	SLC9A1	FKSG30	HPN	PHPT1	CUTC	ADC	PCM1	ZNF536	PEX26
51	CA2									