

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

	PGD	ADGRL2	CFH		MYH11
PTPRB	TALDO1	HECW2	SERPINF1		CNN1
ARHGAP29	BRI3	ARHGEF12	FBXO32		SYNPO2
MGAT4A	TKT	PCDHGC3	CLU		TPM2
ARHGAP31	ATP6V0D1	SPOCK2	MGP	EPAS1	DAB2
NOTCH1	FTH1	ADGRG6	ZHX3	SLC6A4	ITPR1
ZEB1	ATP6V0E1	ACADSB	COL16A1	SLCO2A1	SARAF
ANO6	SQOR	ZC3H7B	THY1	VIPR1	NIBAN1
ATP13A4	ALDOA	NCKAP1	ENAH	ACVRL1	SCARA5
EPB41L5	FTL	TNFAIP1	DPYSL3	SLC9A3R2	PCNX1
KAT6A	FCN3	VNN2	HMGCS1	TEK	FCN3
HMGCS1	HMGCS1	EFNB2	S100A4	GRK5	HMGCS1
EFNB2	SLC39A8	PRX	MSMO1	GIMAP8	MSMO1
FCN1	NPR3	SLC39A8	ANXA3	FOXF1	ANXA3
MGAM	NUPR1	NPR3	MGAM	EFNB2	INSIG1
VSIR	S100A4	ANXA3	FGR	PRX	MNDA
CALCR	MSMO1	MNDA	CALCR	SLC39A8	FCN1
PALM2AKAP2	ANXA3	FCN1	PALM2AKAP2	NUPR1	CALCR
PAPSS2	INSIG1	MGAM	RTKN2	S100A4	TNFSF10
CD36	MNDA	TNFSF10	SPOCK2	ESAM	CHI3L2