

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

	PGD	RGS5	ADGRL2	CFH	MYH11
PTPRB	TALDO1	ANTXR1	HECW2	SERPINF1	CNN1
ARHGAP29	BRI3	FILIP1L	ARHGEF12	FBXO32	SYNPO2
MGAT4A	TKT	IGFBP7	PCDHGC3	CLU	TPM2
ARHGAP31	ATP6V0D1	ROBO1	SPOCK2	MGP	EPAS1
NOTCH1	FTH1	ITGA9	ADGRG6	ZHX3	SLC6A4
ZEB1	ATP6V0E1	ENAH	ACADSB	COL16A1	SLCO2A1
ANO6	SQOR	SH3PXD2A	ZC3H7B	THY1	VIPR1
ATP13A4	ALDOA	NBL1	NCKAP1	ENAH	ACVRL1
EPB41L5	FTL	NRP2	TNFAIP1	DPYSL3	SLC9A3R2
KAT6A	FCN3	VNN2	VNN2	HMGCS1	TEK
HMGCS1	HMGCS1	NPR3	EFNB2	S100A4	GRK5
EFNB2	SLC39A8	MSMO1	PRX	MSMO1	GIMAP8
FCN1	NPR3	MNDA	SLC39A8	ANXA3	FOXF1
MGAM	NUPR1	FCN1	NPR3	MGAM	EFNB2
VSIR	S100A4	VSIR	ANXA3	FGR	PRX
CALCRL	MSMO1	CALCRL	MNDA	CALCRL	SLC39A8
PALM2AKAP2	ANXA3	TNFSF10	FCN1	PALM2AKAP2	NUPR1
PAPSS2	INSIG1	RTKN2	MGAM	RTKN2	S100A4
CD36	MNDA	EPB41L5	TNFSF10	SPOCK2	ESAM
					CHI3L2