

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

TNFAIP3		SELENOW		SCG2
PTGS2	X.2395	CRIP1		CHGA
CSF3	RNA5.8SP6	HSBP1	FCN1	SCN3A
IL6	VEGFA	FXYD3	MNDA	CENPF
LIF	LENG8	CAMK1D	MPEG1	RYR2
SLC7A11	SLC25A6	DYNLL1	PLXNC1	CPE
CXCL1	CEBPB	CHURC1	ARHGDIB	SEC11C
BIRC3	TNS3	PSD3	COTL1	COTL1
PPP1R15A	NTN4	CYB5A	GBP2	PEG10
BTG3	BRI3	EIF4A2	CYBB	ZNF292
NPR3	PATJ	BTNL9	IFITM1	SLC39A8
FADS1	FCN3	NUPR1	FYB1	NUPR1
TNFSF10	TBX3	MGAM	FCN3	S100A4
PRKCE	FADS1	CHI3L2	SLC39A8	ANXA3
MCEMP1	EVI2B	PTPRB	NPR3	EVI2B
PEBP4	ACSS2	CRTAC1	S100A4	BRI3
RALA	CLDN5	HHIP	ANXA3	MYL12A
SHE	RALA	CLDN5	MGAM	HOPX
PCDH17	MYL12A	SHE	FADS1	DBI
ANKRD1	PCDH17	CR1	TNFSF10	DENND3
43	44	50 Significant Latent Factor	51	61