

Genes Associated with Significant Latent Factors	SLIDE Marginal Variables							
	43		50		51		61	
	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		