

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

MT.CO1	KRT8		COL3A1
MT.ND4	KRT18		COL1A1
MT.CO3	COL8A1		COL1A2
MT.ND5	KRT7	IL1RL1	COL6A3
MT.CYB	ITGB6	LDLR	CCDC80
MT.ATP6	RCN1	TXNRD1	MYH11
MT.ND1	CALU	MSMO1	LRP1
MT.ND2	ITGAV	HMGCS1	HSPG2
MT.CO2	LMNA	CYP51A1	CALD1
MT.ATP8	CKAP4	INSIG1	C1S
FCN3	FCN3	HMGCR	FCN3
BTNL9	PRX	IDI1	BTNL9
SLC39A8	SLC39A8	FADS1	PRX
NUPR1	NUPR1	FCN3	SLC39A8
S100A4	S100A4	BTNL9	NUPR1
FCN1	FCN1	PRX	S100A4
VSIR	VSIR	SLC39A8	FCN1
GPX3	GPX3	NUPR1	VSIR
TNFSF10	TNFSF10	S100A4	GPX3
PALM2AKAP2	CHI3L2	STX11	TNFSF10