

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

	c.3.MT1X	c.9.S100A8	c.0.IFI27	c.2.HLADQA1	c.5.APOE
	c.4.MT1X	c.9.S100A9	c.1.IFI27	c.3.COMP	c.14.CD74
	c.5.MT1X	c.9.S100A12	c.11.IFI27	c.12.LGALS1	c.14.HLADRA
	c.6.MT1X	c.9.EREG	c.6.CD74	c.13.TINAGL1	c.14.HLADRB1
	c.8.MT1X	c.9.SERPINB2	c.14.MPZ	c.17.TOB1	c.14.HLADPA1
	c.8.CTGF	c.11.KRT2	c.11.KRT17	c.9.IL23A	c.14.IFITM2
c.3.IGFBP5	c.17.CCL4L2	c.0.KRT2	c.6.CTHRC1	c.13.NDUFA4L2	c.2.APOE
c.3.TPPP3	c.12.AREG	c.9.TNFAIP6	c.15.KRT5	c.12.DUSP14	c.12.CCL4L2
c.3.IGFBP6	c.17.GZMB	c.5.CRYAB	c.4.RGS5	c.9.C1QC	c.4.CCL2
c.3.CD55	c.7.CCL13	c.6.SFRP4	c.9.HLADQB2	c.0.KRT2	c.3.ADH1B
c.3.SLPI	c.3.STMN2	c.3.STMN2	c.3.STMN2	c.10.FOS	c.3.STMN2
c.3.TNXB	c.10.FOS	c.10.FOS	c.10.FOS	c.10.TXNIP	c.10.FOS
c.3.MFAP5	c.10.KLF6	c.10.TXNIP	c.10.TXNIP	c.10.KLF6	c.10.KLF6
c.10.TXNIP	c.3.SAA1	c.10.KLF6	c.10.KLF6	c.3.SAA1	c.13.CCDC80
c.3.SAA1	c.3.TNC	c.3.TNC	c.3.SAA1	c.3.TNC	c.6.CCL19
c.13.CRIP1	c.5.IGFBP5	c.5.IGFBP5	c.3.TNC	c.5.IGFBP5	c.3.STC2
c.3.STMN2	c.13.CCDC80	c.13.CCDC80	c.5.IGFBP5	c.13.CCDC80	c.3.IGFBP2
c.10.FOS	c.6.CCL19	c.6.CCL19	c.13.CCDC80	c.6.CCL19	c.3.GSN
c.10.KLF6	c.3.STC2	c.3.STC2	c.6.CCL19	c.3.STC2	c.3.WIF1
c.3.TNC	c.3.IGFBP2	c.3.IGFBP2	c.3.STC2	c.3.IGFBP2	c.1.APOE

6

26

34
Significant Latent Factor

45

54

62