

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

		c.3.MT1X	c.12.ZNF331	c.14.ISG15	c.16.FOS
c.0.KRT14		c.4.MT1X	c.12.NR4A2	c.14.IFI6	c.16.HES1
c.0.KRT5		c.5.MT1X	c.8.POSTN	c.14.IFI27	c.7.PTGDS
c.0.KRT17		c.6.MT1X	c.10.CCL5	c.14.LY6E	c.17.HSPB1
c.4.FGF7	c.3.SAA1	c.8.MT1X	c.10.CSF2	c.9.HSPH1	c.2.SELE
c.16.SAT1	c.3.STMN2	c.17.CCL4L2	c.12.AP3S1	c.13.SLMAP	c.1.FGFBP1
c.8.ASPN	c.5.IGFBP5	c.3.GPX3	c.8.THBS4	c.13.APOE	c.16.S100A2
c.5.CFH	c.3.TNC	c.17.TOB1	c.14.IRF1	c.3.POSTN	c.9.CXCL1
c.7.CCL2	c.16.KRT75	c.10.CCL5	c.12.BIRC3	c.5.CFD	c.6.SFRP4
c.8.IGFBP5	c.4.RGS5	c.8.THBS4	c.10.CCL4	c.7.HSPD1	c.12.LGALS1
c.13.SLMAP	c.9.CCR7	c.3.SLPI	c.3.SLPI	c.3.SLPI	c.3.SLPI
c.3.SLPI	c.13.TINAGL1	c.3.TPPP3	c.3.TPPP3	c.3.TPPP3	c.3.TPPP3
c.3.TPPP3	c.13.CLU	c.3.DIO2	c.3.DIO2	c.3.DIO2	c.3.DIO2
c.3.DIO2	c.9.AREG	c.10.KLF6	c.10.KLF6	c.10.KLF6	c.10.KLF6
c.10.KLF6	c.3.SLPI	c.3.TNC	c.3.TNC	c.3.TNC	c.3.TNC
c.3.TNC	c.3.TPPP3	c.4.CXCL3	c.4.CXCL3	c.4.CXCL3	c.4.CXCL3
c.4.CXCL3	c.3.DIO2	c.13.CCDC80	c.5.IGFBP5	c.5.IGFBP5	c.5.IGFBP5
c.5.IGFBP5	c.10.KLF6	c.6.CCL19	c.13.CCDC80	c.13.CCDC80	c.13.CCDC80
c.13.CCDC80	c.4.CXCL3	c.3.STC2	c.6.CCL19	c.6.CCL19	c.6.CCL19
c.6.CCL19	c.13.CCDC80	c.3.IGFBP2	c.3.STC2	c.3.STC2	c.3.STC2

1

12

31

90

100

124

Significant Latent Factor