

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

		c.14.ABCA8	c.14.ISG15
		c.14.GADD45B	c.14.IF16
	c.0.LGALS7	c.14.HES1	c.14.IF127
	c.1.LGALS7	c.14.CYR61	c.13.SLMAP
c.3.SAA1	c.11.LGALS7	c.3.TNC	c.14.LY6E
c.3.STMN2	c.16.LGALS7	c.4.SNCG	c.14.IFITM2
c.5.IGFBP5	c.11.KRT15	c.8.COMP	c.3.POSTN
c.13.TINAGL1	c.11.S100A6	c.7.RGS1	c.16.KRT6A
c.3.CCL2	c.9.AC020656.1	c.15.CD164	c.7.PLIN2
c.3.TNC	c.1.APOE	c.10.CCL20	c.4.FGF7
c.3.TPPP3	c.0.CALML5	c.3.TPPP3	c.3.SLPI
c.9.CCR7	c.2.RBP7	c.3.DIO2	c.3.TPPP3
c.9.AREG	c.11.IFITM3	c.10.KLF6	c.3.TNC
c.3.APOE	c.4.IFITM1	c.4.CXCL3	c.4.CXCL3
c.3.SLPI	c.1.KRT17	c.13.CCDC80	c.13.CCDC80
c.3.DIO2	c.10.KLF2	c.6.CCL19	c.3.STC2
c.10.KLF6	c.0.CST6	c.3.STC2	c.3.IGFBP2
c.4.CXCL3	c.1.S100A8	c.3.IGFBP2	c.1.APOE
c.6.CCL19	c.5.CCL19	c.3.WIF1	c.14.APOD