

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

				c.14.ABCA8
				c.14.GADD45B
c.0.KRT14			c.0.LGALS7	c.14.HES1
c.0.KRT5		c.7.APOE	c.1.LGALS7	c.14.CYR61
c.16.SAT1	c.3.SAA1	c.7.APOC1	c.11.LGALS7	c.3.TNC
c.14IFI27	c.3.STMN2	c.7.S100A9	c.16.LGALS7	c.4.SNCG
c.5.APOE	c.5.IGFBP5	c.17.HSPA6	c.11.KRT15	c.8.COMP
c.17.CXCR6	c.13.TINAGL1	c.7.SPP1	c.11.S100A6	c.7.RGS1
c.2CLU	c.3.CCL2	c.10.HSPA6	c.9.AC020656.1	c.15.CD164
c.4.FGF7	c.3.TNC	c.7.G0S2	c.1.APOE	c.10.CCL20
c.3.POSTN	c.3.TPPP3	c.6.CXCL9	c.0.CALML5	c.3.TPPP3
c.12.PPPP1R15A	c.9.CCR7	c.9.CCL20	c.2.RBP7	c.3.DIO2
c.3.TPPP3	c.9.AREG	c.8.HES1	c.11.IFITM3	c.10.KLF6
c.10.KLF6	c.3.APOE	c.4.CXCL3	c.4.IFITM1	c.4.CXCL3
c.4.CXCL3	c.3.SLPI	c.0.S100A2	c.1.KRT17	c.13.CCDC80
c.6.CCL19	c.3.DIO2	c.2.CCL14	c.10.KLF2	c.6.CCL19
c.3.STC2	c.10.KLF6	c.5.CRYAB	c.0.CST6	c.3.STC2
c.3.IGFBP2	c.4.CXCL3	c.7.CCL2	c.1.S100A8	c.3.IGFBP2
c.3.WIF1	c.6.CCL19	c.9.HSPB1	c.5.CCL19	c.3.WIF1