

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

c.0.KRT14			c.11.KRT17	c.14.ISG15
c.0.KRT5		c.3.IGFBP5	c.11.KRT16	c.14.IFI6
c.16.SAT1	c.3.SAA1	c.3.CD55	c.11.KRT6B	c.14.IFI27
c.14.IFI27	c.3.STMN2	c.3.TNXB	c.5.POSTN	c.14.LY6E
c.15.TSC22D1	c.5.IGFBP5	c.3.CCL2	c.15.CLDN10	c.4.FGF7
c.2.HBB	c.3.TNC	c.3.APOE	c.11.FABP5	c.13.EGR1
c.4.FGF7	c.3.CCL2	c.4.RGS5	c.6.CCL19	c.17.KLF2
c.3.TPPP3	c.3.APOE	c.12.AREG	c.3.IGFBP3	c.6.MYOC
c.14.HES1	c.13.TINAGL1	c.12.CCL4L2	c.6.APOD	c.7.PLIN2
c.9.CCR7	c.4.RGS5	c.3.IGFBP6	c.3.SLPI	c.3.SLPI
c.3.SLPI	c.9.AREG	c.5.CCL21	c.3.TPPP3	c.10.KLF6
c.3.DIO2	c.7.HBB	c.3.SLPI	c.10.KLF6	c.4.CXCL3
c.10.KLF6	c.3.SLPI	c.3.TPPP3	c.3.TNC	c.13.CCDC80
c.3.TNC	c.3.TPPP3	c.3.DIO2	c.4.CXCL3	c.3.STC2
c.4.CXCL3	c.10.KLF6	c.10.KLF6	c.13.CCDC80	c.3.IGFBP2
c.13.CCDC80	c.4.CXCL3	c.4.CXCL3	c.3.STC2	c.2.CLU
c.6.CCL19	c.13.CCDC80	c.5.IGFBP5	c.3.IGFBP2	c.3.GSN
c.3.STC2	c.6.CCL19	c.13.CCDC80	c.2.CLU	c.3.WIF1