

Significant Latent Factors – Marginals (bold/italic) and Interactions

Genes Associated with Significant Latent Factors

c.2.IL6		c.0.IFI27	c.13.DUSP1	c.15.SAA1	c.16.FOS
c.2.CSF3	c.5.COL1A1	c.1.IFI27	c.13.CYR61	c.15.SAA2	c.16.HES1
c.13.SCGB2A2	c.5.COL3A1	c.11.IFI27	c.17.GZMB	c.15.LTF	c.2.RBP7
c.9.EREG	c.4.CCL2	c.6.CD74	c.13.EGR1	c.9.AREG	c.6.SFRP4
c.13.DCD	c.9.C1QC	c.14.MPZ	c.13.APOE	c.15.SOD2	c.4.IFIMT1
c.6.SFRP4	c.11.KRT15	c.11.KRT17	c.13.NDUFA4L2	c.9.EREG	c.16.SPRR1B
c.15.TSC22D1	c.6.COMP	c.6.CTHRC1	c.13.SNCG	c.15.MALAT1	c.16.FGFBP1
c.12.DUSP14	c.17.GZMB	c.15.KRT5	c.13.ATF3	c.15.CD59	c.9.CCR7
c.8.COMP	c.15.CD59	c.4.RGS5	c.13.ACTC1	c.17.XCL1	c.7.TNF
c.17.CXCR6	c.15.S100A2	c.9.HLADQB2	c.13.SOCS3	c.15.TSC22D1	c.14.EGR1
c.3.STMN2	c.10.FOS	c.3.STMN2	c.10.FOS	c.3.STMN2	c.3.STMN2
c.10.FOS	c.3.STMN2	c.10.FOS	c.10.TXNIP	c.10.FOS	c.10.FOS
c.10.TXNIP	c.10.TXNIP	c.10.TXNIP	c.10.KLF6	c.10.KLF6	c.10.TXNIP
c.10.KLF6	c.10.KLF6	c.10.KLF6	c.5.IGFBP5	c.3.SAA1	c.10.KLF6
c.3.SAA1	c.3.SAA1	c.3.SAA1	c.13.CCDC80	c.5.IGFBP5	c.3.TNC
c.5.IGFBP5	c.3.TNC	c.3.TNC	c.6.CCL19	c.13.CCDC80	c.5.IGFBP5
c.13.CCDC80	c.5.IGFBP5	c.5.IGFBP5	c.3.STC2	c.6.CCL19	c.13.CCDC80
c.6.CCL19	c.13.CCDC80	c.13.CCDC80	c.2CLU	c.3.STC2	c.6.CCL19
c.3.STC2	c.6.CCL19	c.6.CCL19	c.3.GSN	c.2CLU	c.3.STC2
c.3.IGFBP2	c.3.STC2	c.3.STC2	c.3.WIF1	c.3.GSN	c.3.IGFBP2