

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

Genes Associated with Significant Latent Factors

c.6.PLA2G2A			c.13.DES				c.16.LGALS1	
c.6.CLU			c.13.ACTG2		c.15.AQP5		c.16.S100A2	
c.6.PTGDS	c.3.MT1X	c.0.IFI27	c.13.PCP4		c.15.KRT7		c.16.KRT17	
c.3.GPX3	c.4.MT1X	c.1.IFI27	c.13.RERGL	c.13.DUSP1	c.15.PPP1R1B		c.16.CALML3	c.16.FOS
c.9.HSPA1B	c.5.MT1X	c.11.IFI27	c.13.CNN1	c.13.CYR61	c.5.POSTN		c.16.APOE	c.16.HES1
c.9.DNAJB1	c.6.MT1X	c.16.IFI27	c.13.BCAM	c.17.GZMB	c.5.ACTA2	c.1.CXCL14	c.16.S100A6	c.2.RBP7
c.6.GPX3	c.8.MT1X	c.10.FOSB	c.13.TPM1	c.13.APOE	c.15.CD164	c.16.CXCL14	c.16.IGFBP7	c.8.IGFBP5
c.7.PLTP	c.8.CTGF	c.5.TM4SF1	c.13.SMTN	c.13.ACTC1	c.14.CFD	c.16.KRT15	c.16.KRT75	c.2.SELE
c.10.XCL2	c.3.GPX3	c.3.WIF1	c.13.MYLK	c.13.EGR1	c.15.CD59	c.4.RGS5	c.16.KRT6B	c.4.IFITM1
c.3.PTN	c.17.CCL4L2	c.7.IFI27	c.13.CSRP1	c.4.DCN	c.5.FABP4	c.3.SFRP4	c.16.MYC	c.16.SERPINB2
c.3.GSN	c.7.IFI27	c.15.DCD	c.13.CCDC80	c.14.PRX	c.11.KRT15	c.13.CRIP1	c.10.FOS	c.16.FGFBP1
c.1.APOE	c.12.AREG	c.6.CXCL9	c.3.STC2	c.13.ACTA1	c.10.FOS	c.7.CCL8	c.10.TXNIP	c.5.CFH
c.2.CCL14	c.3.STMN2	c.3.STMN2	c.1.APOE	c.8.TAGLN	c.3.TNC	c.16.KRT6B	c.10.KLF6	c.11.AREG
c.11.KRT17	c.3.SAA1	c.10.FOS	c.10.FOSB	c.3.STC2	c.5.IGFBP5	c.6.SFRP4	c.6.CCL19	c.3.GSN
c.3.IGFBP3	c.3.TNC	c.3.TNC	c.3.POSTN	c.1.APOE	c.13.CCDC80	c.14.CFD	c.3.STC2	c.1.APOE
c.1.POSTN	c.3.STC2	c.13.CCDC80	c.13.RGS5	c.13.TIMP3	c.6.CCL19	c.1.APOE	c.2.CLU	c.3.WISP2
c.3.ADH1B	c.3.WIF1	c.3.STC2	c.4.TCIM	c.13.HES1	c.3.STC2	c.3.WISP2	c.1.APOE	c.3.POSTN
c.15.UQCRH	c.3.POSTN	c.3.IGFBP2	c.7.IFI27	c.17.MTCO1	c.2.CLU	c.7.LYZ	c.3.WISP2	c.7.LYZ
c.5.CCL21	c.10.FOSB	c.3.GSN	c.13.MT1X	c.13.FLNA	c.1.APOE	c.2.SELE	c.16.SAT1	c.17.MTCO1
c.15.S100A1	c.4.TCIM	c.1.APOE	c.13.TIMP3	c.13.CCL2	c.3.WISP2	c.14.PRX	c.3.POSTN	c.3.ADH1B