

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

Genes Associated with Significant Latent Factors

			c.13.DES					
			c.13.ACTG2		c.5.APOE	c.15.AQP5	c.16.S100A8	
	c.3.MT1X	c.0.IFI27	c.13.PCP4		c.14.CD74	c.15.KRT7	c.16.S100A9	
	c.4.MT1X	c.1.IFI27	c.13.RERGL	c.13.DUSP1	c.14.HLADRA	c.15.PPP1R1B	c.9.CCL17	
	c.5.MT1X	c.11.IFI27	c.13.CNN1	c.13.CYR61	c.14.HLADRB1	c.5.POSTN	c.7.CCL18	c.16.FOS
	c.6.MT1X	c.16.IFI27	c.13.BCAM	c.17.GZMB	c.14.HLADPA1	c.5.ACTA2	c.16.S100A7	c.16.HES1
	c.8.MT1X	c.10.FOSB	c.13.TPM1	c.13.APOE	c.14.IFITM2	c.15.CD164	c.5.CCL26	c.2.RBP7
c.7.CCL3L1	c.8.CTGF	c.5.TM4SF1	c.13.SMTN	c.13.ACTC1	c.13.MGP	c.14.CFD	c.11.FABP5	c.8.IGFBP5
c.7.CCL4L2	c.3.GPX3	c.3.WIF1	c.13.MYLK	c.13.EGR1	c.4.IFITM1	c.15.CD59	c.2.FABP5	c.2.SELE
c.7.SOD2	c.17.CCL4L2	c.7.IFI27	c.13.CSRP1	c.4.DCN	c.14.IER3	c.15.FABP4	c.17.JUN	c.4.IFITM1
c.14.CCL2	c.7.IFI27	c.15.DCD	c.13.CCDC80	c.14.PRX	c.4.PHLDA1	c.11.KRT15	c.3.APOD	c.16.SERPINB2
c.14.IFITM2	c.12.AREG	c.6.CXCL9	c.3.STC2	c.13.ACTA1	c.5.IGFBP5	c.10.FOS	c.3.SAA1	c.16.FGFBP1
c.7.RGS1	c.3.STMN2	c.3.STMN2	c.1.APOE	c.8.TAGLN	c.13.CCDC80	c.3.TNC	c.5.CFH	
c.2.APOE	c.3.SAA1	c.10.FOS	c.10.FOSB	c.3.STC2	c.6.CCL19	c.3.TNC	c.6.CCL19	
c.17.TXNIP	c.3.TNC	c.3.TNC	c.3.POSTN	c.1.APOE	c.6.CCL19	c.5.IGFBP5	c.3.GSN	
c.9.TNFAIP6	c.3.STC2	c.13.CCDC80	c.13.RGS5	c.13.TIMP3	c.2.CLU	c.13.CCDC80	c.6.CCL19	
c.13.HES1	c.3.WIF1	c.3.STC2	c.4.TCIM	c.13.HES1	c.2.CLU	c.13.CCDC80	c.2.CLU	
c.9.HSPB1	c.3.POSTN	c.3.IGFBP2	c.4.TCIM	c.13.WIF1	c.3.GSN	c.6.CCL19	c.3.WISP2	
c.7.BIRC3	c.10.FOSB	c.3.GSN	c.4.TCIM	c.13.HES1	c.3.TCIM	c.6.CCL19	c.3.WISP2	
c.1.HSPA6	c.4.TCIM	c.1.APOE	c.4.TCIM	c.13.WIF1	c.3.TCIM	c.13.CCDC80	c.1.APOE	
							c.7.LYZ	
							c.3.POSTN	
							c.1.APOE	
							c.3.WISP2	
							c.17.MTCO1	
							c.3.ADHB1	