

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

		c.15.AQP5	
c.3.MT1X	c.0.IFI27	c.15.KRT7	
c.4.MT1X	c.1.IFI27	c.15.PPP1R1B	
c.5.MT1X	c.11.IFI27	c.5.POSTN	
c.6.MT1X	c.16.IFI27	c.5.ACTA2	c.1.CXCL14
c.8.MT1X	c.10.FOSB	c.15.CD164	c.16.CXCL14
c.8.CTGF	c.5.TM4SF1	c.14.CFD	c.16.KRT15
c.3.GPX3	c.3.WIF1	c.15.CD59	c.4.RGS5
c.17.CCL4L2	c.7.IFI27	c.5.FABP4	c.3.SFRP4
c.7.IFI27	c.15.DCD	c.11.KRT15	c.13.CRIP1
c.12.AREG	c.6.CXCL9	c.10.FOS	c.7.CCL8
c.3.STMN2	c.3.STMN2	c.3.TNC	c.16.KRT6B
c.3.SAA1	c.10.FOS	c.5.IGFBP5	c.6.SFRP4
c.3.TNC	c.3.TNC	c.13.CCDC80	c.14.CFD
c.3.STC2	c.13.CCDC80	c.6.CCL19	c.1.APOE
c.3.WIF1	c.3.STC2	c.3.STC2	c.3.WISP2
c.3.POSTN	c.3.IGFBP2	c.2.CLU	c.7.LYZ
c.10.FOSB	c.3.GSN	c.1.APOE	c.2.SELE
c.4.TCIM	c.1.APOE	c.3.WISP2	c.14.PRX