

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.8.MT1X	c.10.FOSB	c.15.CD164
c.8.CTGF	c.5.TM4SF1	c.14.CFD
c.3.GPX3	c.3.WIF1	c.15.CD59
c.17.CCL4L2	c.7.IFI27	c.5.FABP4
c.7.IFI27	c.15.DCD	c.11.KRT15
c.12.AREG	c.6.CXCL9	c.10.FOS
c.3.STMN2	c.3.STMN2	c.3.TNC
c.3.SAA1	c.10.FOS	c.5.IGFBP5
c.3.TNC	c.3.TNC	c.13.CCDC80
c.3.STC2	c.13.CCDC80	c.6.CCL19
c.3.WIF1	c.3.STC2	c.3.STC2
c.3.POSTN	c.3.IGFBP2	c.2.CLU
c.10.FOSB	c.3.GSN	c.1.APOE
c.4.TCIM	c.1.APOE	c.3.WISP2