

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
	26	45	70
	Significant Latent Factor		