

SLIDE Marginal Variables					
Genes Associated with Significant Latent Factors					
			c.11.KRT17		
	c.0.KRT14		c.11.KRT16	c.14.ABCA8	c.14.ISG15
	c.0.KRT5		c.11.KRT6B	c.14.GADD45B	c.14.IF16
	c.16.SAT1	c.3.SAA1	c.10.CCL20	c.14.HES1	c.13.SLMAP
	c.14.IF127	c.3.STMN2	c.5.POSTN	c.14.CYR61	c.14.LY6E
	c.15.TSC22D1	c.5.IGFBP5	c.15.CLDN10	c.4.SNCG	c.4.FGF7
	c.2.HBB	c.3.TNC	c.11.FABP5	c.3.TNC	c.13.EGR1
	c.4.FGF7	c.3.CCL2	c.6.CCL19	c.7.RGS1	c.17.KLF2
	c.3.TPPP3	c.3.APOE	c.3.IGFBP3	c.4.RGS16	c.6.MYOC
	c.14.HES1	c.13.TINAGL1	c.6.APOD	c.3.TPPP3	c.7.PLIN2
	c.9.CCR7	c.4.RGS5	c.3.SLPI	c.15.NEAT1	c.3.SLPI
	c.3.SLPI	c.9.AREG	c.3.TPPP3	c.3.SLPI	c.10.KLF6
	c.3.DIO2	c.7.HBB	c.10.KLF6	c.3.DIO2	c.4.CXCL3
	c.10.KLF6	c.3.SLPI	c.3.TNC	c.4.CXCL3	c.13.CCDC80
	c.3.TNC	c.3.TPPP3	c.4.CXCL3	c.5.IGFBP5	c.3.STC2
	c.4.CXCL3	c.10.KLF6	c.13.CCDC80	c.13.CCDC80	c.3.IGFBP2
	c.13.CCDC80	c.4.CXCL3	c.3.STC2	c.6.CCL19	c.2.CLU
	c.6.CCL19	c.13.CCDC80	c.3.IGFBP2	c.3.STC2	c.3.GSN
	c.3.STC2	c.6.CCL19	c.2.CLU	c.3.IGFBP2	c.3.WIF1
Significant Latent Factor					
1127699100					