

SLIDE Marginal Variables					
Genes Associated with Significant Latent Factors	c.3.MT1X	c.12.ZNF331	c.14.ISG15	c.16.FOS	
	c.0.KRT14	c.4.MT1X	c.12.NR4A2	c.14.IFI6	c.16.HES1
	c.0.KRT5	c.5.MT1X	c.8.POSTN	c.14.IFI27	c.7.PTGDS
	c.0.KRT17	c.6.MT1X	c.10.CCL5	c.14.LY6E	c.17.HSPB1
	c.4.FGF7	c.3.SAA1	c.8.MT1X	c.10.CSF2	c.9.HSPH1
	c.16.SAT1	c.3.STMN2	c.17.CCL4L2	c.12.AP3S1	c.1.FGFBP1
	c.8.ASPN	c.5.IGFBP5	c.3.GPX3	c.8.THBS4	c.13.APOE
	c.5.CFH	c.3.TNC	c.17.TOB1	c.14.IRF1	c.3.POSTN
	c.7.CCL2	c.16.KRT75	c.10.CCL5	c.12.BIRC3	c.5.CFD
	c.8.IGFBP5	c.4.RGS5	c.8.THBS4	c.10.CCL4	c.7.HSPD1
	c.13.SLMAP	c.9.CCR7	c.3.SLPI	c.3.SLPI	c.3.SLPI
	c.3.SLPI	c.13.TINAGL1	c.3.TPPP3	c.3.TPPP3	c.3.TPPP3
	c.3.TPPP3	c.13CLU	c.3.DIO2	c.3.DIO2	c.3.DIO2
	c.3.DIO2	c.9.AREG	c.10.KLF6	c.10.KLF6	c.10.KLF6
	c.10.KLF6	c.3.SLPI	c.3.TNC	c.3.TNC	c.3.TNC
c.3.TNC	c.3.TPPP3	c.4.CXCL3	c.4.CXCL3	c.4.CXCL3	c.4.CXCL3
c.4.CXCL3	c.3.DIO2	c.13.CCDC80	c.5.IGFBP5	c.5.IGFBP5	c.5.IGFBP5
c.5.IGFBP5	c.10.KLF6	c.6.CCL19	c.13.CCDC80	c.13.CCDC80	c.13.CCDC80
c.13.CCDC80	c.4.CXCL3	c.3.STC2	c.6.CCL19	c.6.CCL19	c.6.CCL19
c.6.CCL19	c.13.CCDC80	c.3.IGFBP2	c.3.STC2	c.3.STC2	c.3.STC2
1	12	31	90	100	124
		Significant Latent Factor			