

SLIDE Marginal Variables			
Genes Associated with Significant Latent Factors		c.15.AQP5	
		c.15.KRT7	
	c.9.S100A8	c.15.PPP1R1B	
	c.9.S100A9	c.15.S100A1	c.1.CXCL14
	c.9.S100A12	c.15.CD164	c.16.CXCL14
	c.9.SERPINB2	c.15.CD59	c.16.KRT15
	c.9.EREG	c.14.CFD	c.16.KRT6B
	c.0.KRT2	c.0.S100A2	c.4.RGS5
	c.9.TNFAIP6	c.5.FABP4	c.14.CFD
	c.11.KRT2	c.16.MYC	c.3.SFRP4
	c.5.CRYAB	c.13.CCDC80	c.13.CRIP1
	c.9.CCL20	c.6.CCL19	c.7.CCL8
	c.1.APOE	c.2.CLU	c.5.SOD2
	c.3.WISP2	c.3.MMP11	c.8.CXCL2
	c.7.LYZ	c.3.MFAP5	c.13.SYNPO2
	c.4.CCL19	c.14.ABCA8	c.6.SFRP4
	34	70	77
	Significant Latent Factor		