c.13.DUSP1

c.13.CYR61

c.13.APOE

c.13.ACTC1

c.17.GZMB

c.14.PRX

c.4.DCN

c.13.EGR1

c.13.ACTA1

c.13.FLNA

c.2.FN1

Significant Latent Factors – Margin		
Genes Associated with Significant Latent Factors		c.13.JUNB
	c.9.S100A8	c.13.ZFP36
	c.9.S100A9	c.13.JUN
	c.9.S100A12	c.13.FOS
	c.9.SERPINB2	c.13.HES1
	c.9.EREG	c.13.EGR1
	c.0.KRT2	c.13.SOCS3
	c.9.TNFAIP6	c.13.ATF3
	c.11.KRT2	c.13.RGS16
	c.5.CRYAB	c.13.CEBPD
	c.9.CCL20	c.10.TXNIP
	c.1.APOE	c.10.KLF2
	c.3.WISP2	c.1.POSTN
	c.7.LYZ	c.1.SYT8
	c.4.CCL19	c.13.CCL2
	34	58

 c.15.S100A1
 c.1.0

 c.15.CD164
 c.16.

 c.15.CD59
 c.16.

 c.14.CFD
 c.16.

 c.0.S100A2
 c.4

 c.5.FABP4
 c.1

 c.16.MYC
 c.3.

c.15.AQP5

c.15.KRT7

c.15.PPP1R1B

c.13.CCDC80

c.6.CCL19

c.2.CLU

c.3.MMP11

c.3.MFAP5

c.14.ABCA8

Significant Latent Factor

c.1.CXCL14 c.16.CXCL14 c.16.KRT15 c.16.KRT6B c.4.RGS5 c.14.CFD c.3.SFRP4 c.13.CRIP1 **c.7.CCL8** c.5.SOD2 c.8.CXCL2 c.13.SYNPO2 c.6.SFRP4

77

c.16.S100A2 c.16.KRT17 c.16.CALML3 c.16.APOE c.16.S100A6 c.16.IGFBP7 c.16.KRT6B c.16.MYC c.16.KRT75 c.10.KLF6 c.16.SAT1 c.15.CD164 c.5.CCL21 c.13.FLNA

80

c.16.LGALS1