

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

			c.0.HBB	c.0.HSPA1A	
			c.0.HBA2	c.0.HSPA1B	c.17.KLRB1
c.0.KRT6B	c.0.IFI27	c.16.KRT1	c.1.HBB	c.1.HSPA1A	c.17.LTB
c.0.KRT6C	c.1.IFI27	c.16.KRTDAP	c.1.HBA2	c.1.HSPA1B	c.10.TSC22D3
c.3.CFD	c.11.IFI27	c.16.KRT10	c.2.HBB	c.2.HSPA1A	c.12.UTS2
c.4.PHLDA1	c.16.IFI27	c.2.CYP1B1	c.2.HBA2	c.2.HSPA1B	c.5.ACTA2
c.5.POSTN	c.10.FOSB	c.15.ZG16B	c.3.HBB	c.2.DNAJB1	c.17.CCL4
c.5.ACTA2	c.5.TM4SF1	c.0.CST6	c.3.HBA2	c.2.HSP90AA1	c.5.CRYAB
c.11.KRT6B	c.3.WIF1	c.16.LY6D	c.4.HBB	c.4.HSPA6	c.8.TAGLN
c.1.KRT15	c.7.IFI27	c.13.MGP	c.4.HBA2	c.4.HSPA1B	c.10.DUSP1
c.13.CCDC80	c.15.DCD	c.9.CXCL1	c.3.STMN2	c.10.TXNIP	c.15.NEAT1
c.12.PPP1R15A	c.6.CXCL9	c.9.HSPB1	c.10.FOSB	c.10.KLF6	c.3.TNC
c.3.STMN2	c.3.STMN2	c.3.WIF1	c.3.POSTN	c.1.APOE	c.1.APOE
c.10.TXNIP	c.10.FOS	c.3.POSTN	c.9.S100A12	c.10.KLF2	c.3.WISP2
c.10.KLF6	c.3.TNC	c.3.ADH1B	c.5.FABP4	c.1.POSTN	c.0.S100A2
c.3.SAA1	c.13.CCDC80	c.15.UQCRH	c.0.CST6	c.7.LYZ	c.5.FABP4
c.3.IGFBP2	c.3.STC2	c.6.POSTN	c.14.HBB	c.17.MTCO1	c.1.POSTN
c.2.CLU	c.3.IGFBP2	c.13.FLNA	c.15.MALAT1	c.10.TSC22D3	c.7.LYZ
c.3.WIF1	c.3.GSN	c.5.CFH	c.3.ADH1B	c.5.CCL21	c.5.CCL21
c.3.POSTN	c.1.APOE	c.16.SPRR1B	c.15.UQCRH	c.11.HSPA1B	c.16.MYC

1

45

79

81

85

86

Significant Latent Factor