

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

				c.0.S100A8
				c.0.KRT6A
				c.0.IFI27
				c.0.S100A7
				c.0.KRT16
	c.9.S100A8			c.0.SERPINB2
c.2.HLADRB5	c.9.S100A9		c.2.KRT14	c.0.KRT14
c.7.HLADRB5	c.9.SERPINB2		c.10.KRT14	c.0.S100A9
c.9.HLADRB5	c.9.THBS1		c.15.KRT14	c.0.HES1
c.13.CFD	c.9.S100A12		c.15.S100A2	c.0.FOS
c.15.CXCL2	c.9.TNFAIP6		c.15.KRT5	c.4.NNMT
c.3.PLAG2A	c.0.KRT2		c.15.LY6D	c.3.IGFBP5
c.16.LYPD3	c.9.EREG		c.5.CCL2	c.3.STMN2
c.16.TM4SF1	c.9.CCL20		c.15.HSPA1A	c.0.KRT5
c.2.IER3	c.12.PLIN2	c.2.HLADQA1	c.13.CLU	c.11.KRT6A
c.16.SERPINB2	c.14.KRT1	c.9.IL23A	c.4.TCIM	c.3.CD55
c.14.CFD	c.9.AREG	c.3.COMP	c.2.CLU	c.3.IFI27
c.13.CRIP1	c.12.JUN	c.17.TOB1	c.1.KRT15	c.3.SLPI
c.9.CXCL3	c.16.LYPD3	c.12.LGALS1	c.6.ADAMDEC1	c.3.TPPP3
c.17.LTB	c.15.SLC12A2		c.5.CFD	c.4.C1R