

## SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

c.3.MT1X	c.9.S100A8	c.10.CCL20	c.2.HLADQA1	c.5.APOE	
c.4.MT1X	c.9.S100A9	c.11.KRT16	c.3.COMP	c.14.CD74	c.2.IFIT1
c.5.MT1X	c.9.S100A12	c.17.GZMB	c.12.LGALS1	c.14.HLADRA	c.2.ISG15
c.6.MT1X	c.9.EREG	c.7.CCL13	c.13.TINAGL1	c.14.HLADRB1	c.2.IFI44L
c.8.MT1X	c.9.SERPINB2	c.8.CTGF	c.17.TOB1	c.14.HLADPA1	c.2.IFI6
c.8.CTGF	c.11.KRT2	c.11.FABP5	c.9.IL23A	c.14.IFIM2	c.3.IFI27
c.17.CCL4L2	c.0.KRT2	c.14.MBP	c.13.NDUFA4L2	c.2.APOE	c.3.IFI6
c.12.AREG	c.9.TNFAIP6	c.11.KRT15	c.12.DUSP14	c.12.CCL4L2	c.4.ISG15
c.17.GZMB	c.5.CRYAB	c.5.POSTN	c.9.C1QC	c.4.CCL2	c.5.IFI27
c.7.CCL13	c.6.SFRP4	c.4.PHLDA1	c.0.KRT2	c.3.ADHB1	c.5.IFI6
c.3.STMN2	c.3.STMN2	c.3.STMN2	c.10.FOS	c.3.STMN2	c.6.IFI27
c.10.FOS	c.10.FOS	c.10.FOS	c.10.TXNIP	c.10.FOS	c.3.STMN2
c.10.KLF6	c.10.TXNIP	c.10.TXNIP	c.10.KLF6	c.10.KLF6	c.10.FOS
c.3.SAA1	c.10.KLF6	c.10.KLF6	c.3.SAA1	c.13.CCDC80	c.10.TXNIP
c.3.TNC	c.3.TNC	c.3.SAA1	c.3.TNC	c.6.CCL19	c.10.KLF6
c.5.IGFBP5	c.5.IGFBP5	c.3.TNC	c.5.IGFBP5	c.3.STC2	c.3.SAA1
c.13.CCDC80	c.13.CCDC80	c.5.IGFBP5	c.13.CCDC80	c.3.IGFBP2	c.3.TNC
c.6.CCL19	c.6.CCL19	c.13.CCDC80	c.6.CCL19	c.3.GSN	c.5.IGFBP5
c.3.STC2	c.3.STC2	c.6.CCL19	c.3.STC2	c.3.WIF1	c.13.CCDC80
c.3.IGFBP2	c.3.IGFBP2	c.3.STC2	c.3.IGFBP2	c.1.APOE	c.6.CCL19