

SLIDE Marginal Variables

Genes Associated with Significant Latent Factors

c.2.ISG15		c.3.PRSS23	c.3.MT1X	c.7.APOE	c.2.HLADRB5	c.10.HSPA1B	c.4.TCIM	c.15.SAA1
c.2.IF16		c.3.SERPINE2	c.6.MT1X	c.7.APOC1	c.7.HLADRB5	c.10.DNAJB1	c.13.FLNA	c.15.SAA2
c.5.TM4SF1		c.3.MFAP5	c.12.AREG	c.7.S100A9	c.9.HLADRB5	c.10.HSPA1A	c.9.CCR7	c.15.LTF
c.2.IF144L	c.3.IGFBP5	c.3.IGFBP2	c.5.MT1X	c.10.HSPA6	c.15.CXCL2	c.15.TSC22D1	c.16.SPINK5	c.15.SOD2
c.14.IF127	c.3.CD55	c.10.LPAR6	c.17.TOB1	c.17.HSPA6	c.1.CXCL14	c.7.TNF	c.17.HSPA6	c.15.SLC12A2
c.1.NEAT1	c.12.AREG	c.8.DPEP1	c.4.MT1X	c.12.ID2	c.3.PLA2G2A	c.12.HSPE1	c.3.PTN	c.15.MALAT1
c.13.PTGDS	c.9.CCR7	c.6.GPC3	c.6.CLU	c.8.HES1	c.17.IFNG	c.6.CXCL2	c.4.PLN	c.10.CSF2
c.16.IF127	c.17.CXCR6	c.3.LUM	c.3.MFAP5	c.7.SPP1	c.13.CFD	c.8.HES1	c.15.TSC22D1	c.8.TAGLN
c.14.FOS	c.10.TXNIP	c.17.CCL4L2	c.3.WISP2	c.9.CCR7	c.0.HES1	c.7.S100A9	c.15.SLC12A2	c.8.POSTN
c.0.S100A7	c.3.APOE	c.5.ACTA2	c.7.PLTP	c.2.HSPA1A	c.5.HSPA6	c.8.APOD	c.16.KRT6B	c.12.AP3S1
c.3.SLPI	c.9.SGK1	c.3.SLPI	c.3.SLPI	c.3.SLPI	c.3.SLPI	c.3.SLPI	c.3.SLPI	c.3.SLPI
c.3.TPPP3	c.10.XCL2	c.3.TPPP3	c.3.TPPP3	c.3.TPPP3	c.3.TPPP3	c.3.TPPP3	c.3.TPPP3	c.3.TPPP3
c.10.FOS	c.5.MT1X	c.10.FOS	c.10.FOS	c.10.FOS	c.10.FOS	c.10.FOS	c.10.FOS	c.10.FOS
c.10.TXNIP	c.3.SLPI	c.10.TXNIP	c.10.TXNIP	c.10.TXNIP	c.10.TXNIP	c.10.TXNIP	c.10.TXNIP	c.10.TXNIP
c.3.DIO2	c.3.TPPP3	c.3.DIO2	c.3.DIO2	c.3.DIO2	c.3.DIO2	c.3.DIO2	c.3.DIO2	c.3.DIO2
c.10.KLF6	c.10.FOS	c.10.KLF6	c.10.KLF6	c.10.KLF6	c.10.KLF6	c.10.KLF6	c.10.KLF6	c.10.KLF6
c.3.TNC	c.3.DIO2	c.3.TNC	c.3.TNC	c.3.TNC	c.3.TNC	c.3.TNC	c.3.TNC	c.3.TNC
c.4.CXCL3	c.10.KLF6	c.4.CXCL3	c.4.CXCL3	c.4.CXCL3	c.4.CXCL3	c.4.CXCL3	c.4.CXCL3	c.4.CXCL3
c.3.TNXB	c.3.TNC	c.3.TNXB	c.3.TNXB	c.3.TNXB	c.3.TNXB	c.3.TNXB	c.3.TNXB	c.3.TNXB
c.5.IGFBP5	c.4.CXCL3	c.5.IGFBP5	c.5.IGFBP5	c.5.IGFBP5	c.5.IGFBP5	c.5.IGFBP5	c.5.IGFBP5	c.5.IGFBP5