Up in sLC vs. vLC (58) SLC9A2 C100RF35 PDLIM3 PDLIM3 C110RF41 NRCAM FLJ10986 C19ORF64 TSPAN13 MLLT11 СОСН KIAA0363 PON2 TMEM200A FGGY PDE9A NINJ2 CD9 HS.561747 ALDH1B1 ANXA3 KIFAP3 RTKN SERPINE2 AXIN2 SWAP70 TNFSF4 WRN TBC1D4 ABCB10 TUBB2B MALT1 INSM1 SQLE

SLC25A17

HS.371609

LIMCH1

IGSF3

PHF16

CLINT1

HLA-DPB1

FAM129A

FOXD4L1

LOC286512

LOC652479

SAMD9L

ATP11C

BUB3

CSDE1

FYN

sLC sCD14- DC sCD14+ DC vLC vCD14- DC vCD14+ DC

GALC

N4BP2L1

EPM2AIP1

HMGN4

ST7

RAB23

FYTTD1

GCLC



Up in vLC vs. sLC (107)

DAB2

CD93

CLEC4A

PTPRE

MS4A7

SOD2

PTPRE

CASP1

ADRB2

CXCR7

ATP8B4

NDRG1

RHOQ

FOXO3

NBPF10

NPC1

SGK

IL1RN

LOC653158

FAM108C1

ZNF148

ITPRIP

RIOK3

ECHDC2

GSTM2

GSTM1

A2ML1

DENND2C

SERPINB3

FGFBP1

IGLL1

KLK10

RDH13

MALL

SEN

PRDM1

HSPB8

SPINK7

IGFBP3

PLCXD1

ABLIM1

C140RF78

DEFB103B

TMPRSS11D

TMEM156

METRNL

KLF2

Figure S8 - Hierarchical clustering of the 165 transcripts differentially expressed between sLC and vLC. Left panel: 58 transcriptsover-expressed in sLC. Right panel: 107 transcripts over-expressed in vLC. Data are normalized to the median of all samples.