39_treated_v1, Cytokine_genes, genes=31

species

stage

hippo_group_untreated

Row labels (top to bottom):
IL16
CXCL16
CX3CL1
CXCL1
CCL20
IL1B
CCL2
IL6
CXCL2
CCL5
CXCL10
IL4
CCL25
CCL17
CCL3
CCL27
IL10
CXCL11
CXCL5
CCL22
CSF2
CCL24
CCL4
IFNG
IL2
CCL19
TNF
CXCL12
CCL7
CXCL13
CCL11

Column labels:
Hs_WM45.1.G007.LK
Hs_WM266.G007.LK
Hs_WM239.G007.LK
Hs_MeWo.G007.LK
Hs_FemxI.G007.LK
Hs_FemxV.G007.LK
Hs_WM35.G007.LK
Hs_WM1341.G007.LK
Hs_Skmel28.G007.LK
Hs_WM115.G007.LK
Hs_WM1382.G007.LK
Hs_WM1366.G007.LK
Hs_A375.G007.LK
Mm_WNT_G007LK
Mm_G007LK
Hs_WM983.G007.LK
Hs_WM852.G007.LK
Hs_WM9.G007.LK
Hs_WM793.G007.LK
Hs_LOX.G007.LK

species
human
mouse

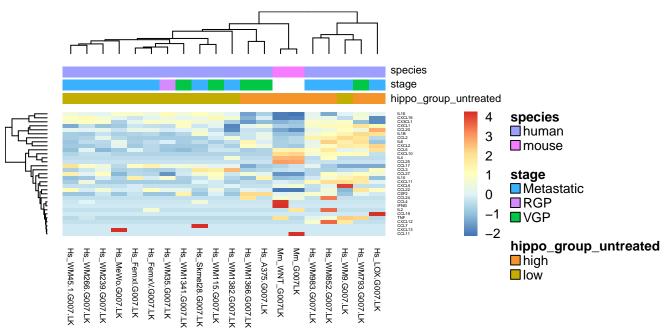**stage**
Metastatic
RGP
VGP

hippo_group_untreated
high
low

Legend scale: 4, 3, 2, 1, 0, −1, −2

samples: G007LK and also WNT_G007LK for mouse
genes: 32 genes described as cytokins.
value: log2 of TPM
zscorecolors: row