## PCA plot of dataset (SCP1288) with method: deseq2 41BB-Hi\_CD8+\_T\_cell 41BB-Lo\_CD8+\_T\_cell CD16+\_Monocyte B\_cell CD16-\_Monocyte CD1C+\_DC 20 10-0 -10 --20· -30 -CLEC9A+\_DC CXCL10-Hi\_TAM Cycling\_CD8+\_T\_cell Cycling\_TAM Cycling\_Tumor Effector\_T-Helper 20 -10 -0 --10 --20 **-**-30 -Endothelial FGFBP2-\_NK FGFBP2+\_NK Fibroblast FOLR2-Hi\_TAM GPNMB-Hi\_TAM 20 -10-0 -10· -20· 25 -30 J ICB\_Exposed ICB LowLibSize\_Macrophage Mast\_cell Memory\_T-Helper MitoHigh\_CD8+\_T\_cell Misc\_Undetermined MitoHigh\_Myeloid NoICB 10 0 -10· -20 **-**-30 -MitoHigh\_T-Helper MitoHigh\_NK MX1-Hi\_CD8+\_T\_cell NKT Plasma\_cell T-Reg 20 -10 0 -10· -20· -30 --25 -25 25 -25 25 25 TP2 VSIR-Hi\_TAM TP1 20 -10 --10

-20 -30

-25

0

-25

25

-25

25

PC1

25