## PCA plot of dataset (SCP1288) with method: edger\_robust\_likelihood\_ratio 41BB-Hi\_CD8+\_T\_cell 41BB-Lo\_CD8+\_T\_cell B\_cell CD16+\_Monocyte 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· 22 -30 J ICB\_Exposed ICB LowLibSize\_Macrophage Mast\_cell Memory\_T-Helper MitoHigh\_CD8+\_T\_cell Misc\_Undetermined MitoHigh\_Myeloid NoICB 10 -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 TP2 VSIR-Hi\_TAM TP1 20 · 10 --10 -20 · -30

-25

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0

-25

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PC1

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