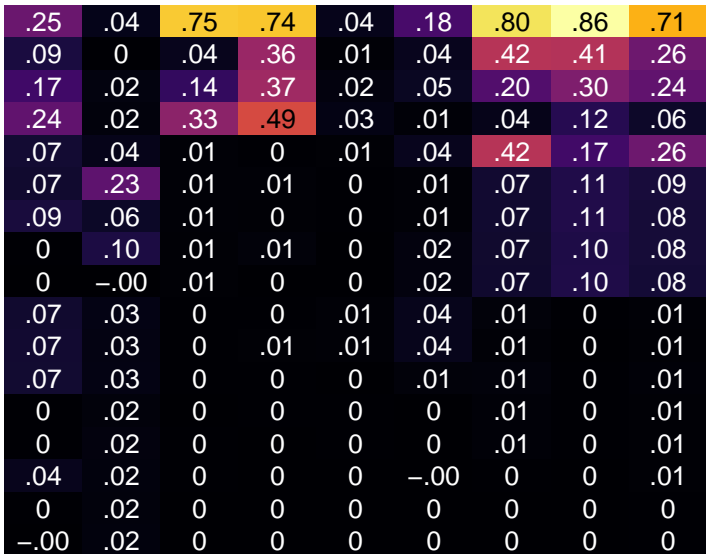
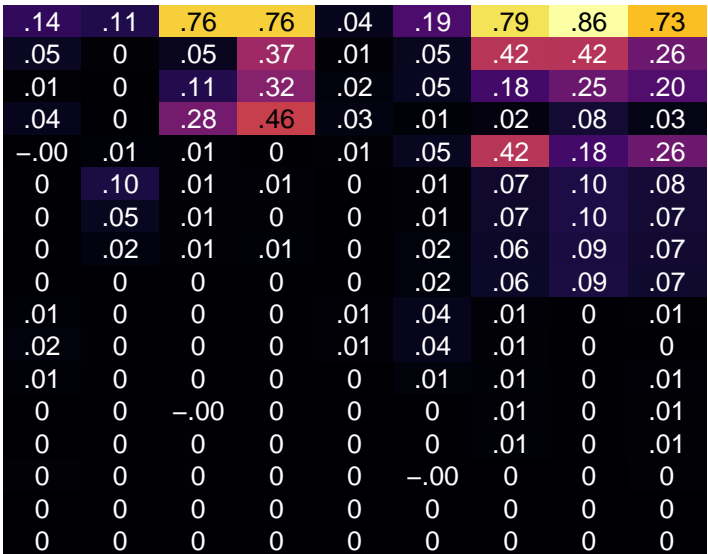


A

var explained by
log10_total_counts

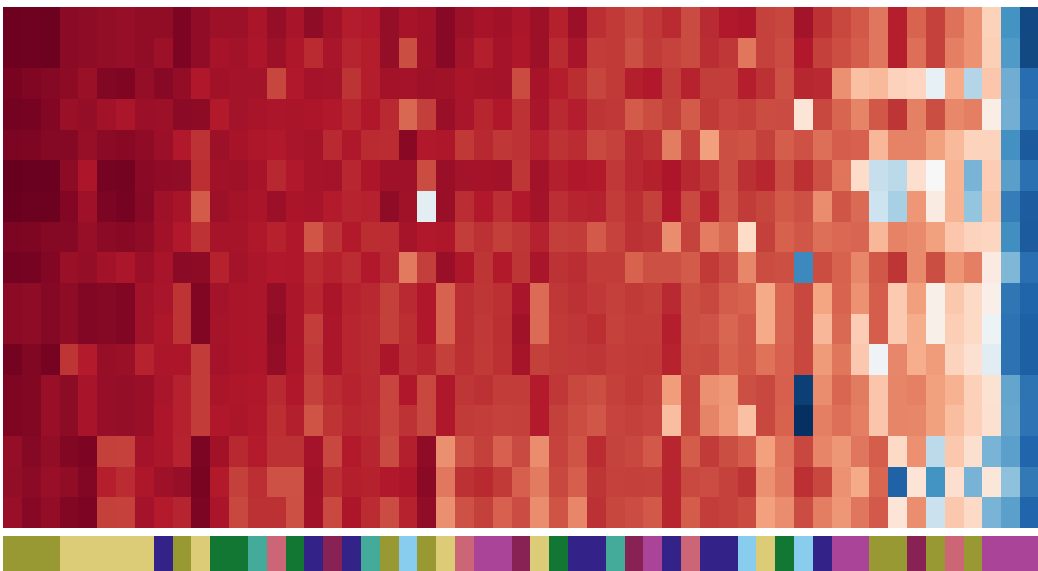
var explained by
log10_total_features



none
none.scaled
seurat (no scaling)
scran
scnorm.scaled
seurat.feats_mt_regress
seurat.feats_regress
sctransform.feats_mt_regress
sctransform.feats_regress
stableG
stableG.nucleus
stableGsum
seurat.mt_regress
seurat
scran.scaled
sctransform
sctransform.mt_regress

B

Subpopulation silhouette



sctransform
sctransform.mt_regress
seurat (no scaling)
sctransform.feats_regress
seurat
scran
scran.scaled
seurat.mt_regress
sctransform.feats_mt_regress
stableG.nucleus
stableG
stableGsum
seurat.feats_regress
seurat.feats_mt_regress
scnorm.scaled
none
none.scaled

silhouette
width
1
0.5
0
-0.5

dataset
Koh
Kumar
mixology10x3cl
mixology10x5cl
simMix1
simMix2
Zhengmix4eq
Zhengmix4uneq
Zhengmix8eq

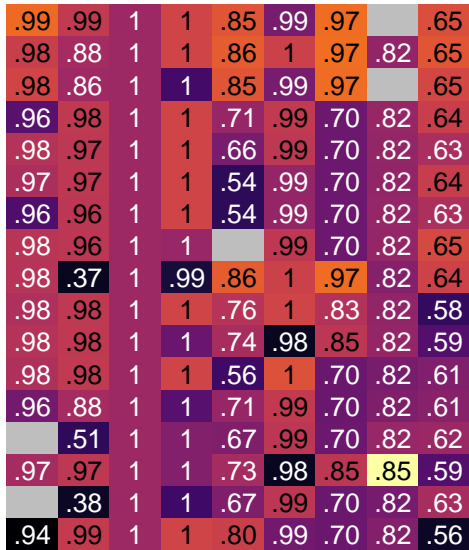
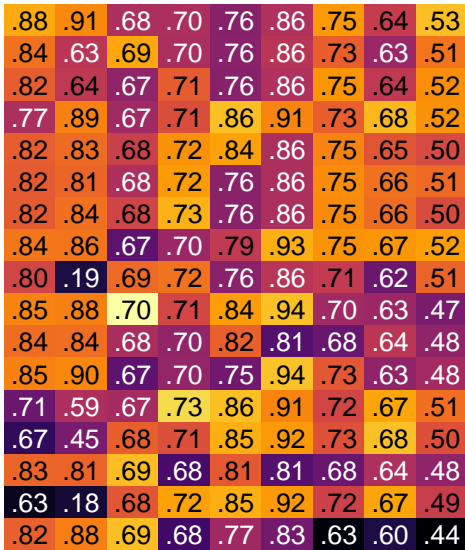
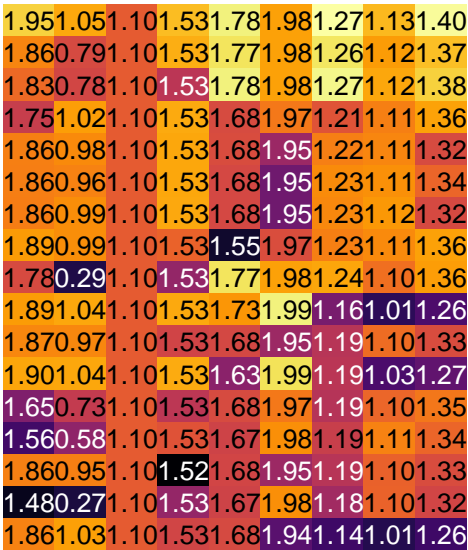
C

MI

ARI

ARI at true

of clusters



sctransform
sctransform.feats_regress
sctransform.mt_regress
seurat
stableGsum
stableG.nucleus
stableG
scran.scaled
sctransform.feats_mt_regress
seurat (no scaling)
scnorm.scaled
scran
seurat.mt_regress
seurat.feats_regress
none.scaled
seurat.feats_mt_regress
none