

18 head and neck cancer samples, 5,902 cells

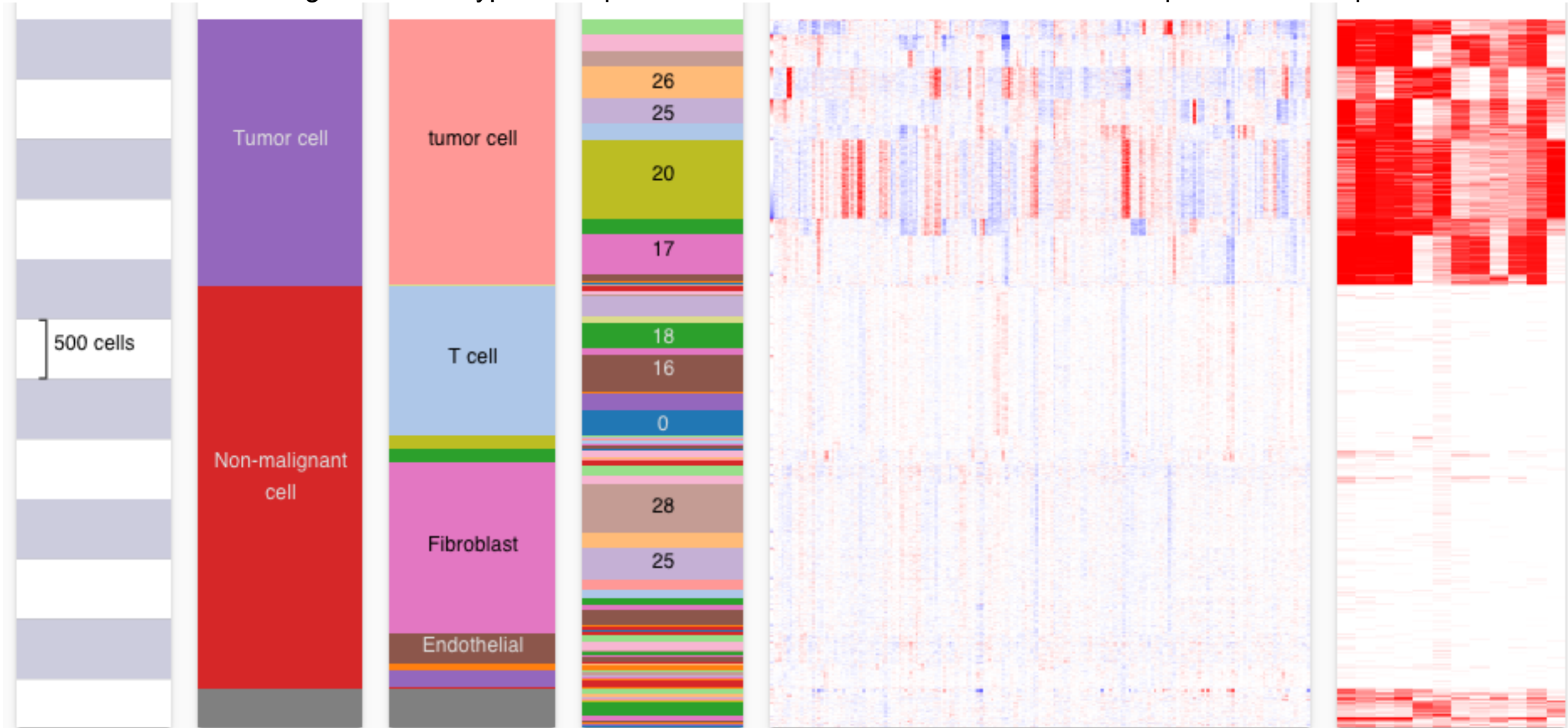
malignant vs.
non-malignant

cell type

patient ID

whole genome large-scale CNV
estimated from RNAseq data

gene expression
epithelial markers



-0.35 0.35
z-score

low high
TPM