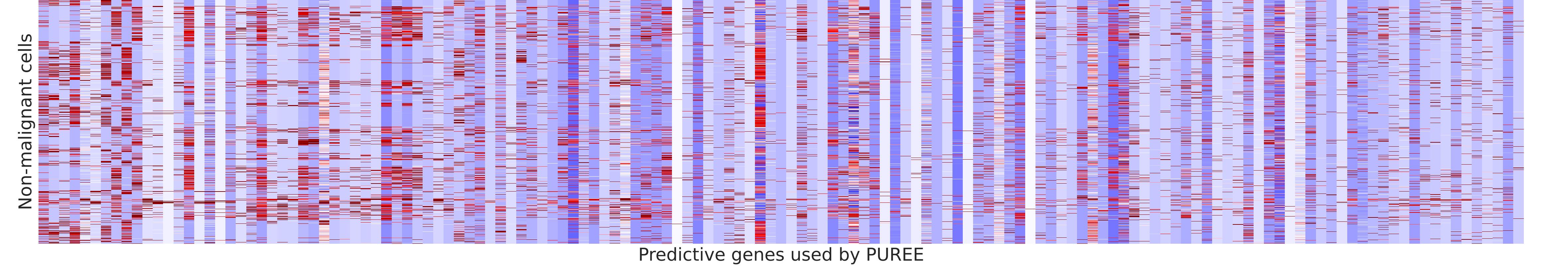
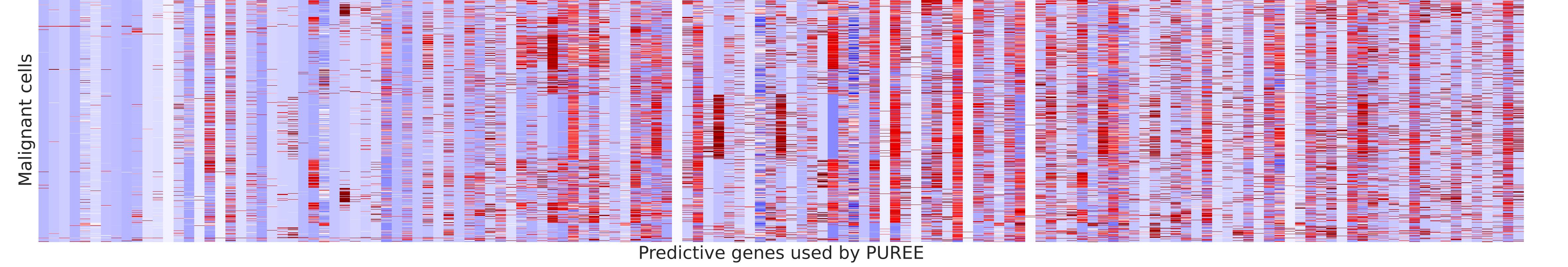


Head and neck cancers single-cell dataset, 5902 cells

Z-scores of NON-MALIGNANT cells' expression, 2539 cells



Z-scores of MALIGNANT cells' expression, 3363 cells



Correlation of genes' expressions with tumor purity on TCGA train set, average of mean across 20 cancer types

