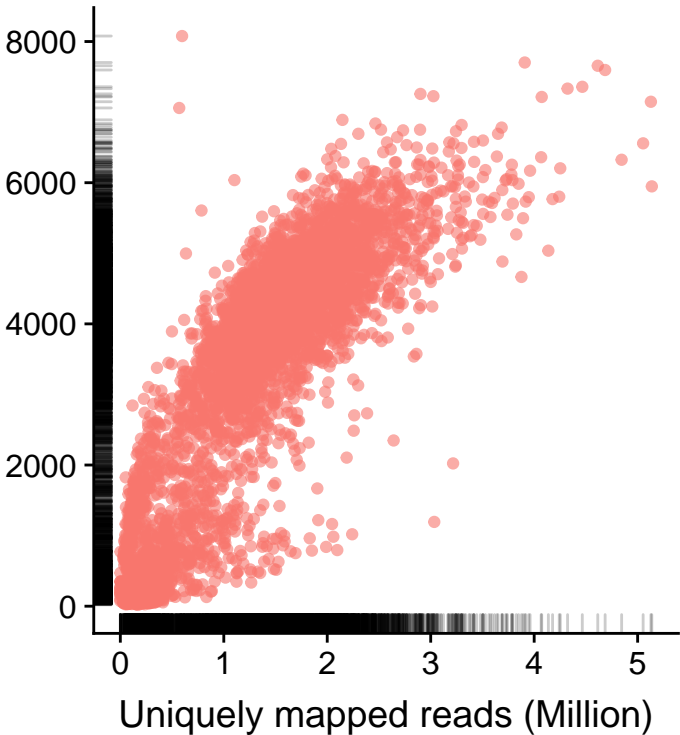


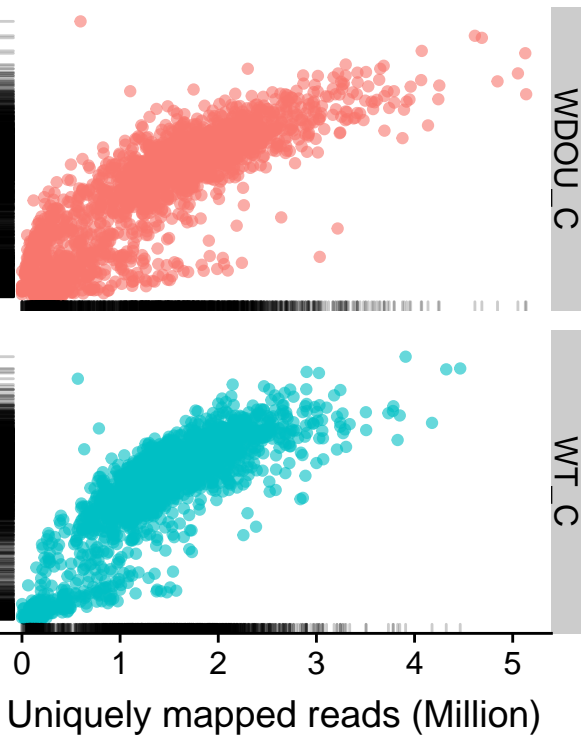
Number of detected genes

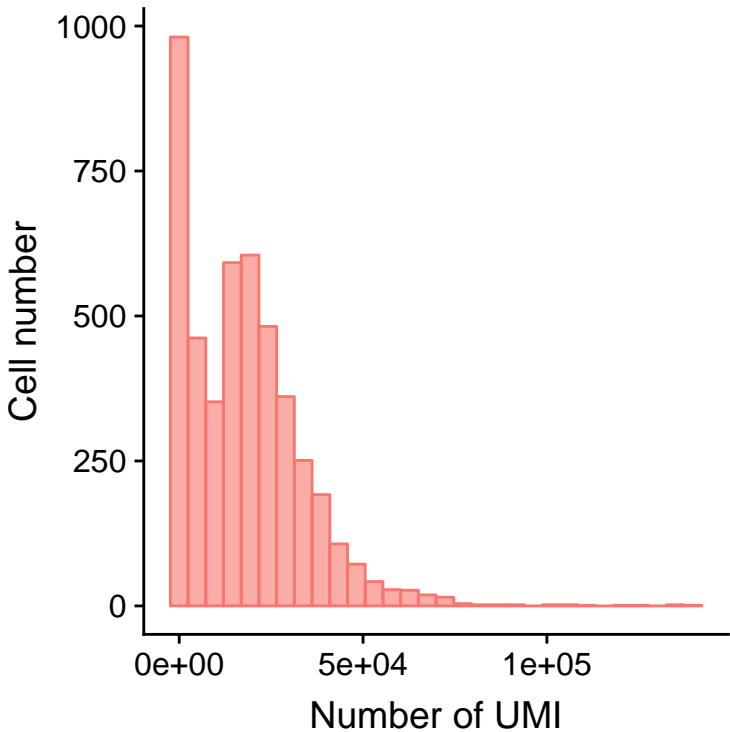


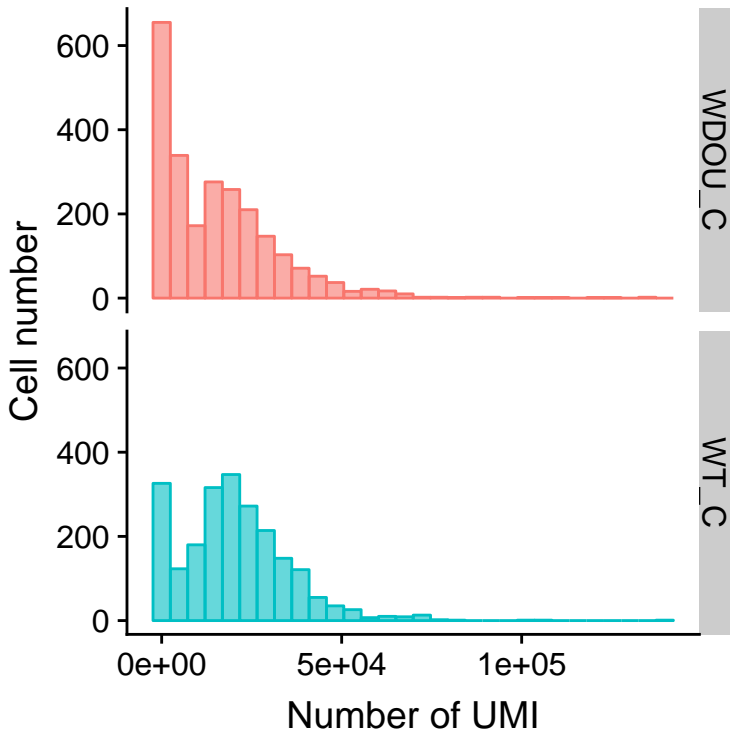
Number of detected genes

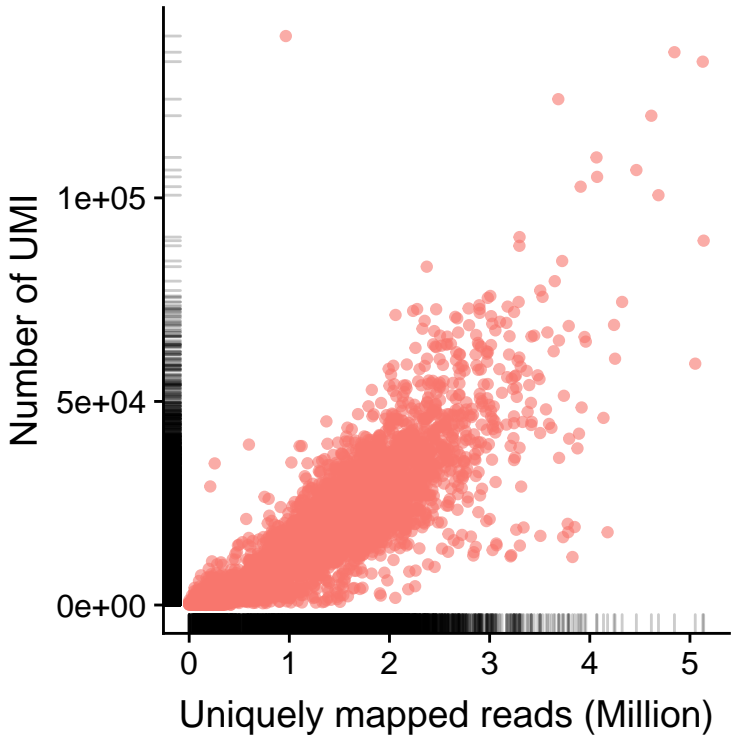
WDOU_C

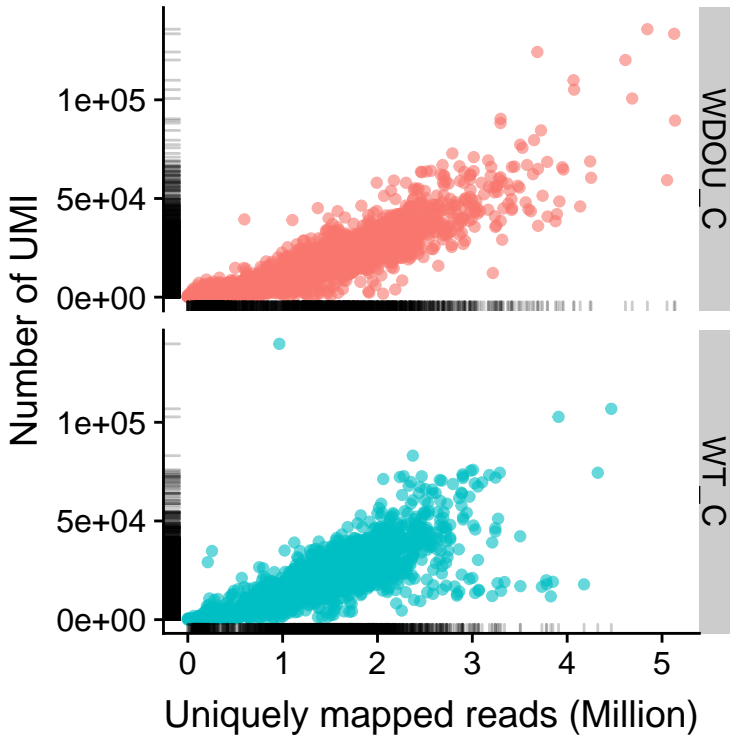
WT_C



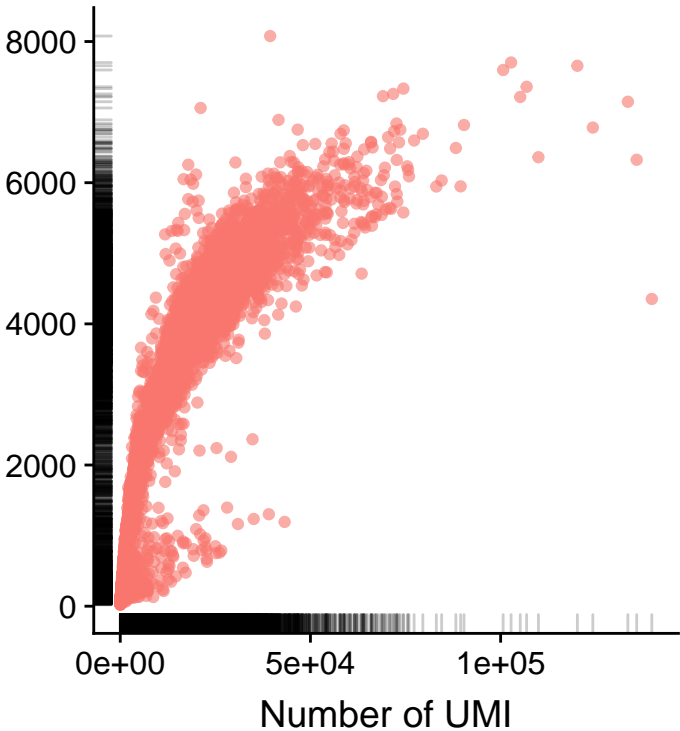








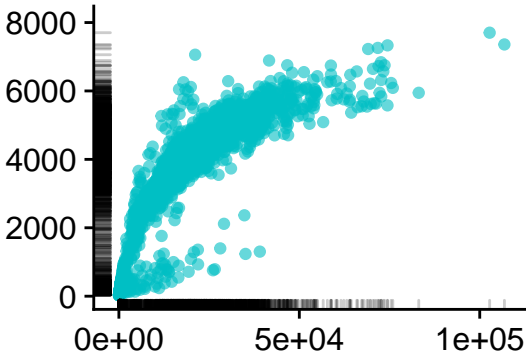
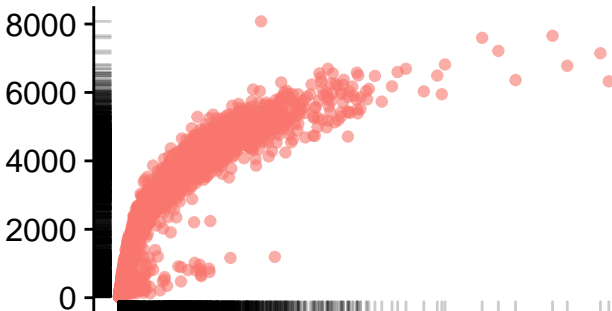
Number of detected genes



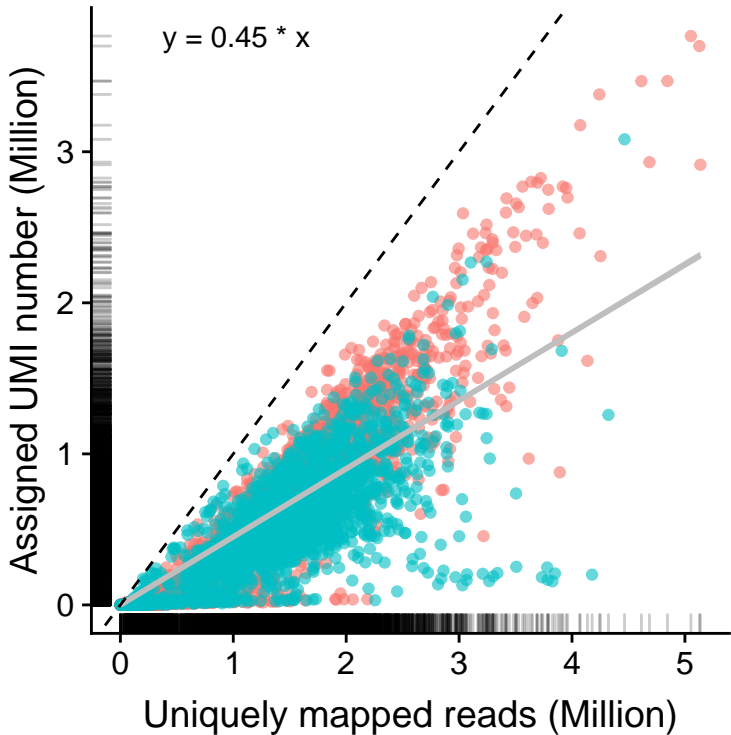
Number of detected genes

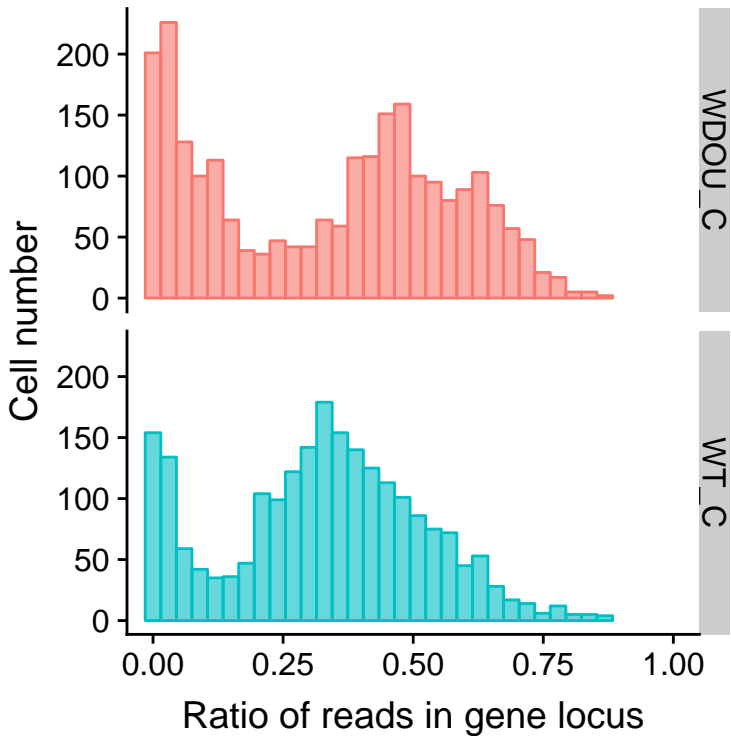
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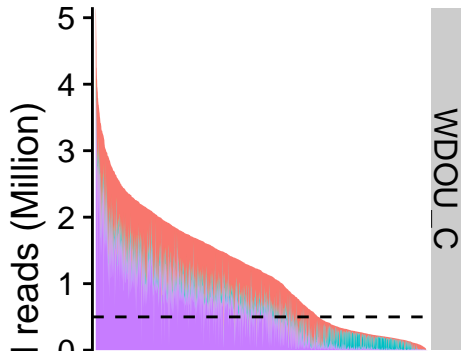
WT_C



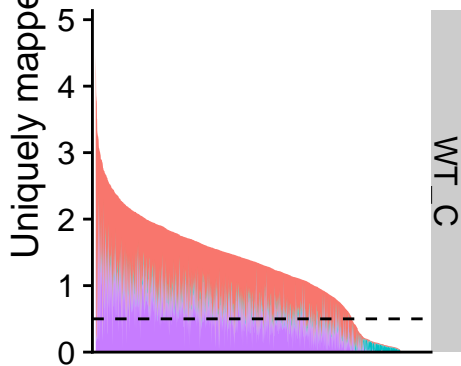
Number of UMI

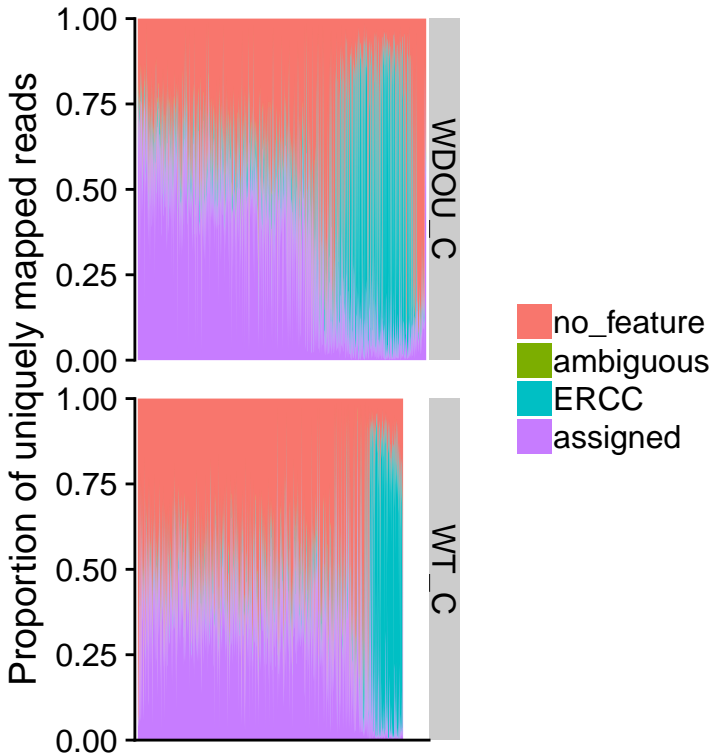


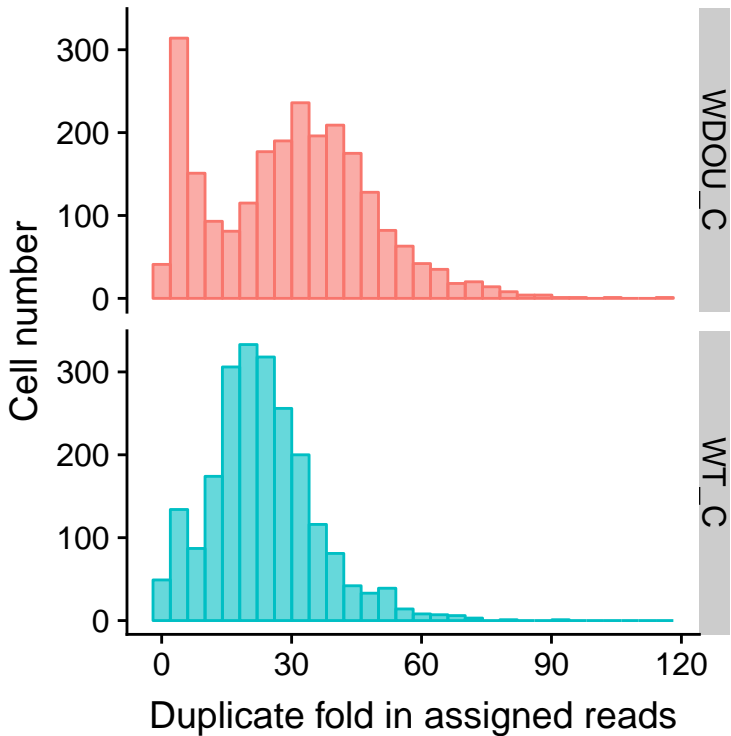




no_feature
ambiguous
ERCC
assigned



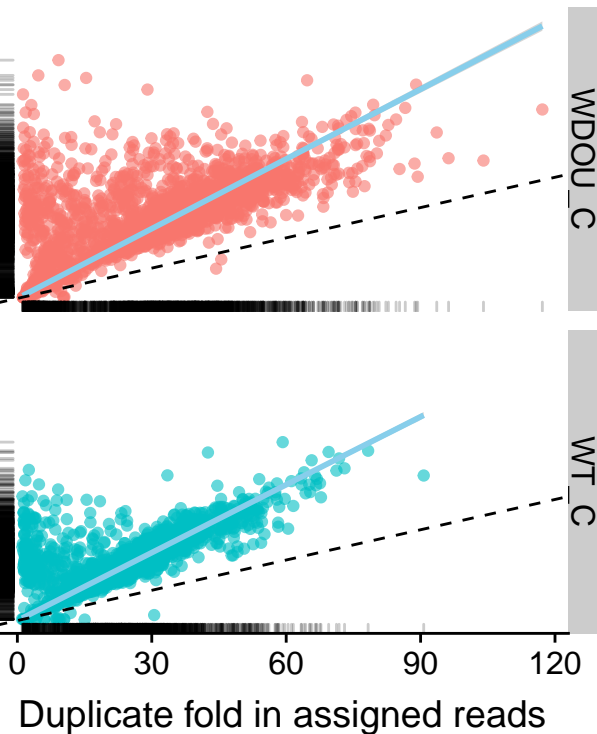




Duplicate fold in ERCC reads

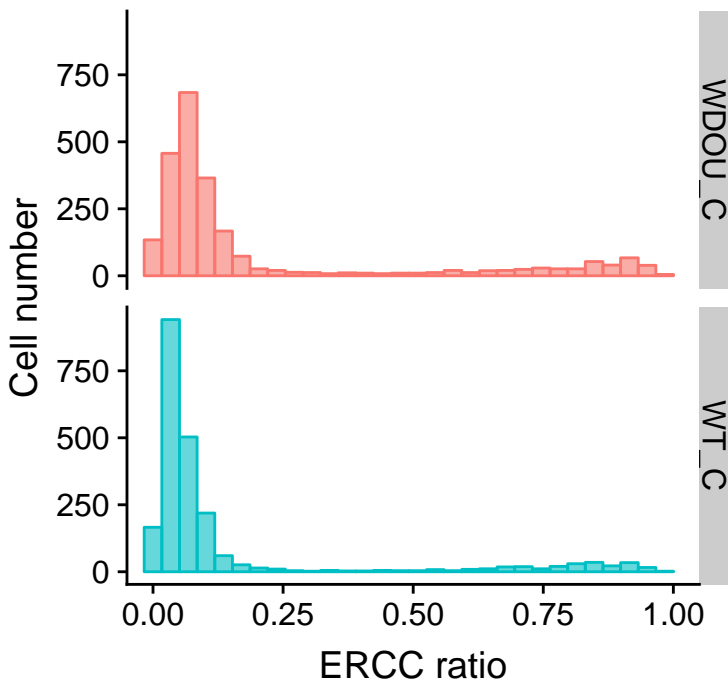
WDOU_C

WT_C

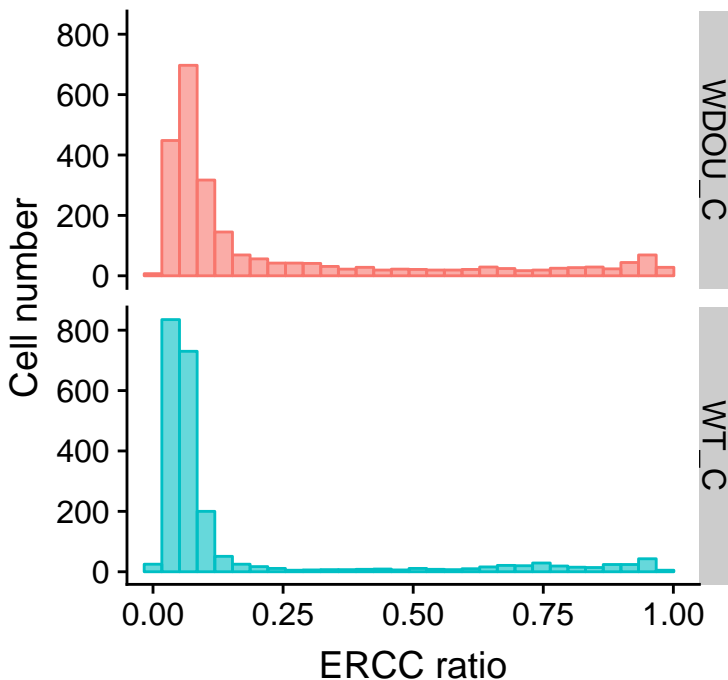


Duplicate fold in assigned reads

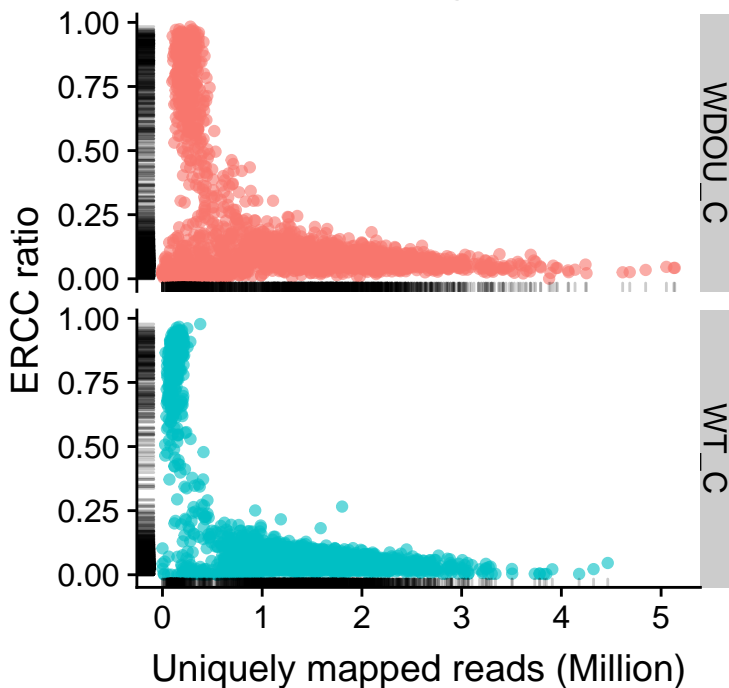
Before removing duplicates



After removing duplicates



Before removing duplicates



After removing duplicates

