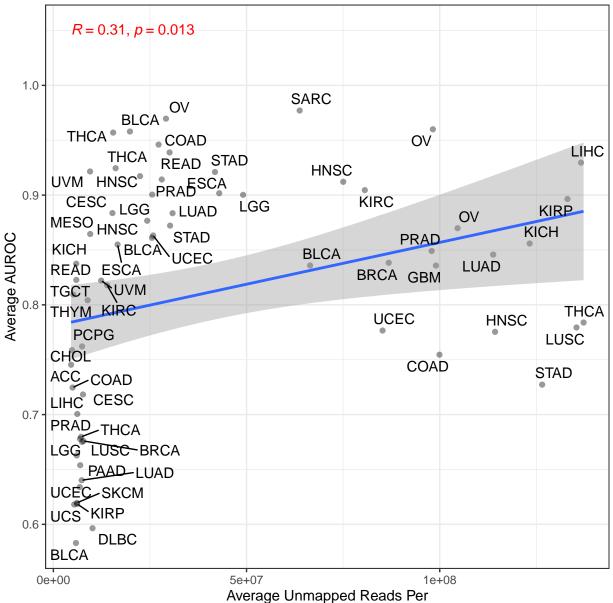
Correlating AUROC and unmapped reads for primary tumor TCGA models (Raw data)



Seq Center & Cancer Type (Primary Tumor)