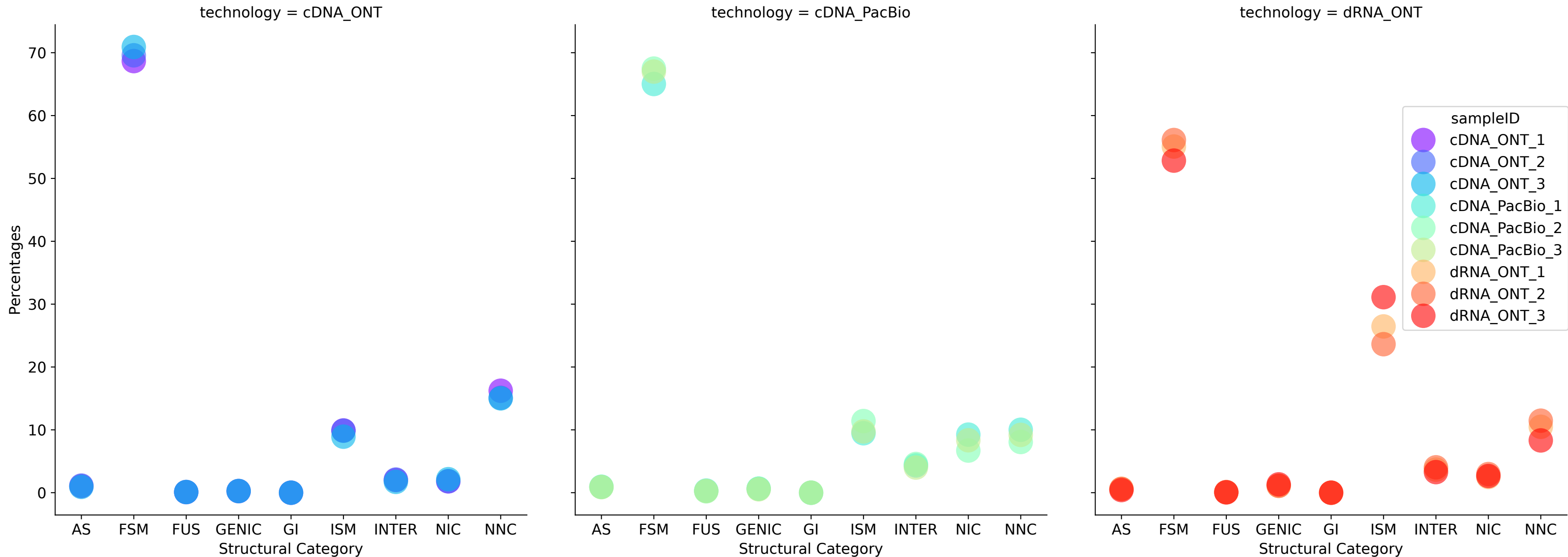
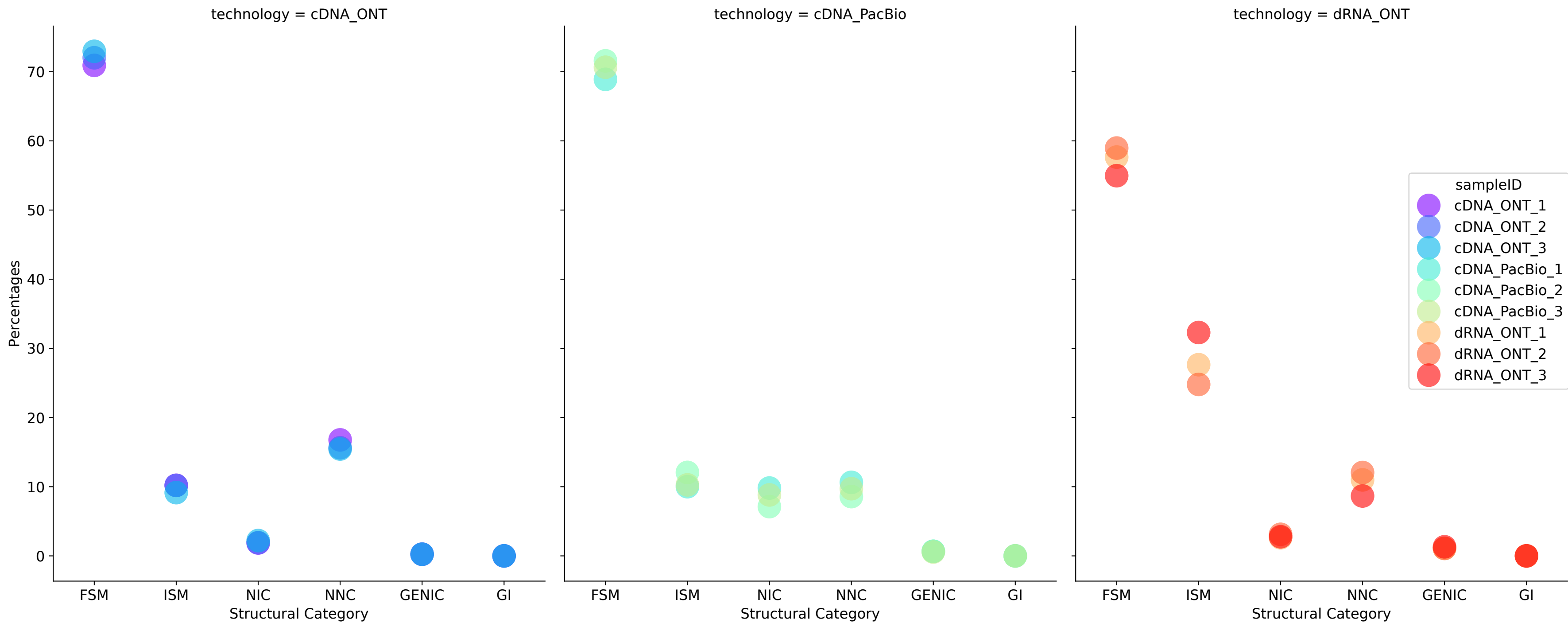


SQANTI-reads report

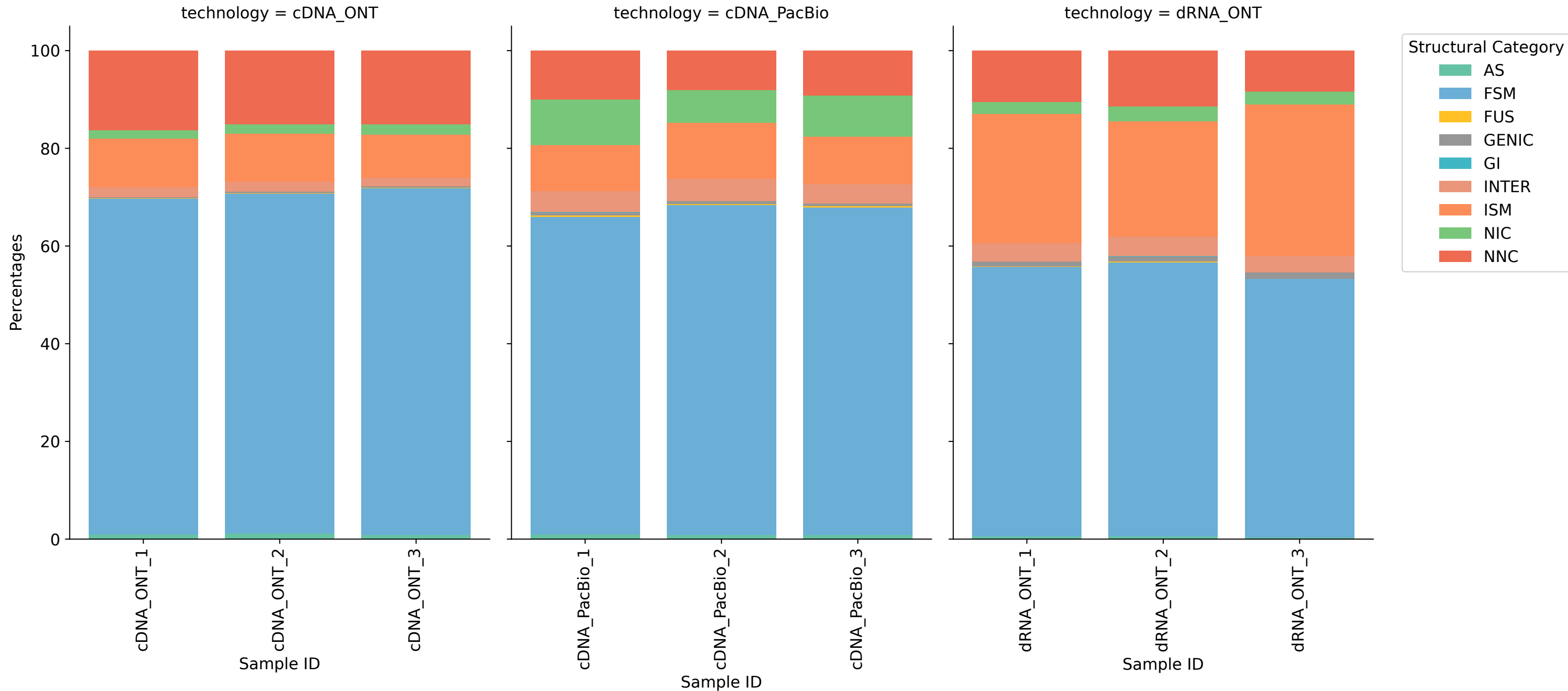
Percent Reads in Each Structural Category - All Genes



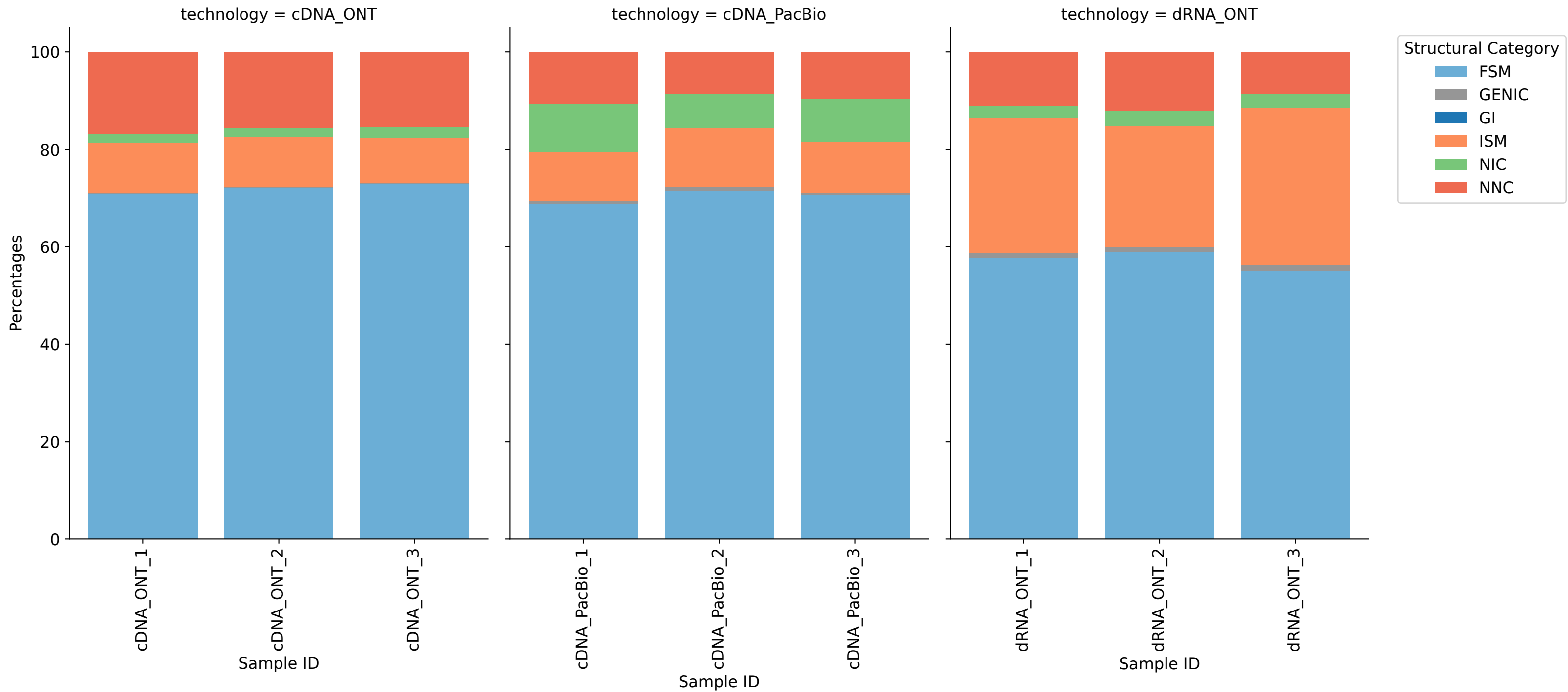
Percent Reads in Each Structural Category - Annotated Genes



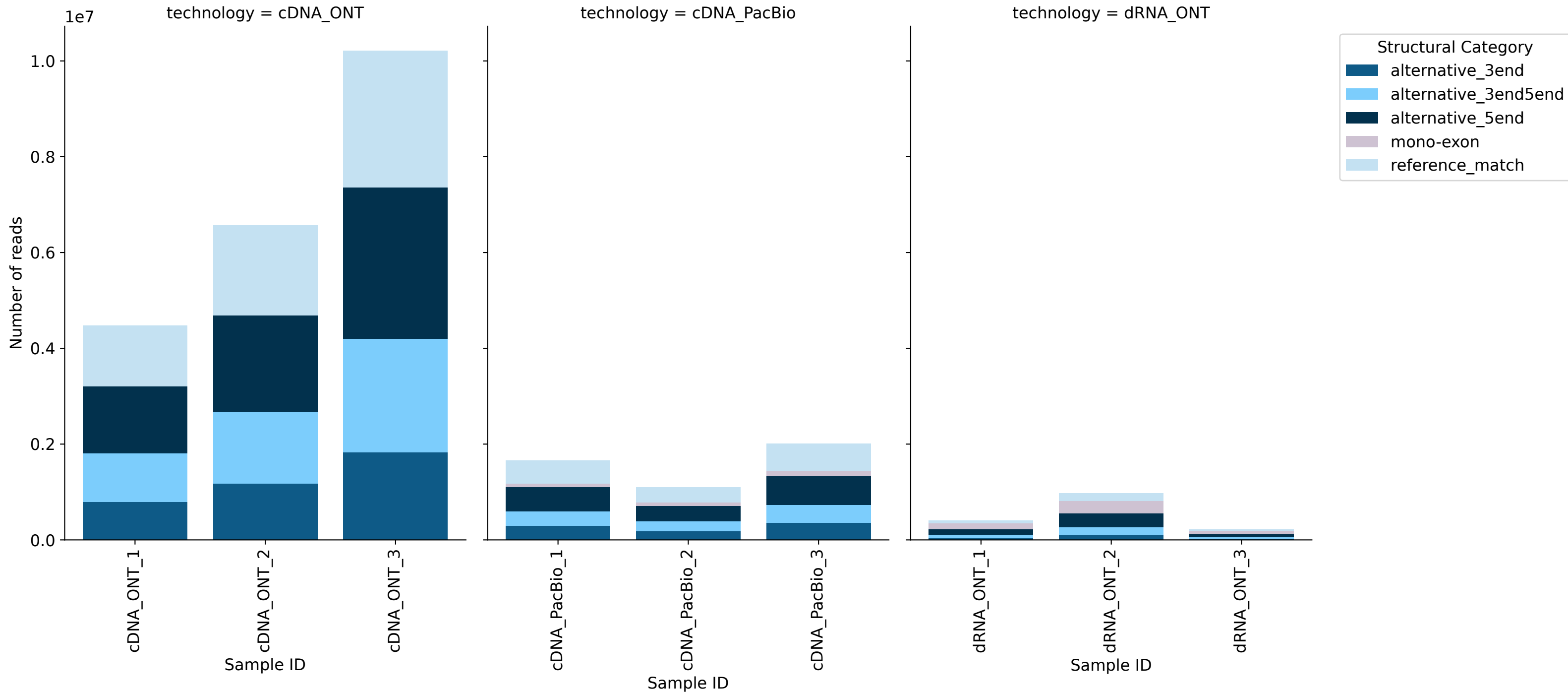
Percentage Reads in Each Structural Category - All Genes



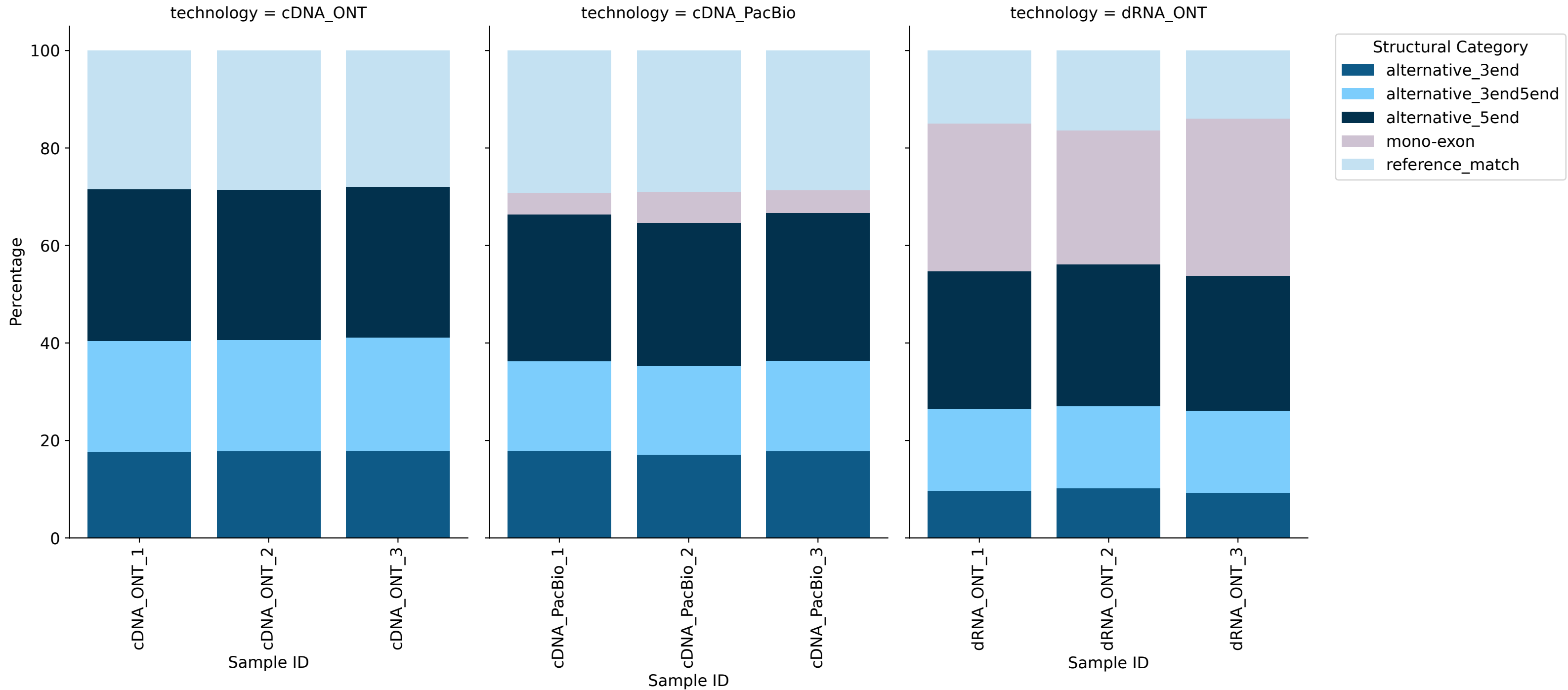
Percentage Reads in Each Structural Category - Annotated Genes



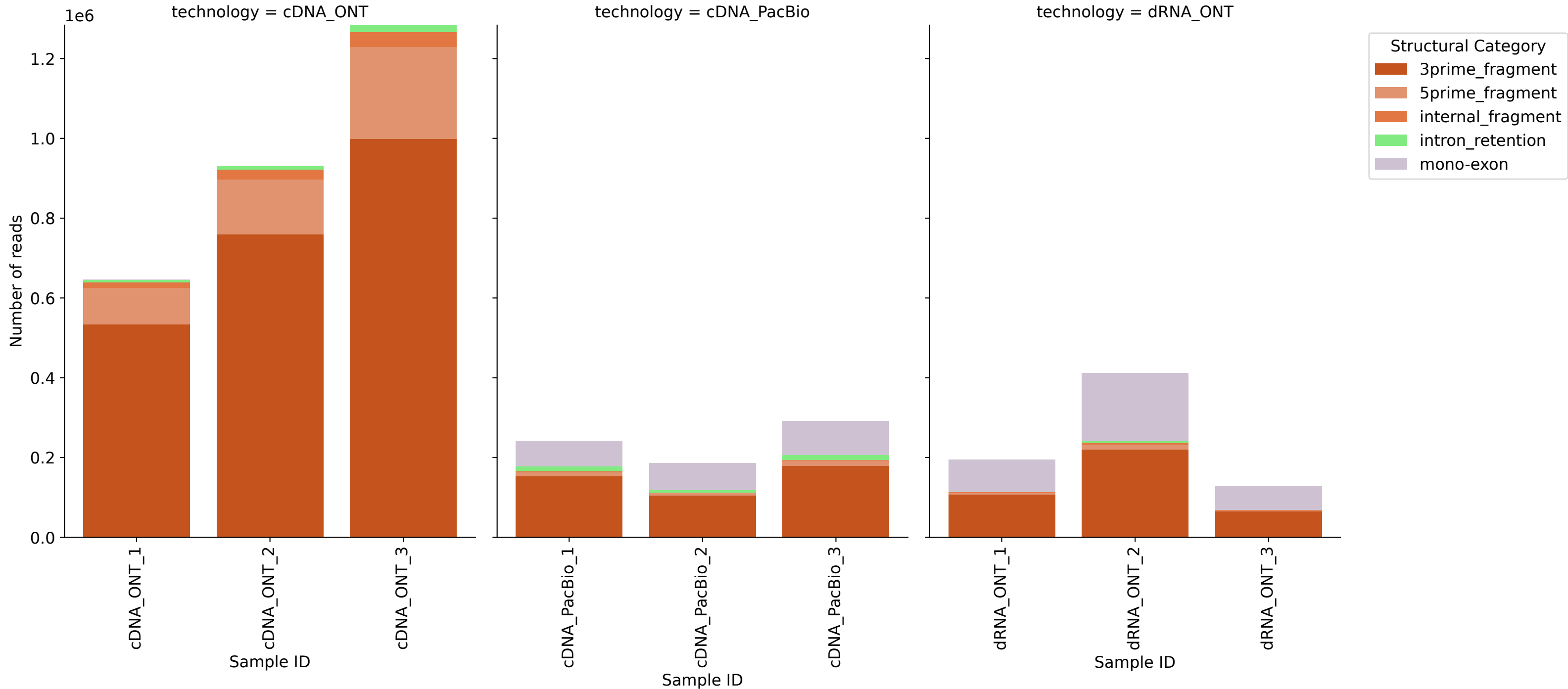
Number of Reads in Each Sub-Category - FSM



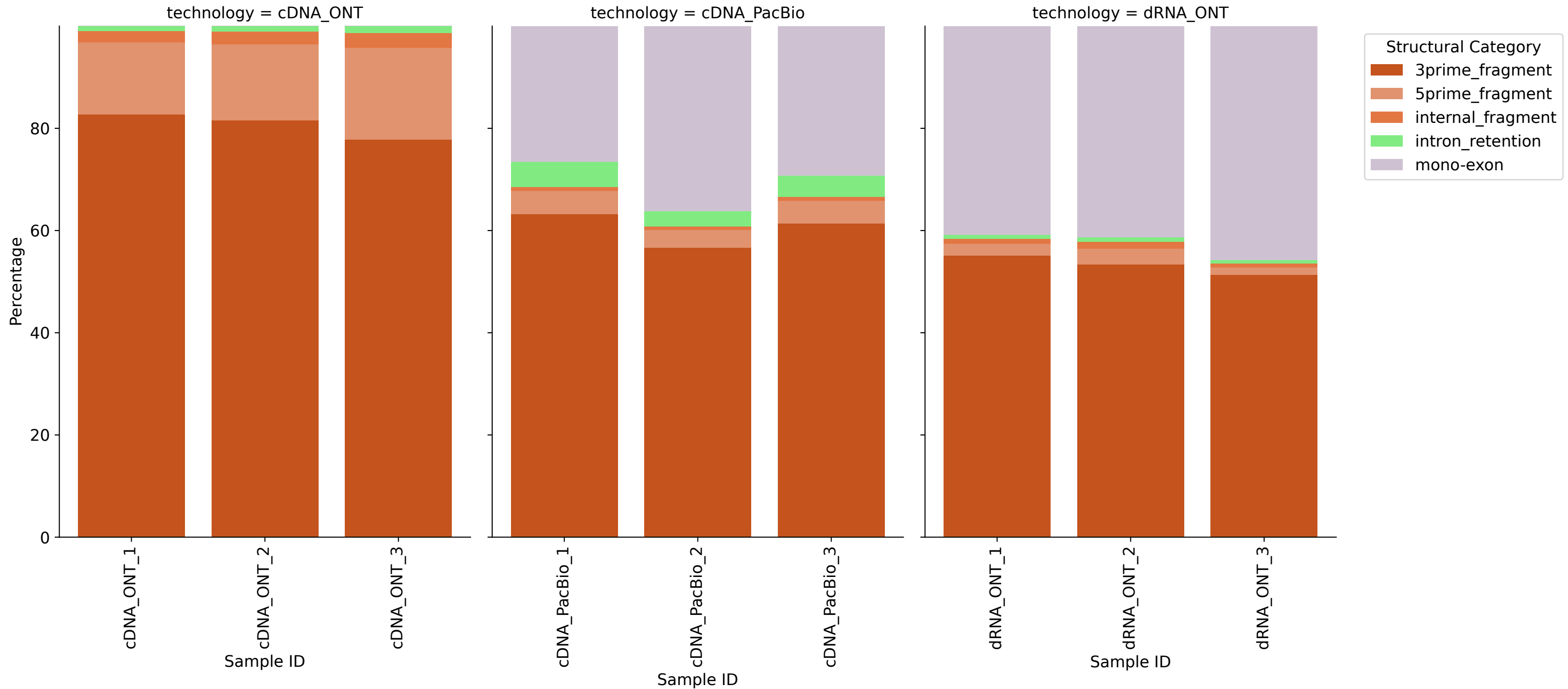
Percentage of FSM Reads in Each Sub-Category



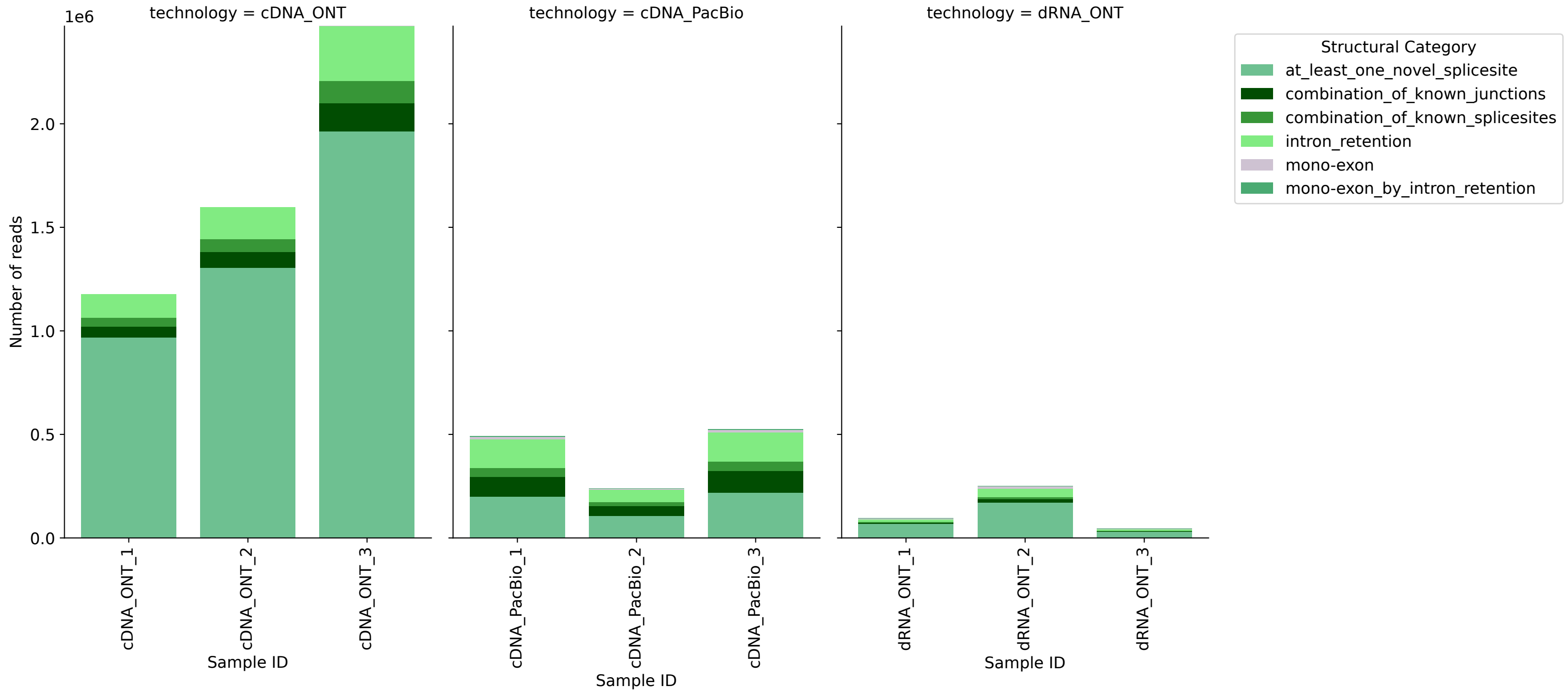
Number of Reads in Each Sub-Category - ISM



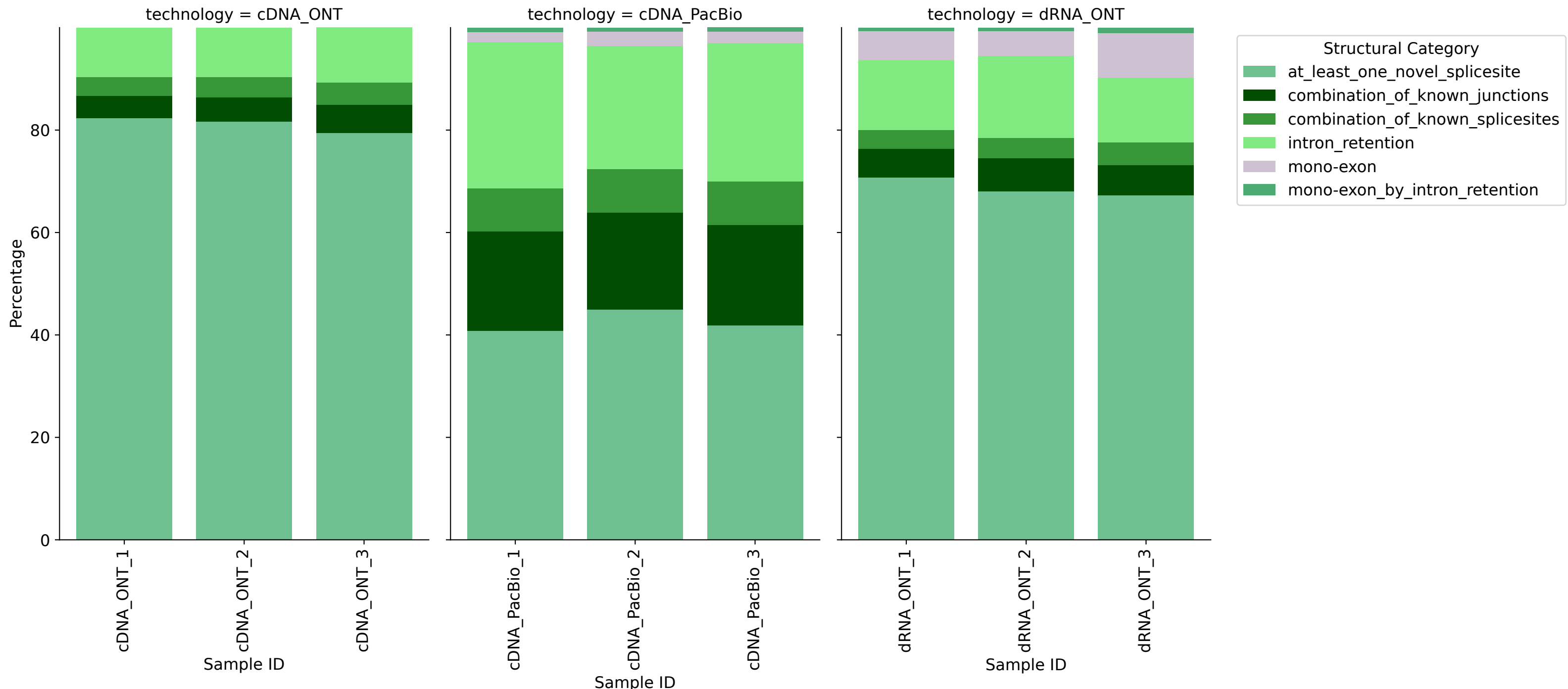
Percentage of ISM Reads in Each Sub-Category



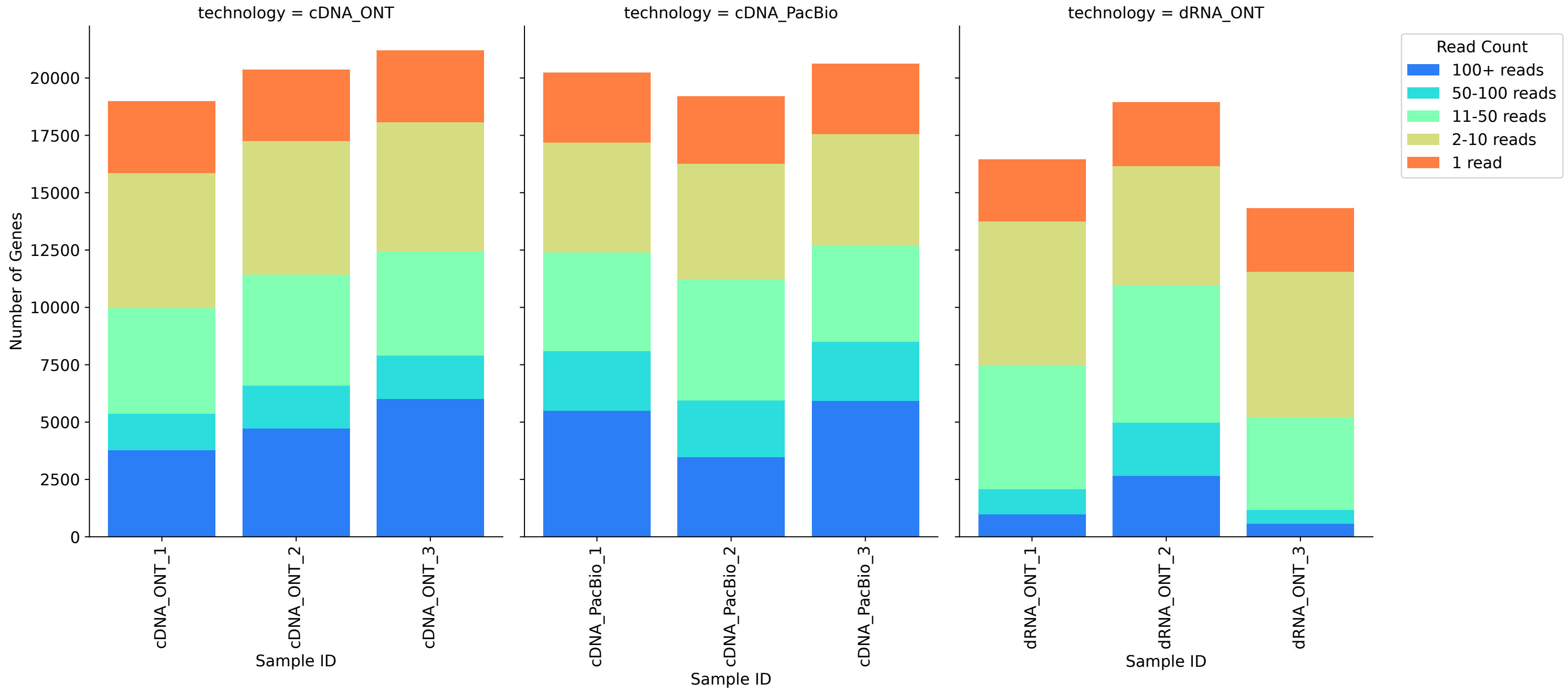
Number of Reads in Each Sub-Category - NIC and NNC



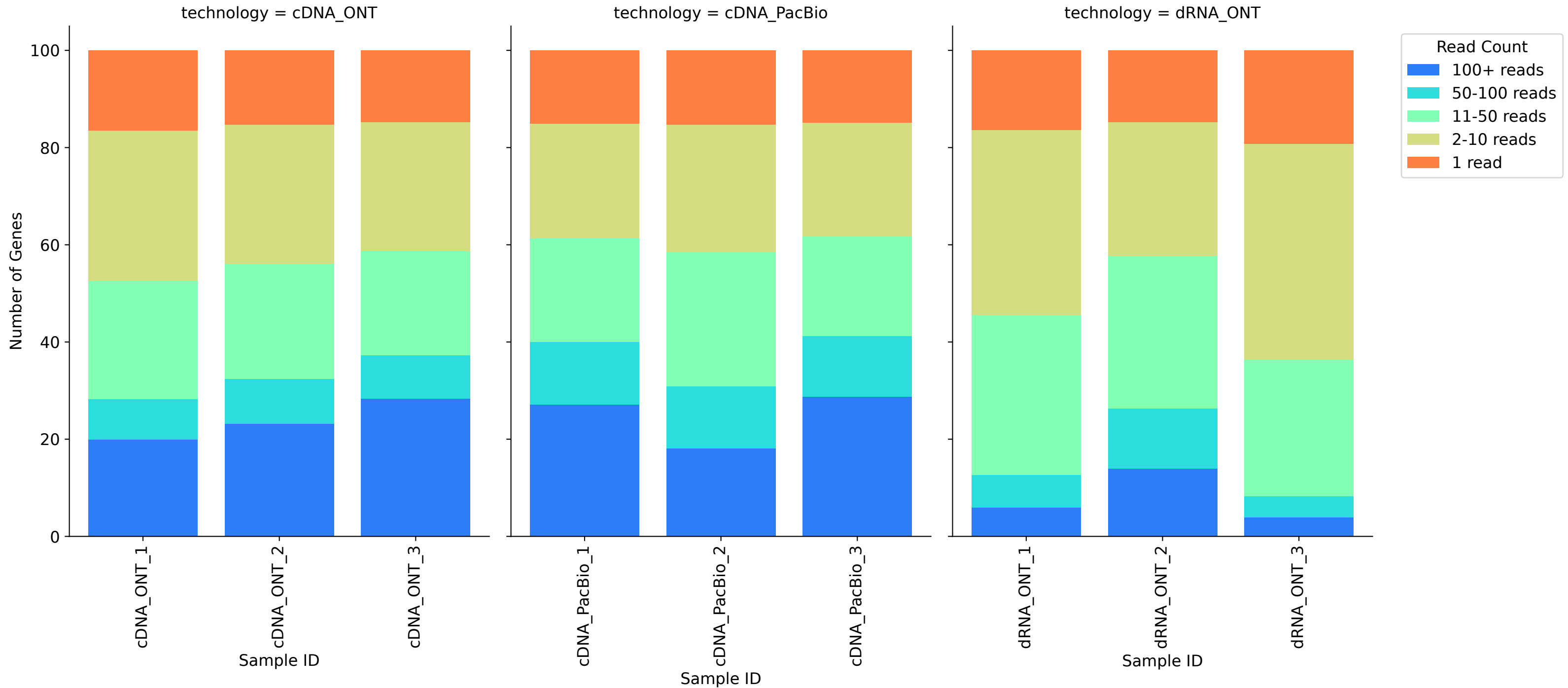
Percentage of NIC/NNC Reads in Each Sub-Category



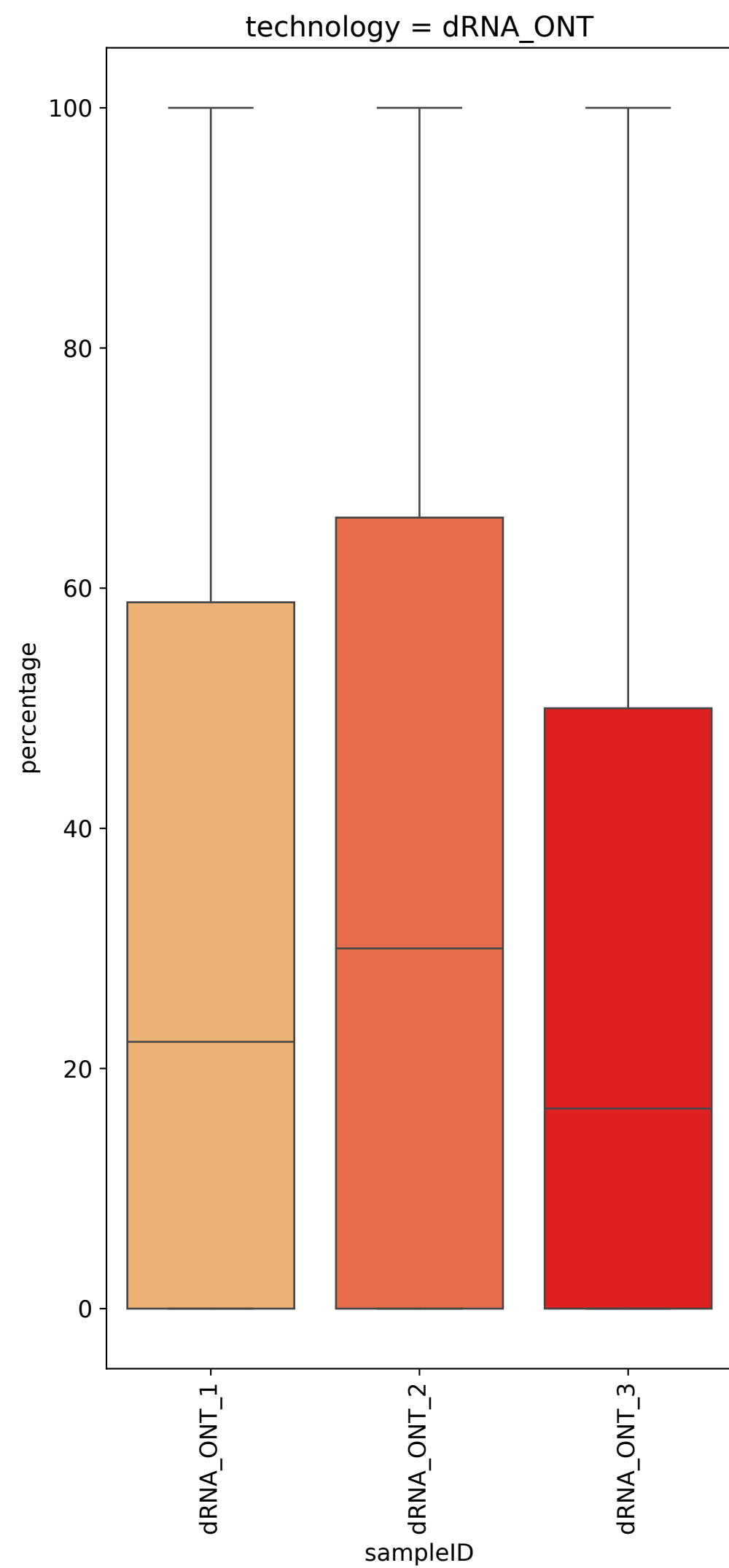
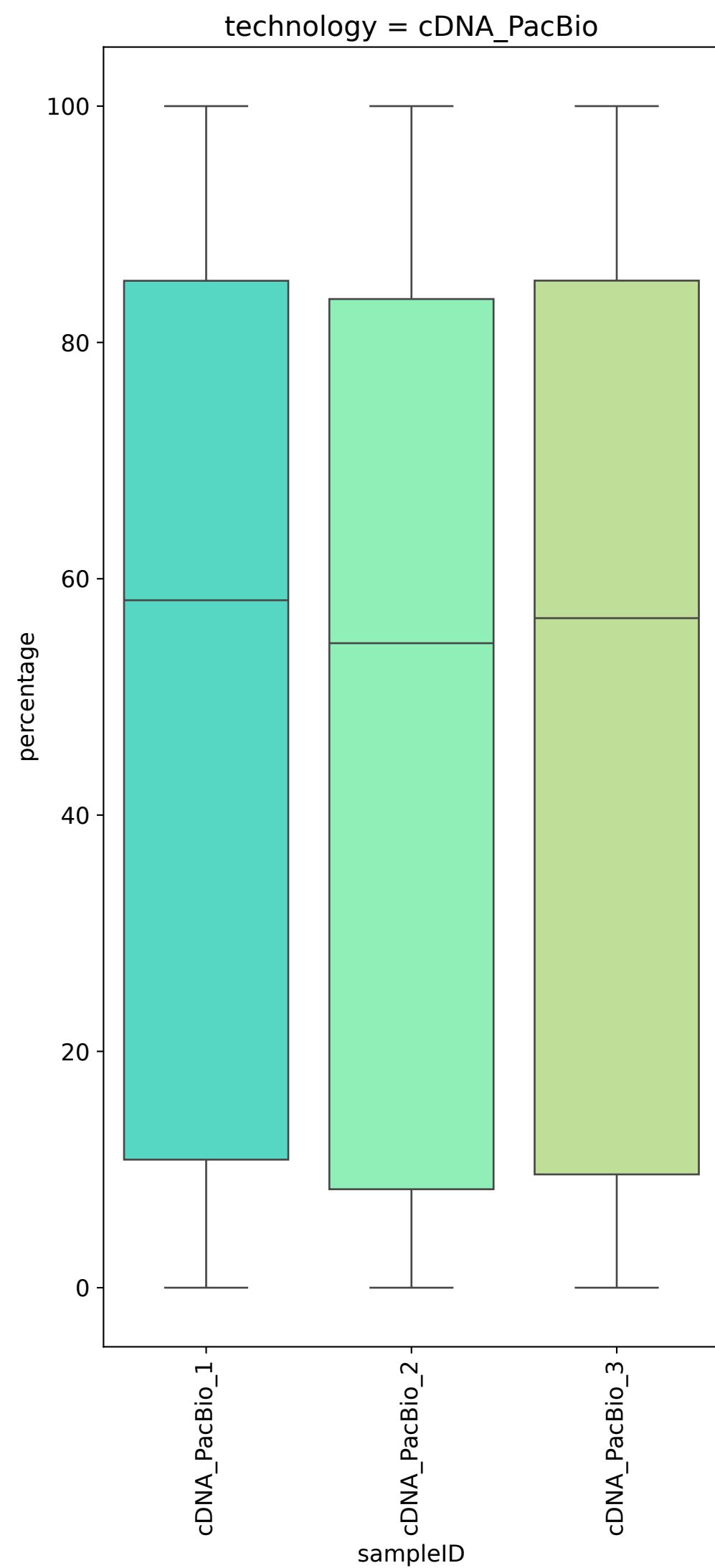
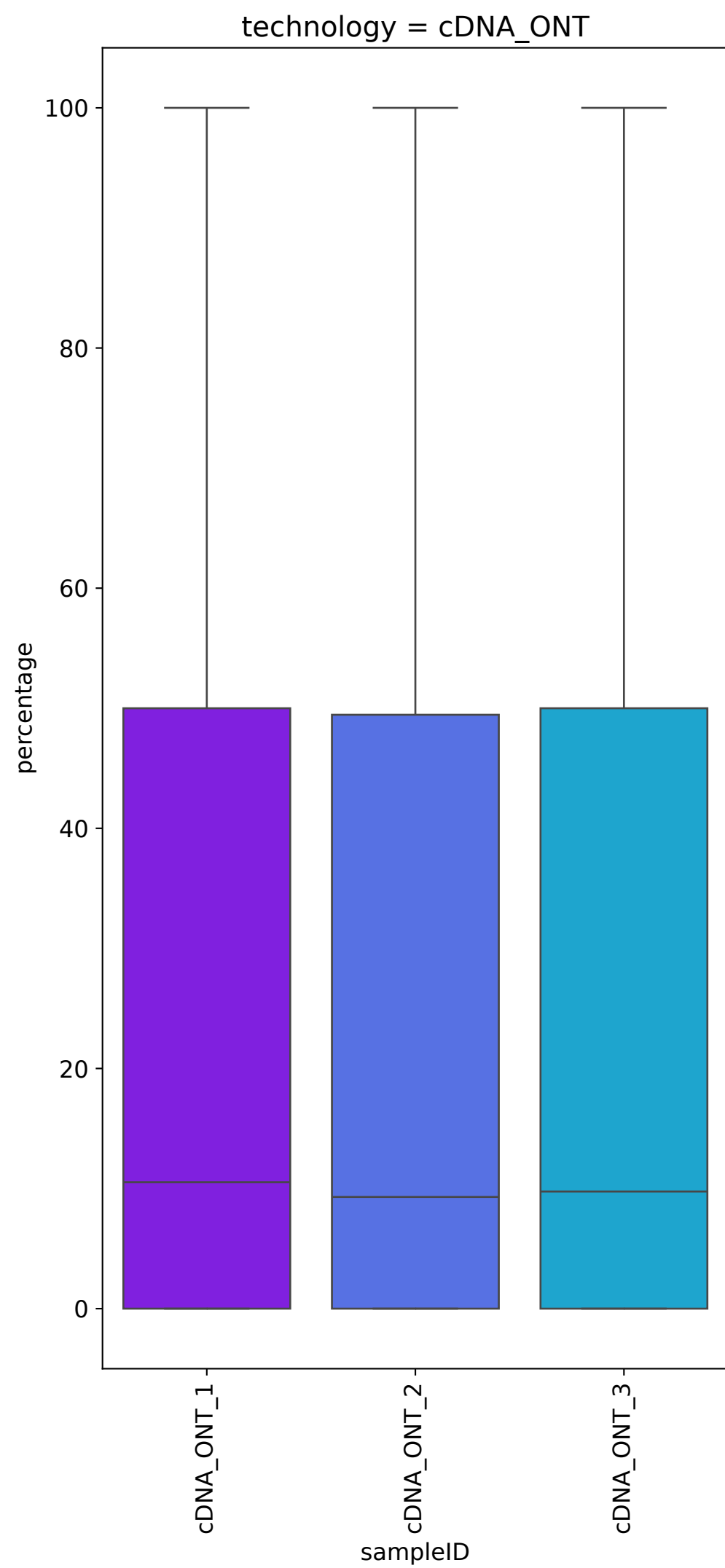
Number of Genes Detected



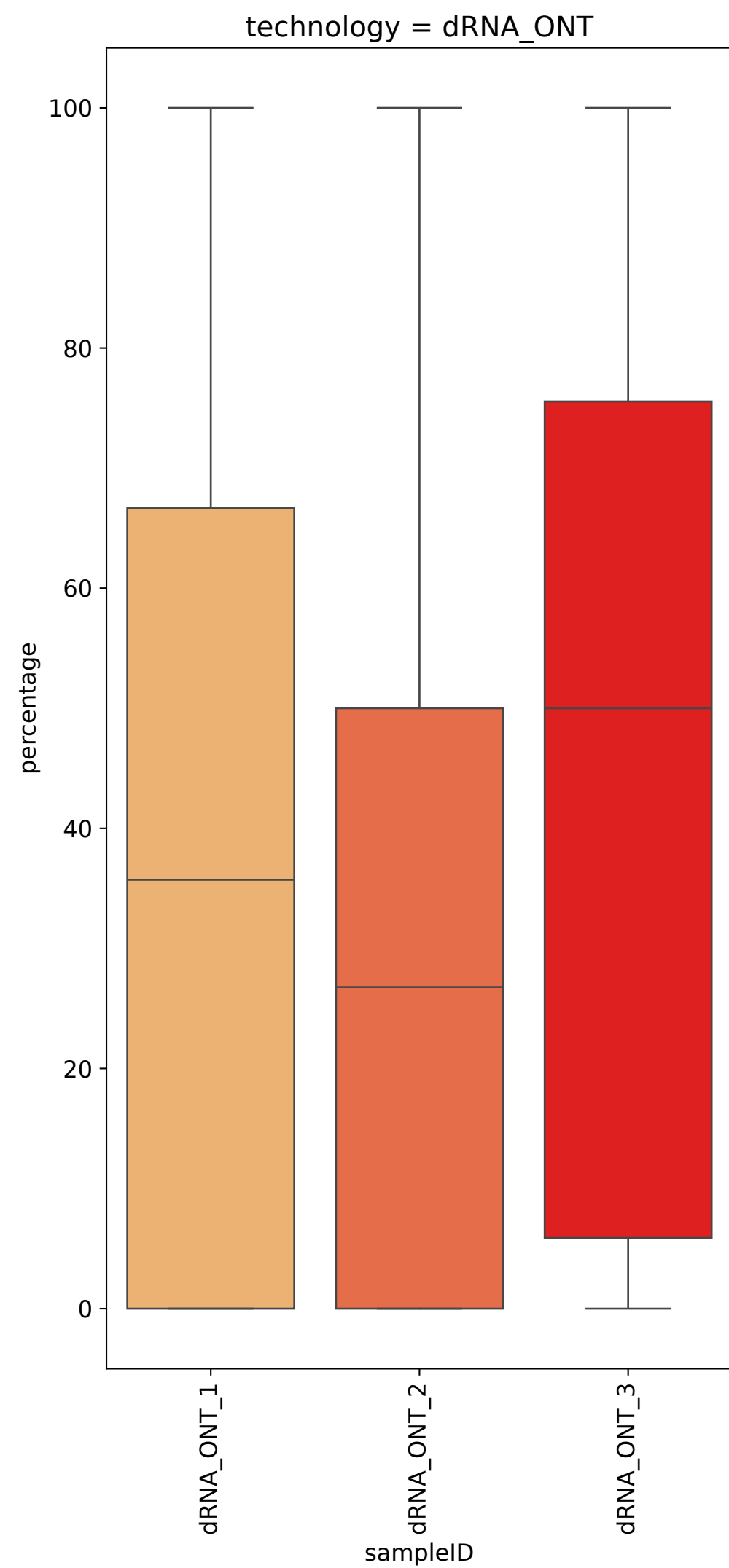
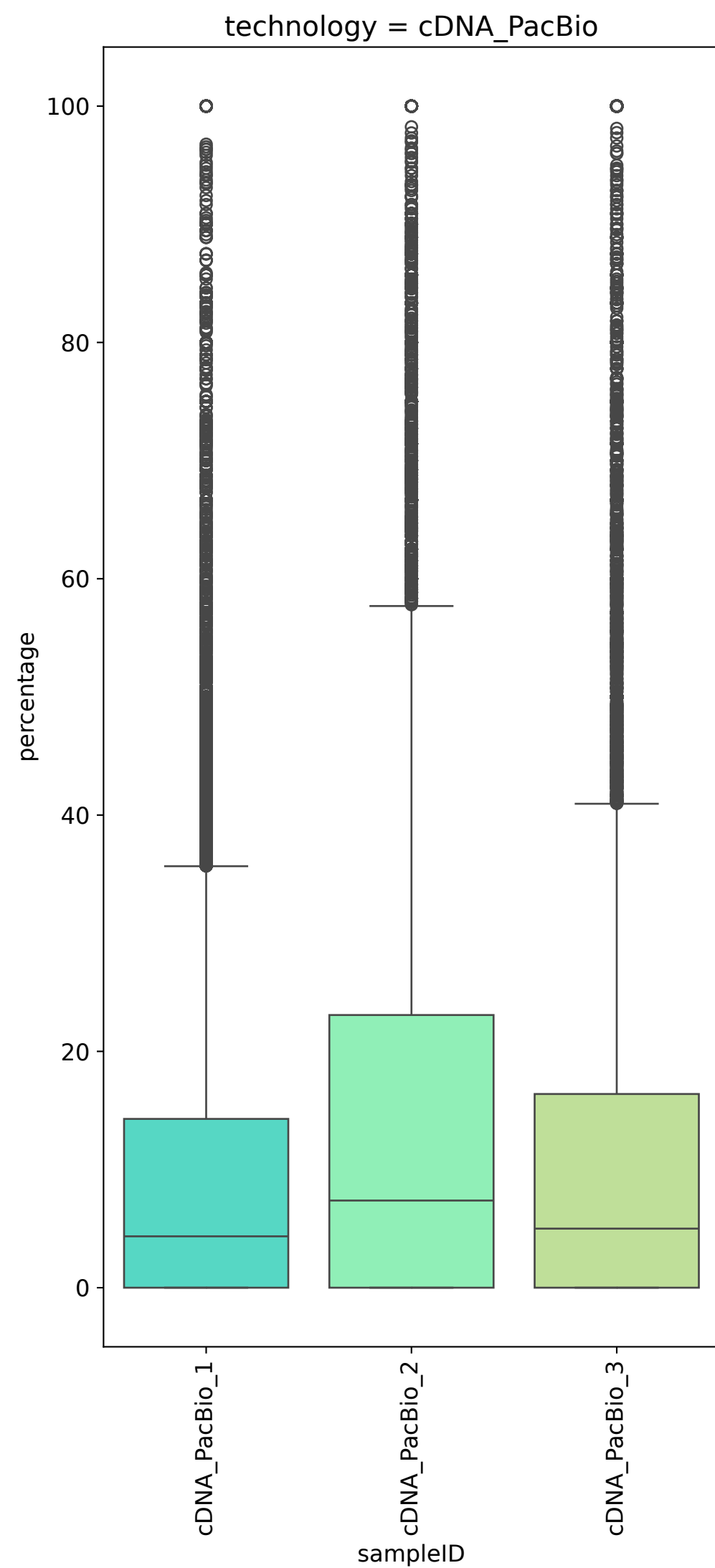
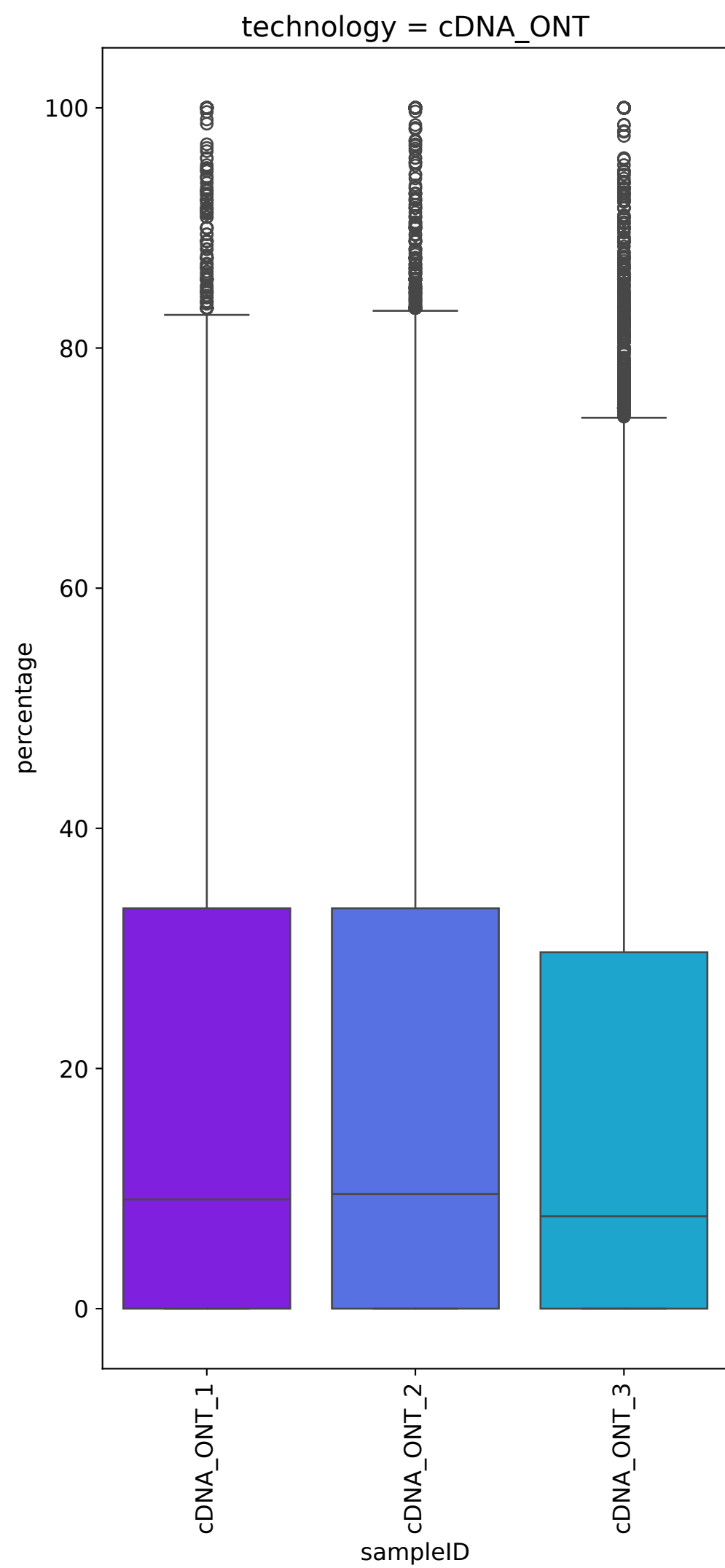
Number of Genes Detected



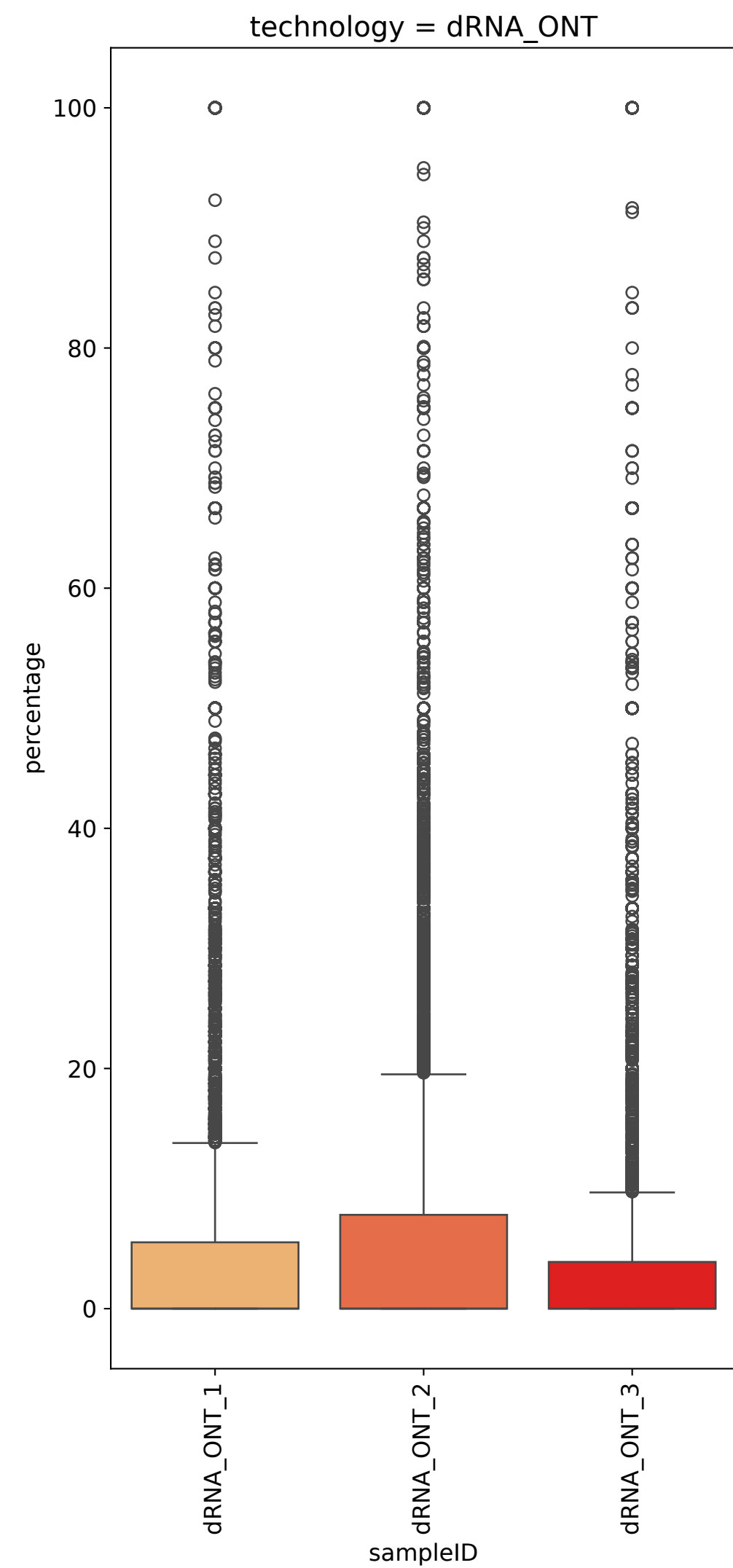
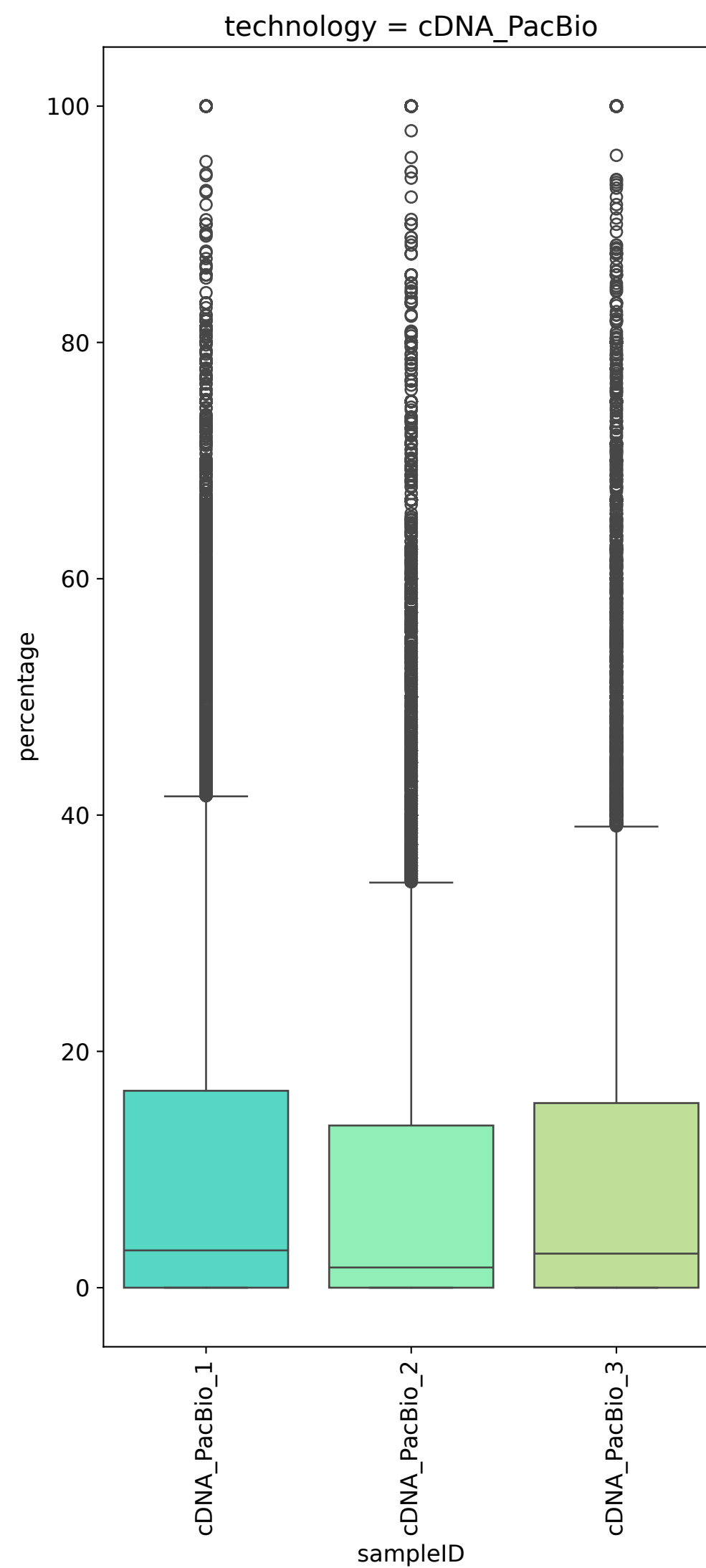
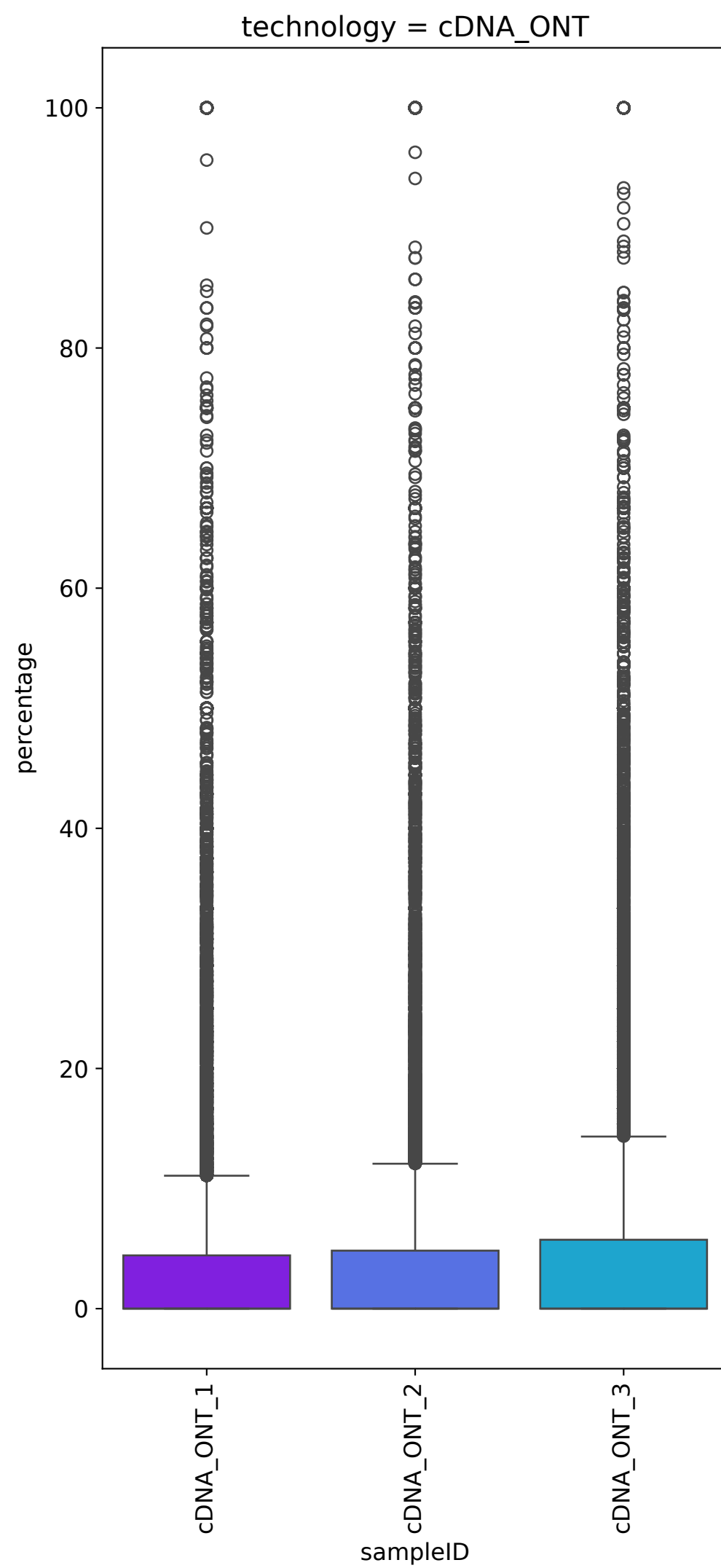
Gene distribution - percent_full-splice_match



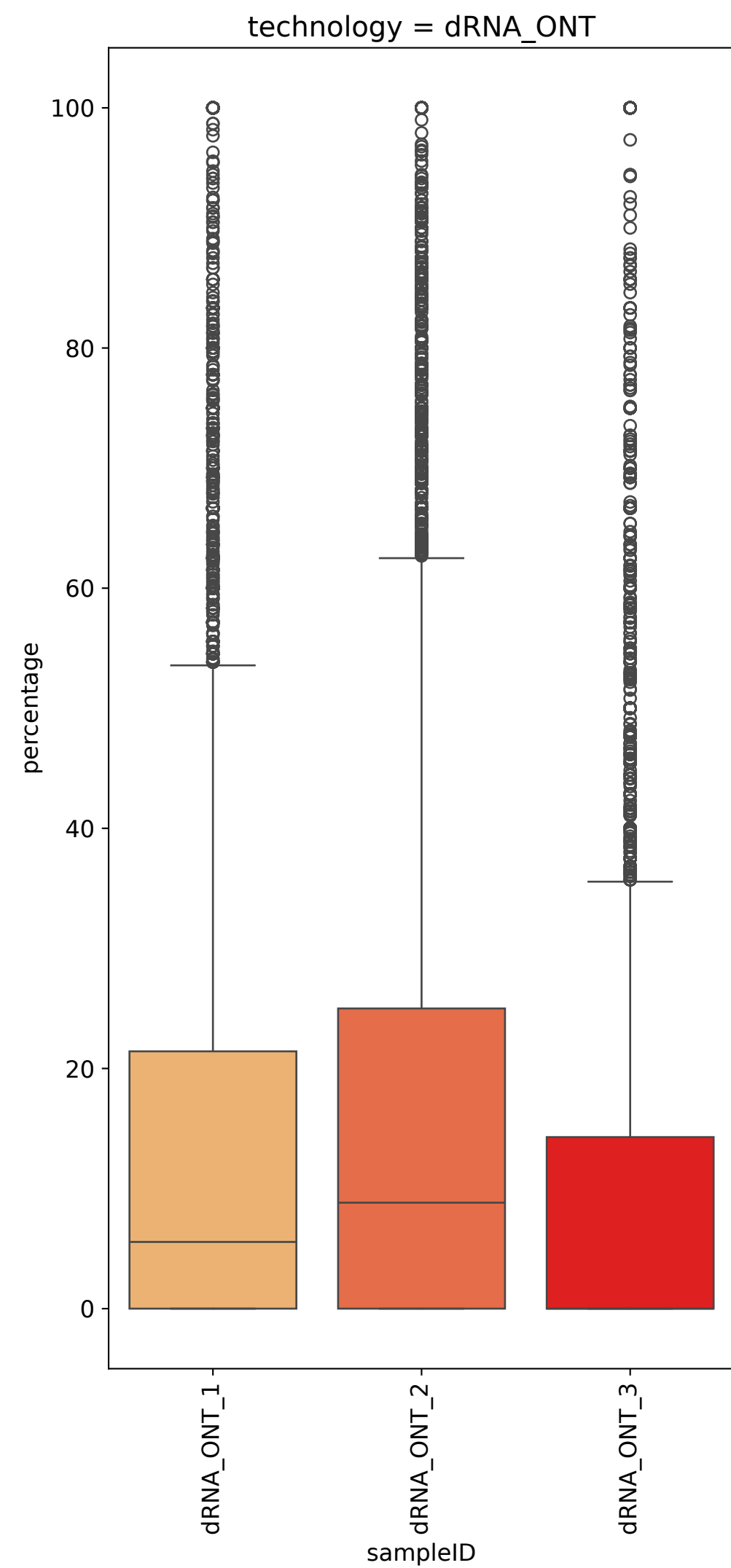
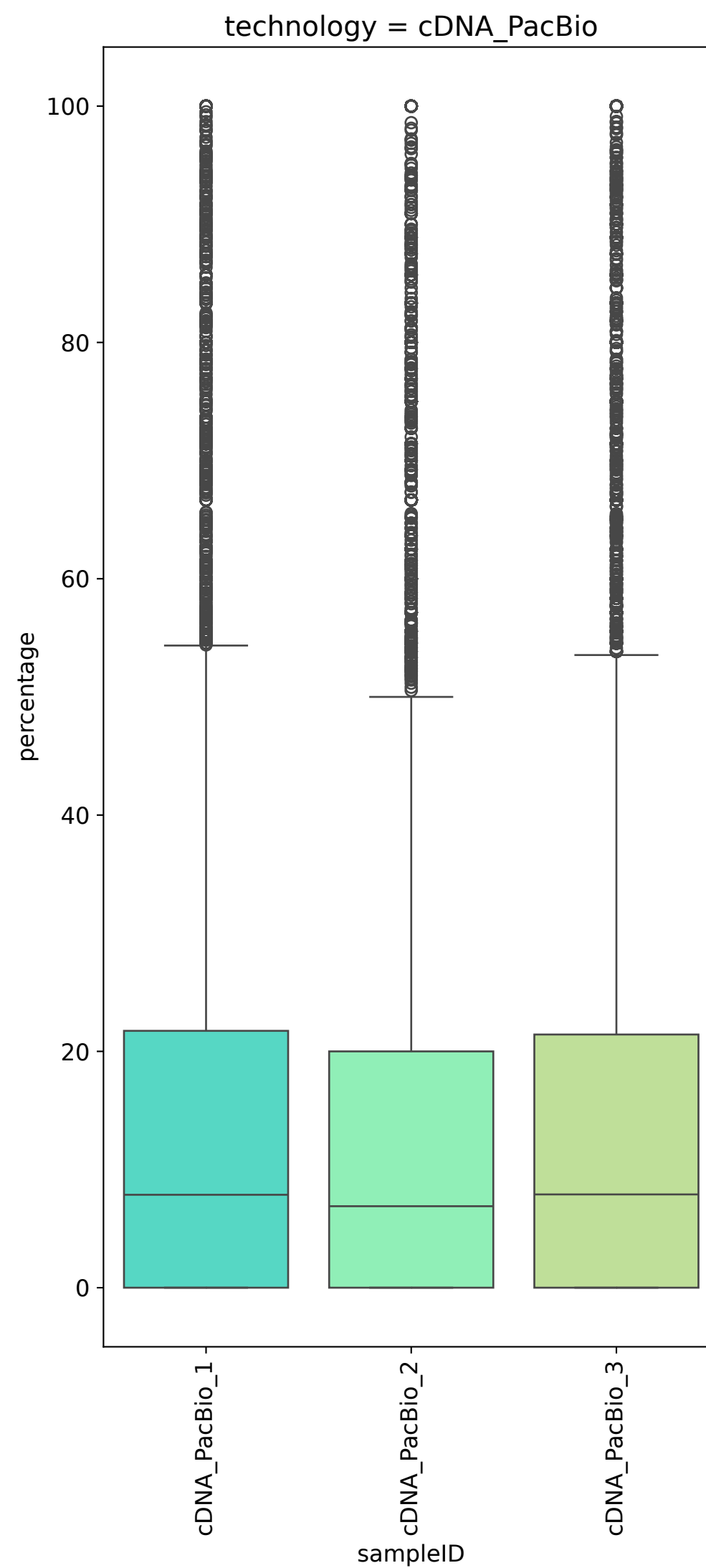
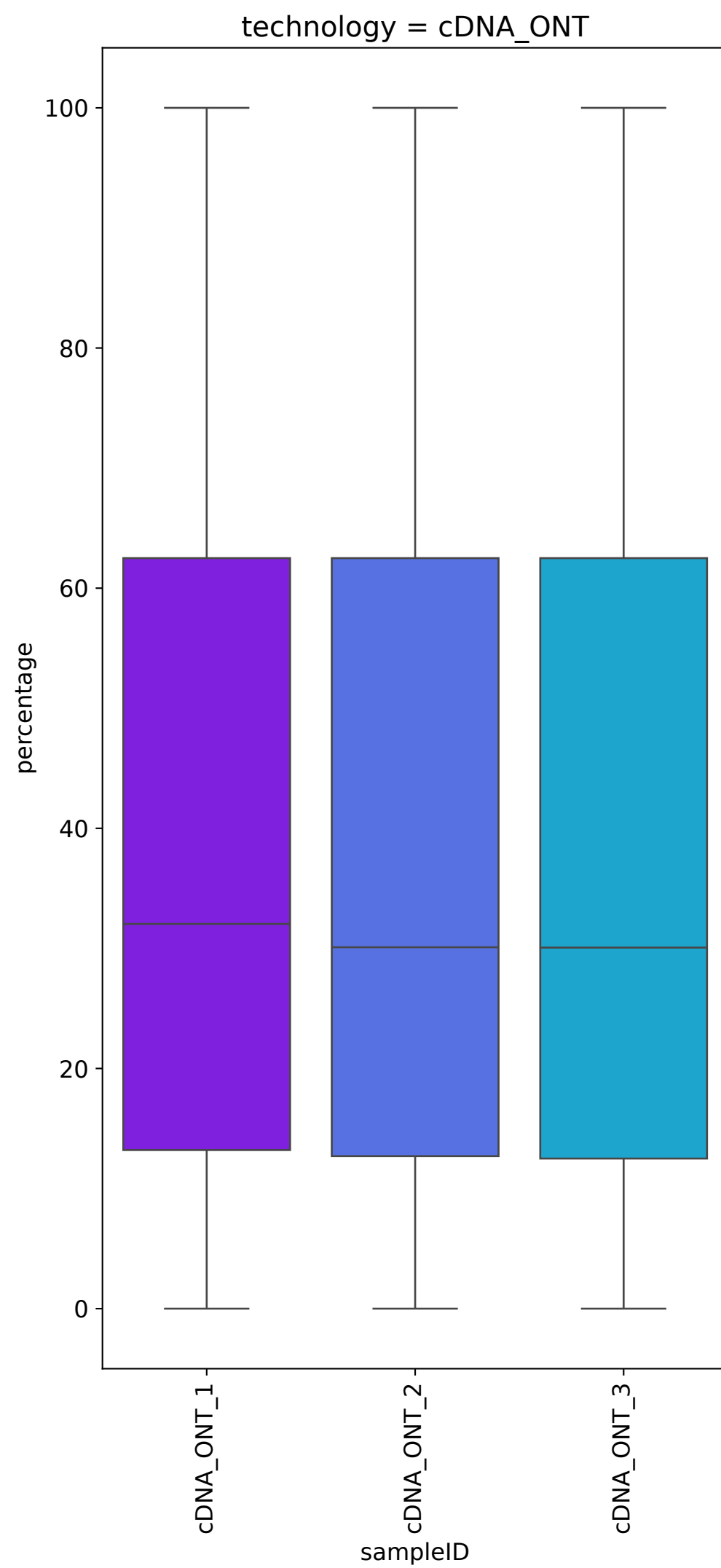
Gene distribution - percent_incomplete-splice_match



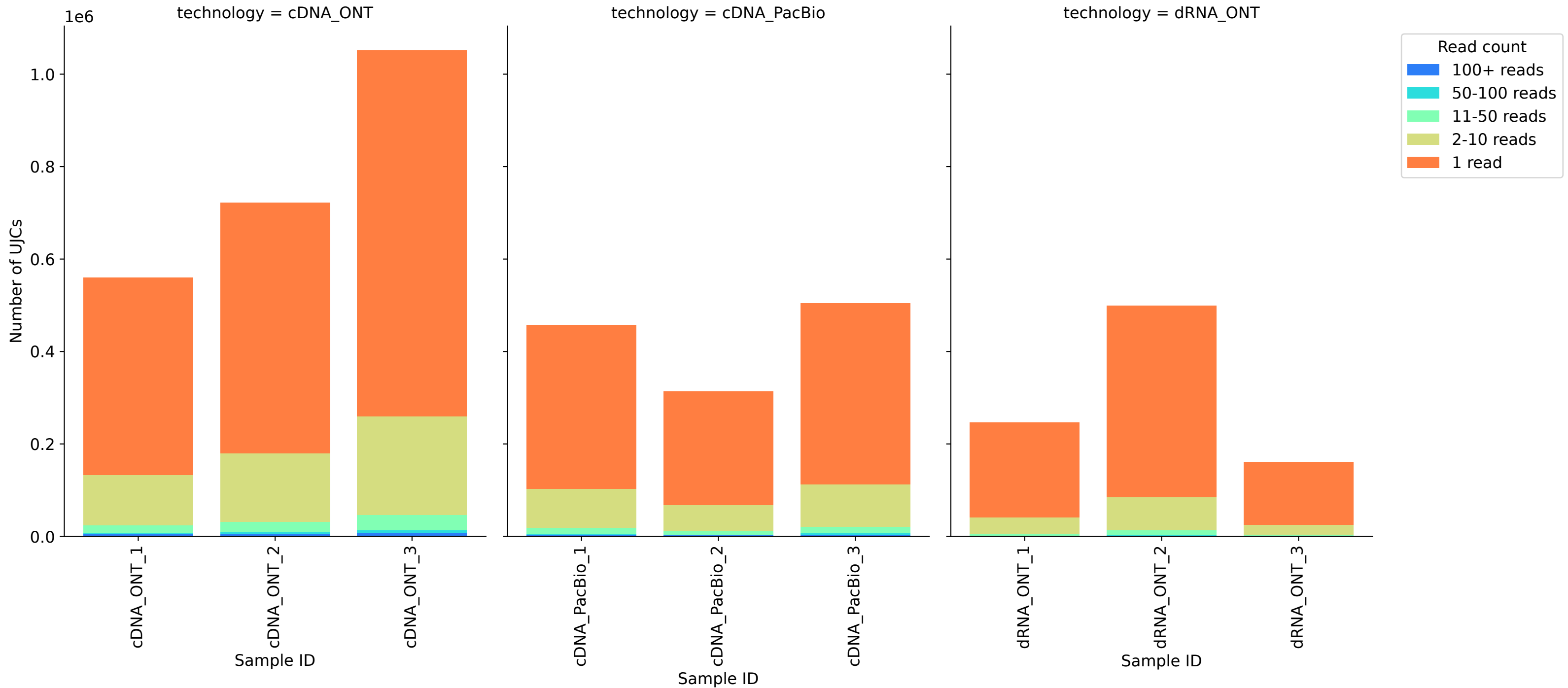
Gene distribution - percent_novel_in_catalog



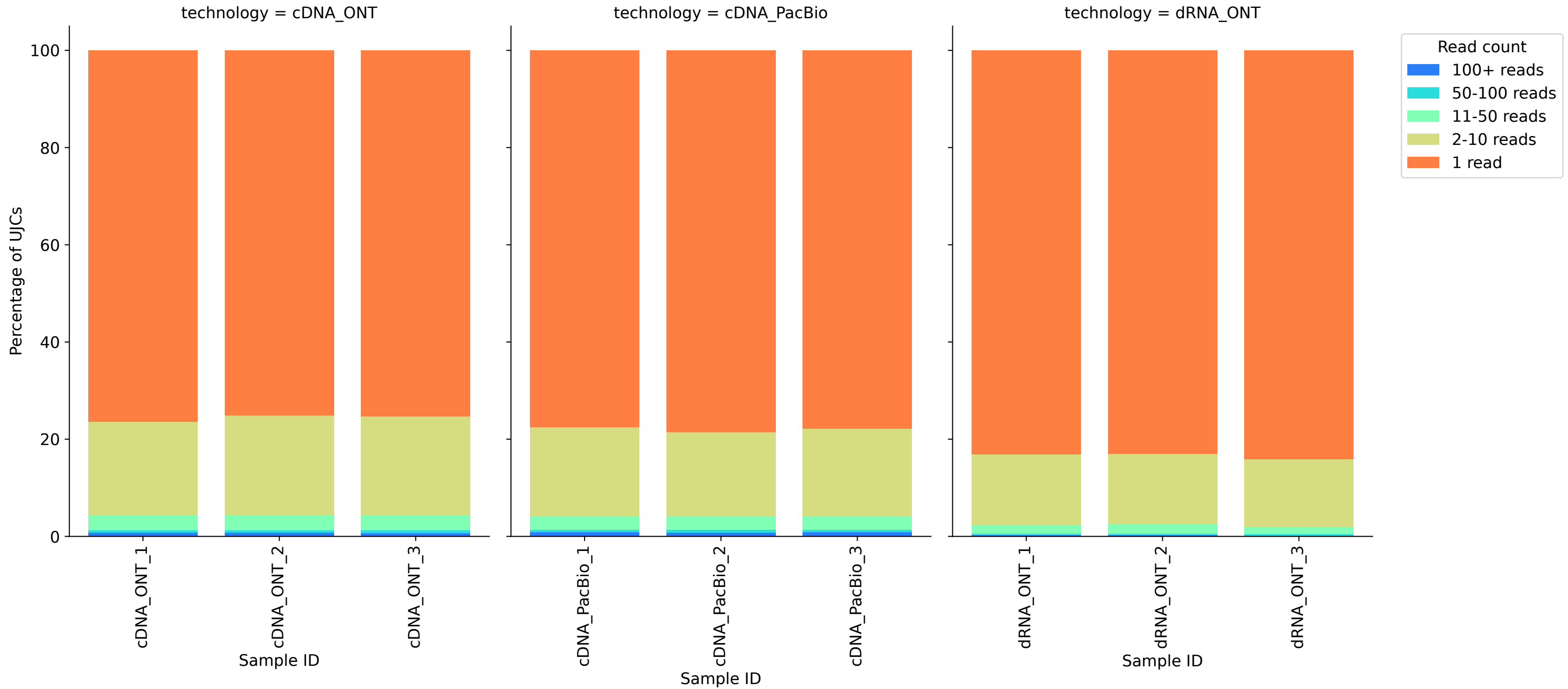
Gene distribution - percent_novel_not_in_catalog



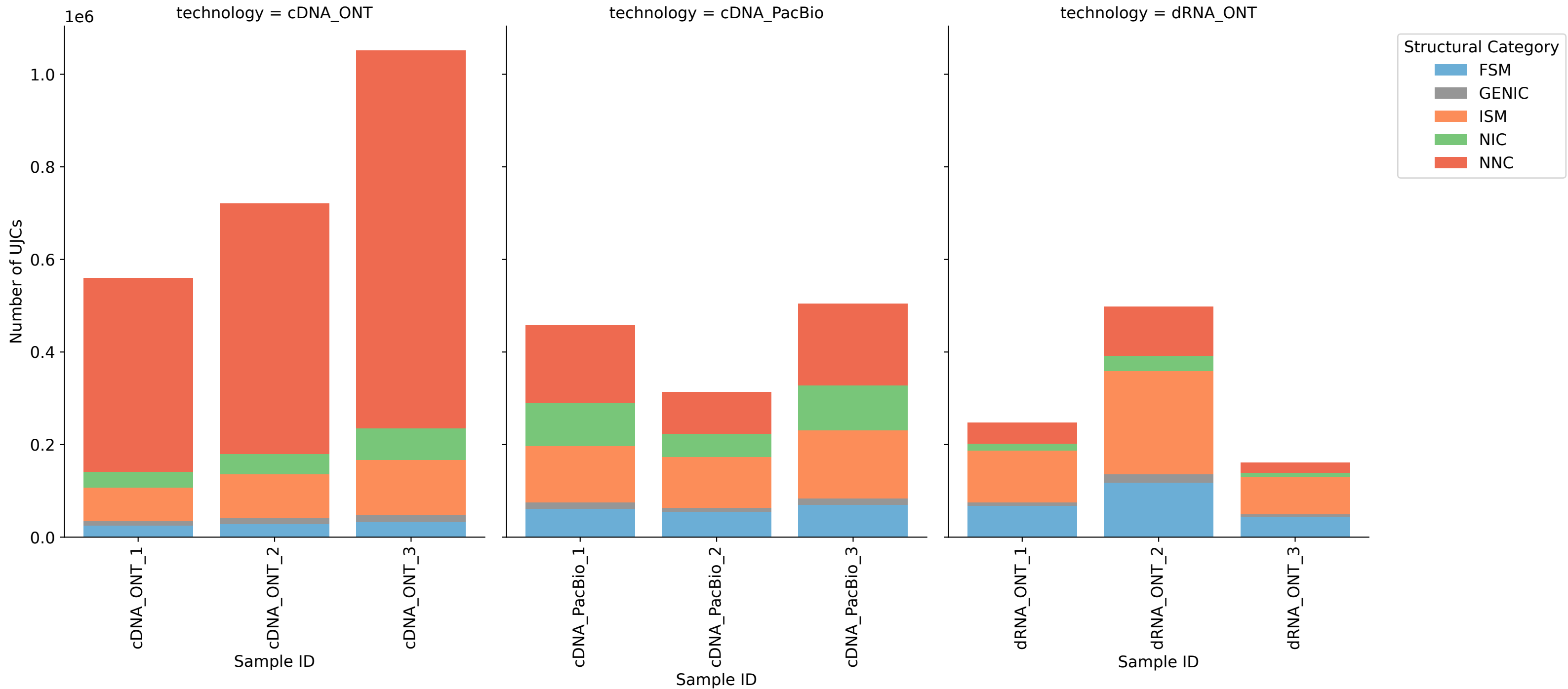
Number of UJCs Detected



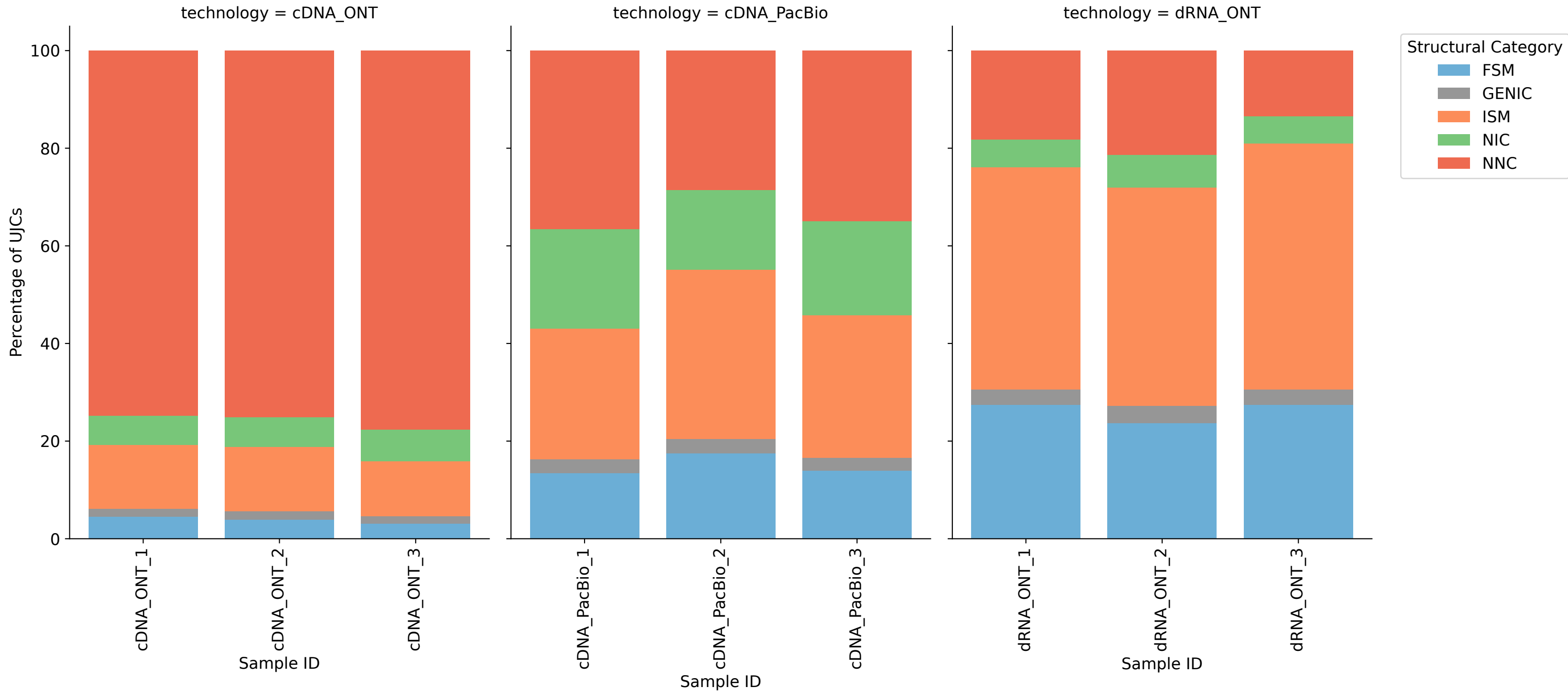
UJCs Detected



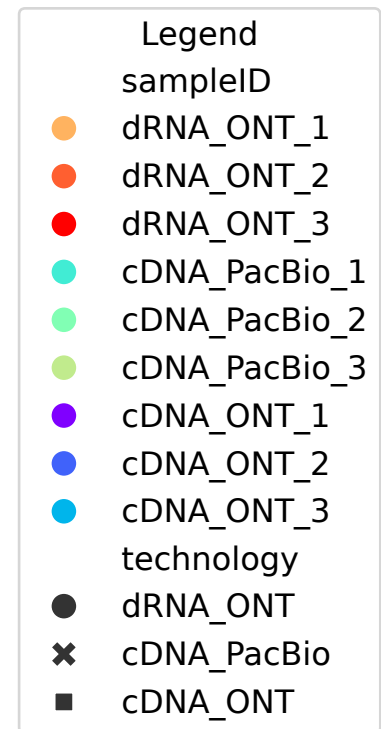
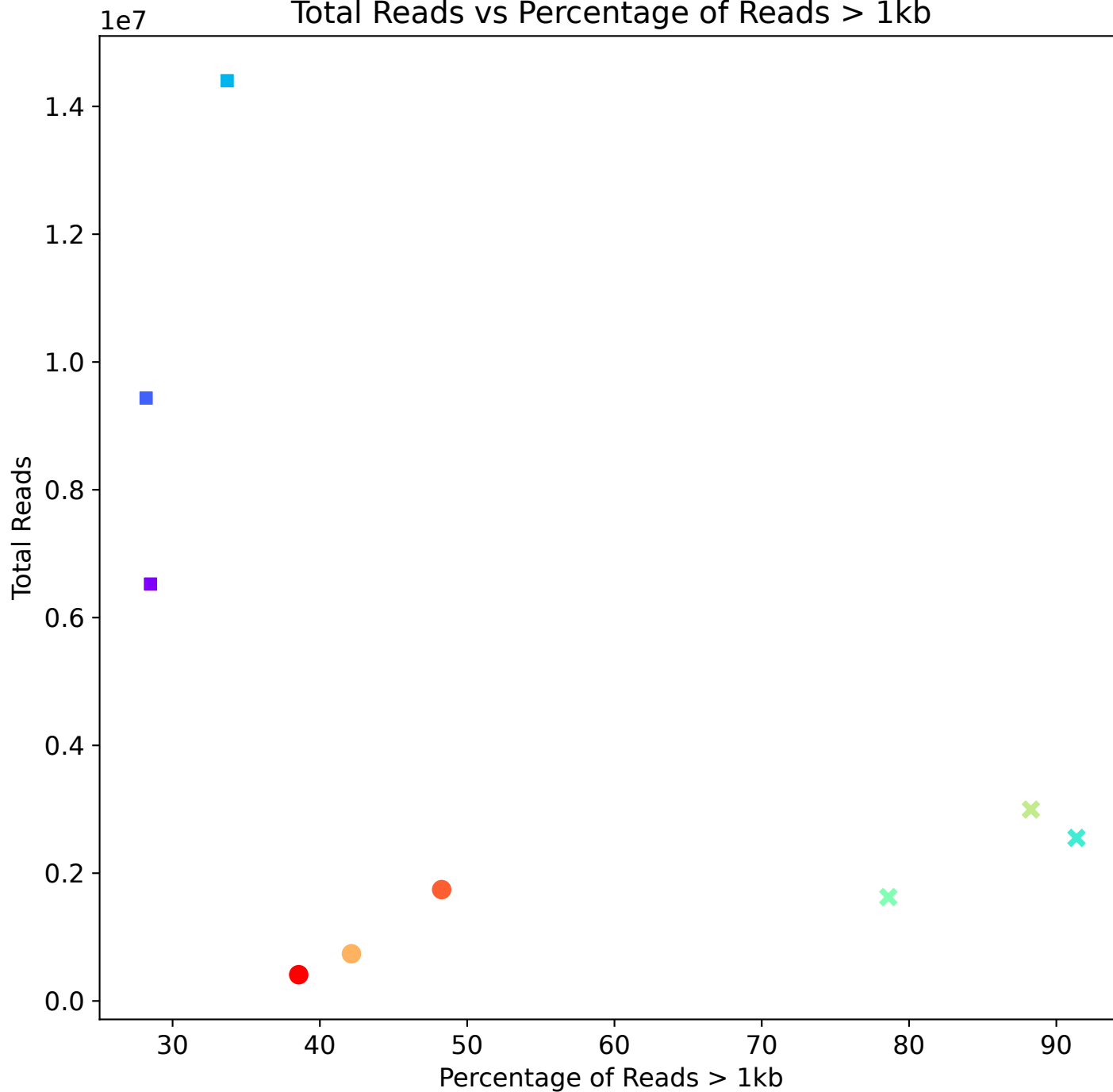
Number of UJCs detected



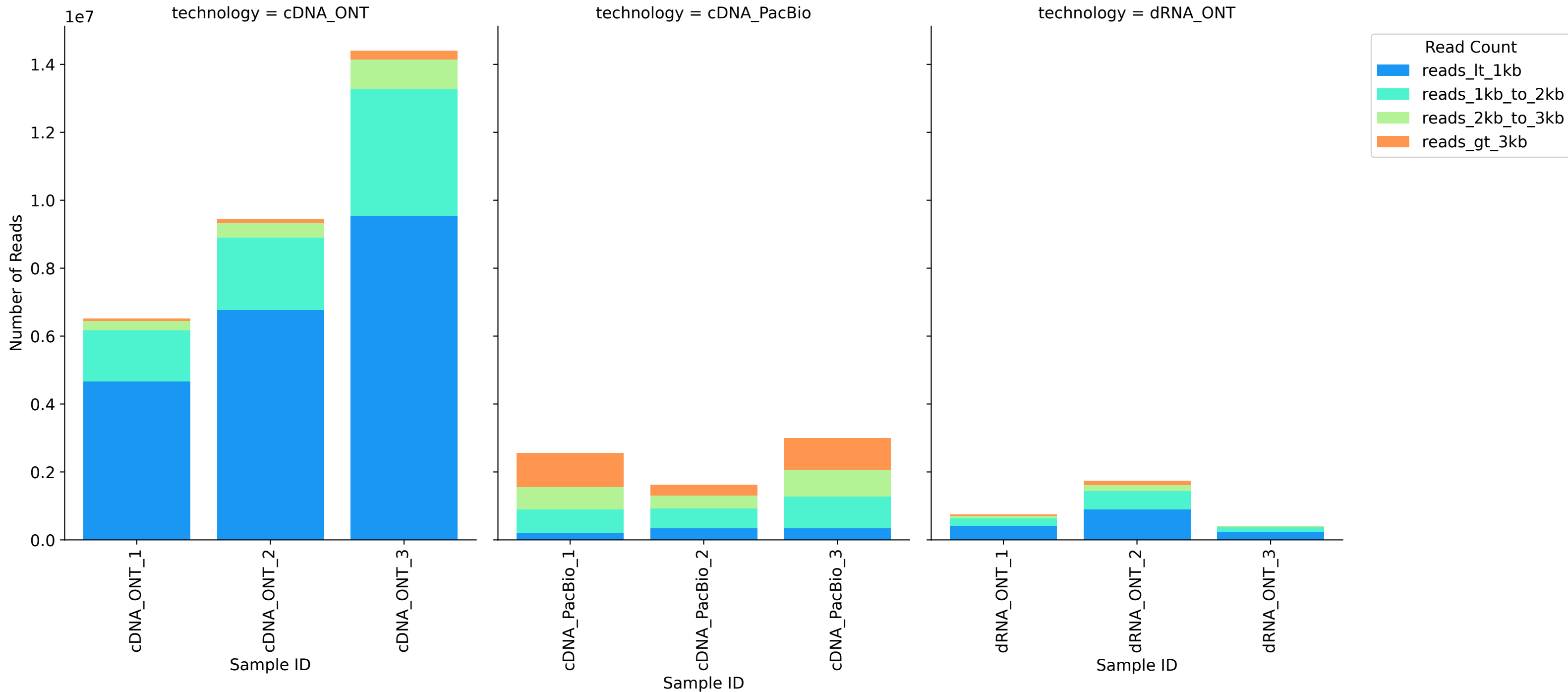
UJCs detected



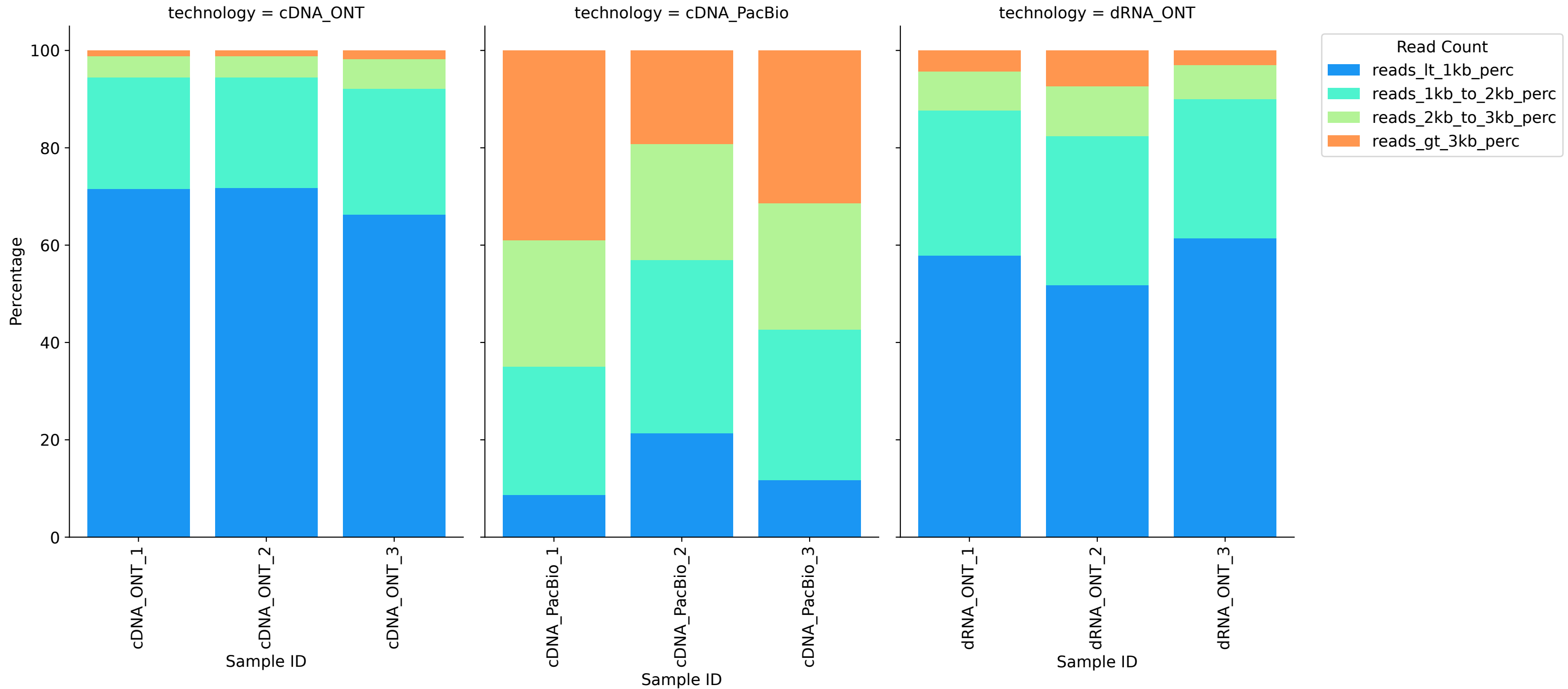
Total Reads vs Percentage of Reads > 1kb



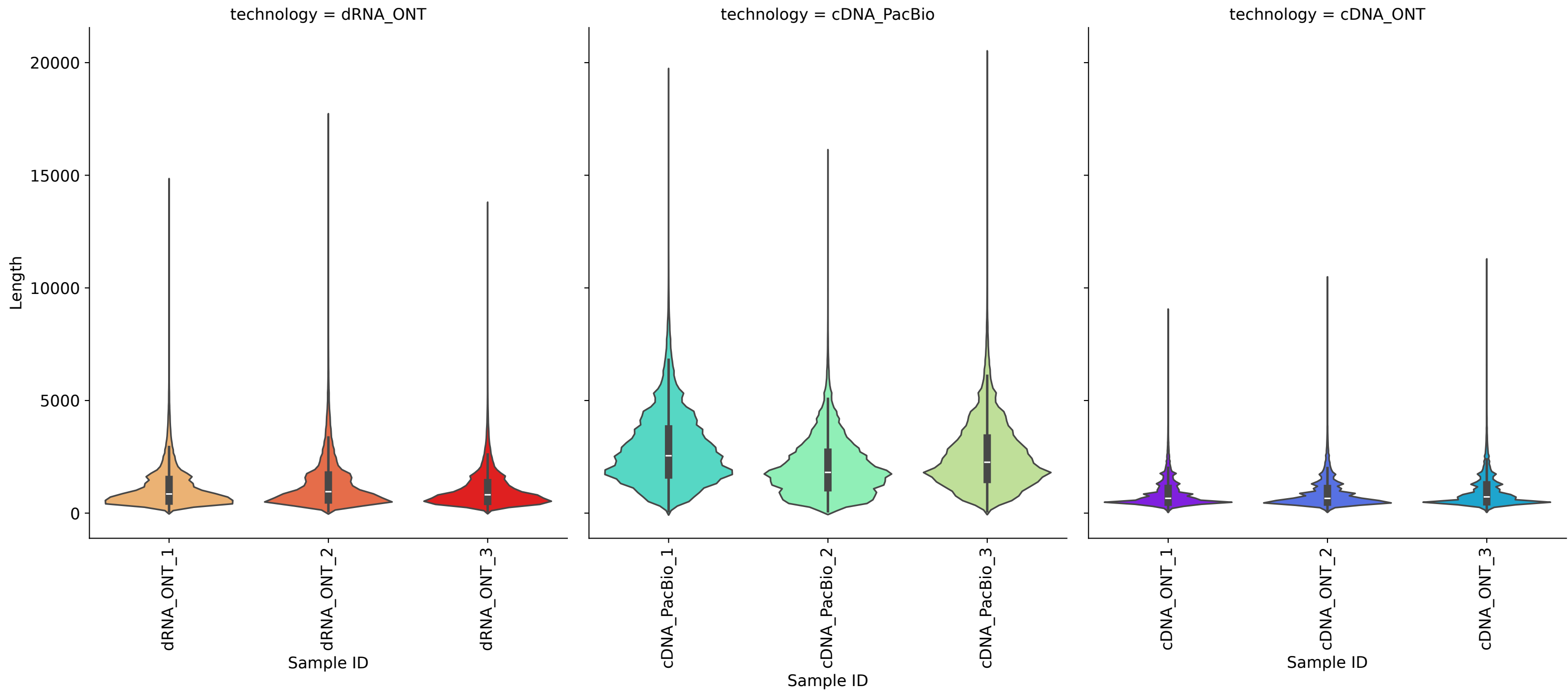
Lengths of All Mapped Reads



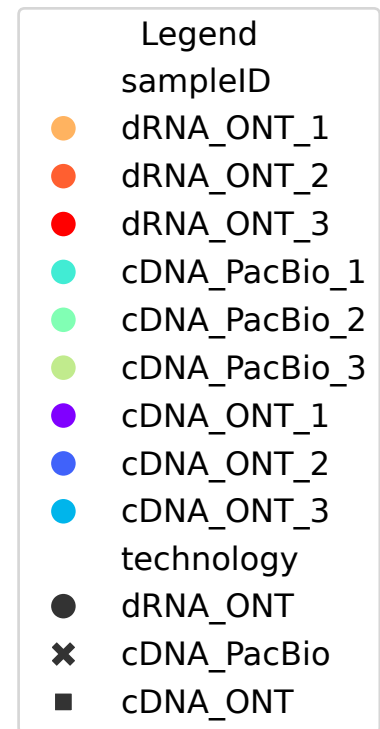
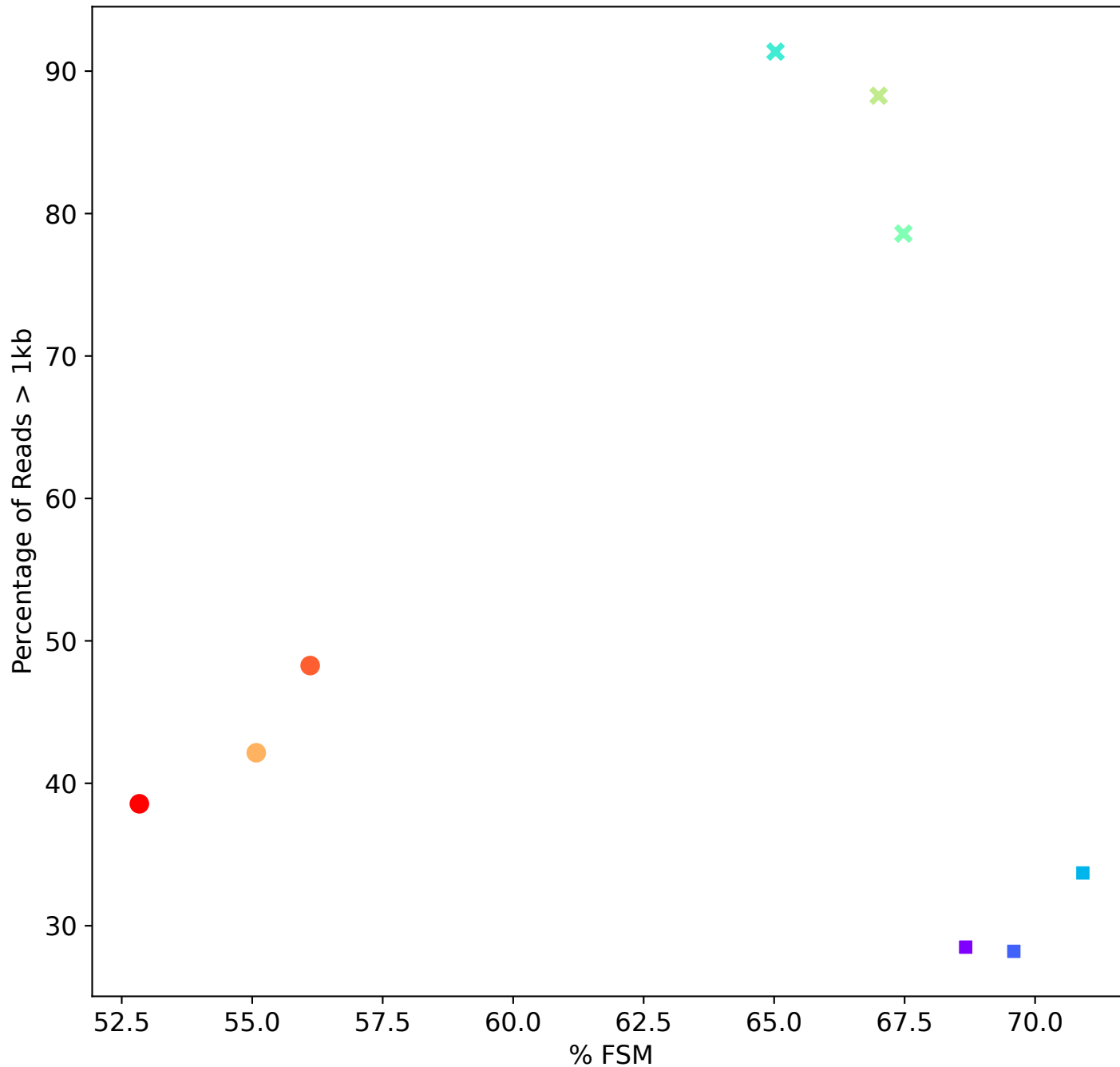
Lengths of All Mapped Reads



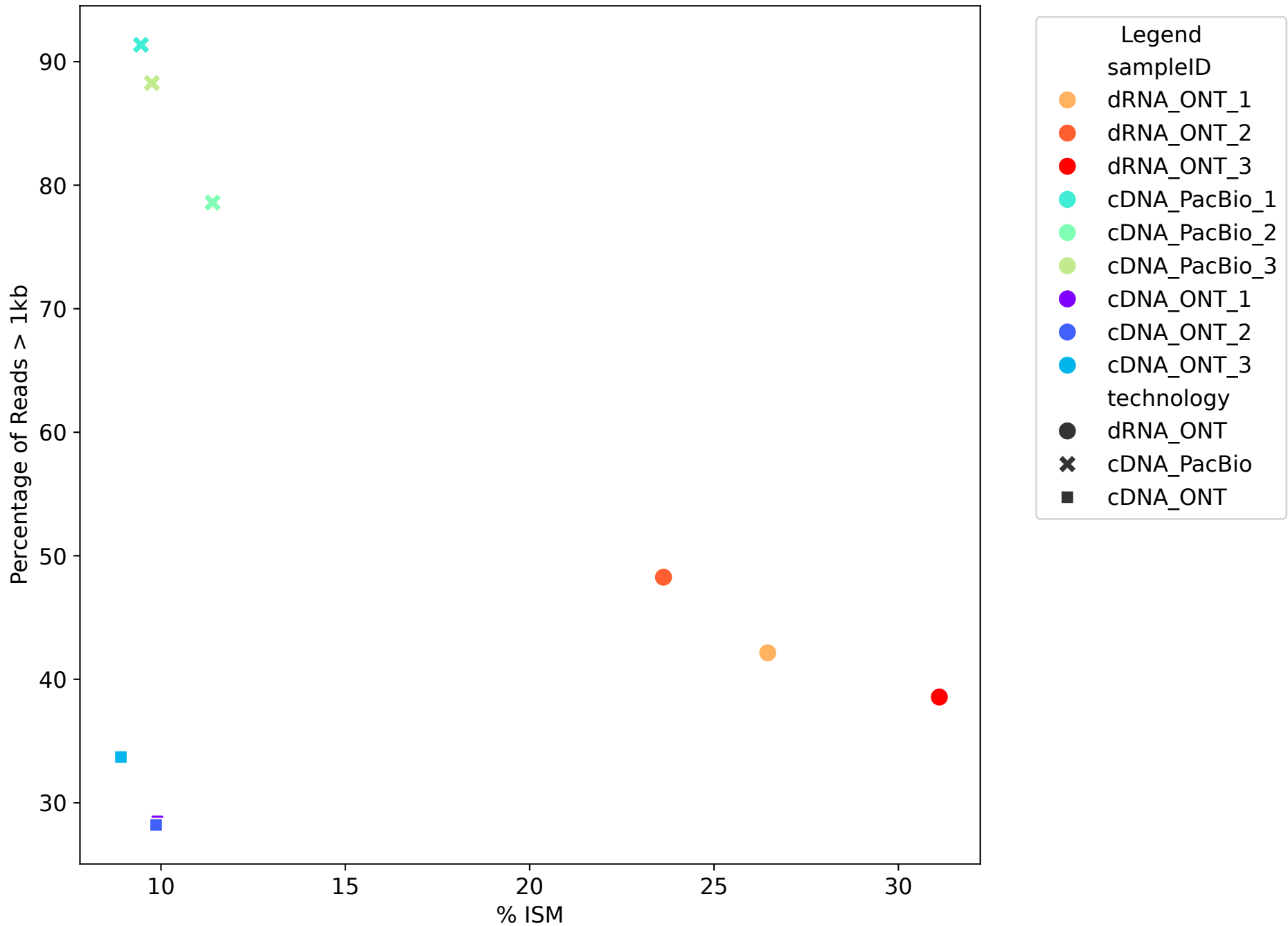
Read Length Distribution



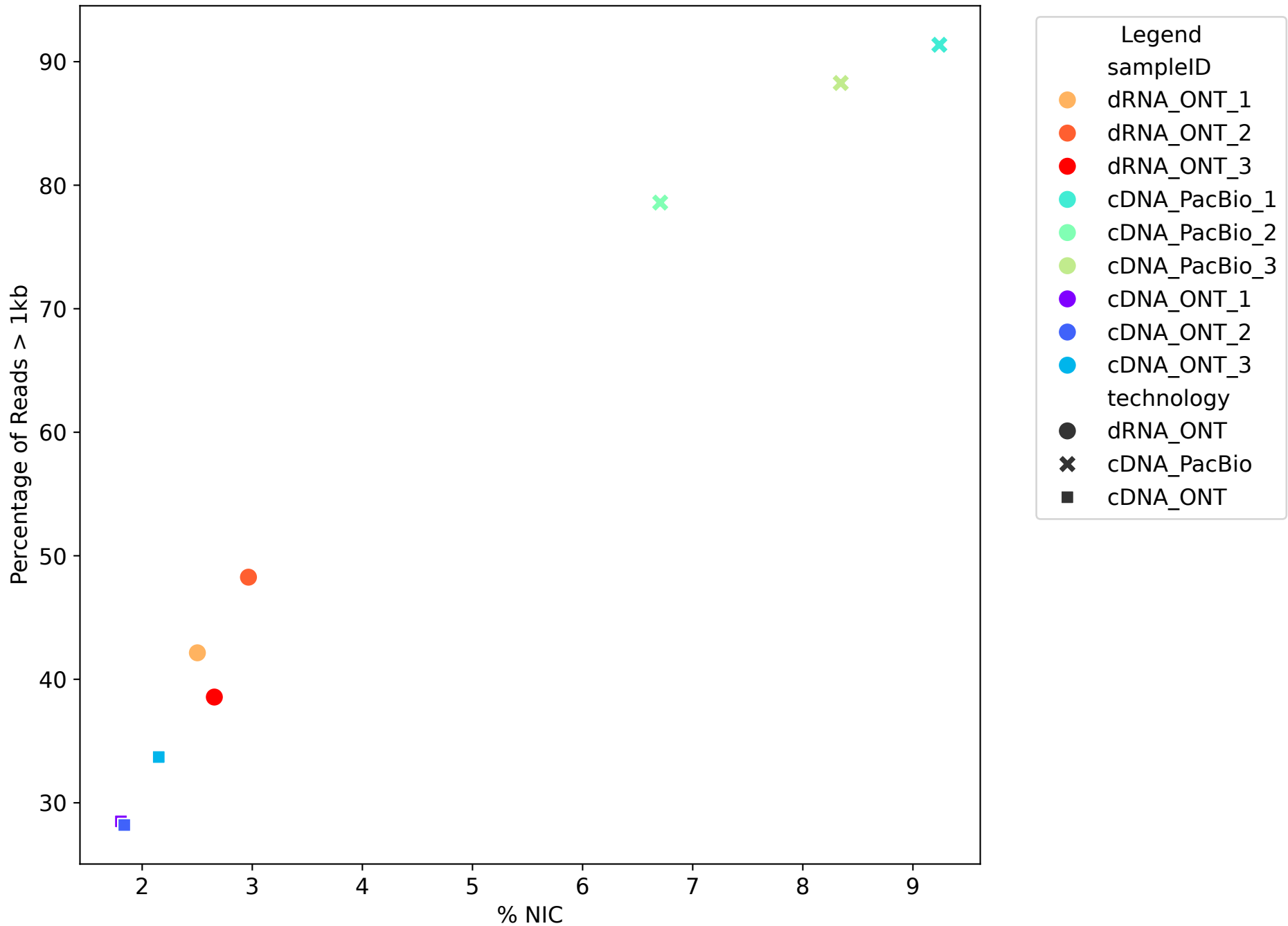
Percentage of Reads > 1kb vs %FSM



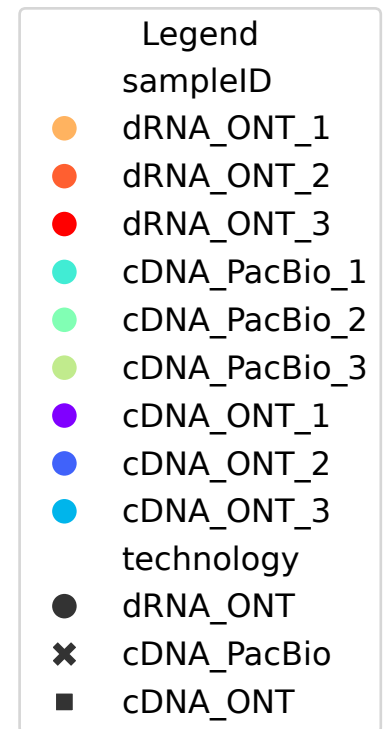
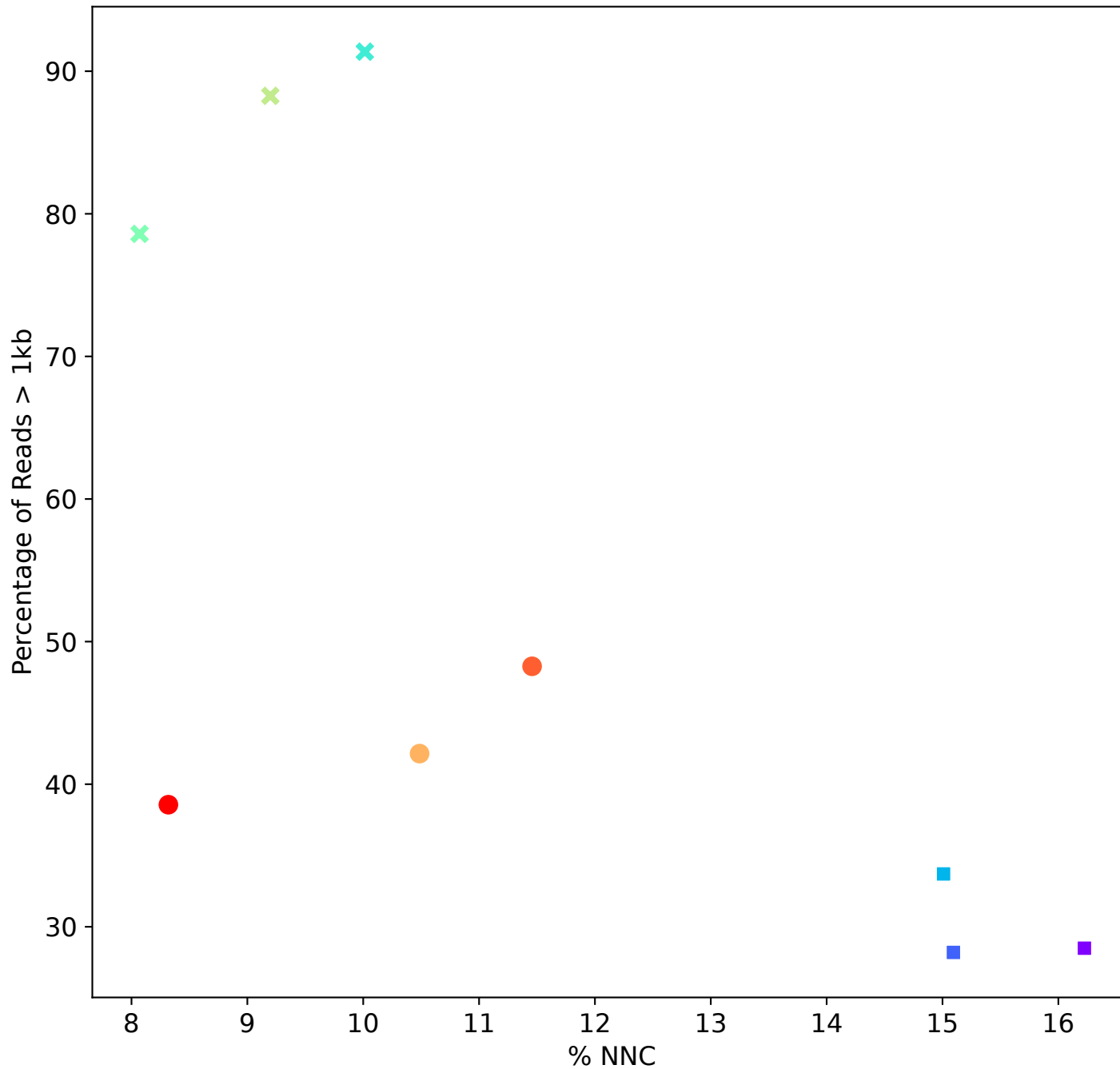
Percentage of Reads > 1kb vs %ISM



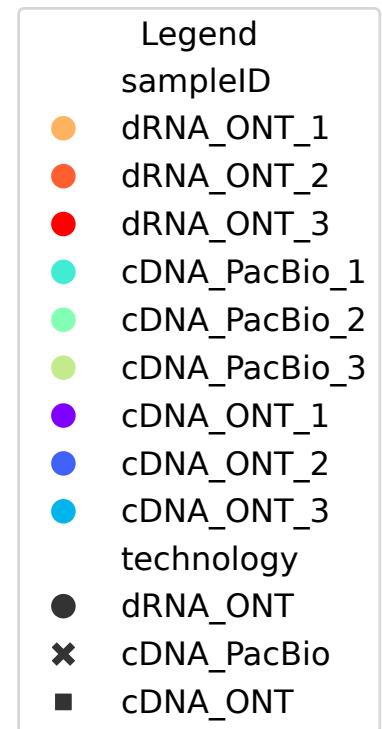
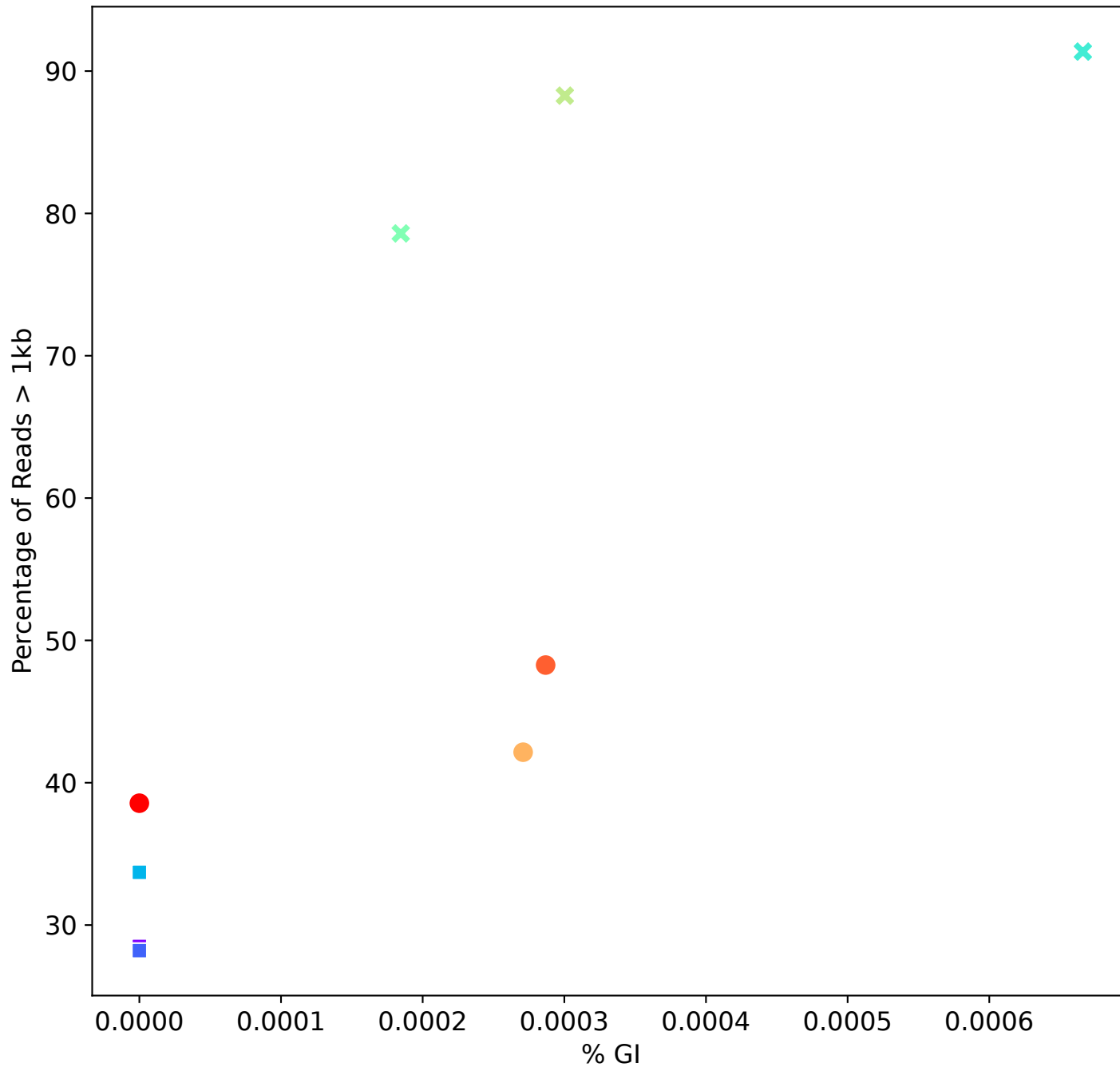
Percentage of Reads > 1kb vs %NIC



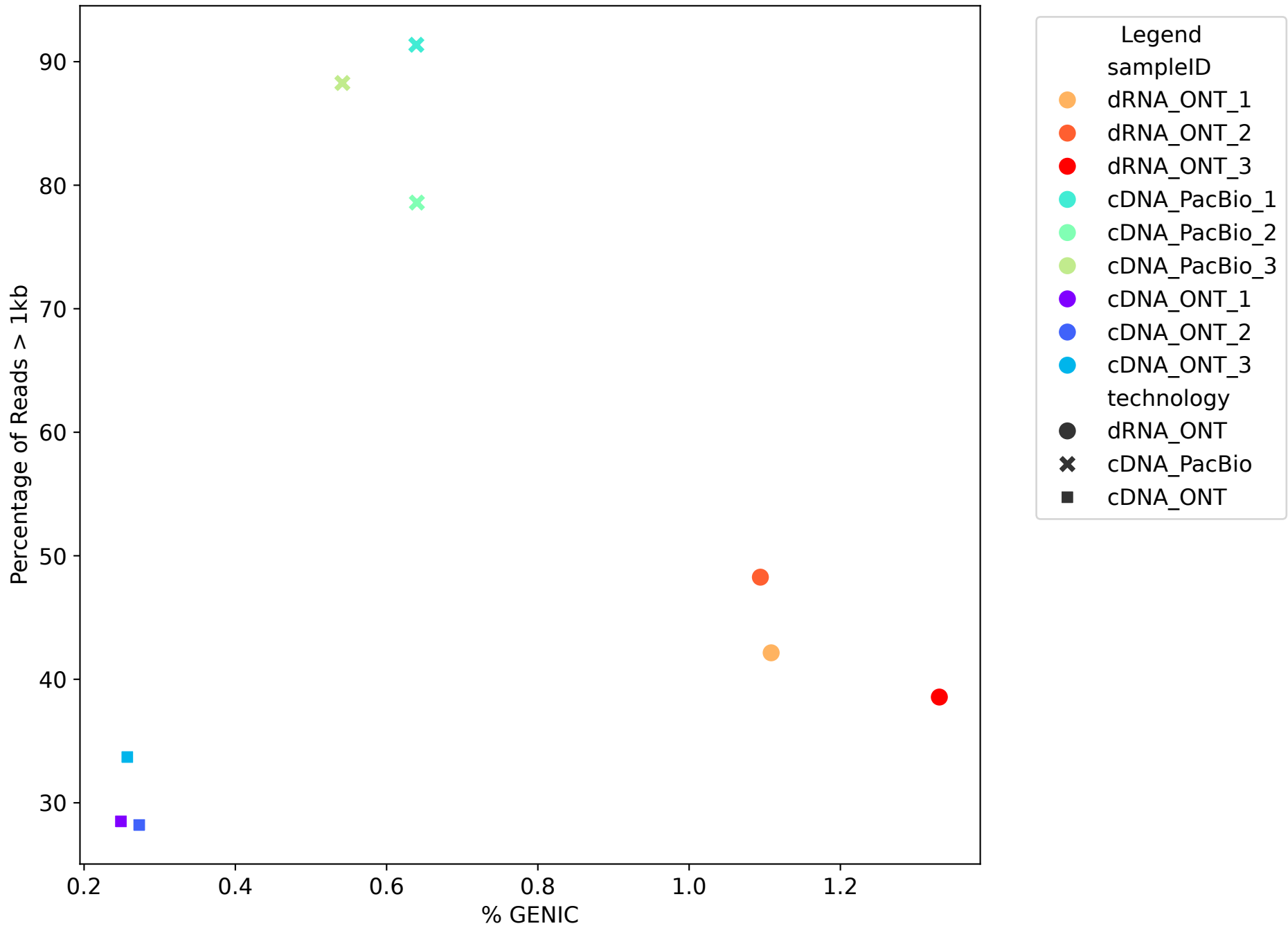
Percentage of Reads > 1kb vs %NNC



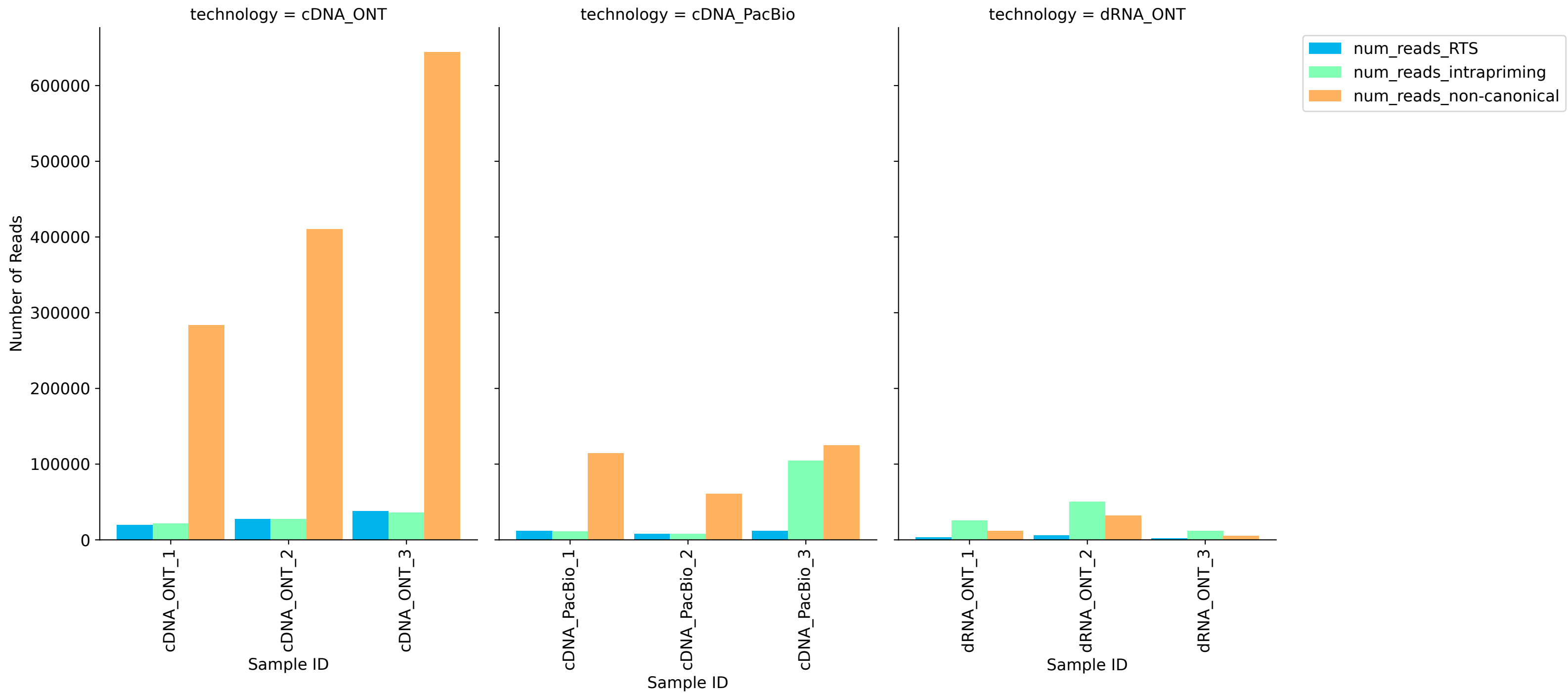
Percentage of Reads > 1kb vs %GI



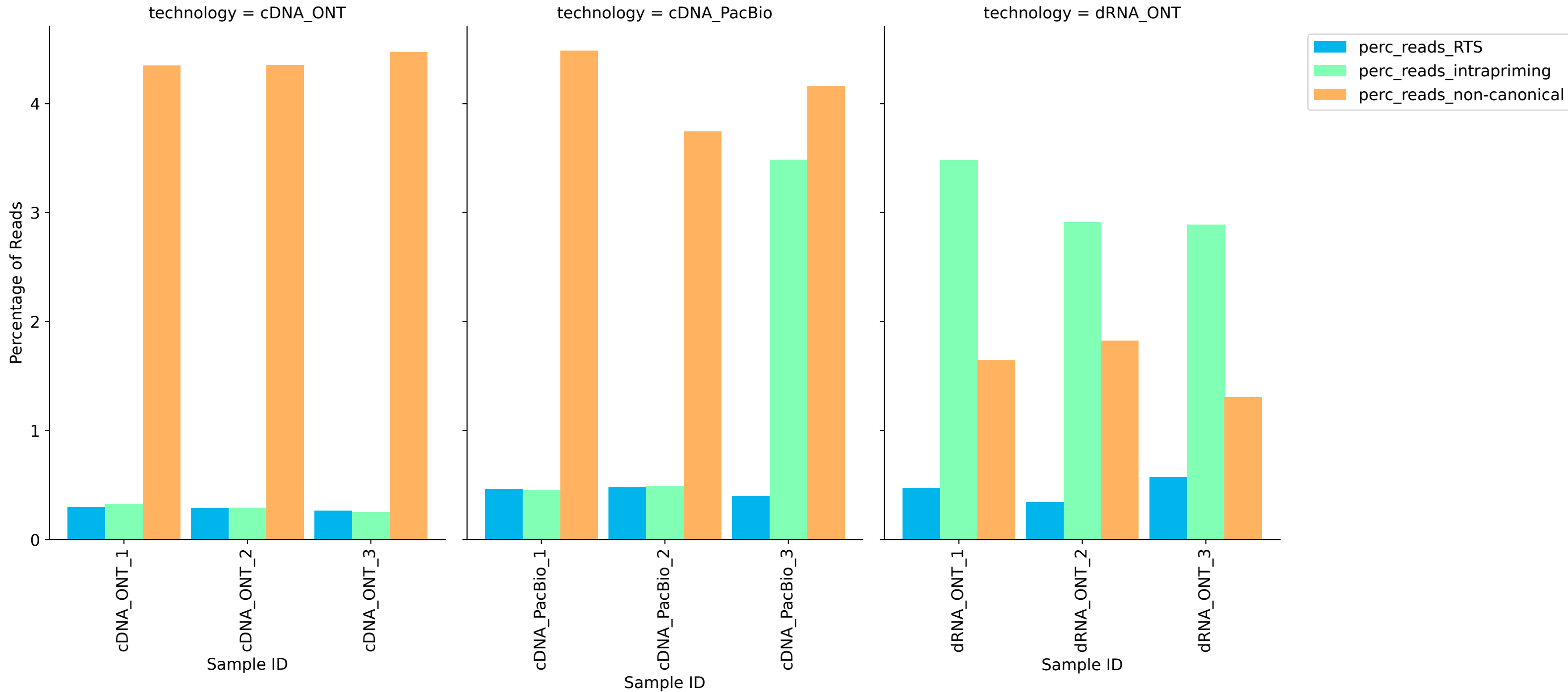
Percentage of Reads > 1kb vs %GENIC



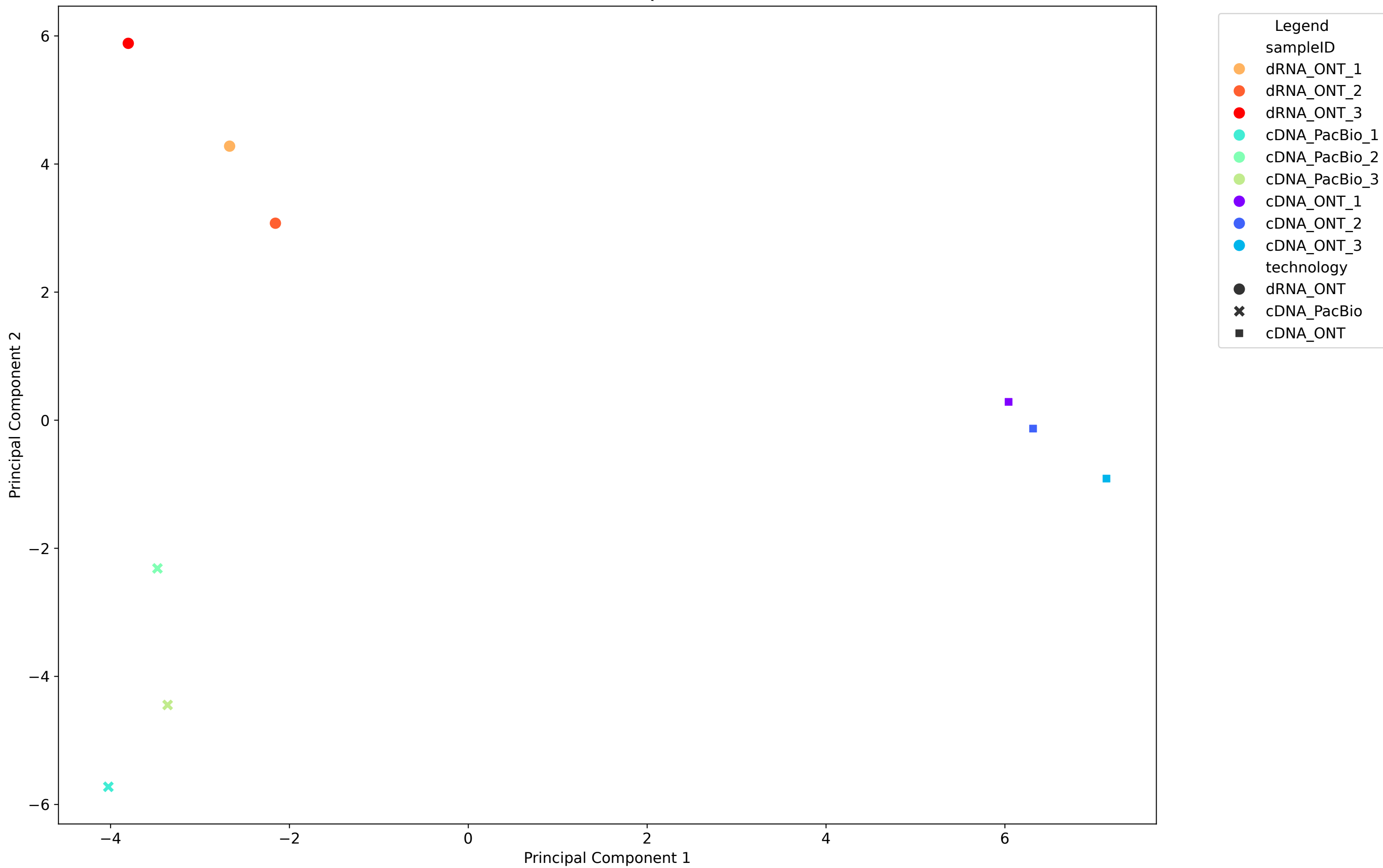
Number of Artefact Reads



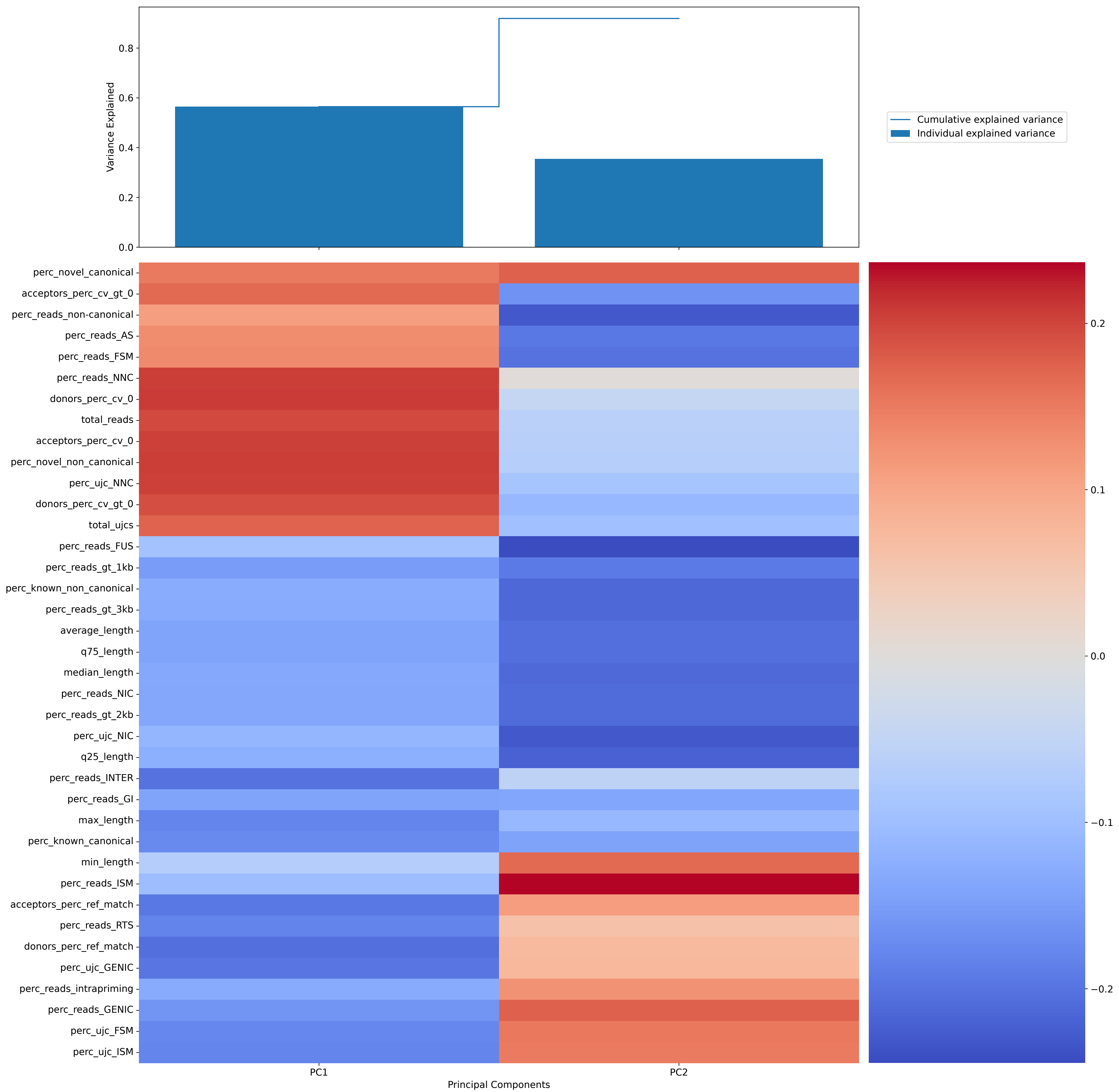
Percentage of Artefact Reads



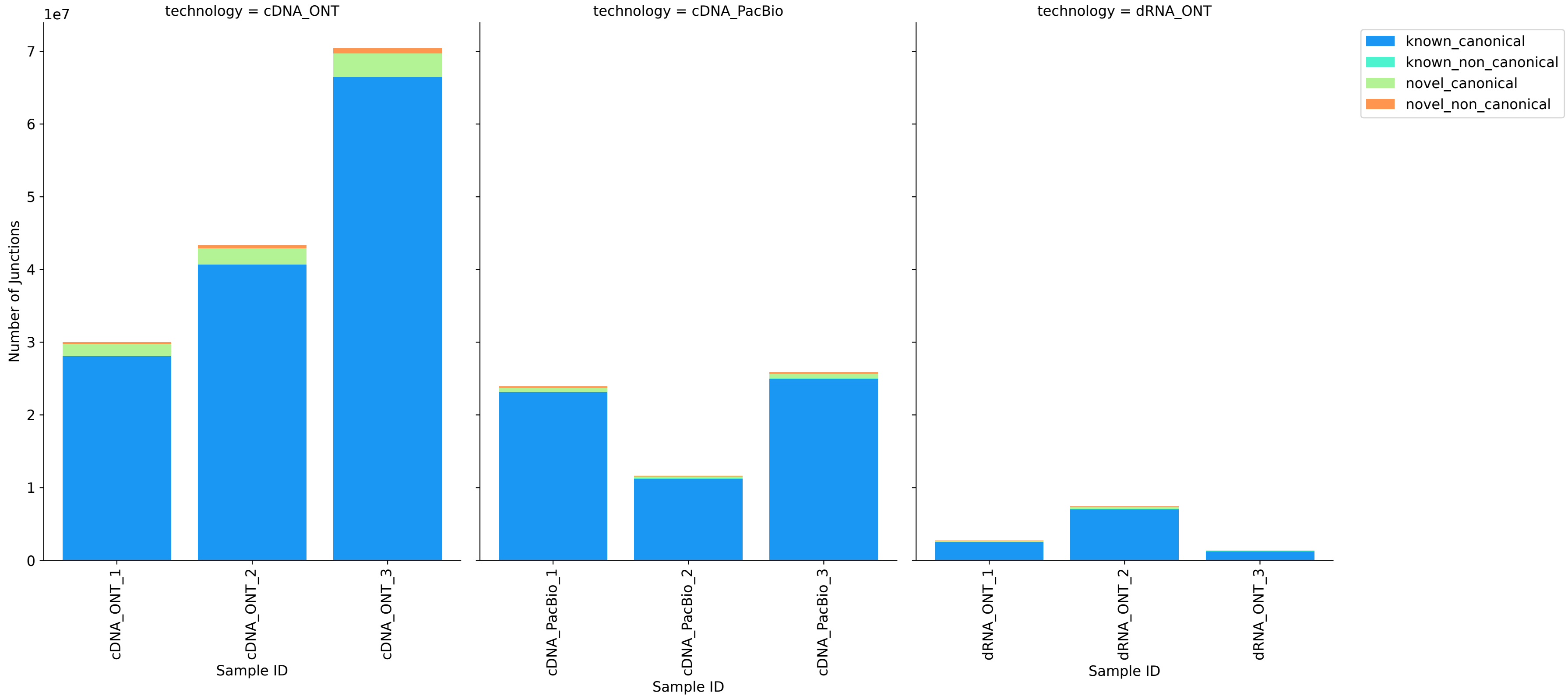
PCA Plot Based on QC metrics



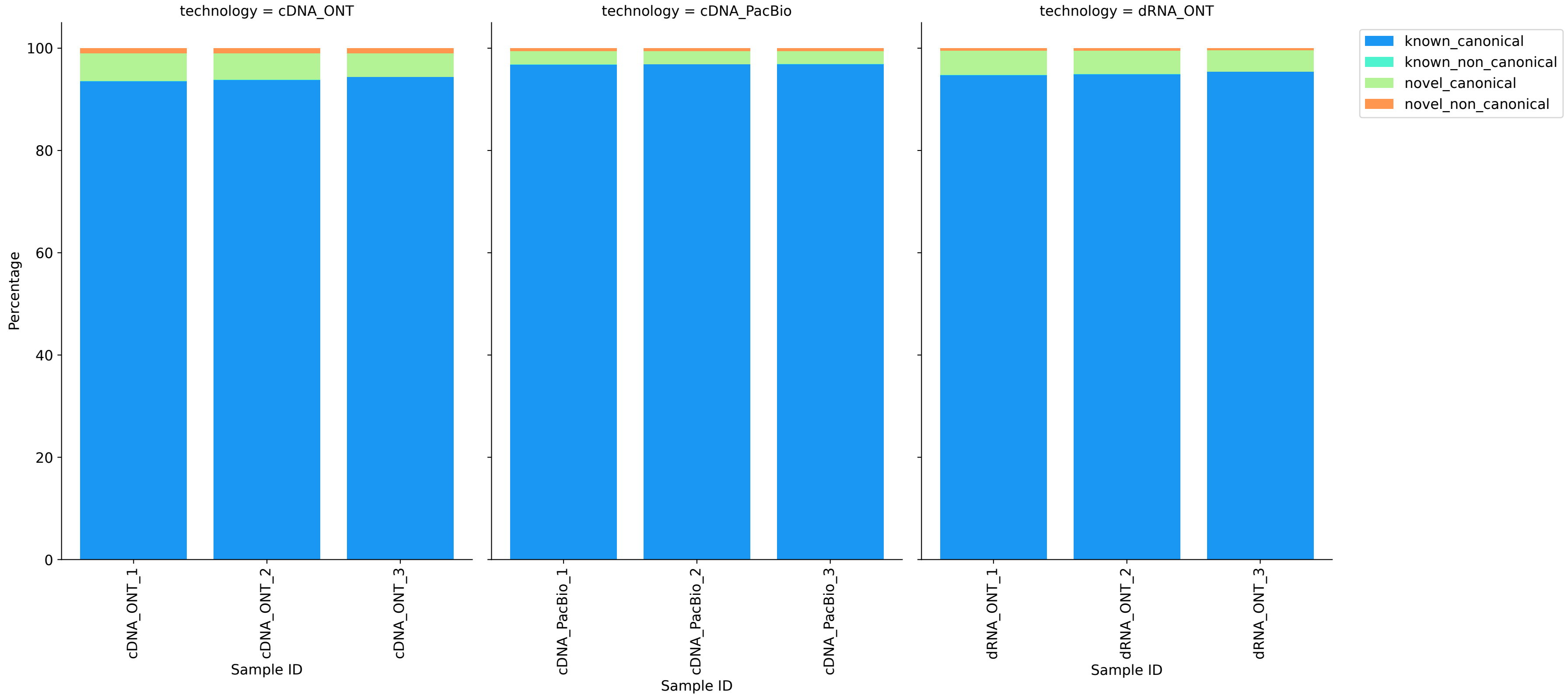
Variance and Heatmap of PC loadings



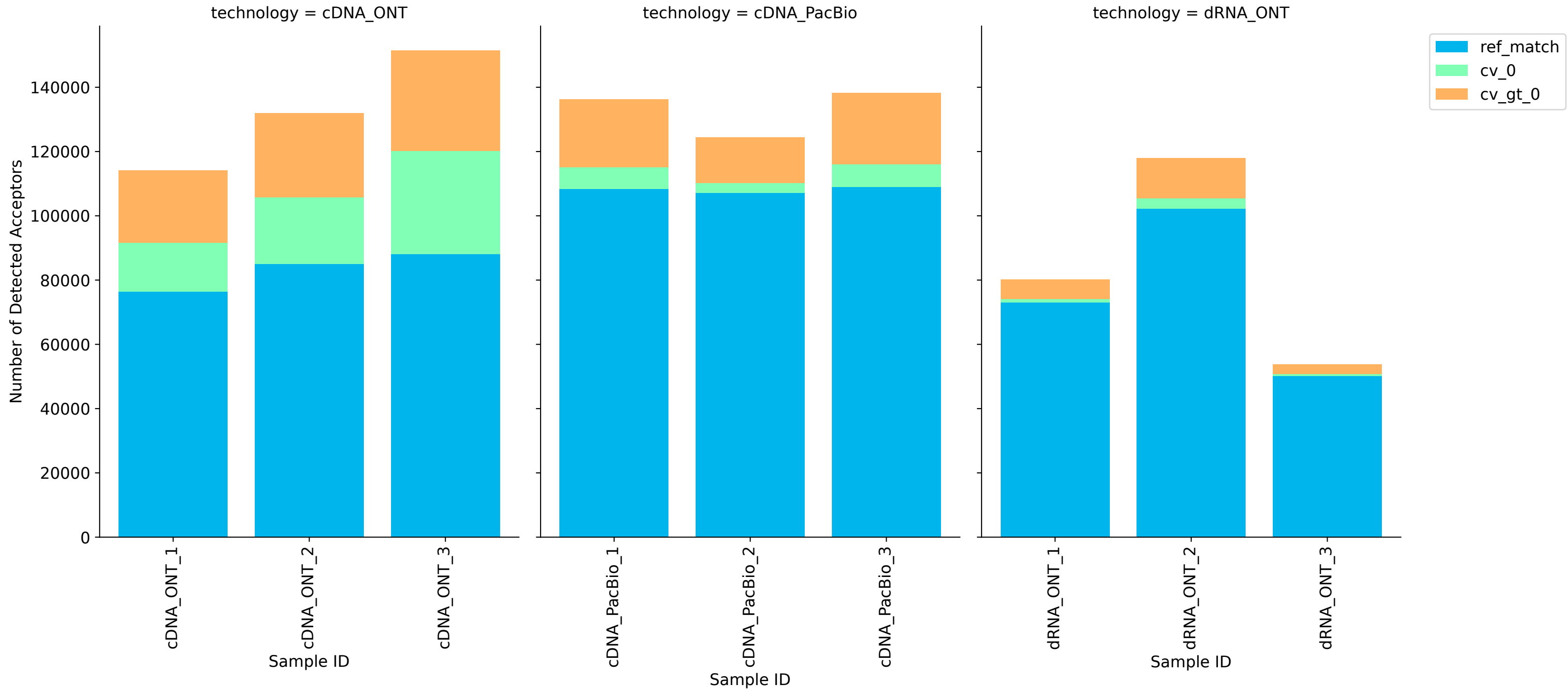
Junctions by Category



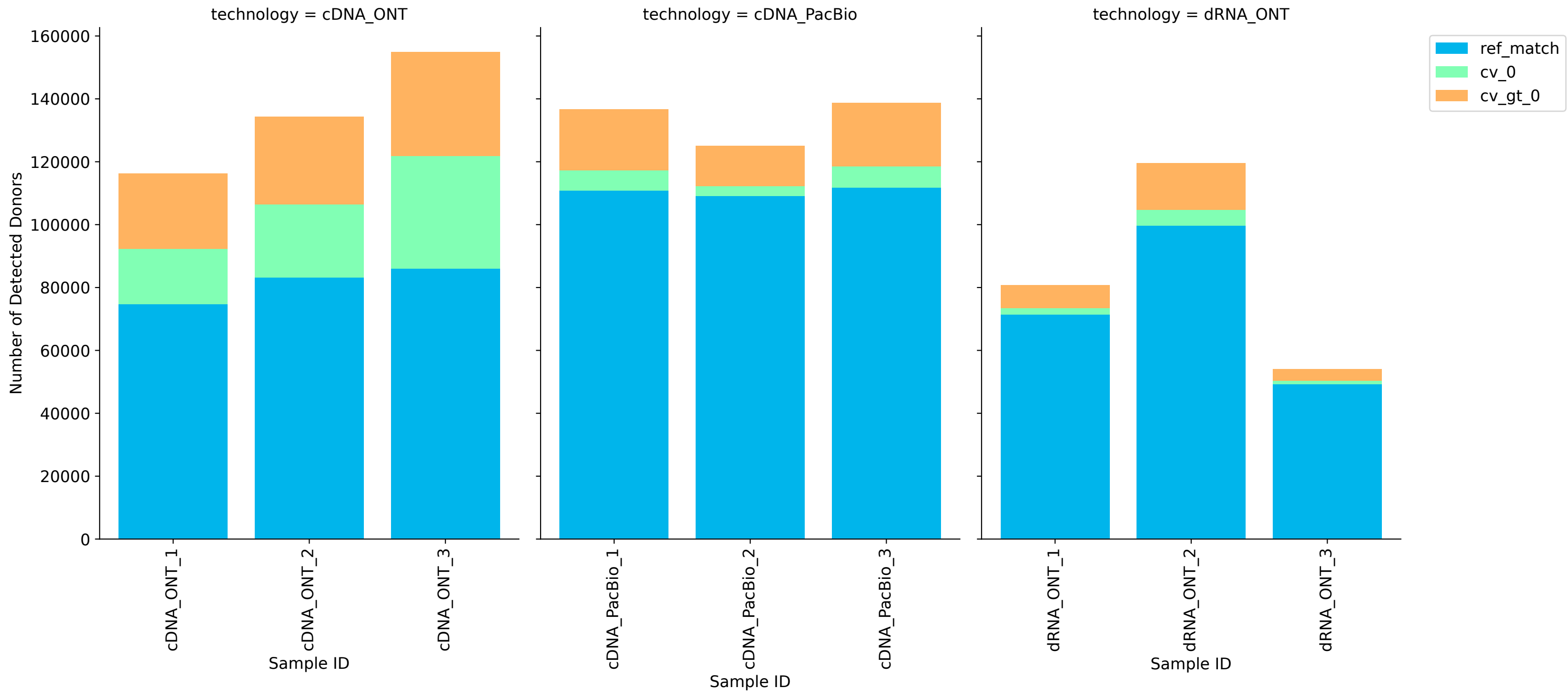
Junctions by Category



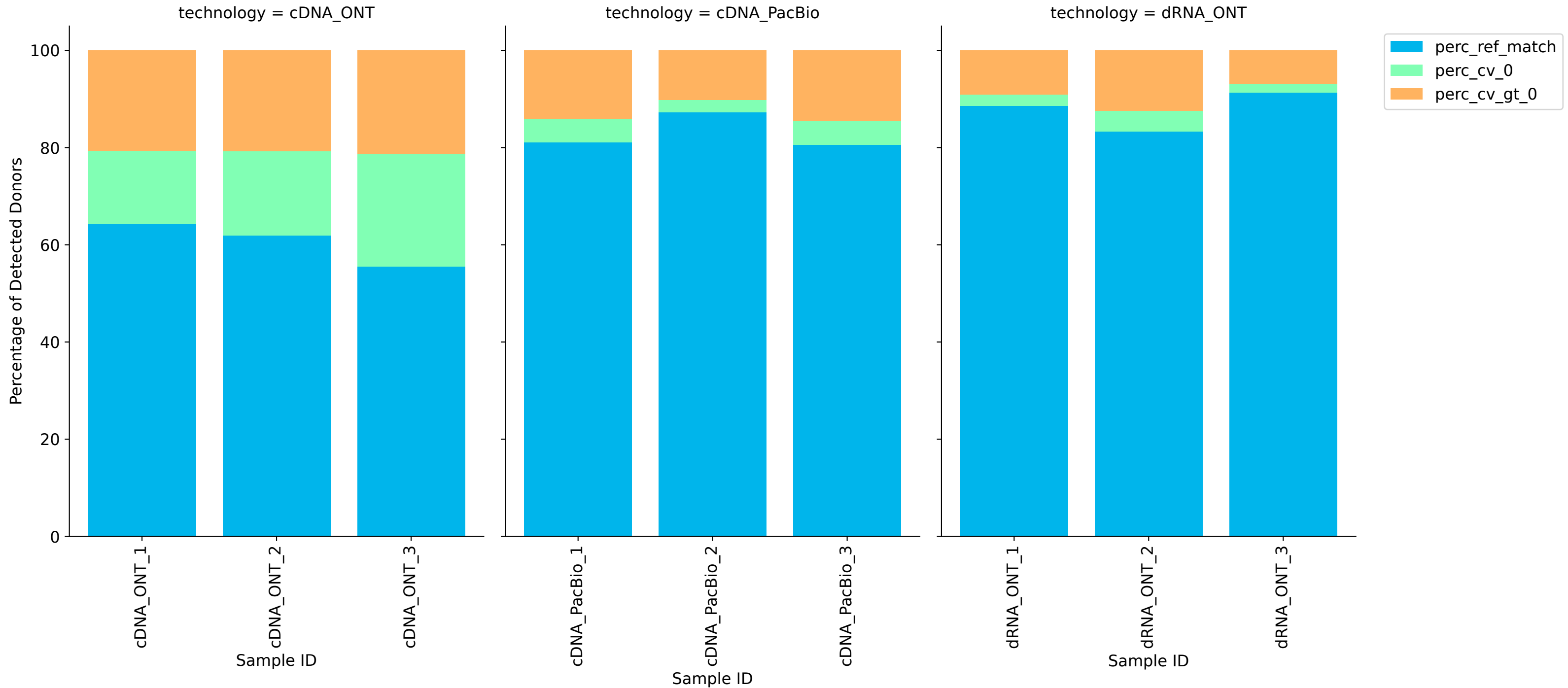
Number of Detected Acceptors



Number of Detected Donors



Percentage of Detected Donors



Percentage of Detected Acceptors

