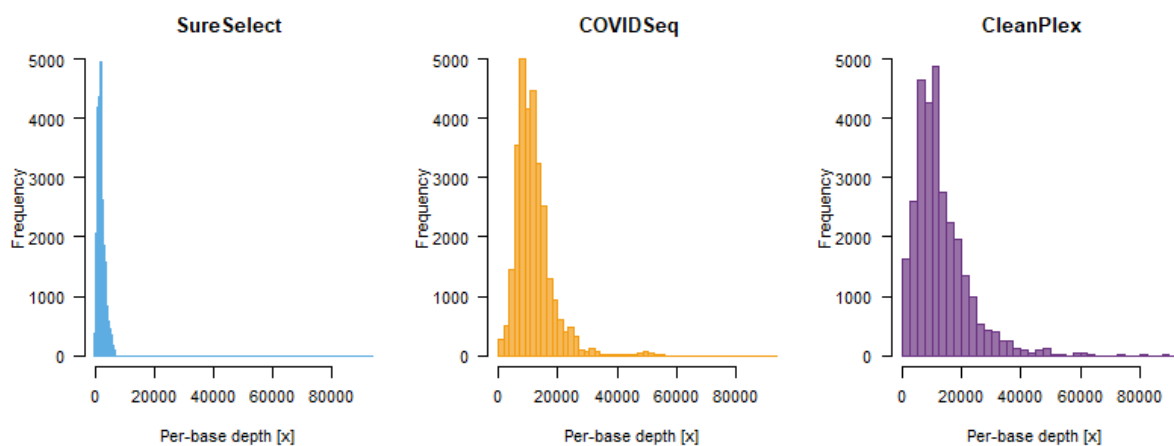


A comparison of high-throughput SARS-CoV-2 sequencing methods from nasopharyngeal samples

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Supplementary Figures



Suppl. Figure 1. A histogram of per-base depth of a typical library. A wider distribution confirms heterogeneous coverage for CleanPlex. Furthermore, higher frequency in the leftmost bin indicates the presence of poorly covered regions in case of CleanPlex.