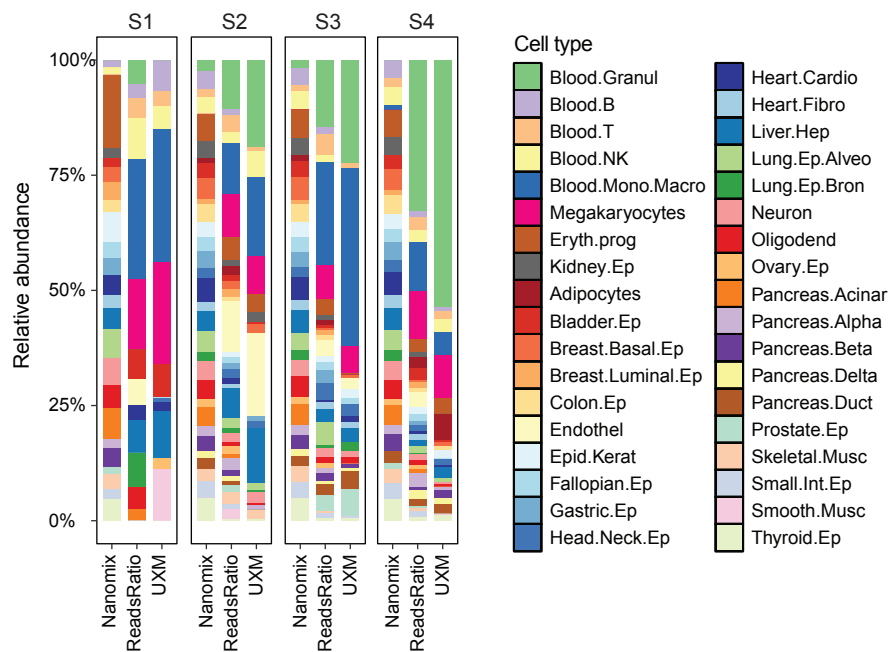


A



B

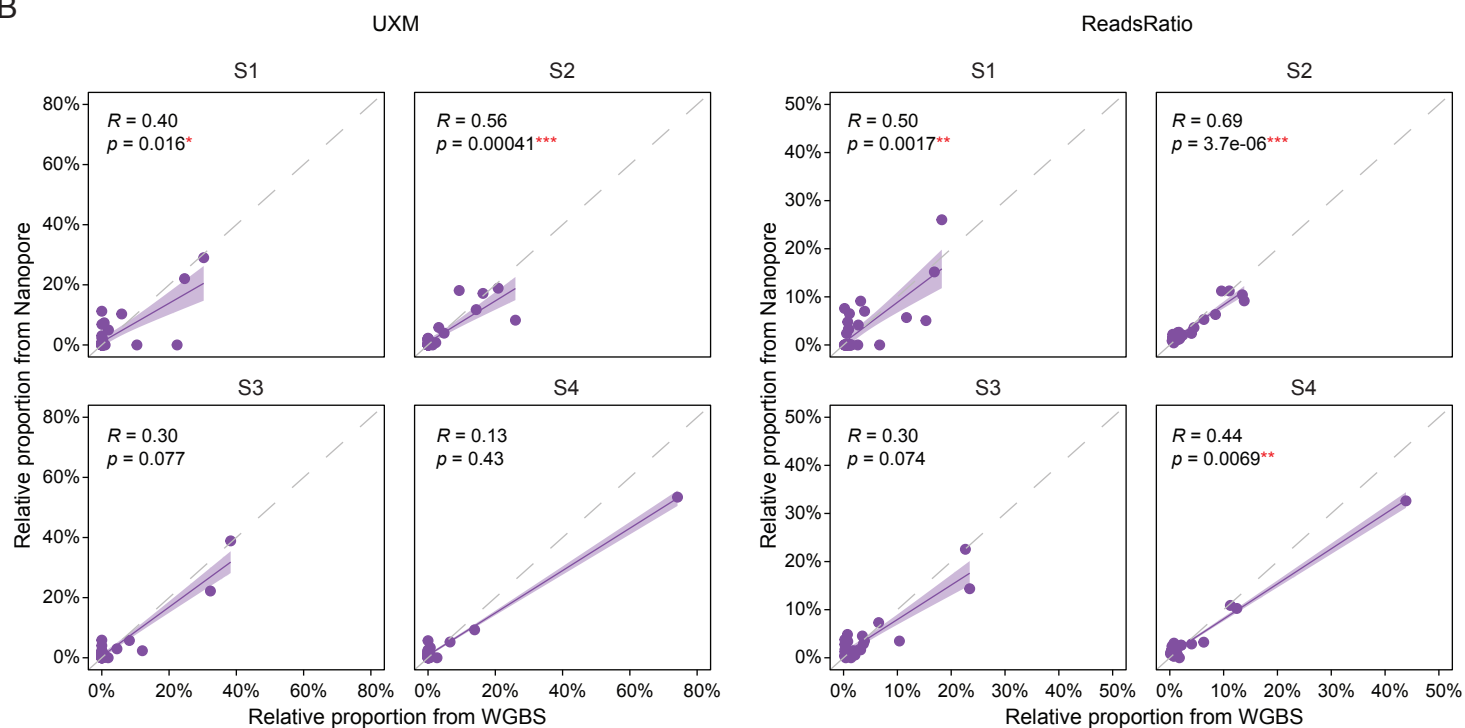


Fig S16. Performance of ReadsRatio method in Nanopore cfDNA sequencing data.

A. Cell type composition in Nanopore sequencing samples deconvoluted by different methods. B. Correlation of cell type composition between paired WGBS and Nanopore data for each individual sample. $p < 0.05^*$, $p < 0.01^{**}$, $p < 0.001^{***}$.