Description of additional supplementary files

File Name: **Supplementary Data 1**
Description: Comparing ccsmeth and primrose/pb-CpG-tools against BS-seq and nanopore sequencing under different coverages of HG002 CCS reads (71.0× in total). Values for coverage 5×-70× are the average and standard deviation of 5 repeated tests in "average ± std" format. r: Pearson correlation; r^2: the coefficient of determination; ρ: Spearman correlation; RMSE: root mean square error.