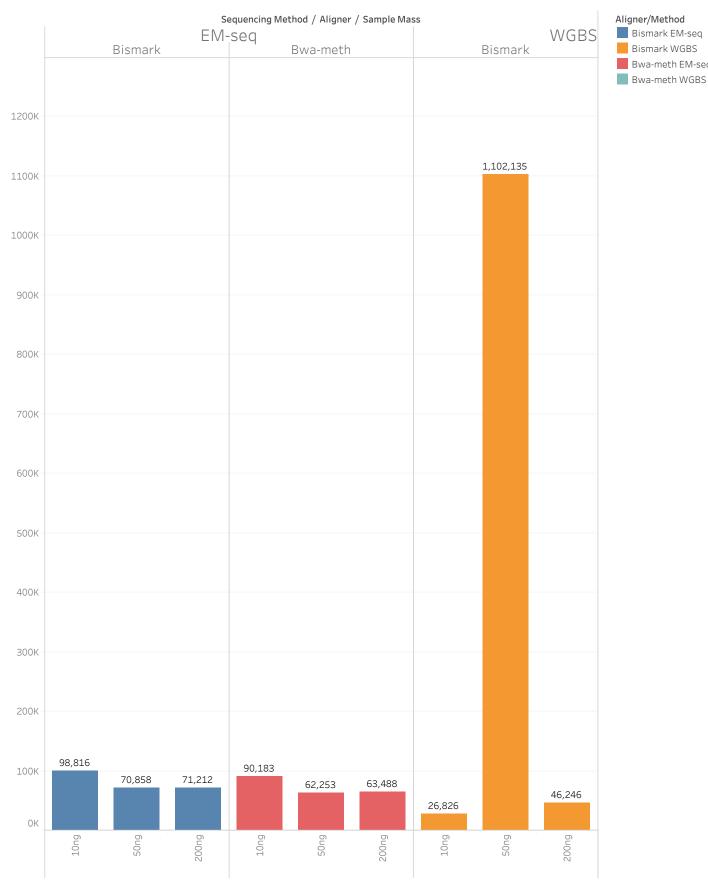
Resolved Calls

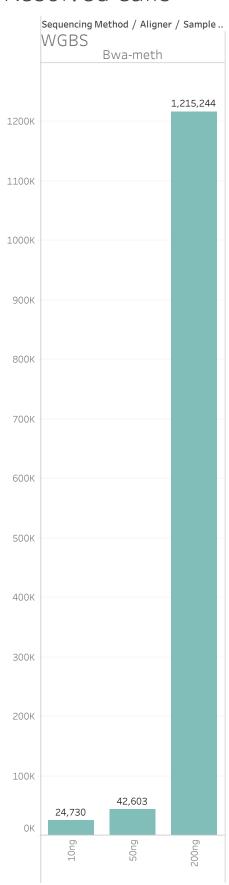


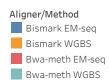
Bismark WGBS

Bwa-meth EM-seq

Sum of Miscalled Cs for each Sample Mass broken down by Sequencing Method and Aligner. Color shows details about Aligner/Method. $The \ marks \ are \ labeled \ by \ sum \ of \ Miscalled \ Cs. \ The \ data \ is \ filtered \ on \ Analysis \ and \ Reference. \ The \ Analysis \ filter \ keeps \ resolved_calls. \ The \ Miscalled \ Cs. \ The \ data \ is \ filtered \ on \ Analysis \ and \ Reference.$ Reference filter keeps grch38.

Resolved Calls





Sum of Miscalled Cs for each Sample Mass broken down by Sequencing Method and Aligner. Color shows details about Aligner/Method. The marks are labeled by sum of Miscalled Cs. The data is filtered on Analysis and Reference. The Analysis filter keeps resolved_calls. The Reference filter keeps grch38.