



Methylation Detection System

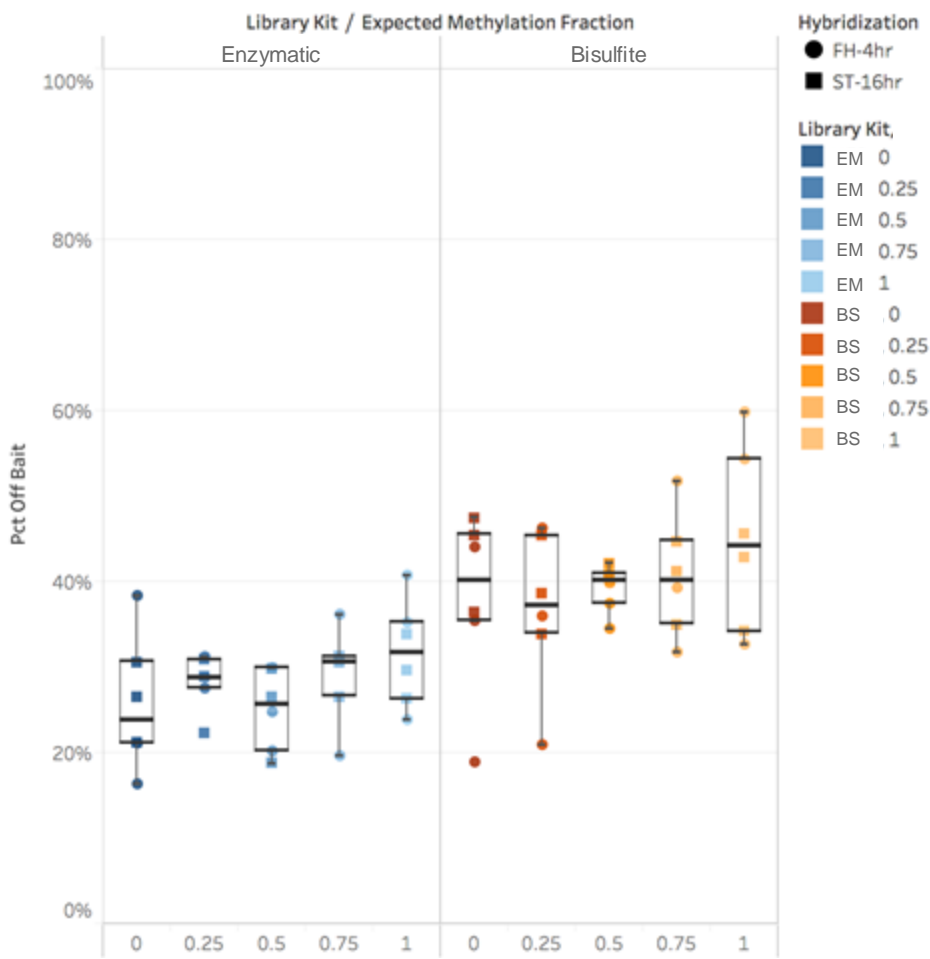
Tim Farinholt, PhD
NGS Applications Scientist



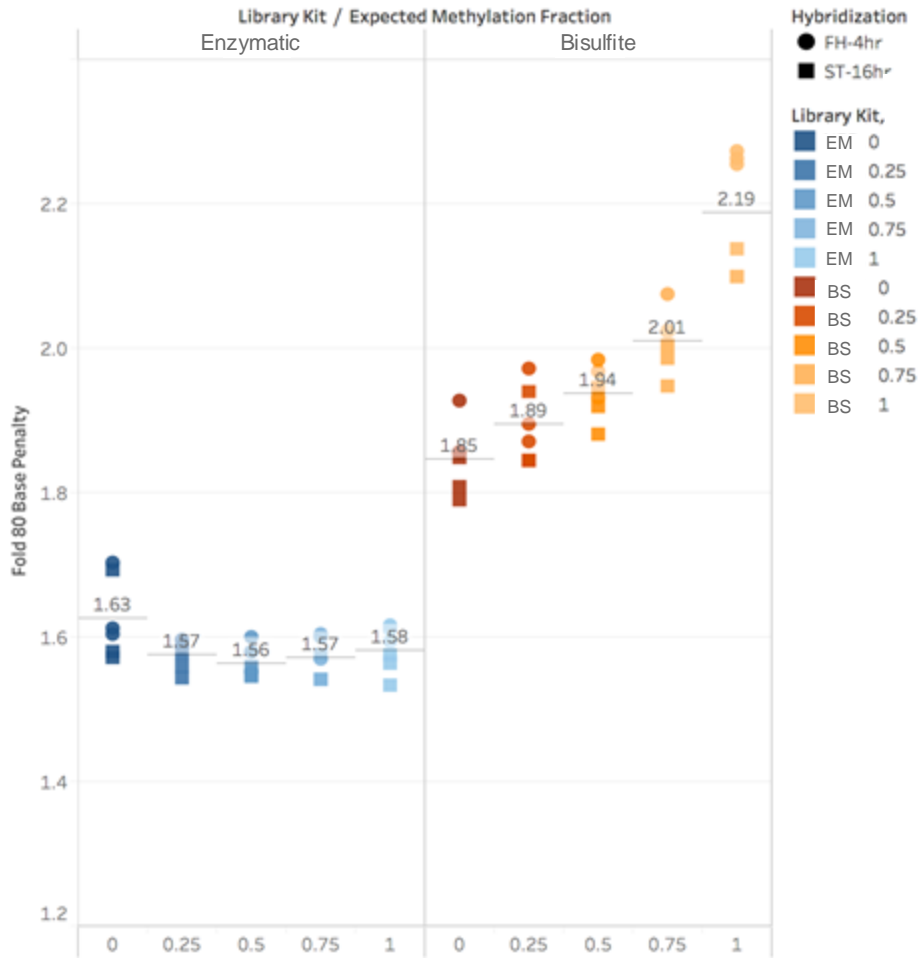
Increased Specificity and Uniformity with Enzymatic Conversion



Percent Off Bait



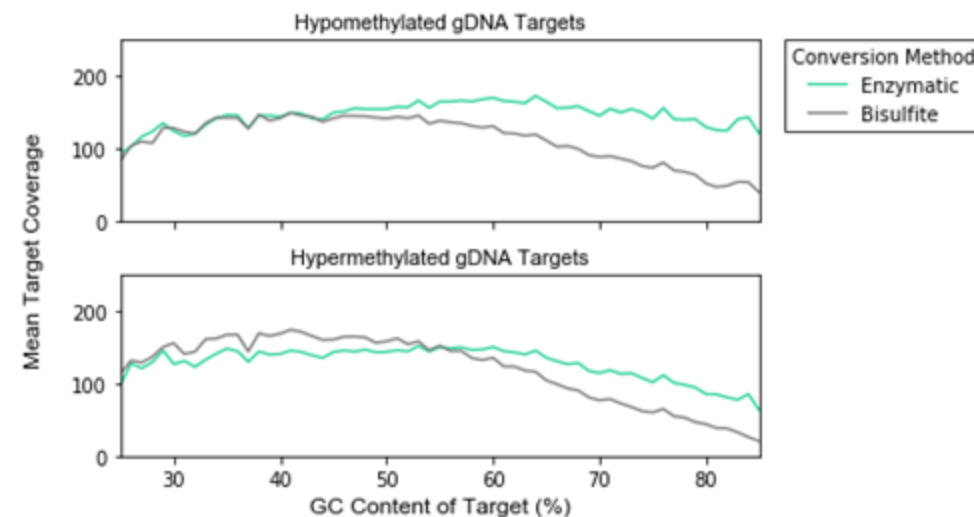
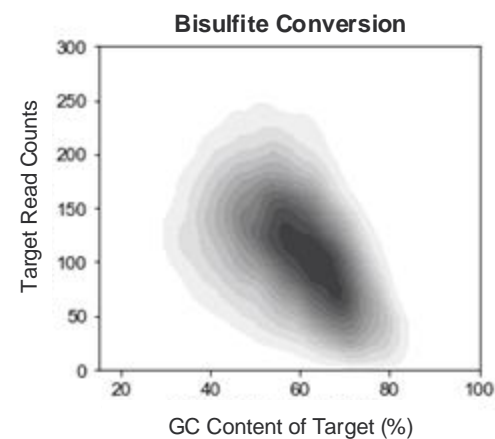
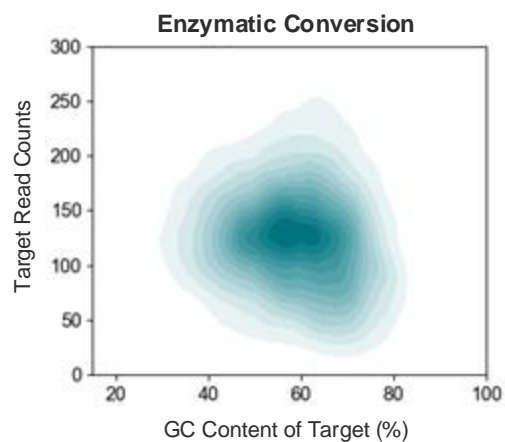
Fold-80 Base Penalty



Enzymatic Method Provides Better GC Representation



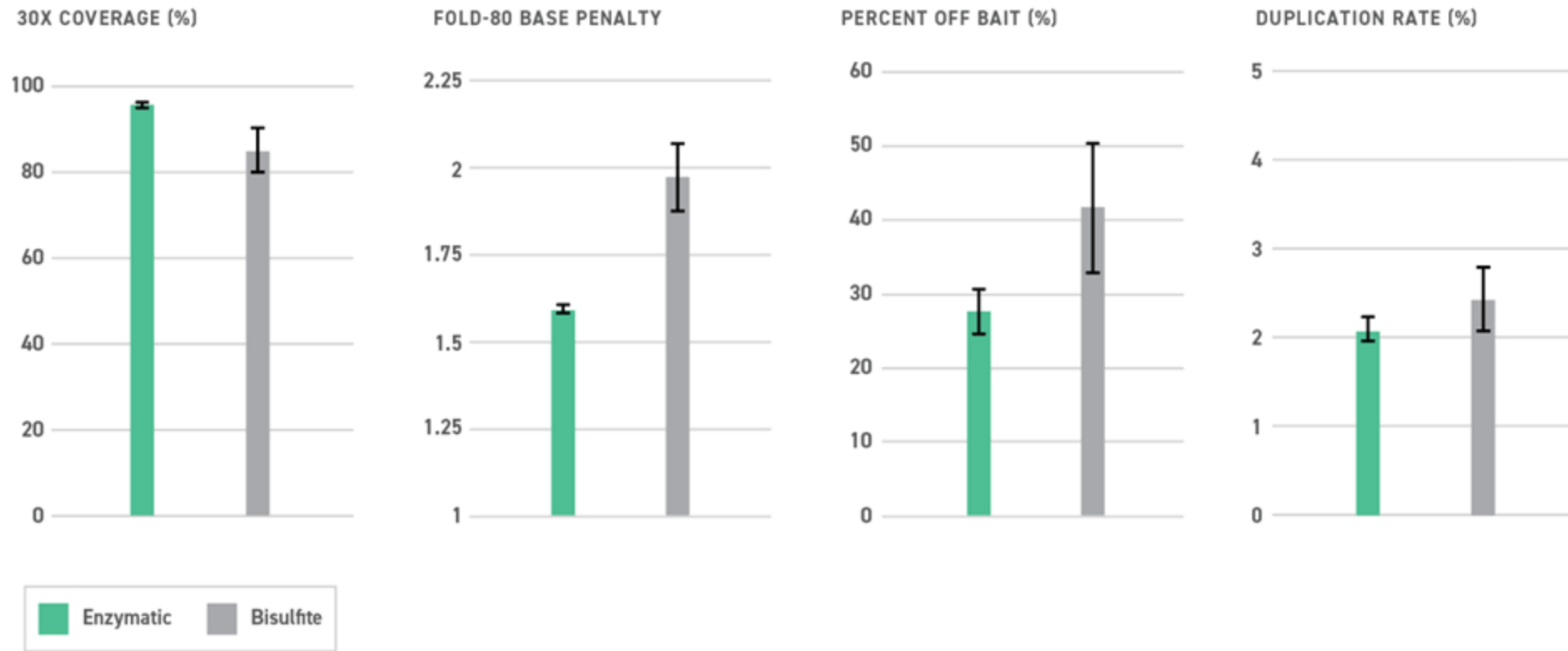
The enzymatic conversion method shows significantly less bias in high GC target regions



Library Preparation - Bisulfite vs Enzymatic Conversion



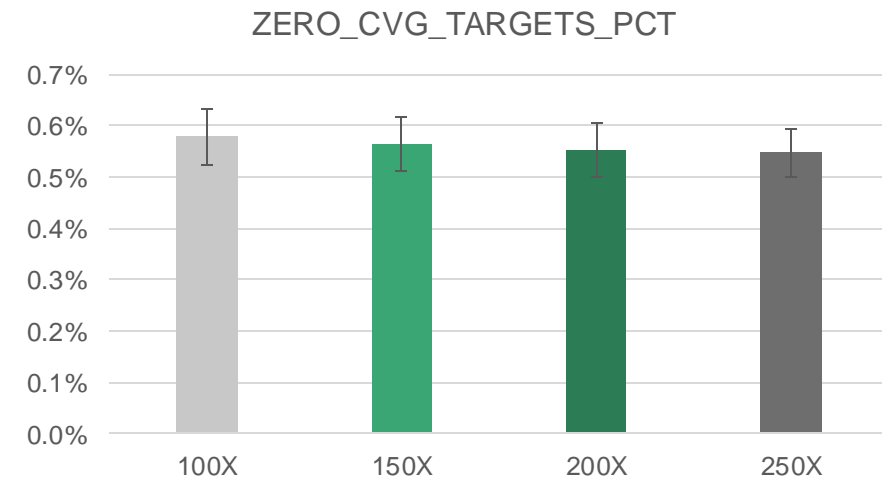
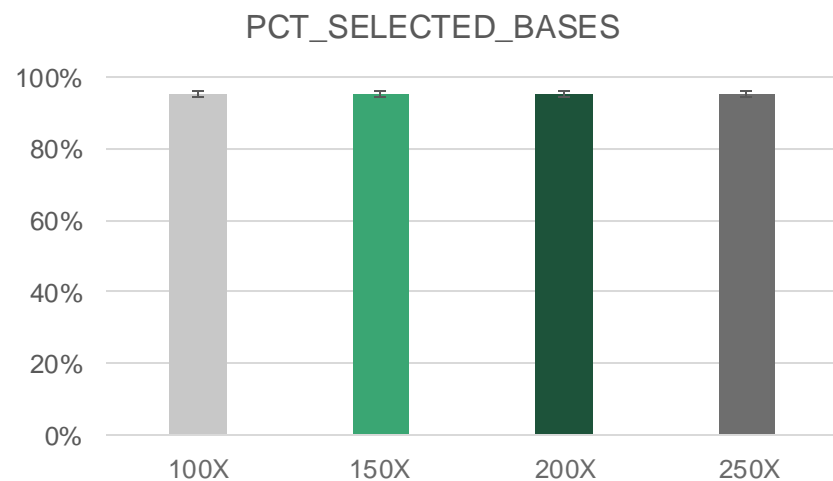
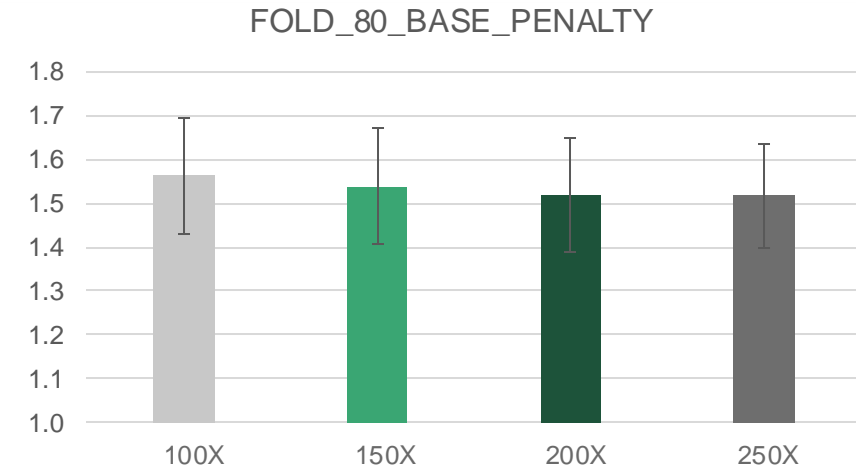
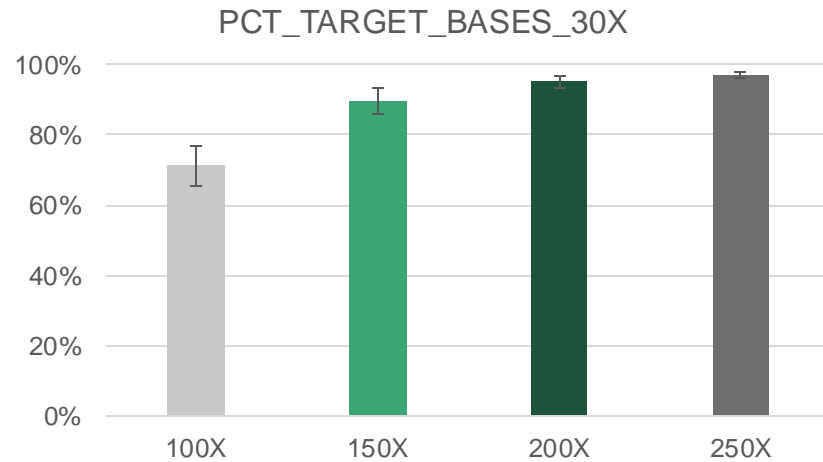
Enzymatic Conversion is Superior to Bisulfite Conversion



Key Performance Metrics



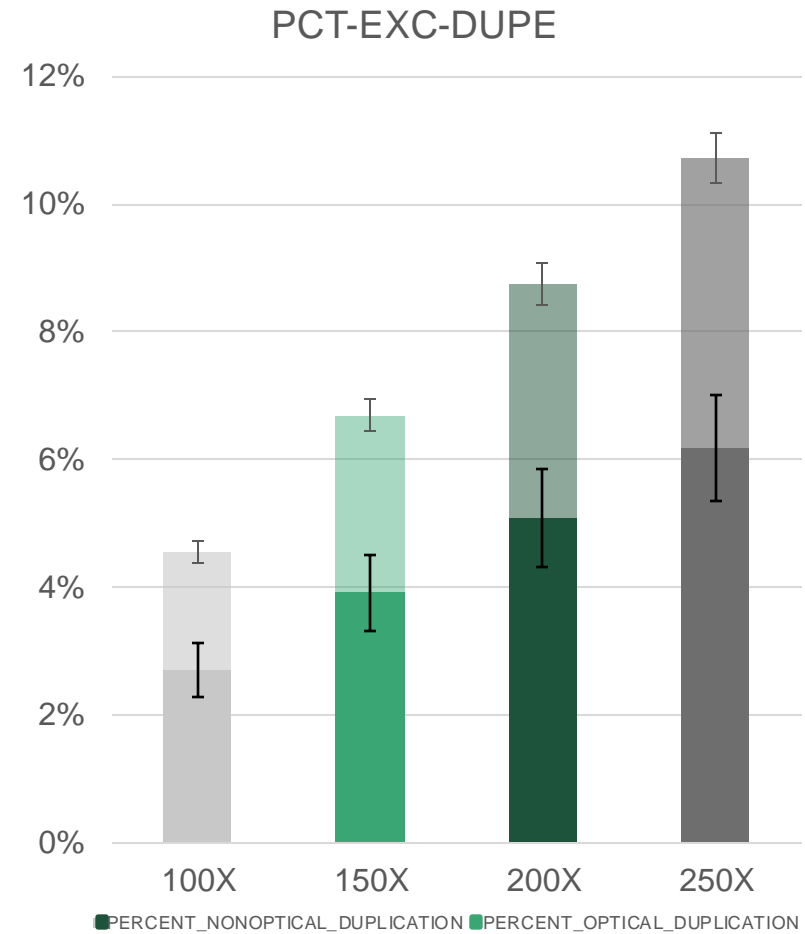
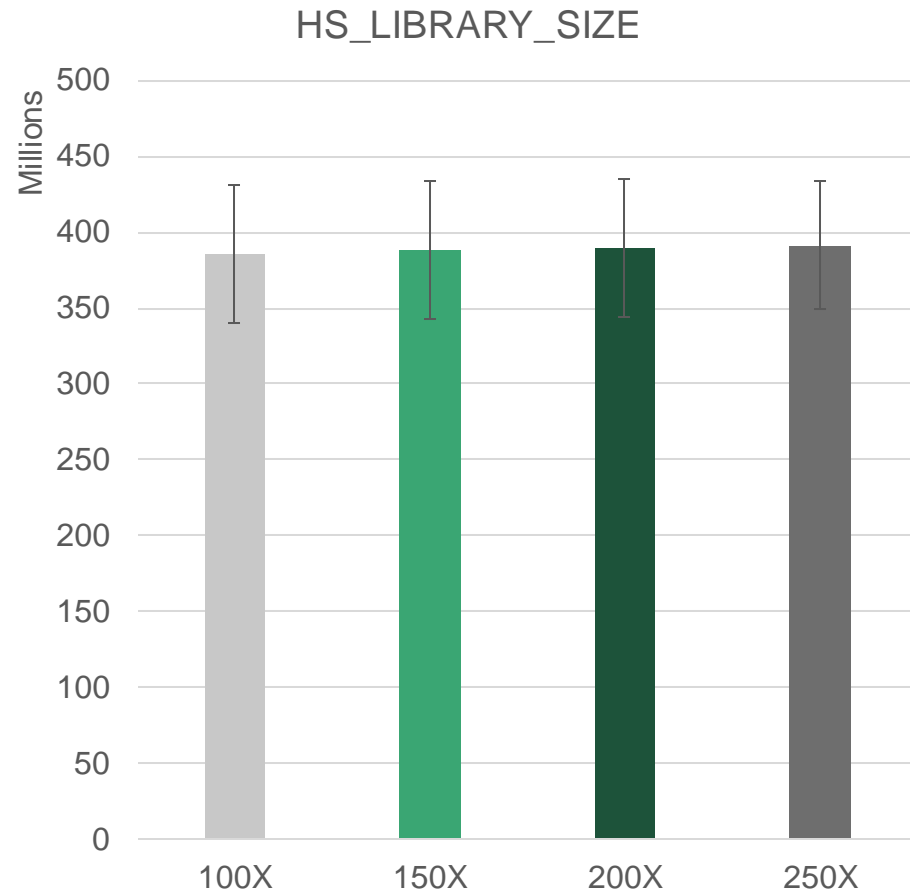
150x raw coverage = 90% at 30x bases, Fold 80 1.54, On-target 95%, Zero coverage 0.56%



Library Complexity



Full saturation across all coverage values



Methylation Specific Metrics

