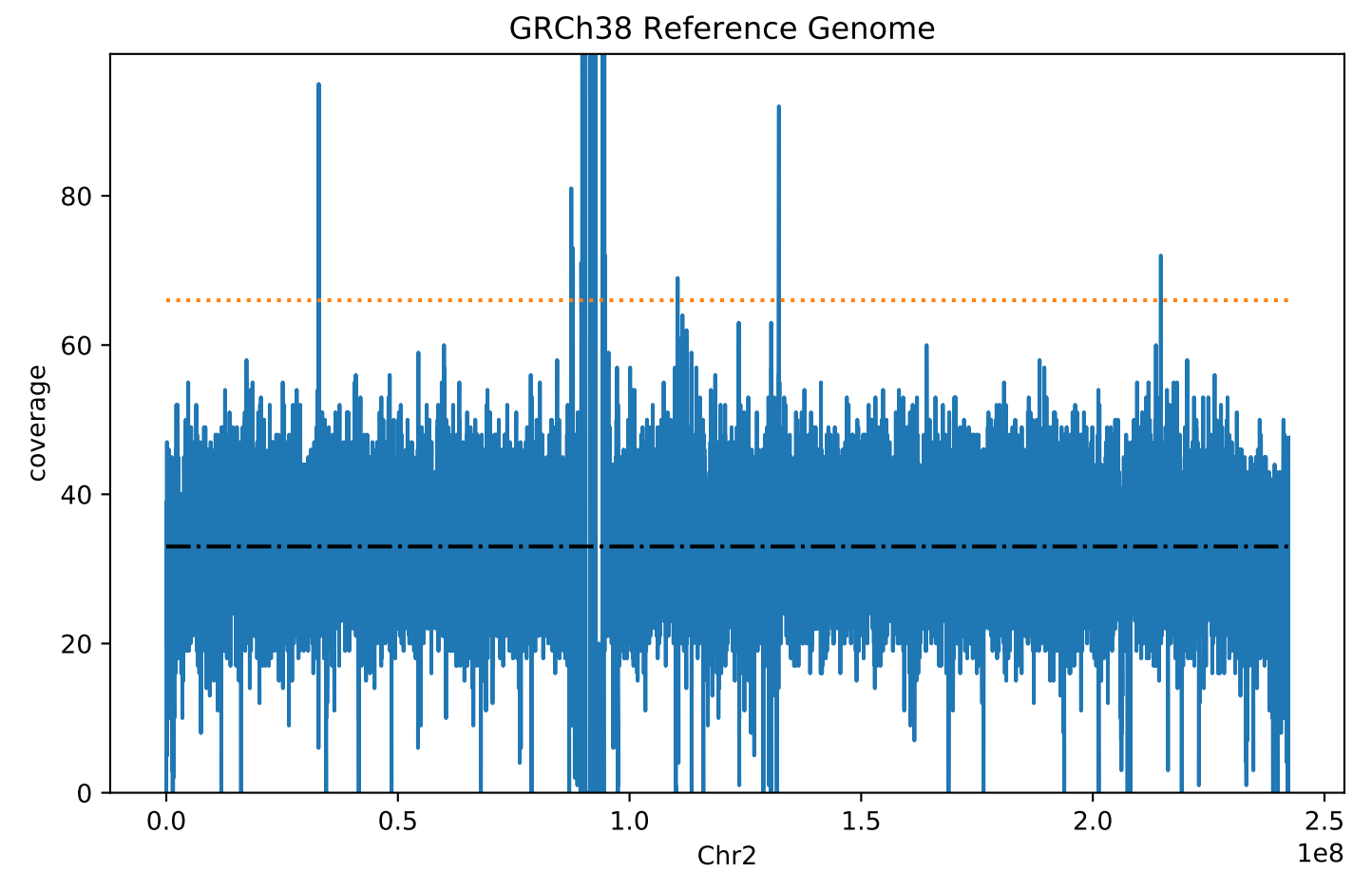
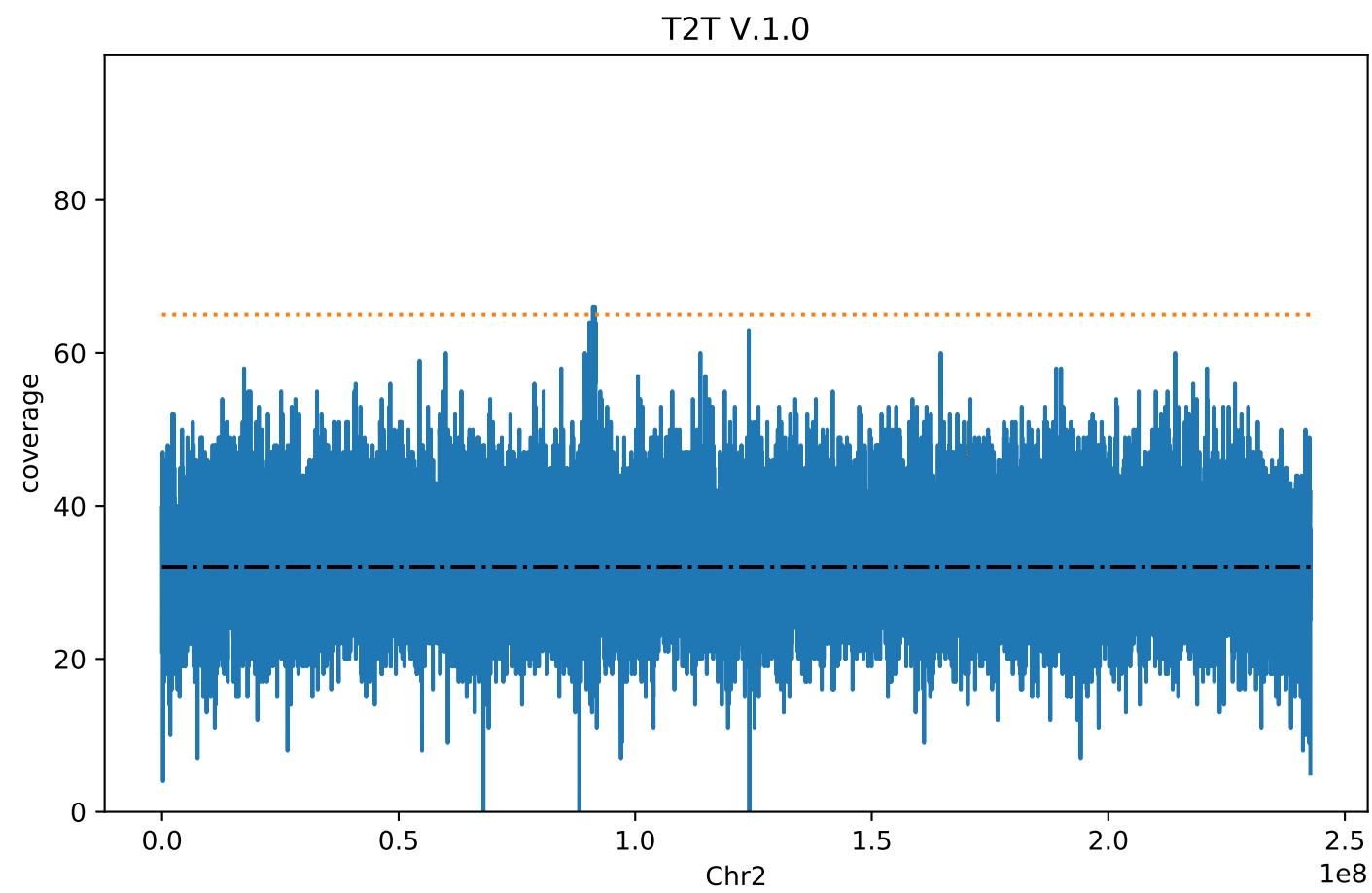
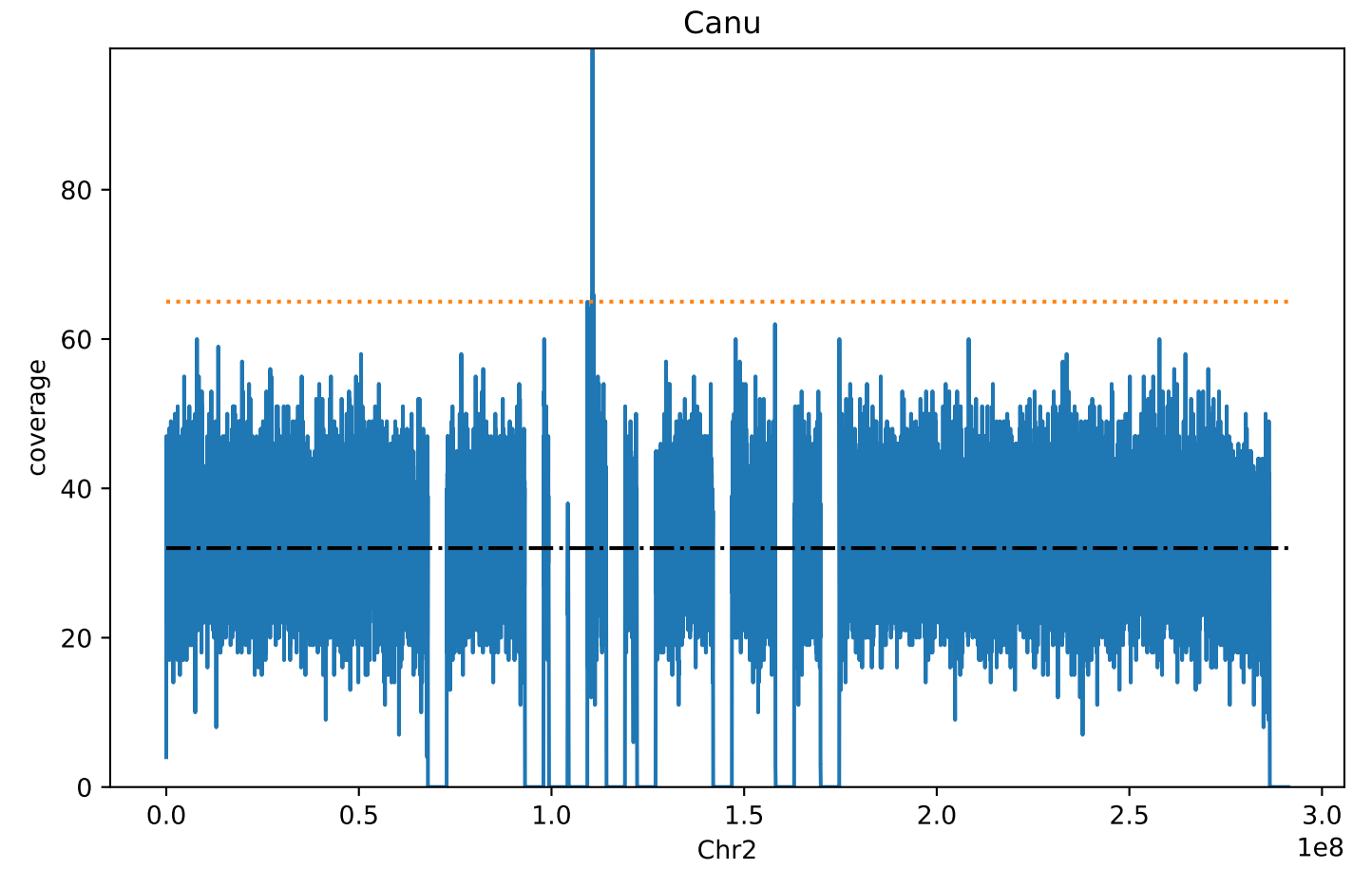
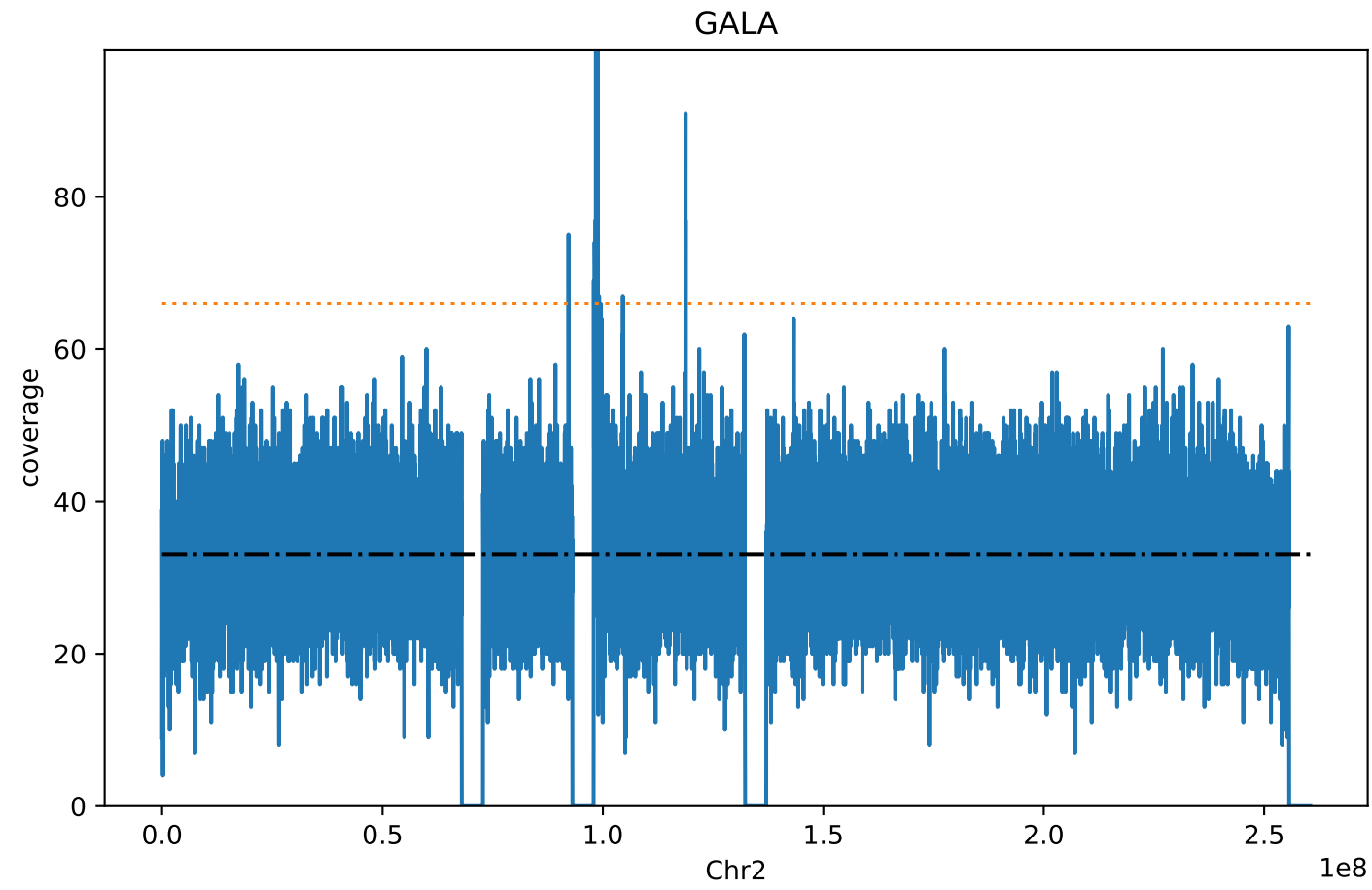
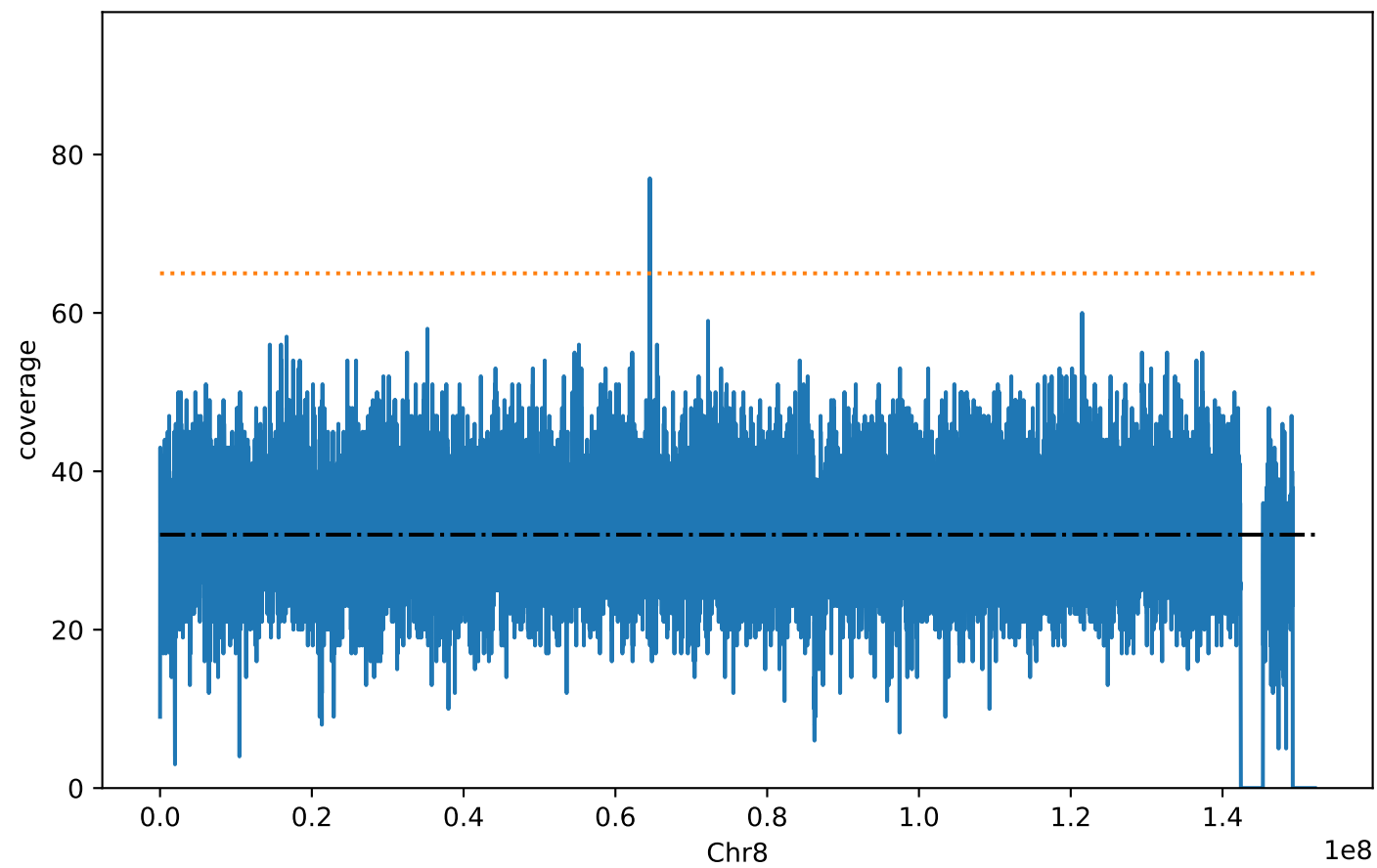


Human Coverage plot

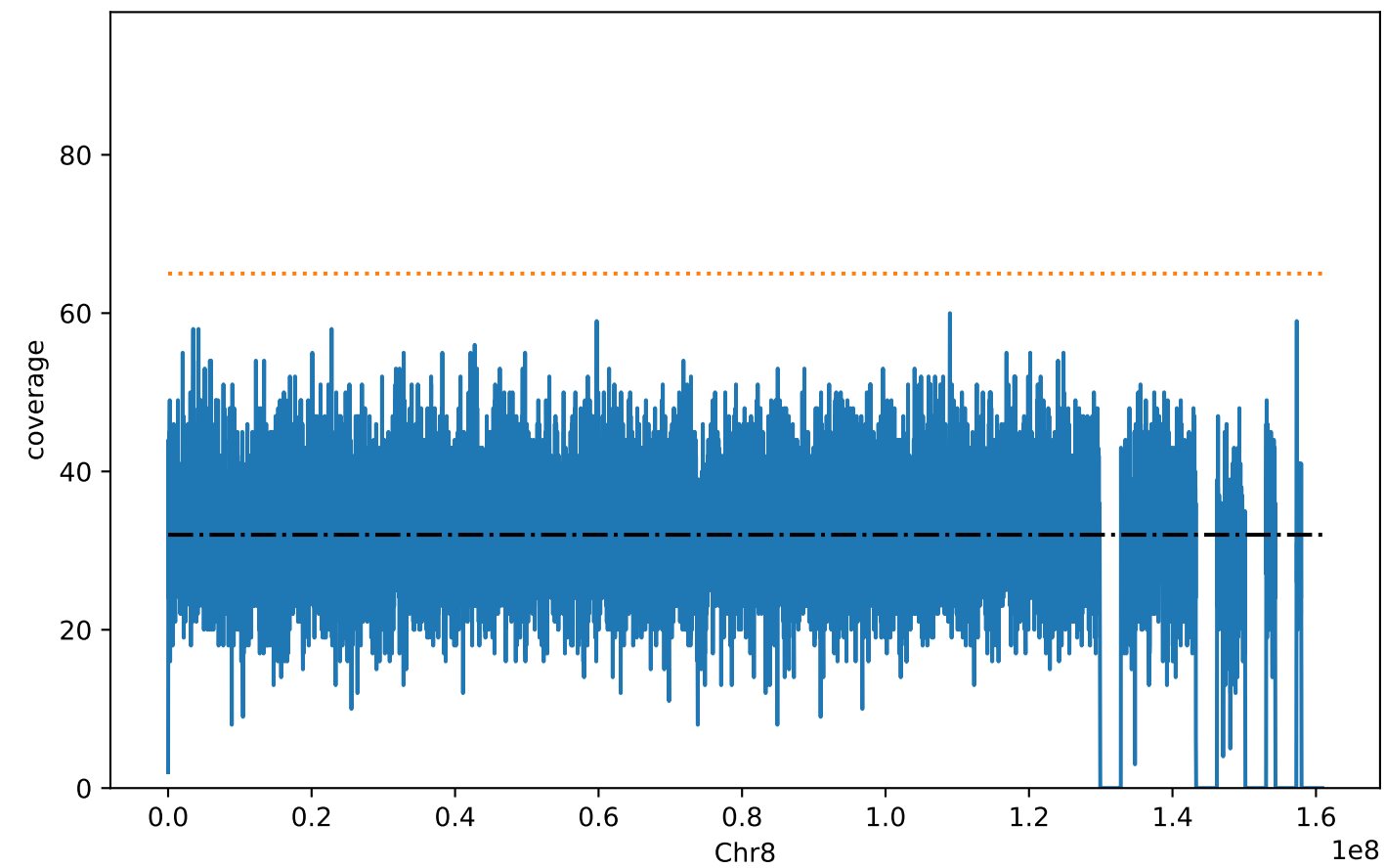
----- Average depth * 2
-.- Average depth



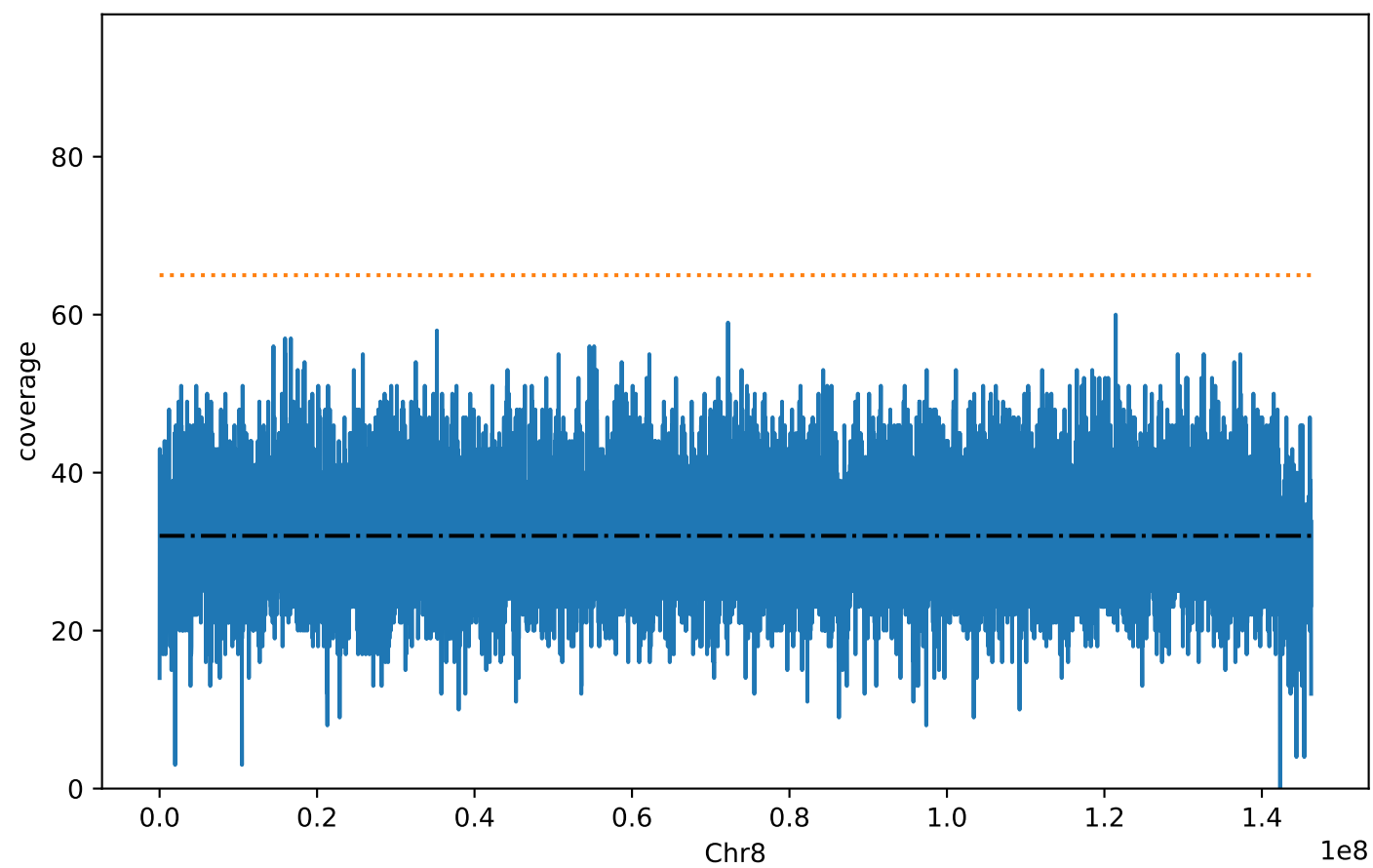
GALA



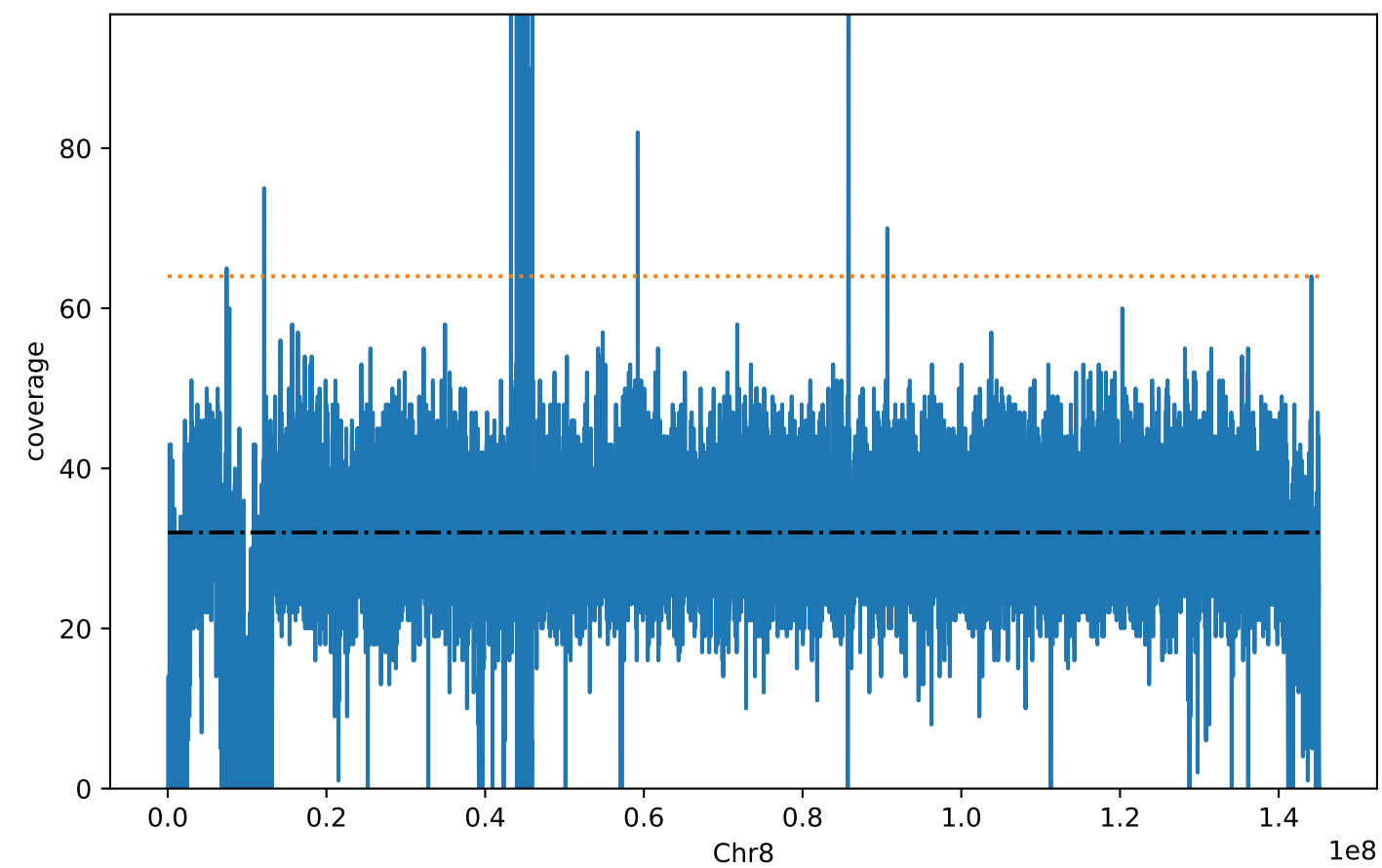
Canu



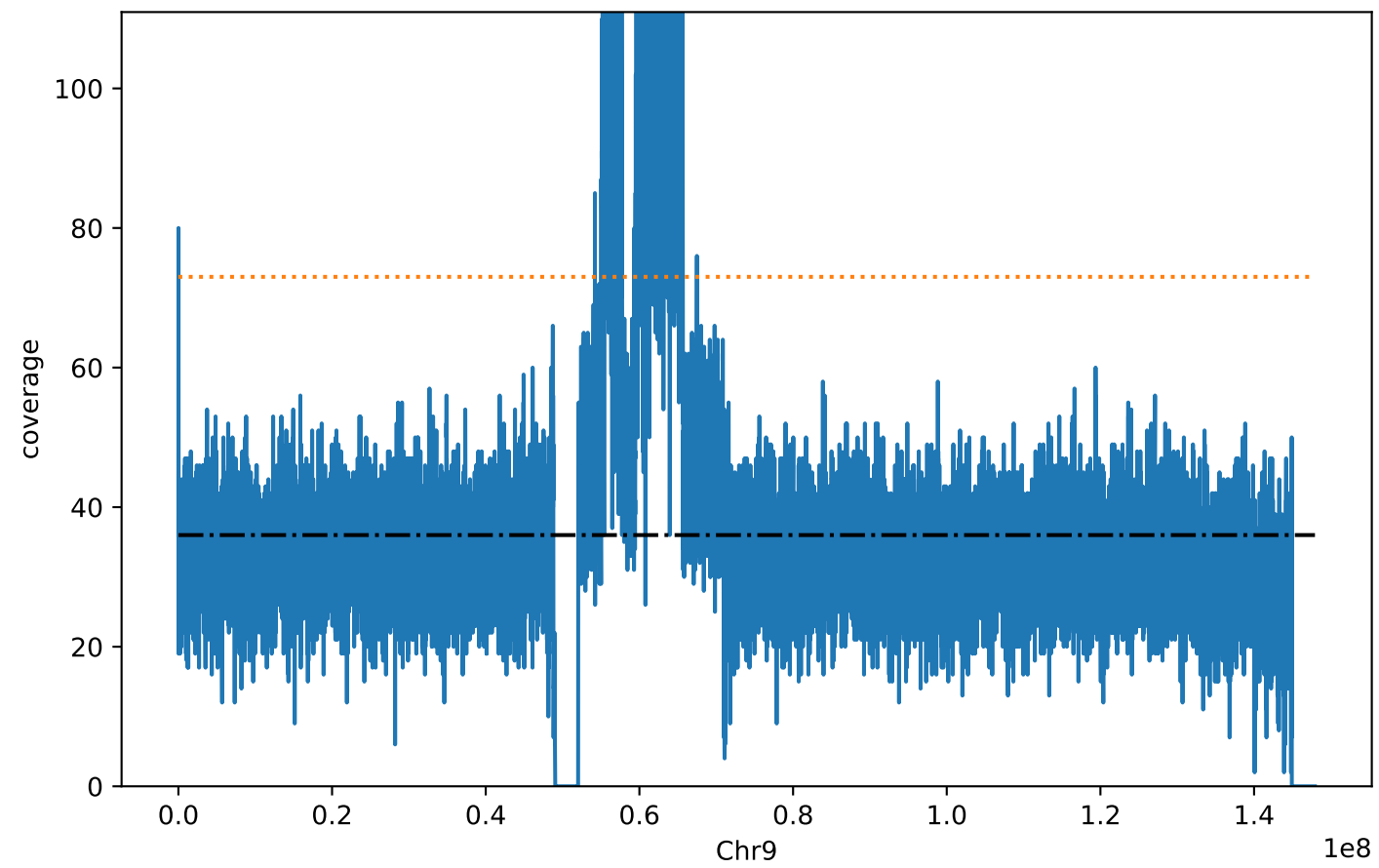
T2T V.1.0



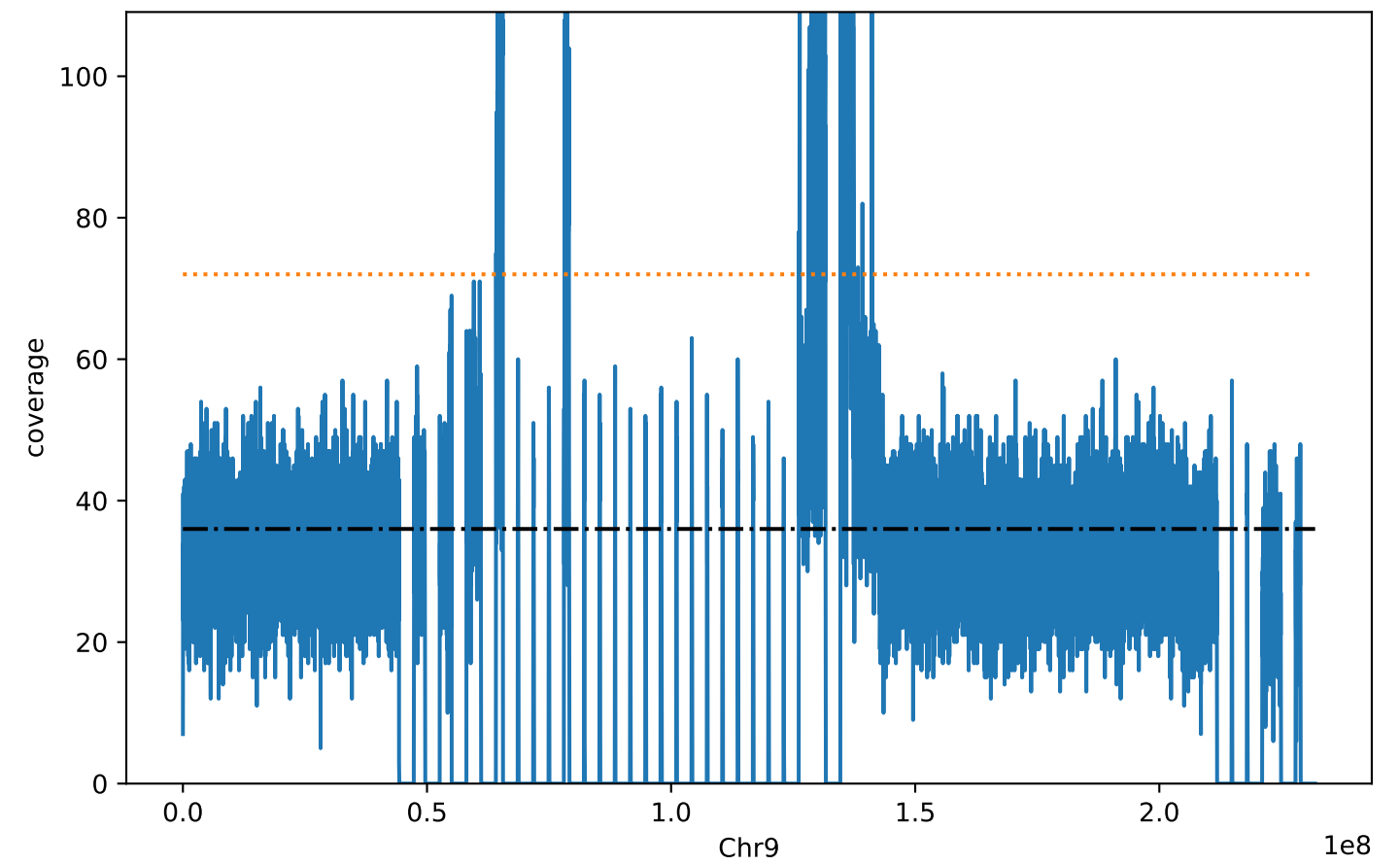
GRCh38 Reference Genome



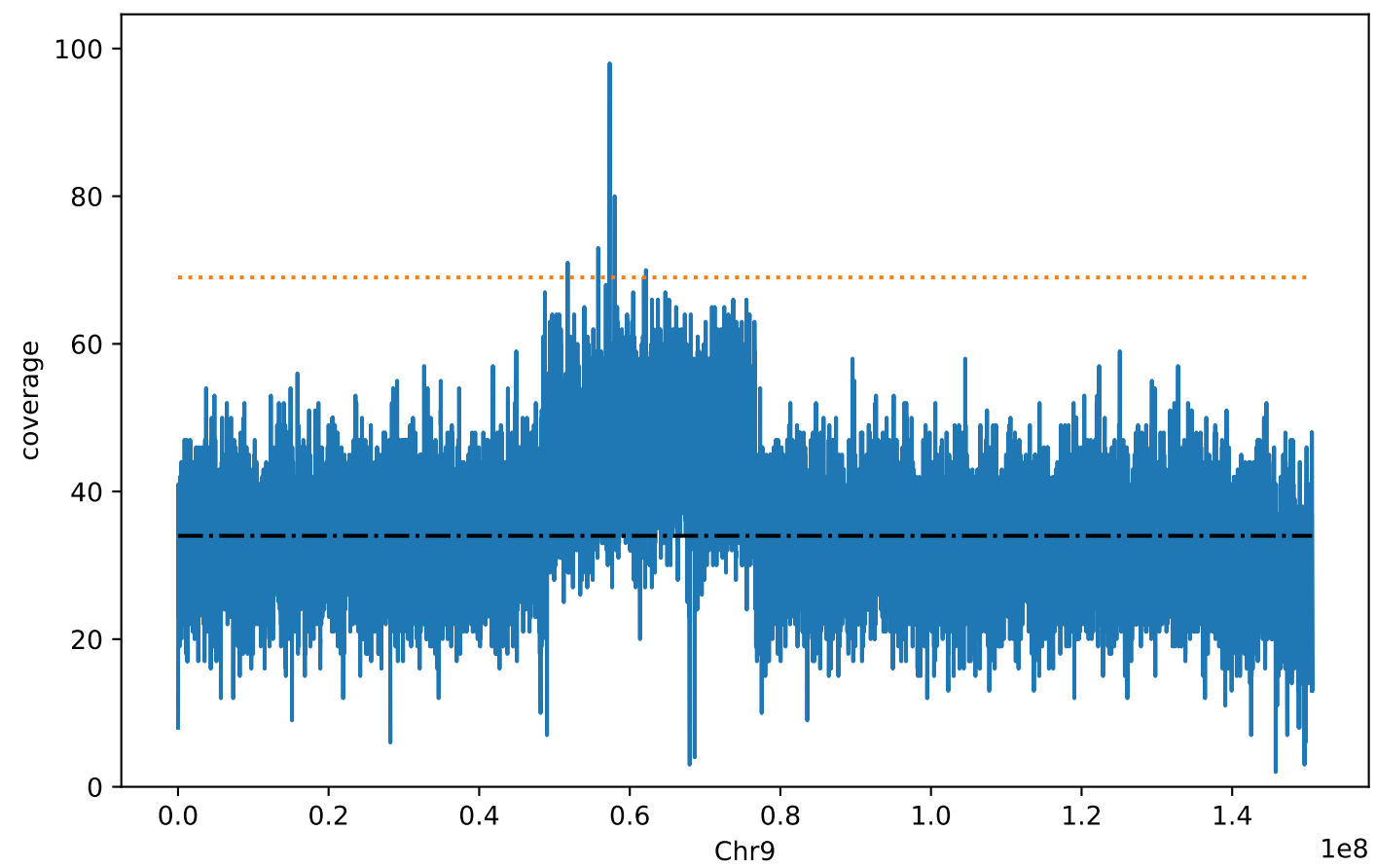
GALA



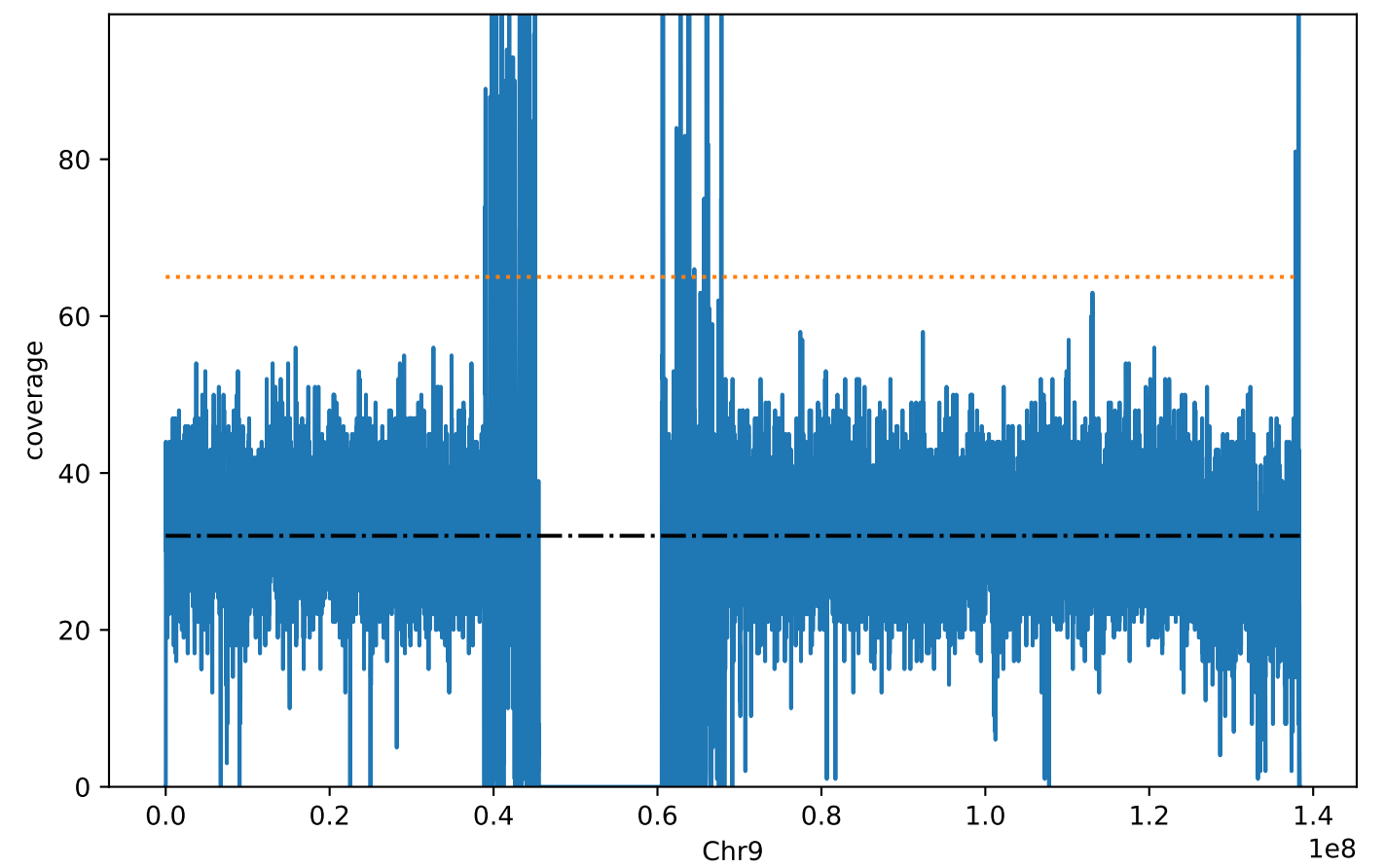
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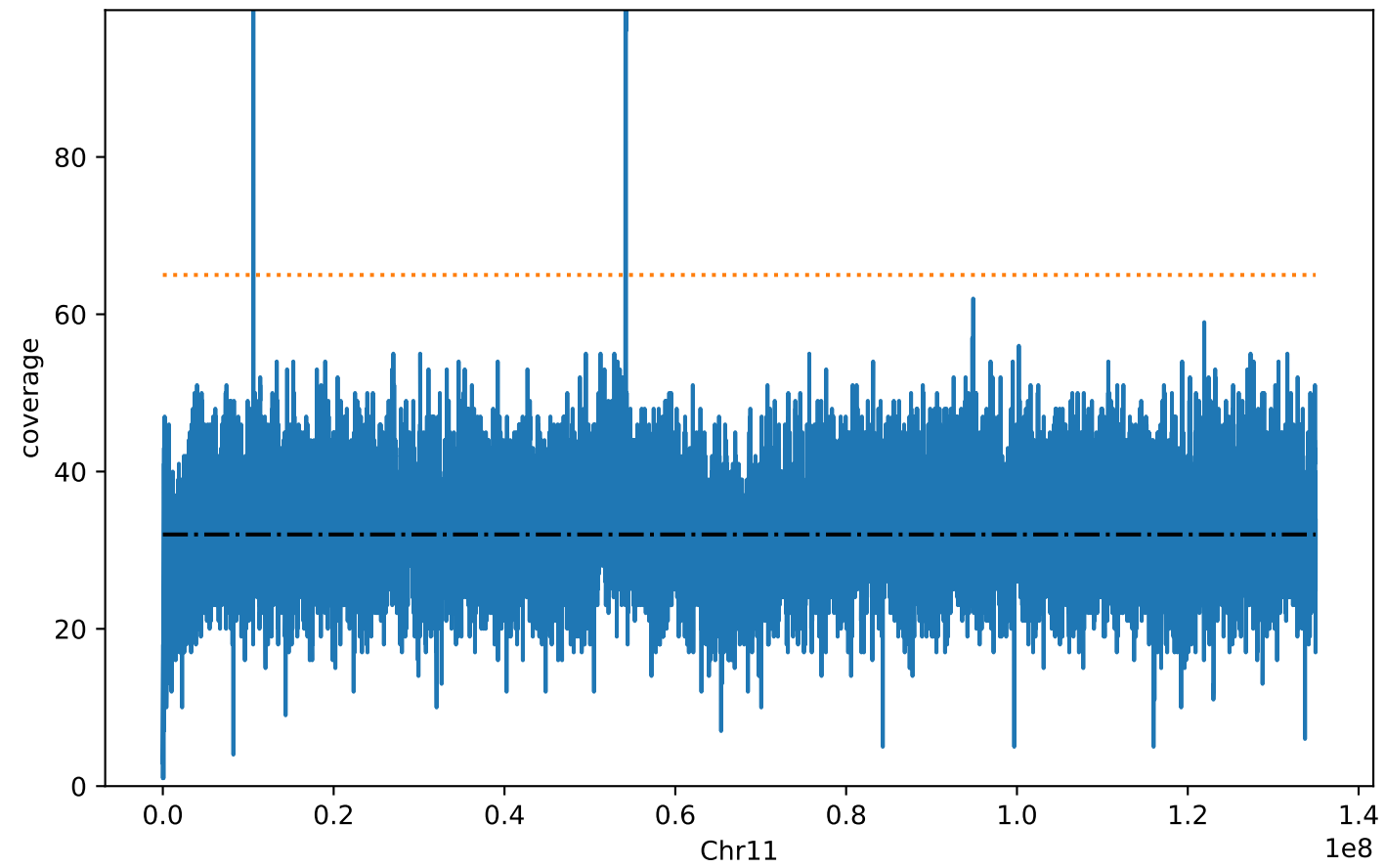
T2T V.1.0



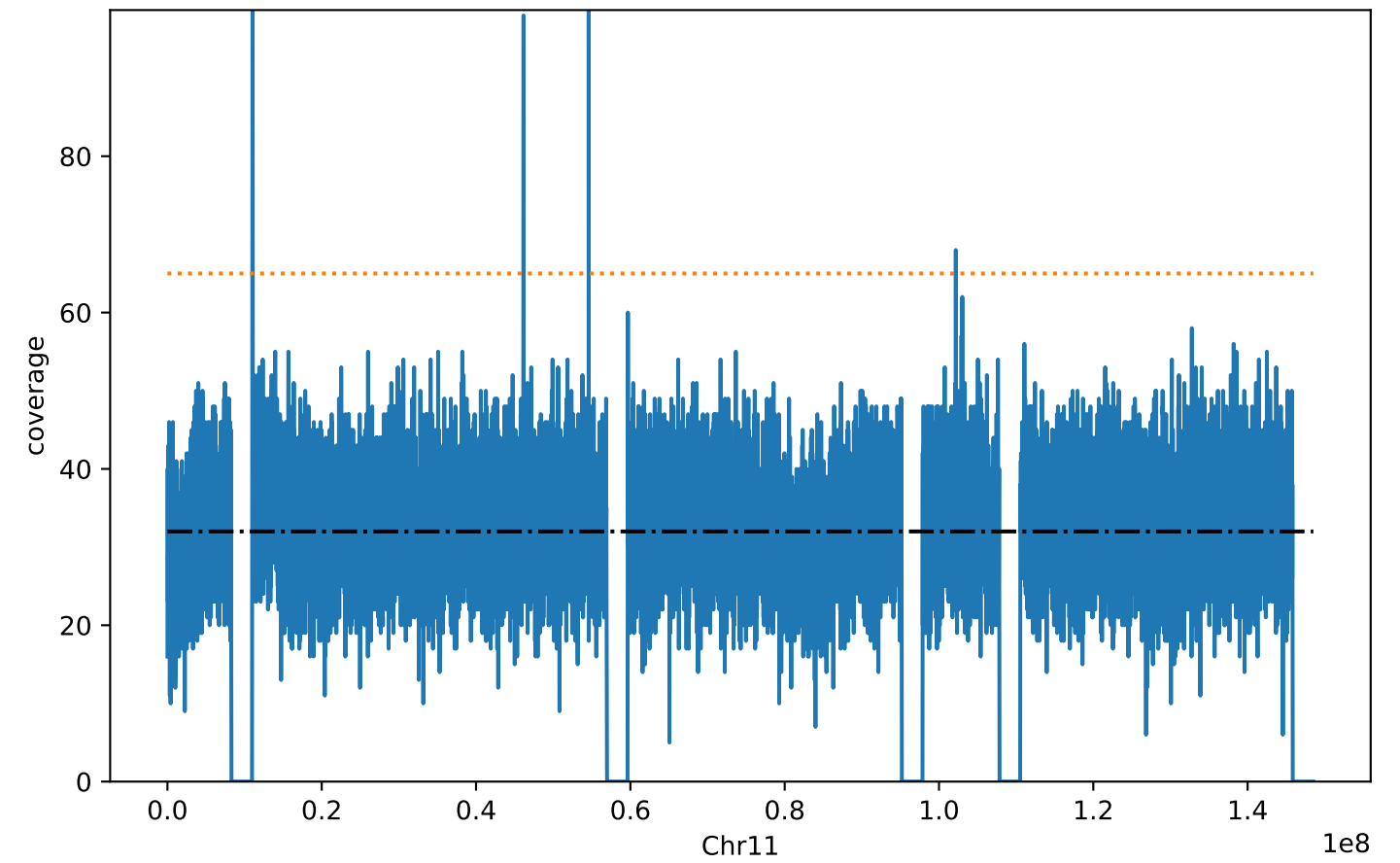
GRCh38 Reference Genome



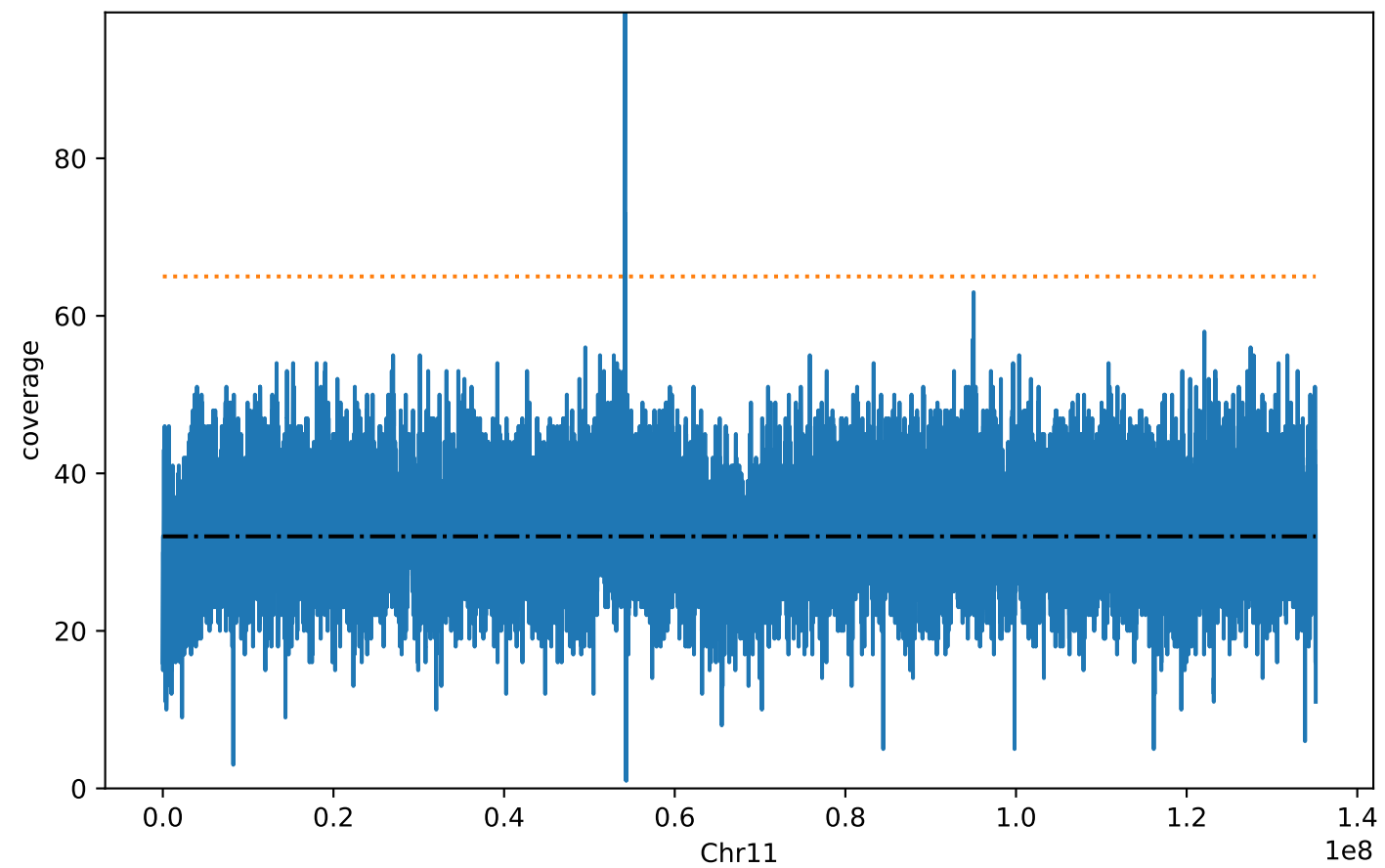
GALA



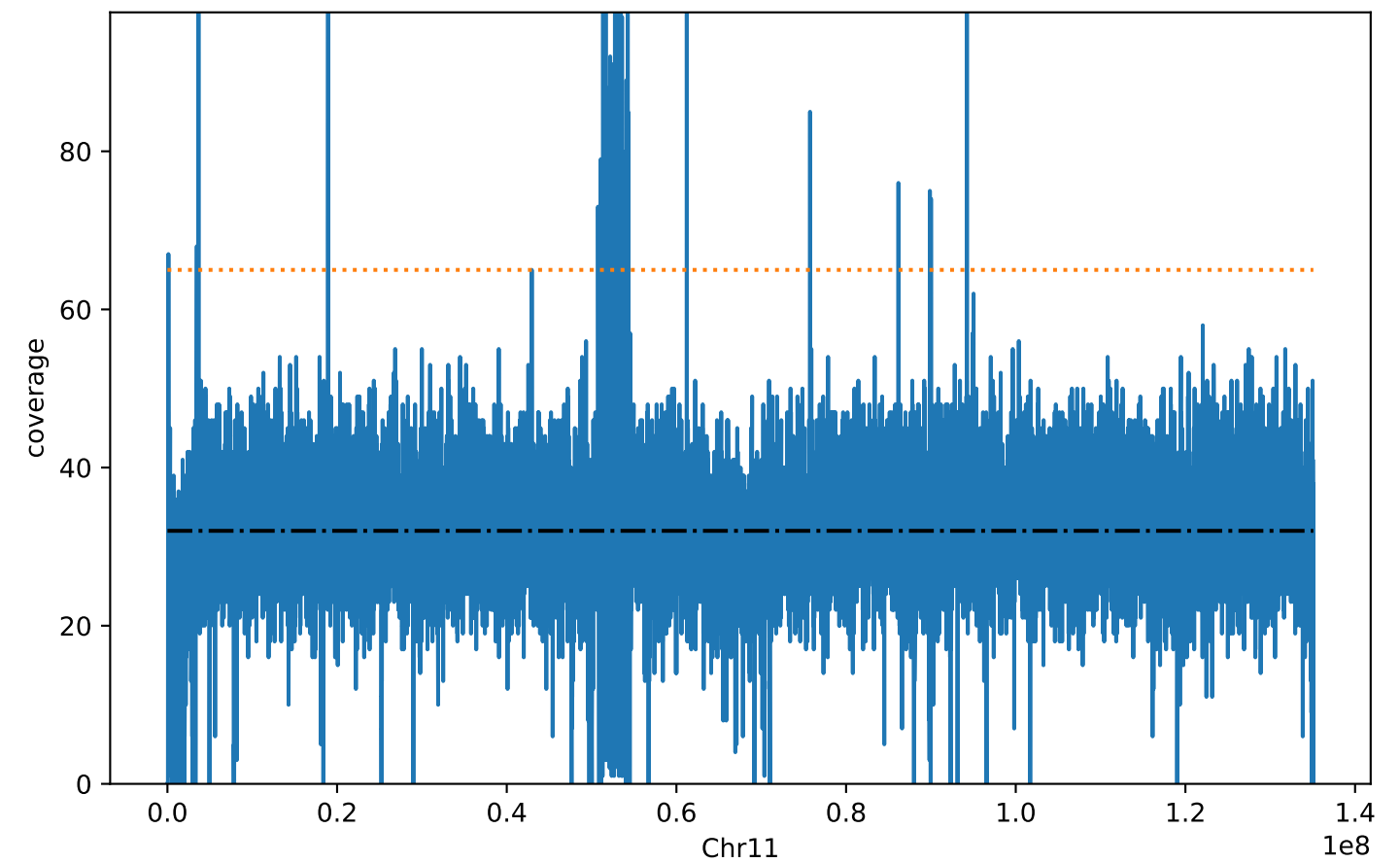
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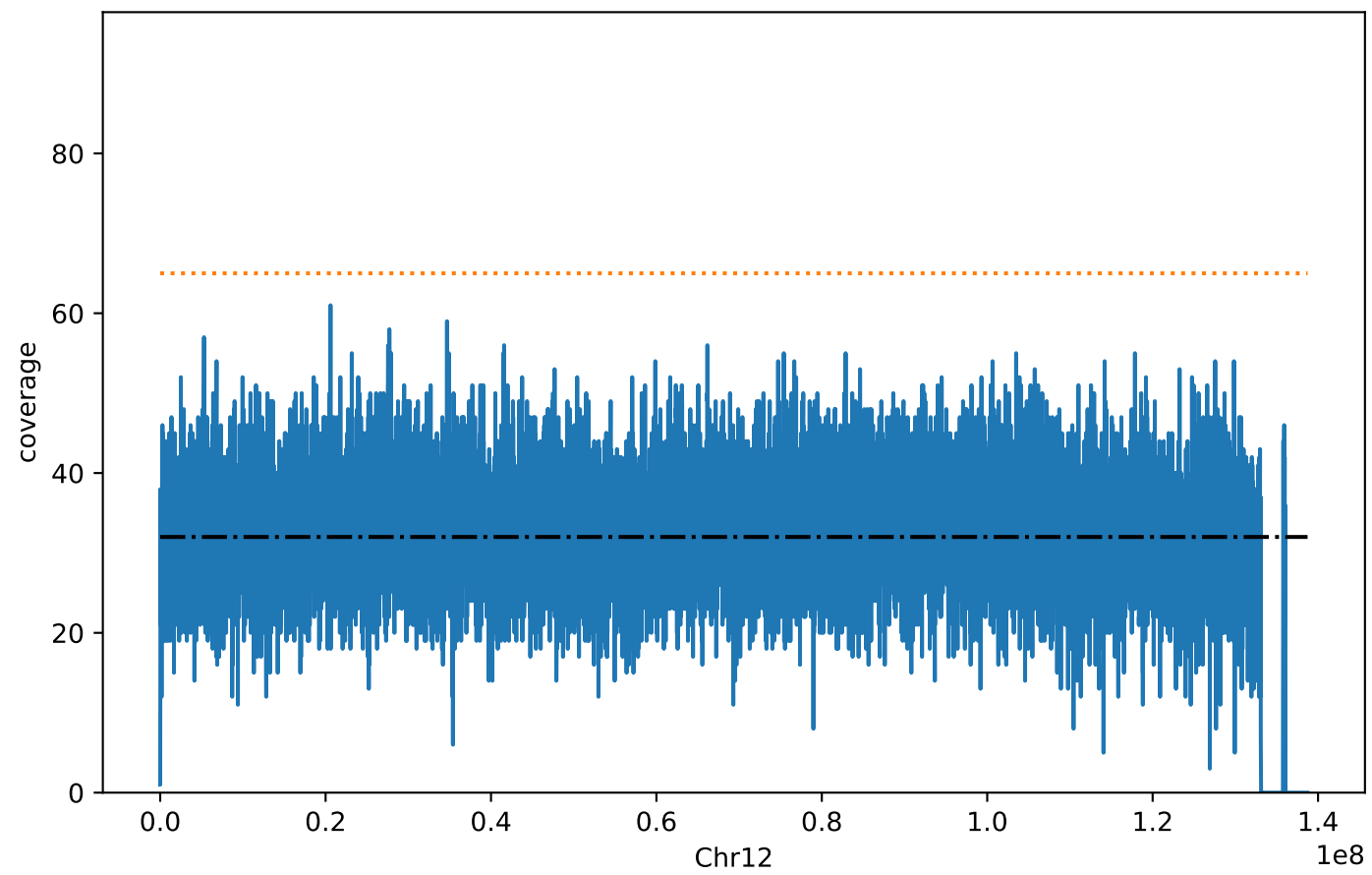
T2T V.1.0



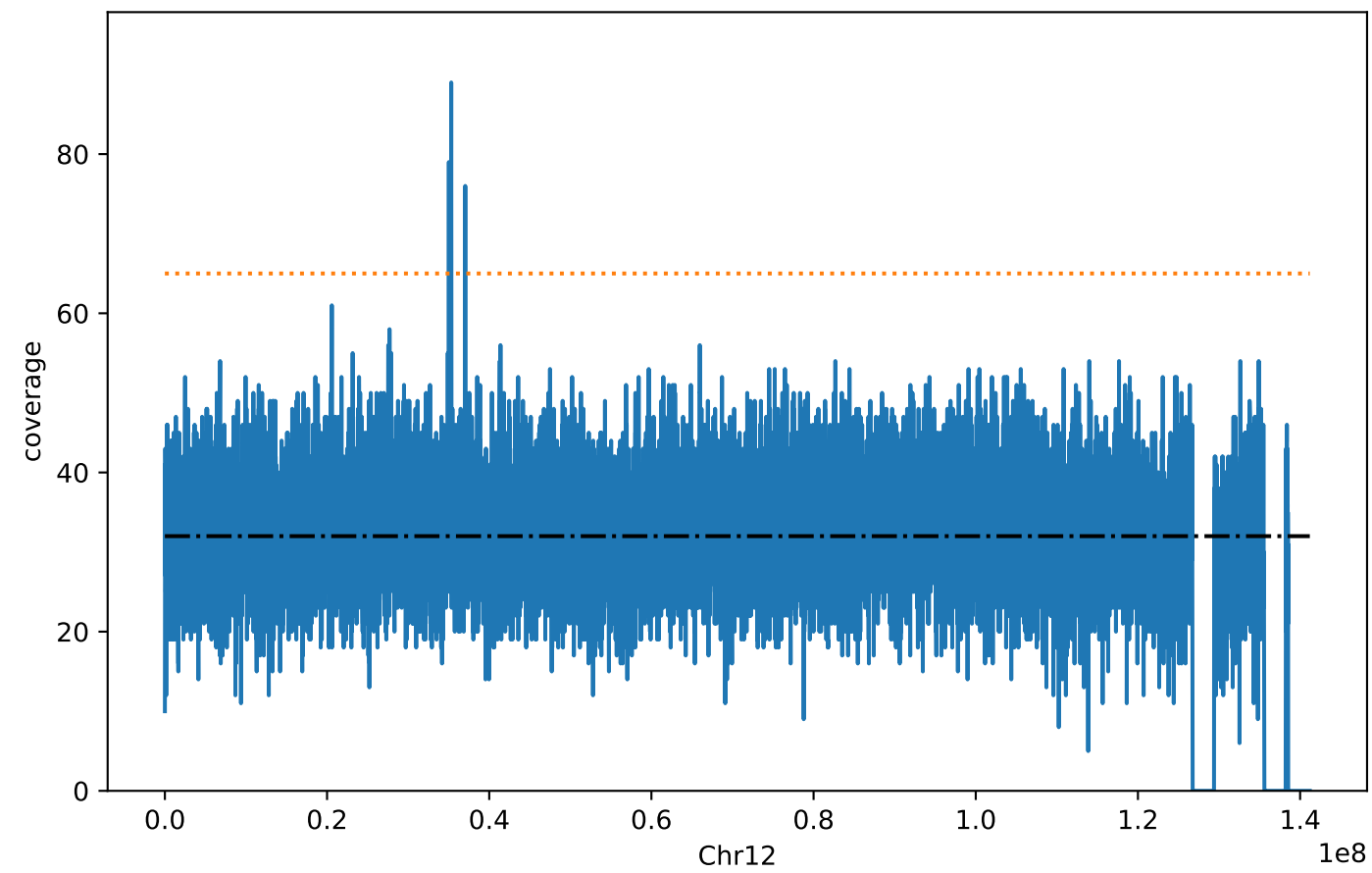
GRCh38 Reference Genome



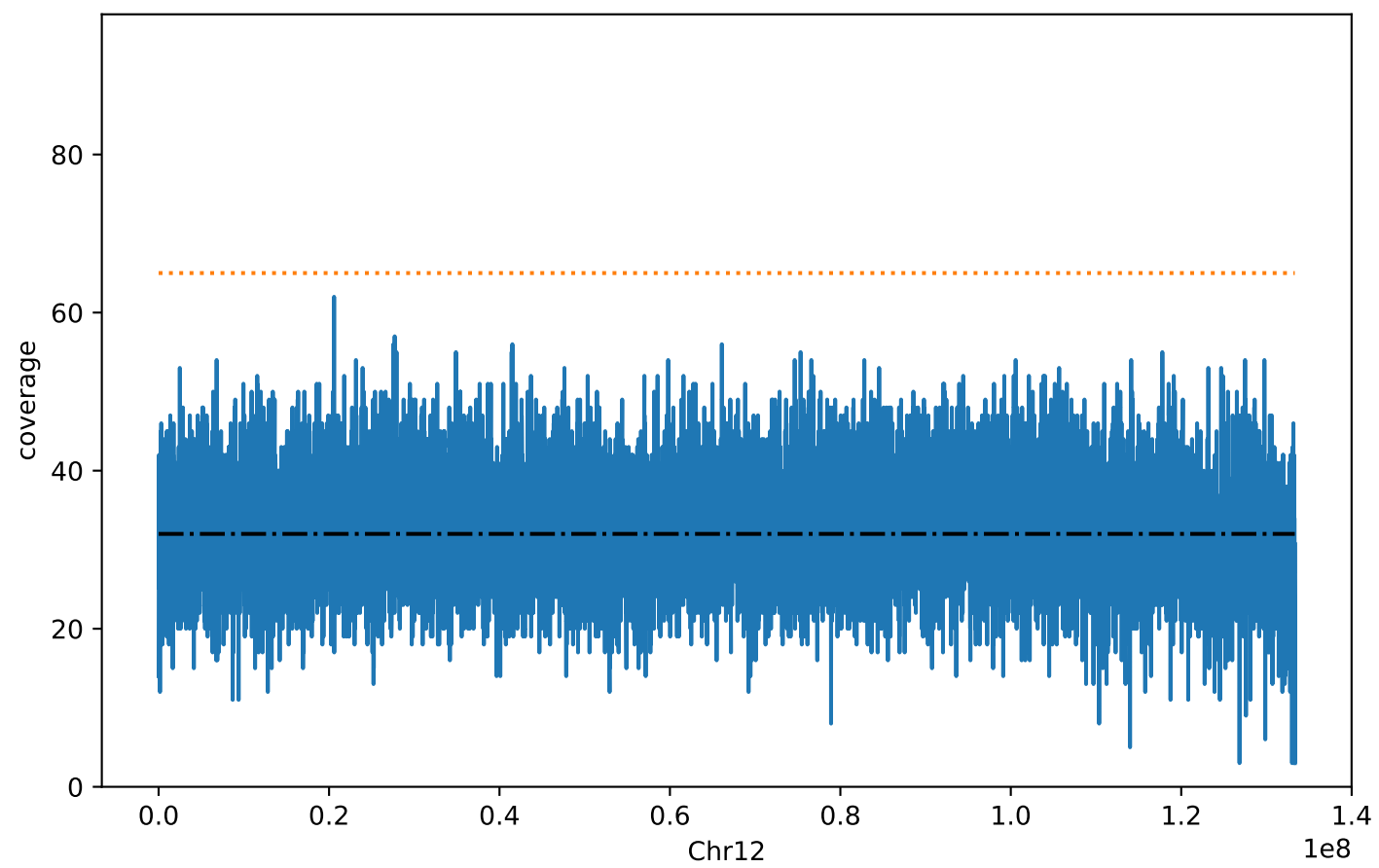
GALA



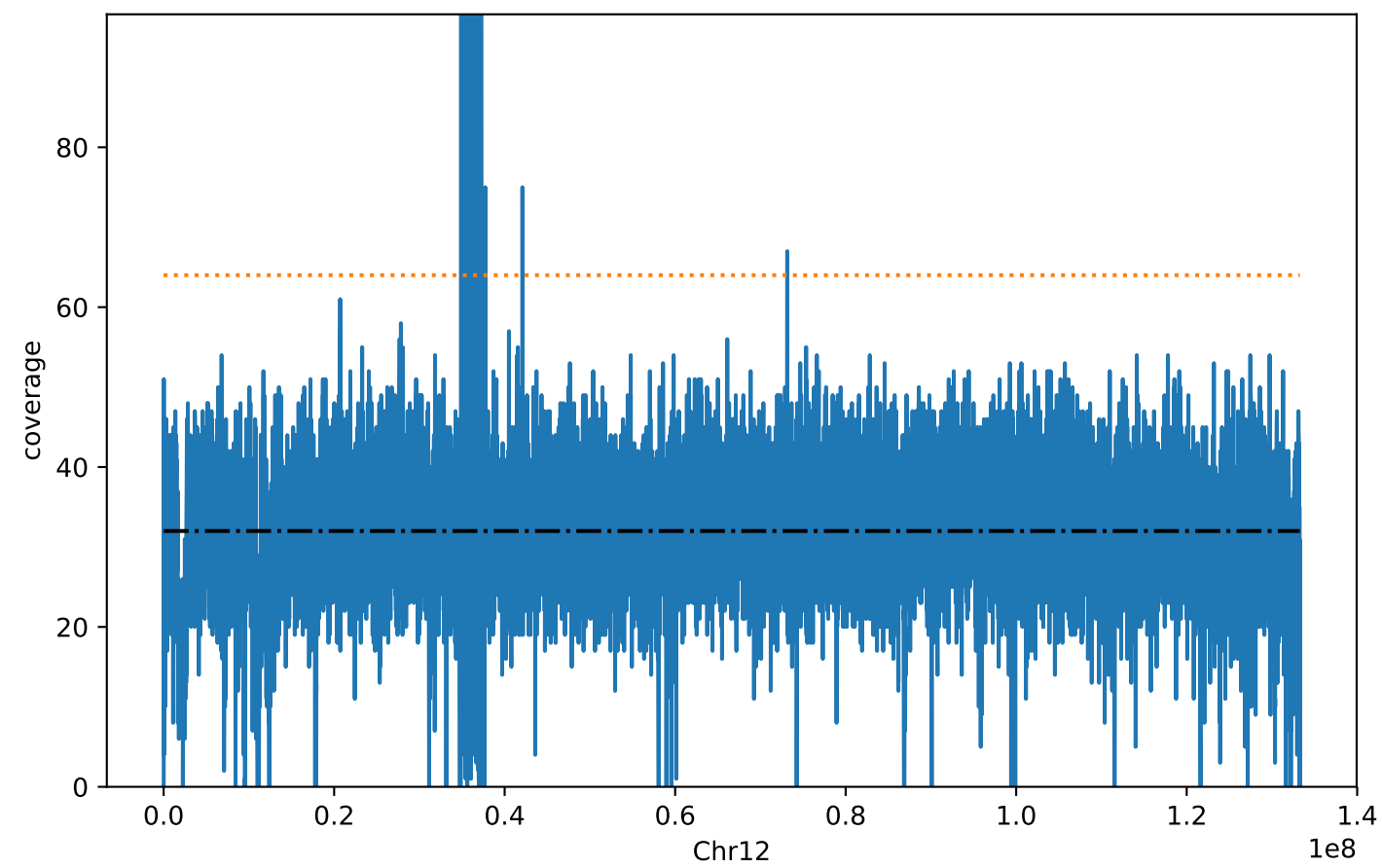
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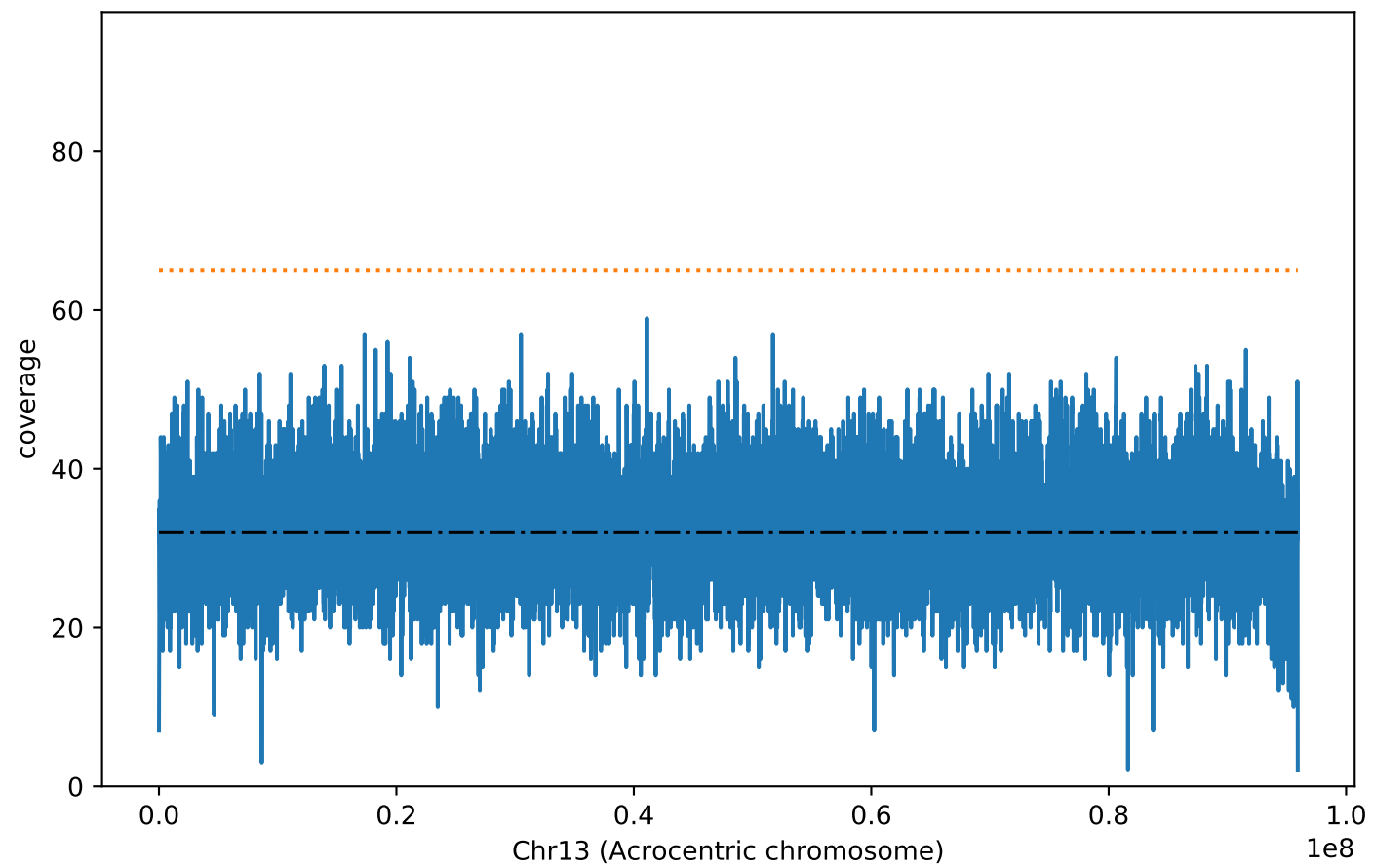
T2T V.1.0



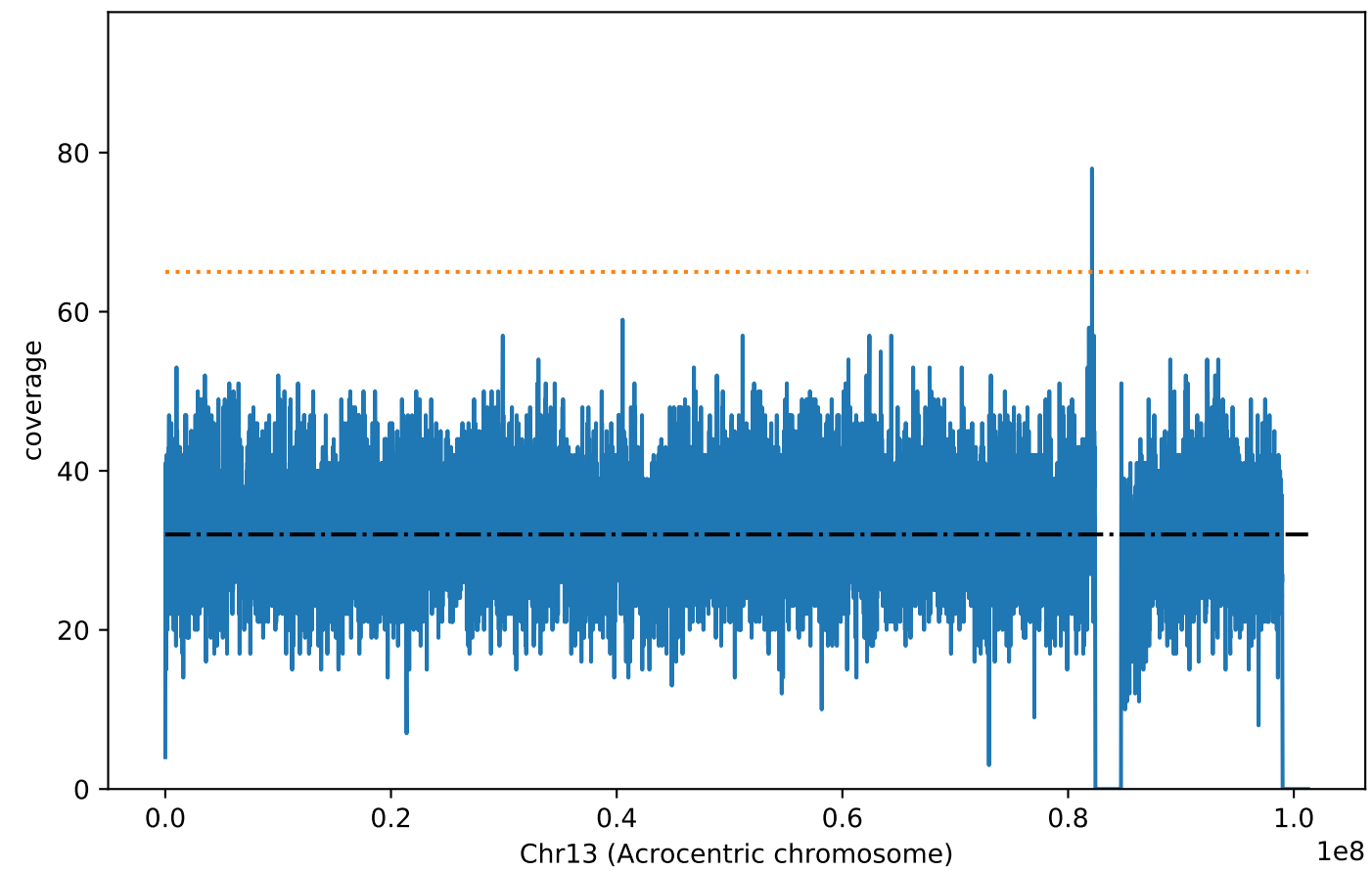
GRCh38 Reference Genome



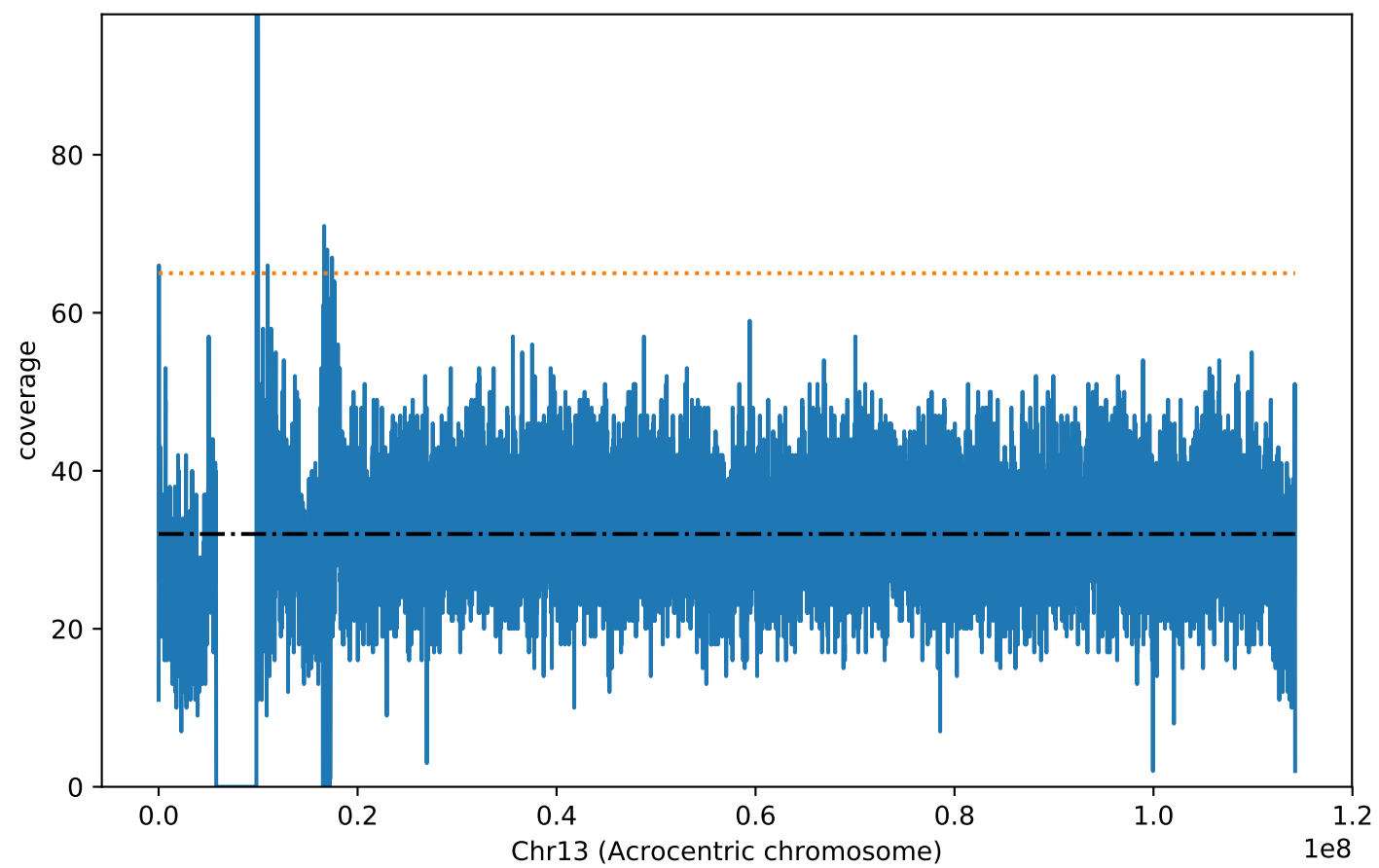
GALA



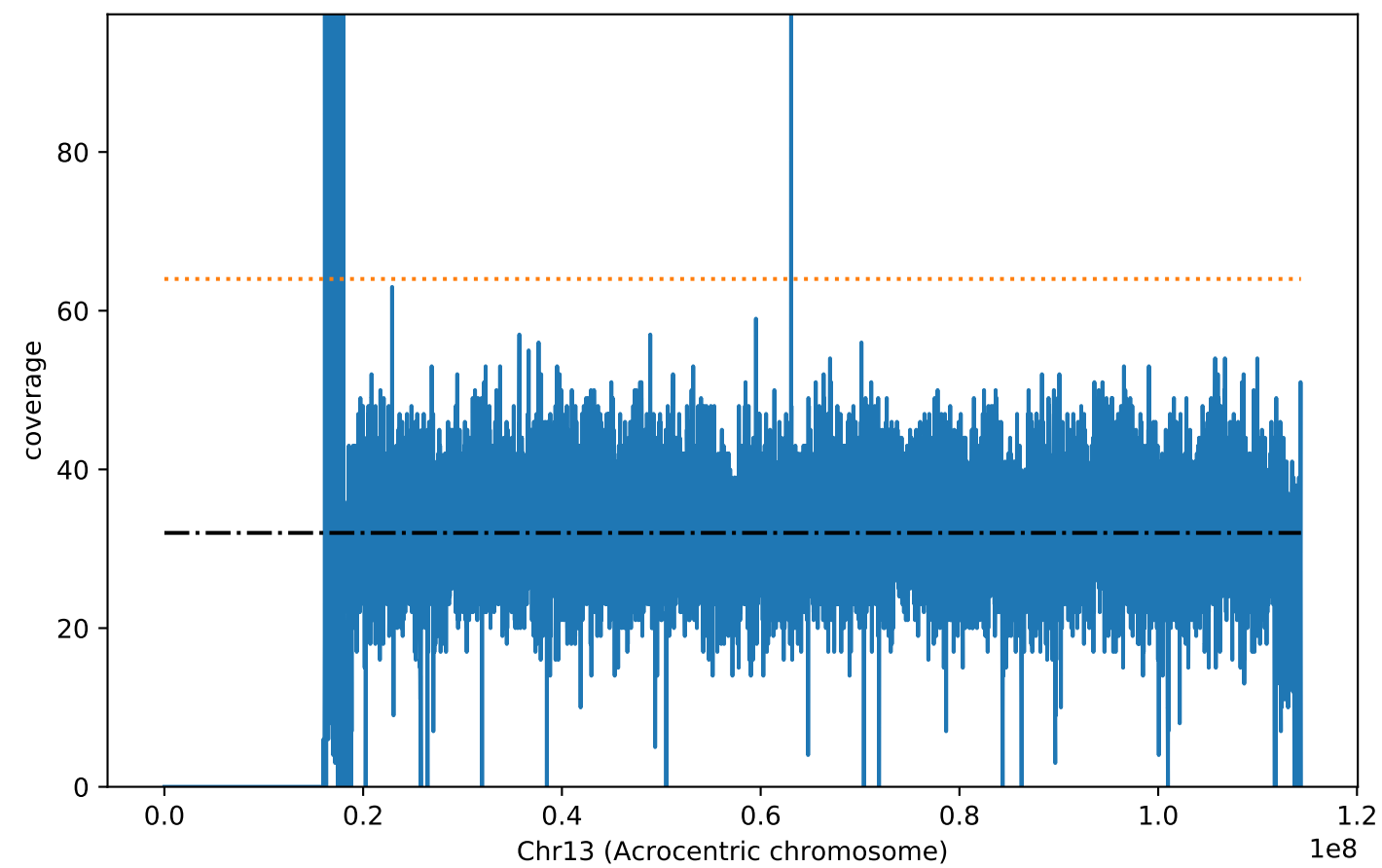
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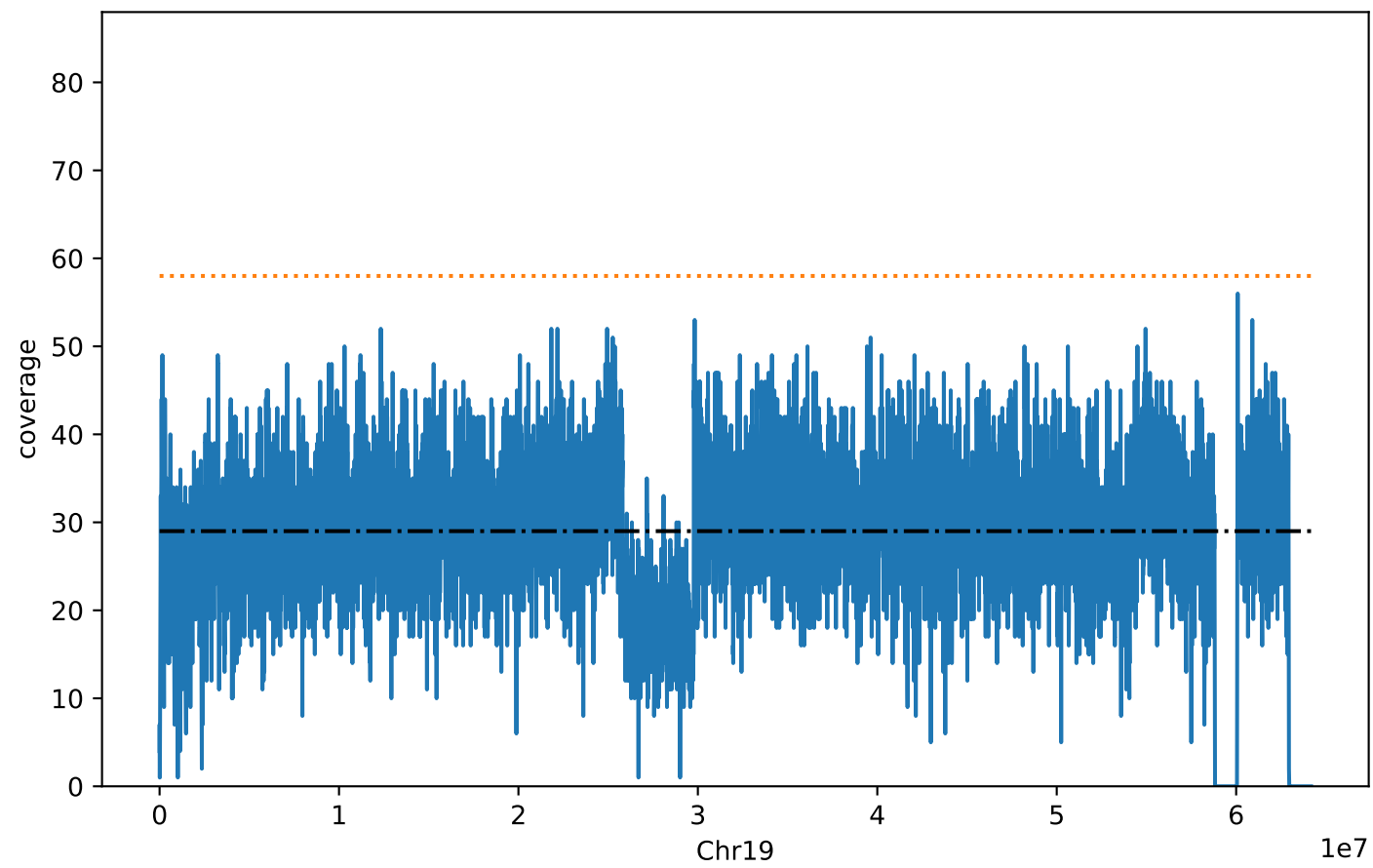
T2T V.1.0



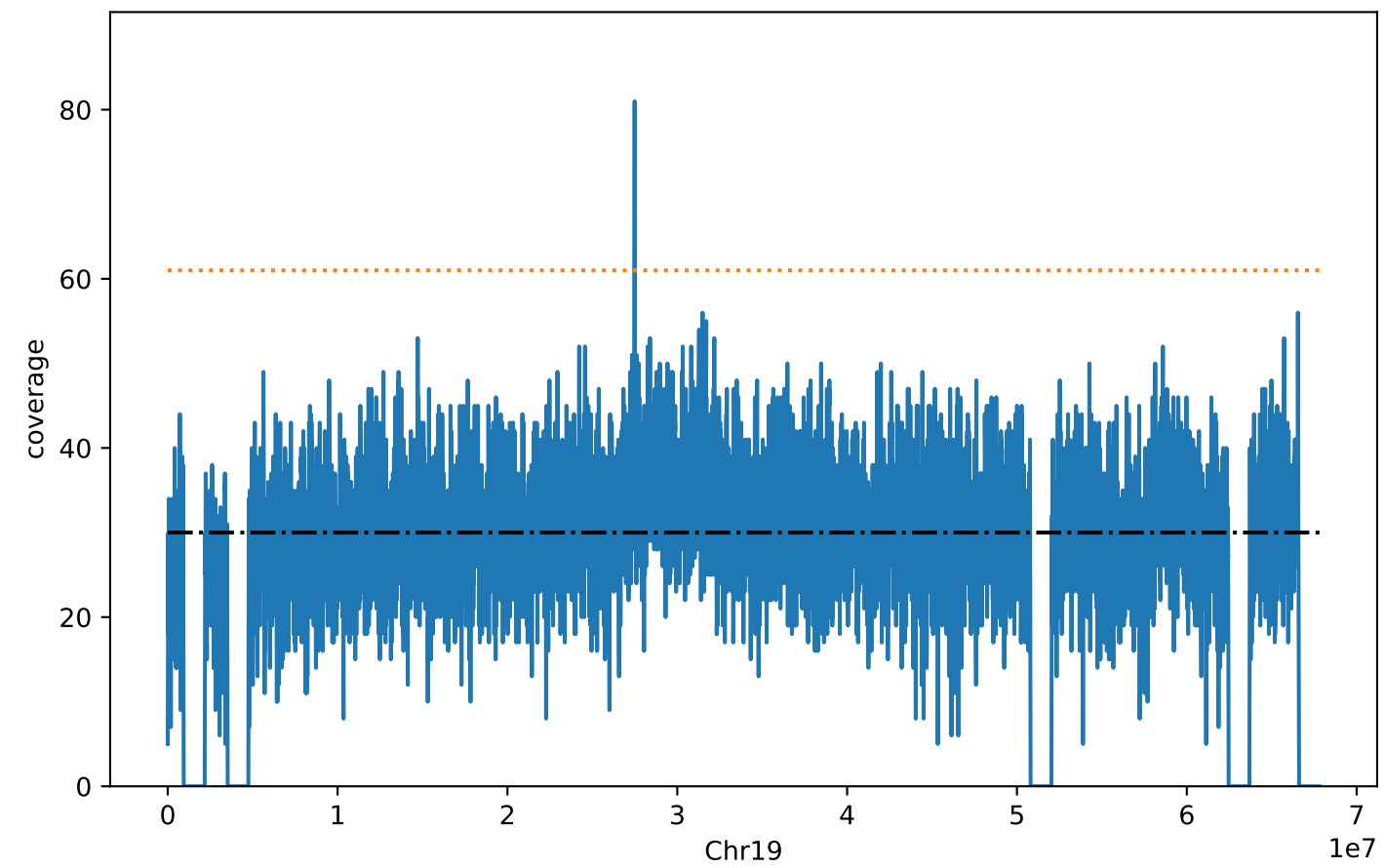
GRCh38 Reference Genome



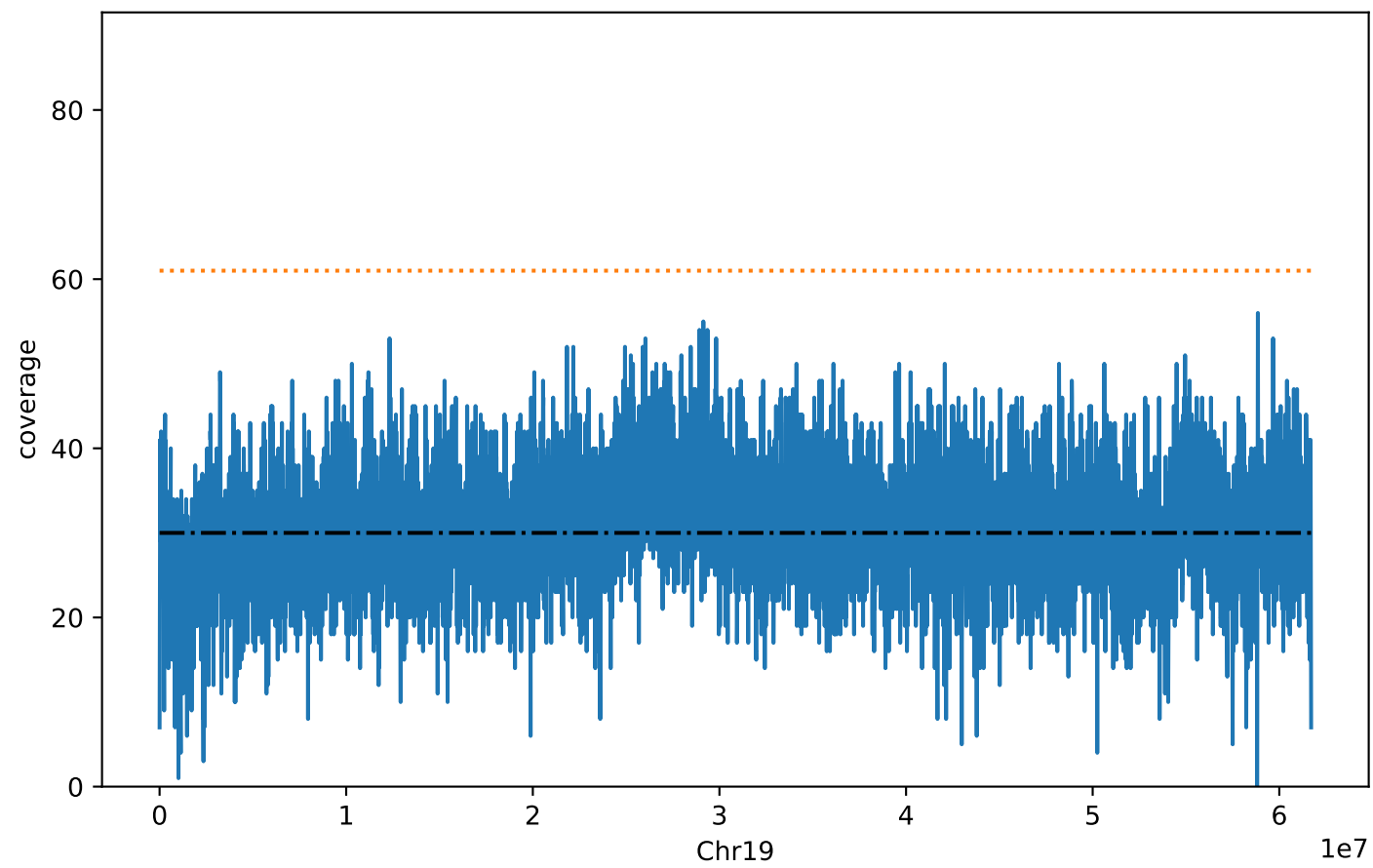
GALA



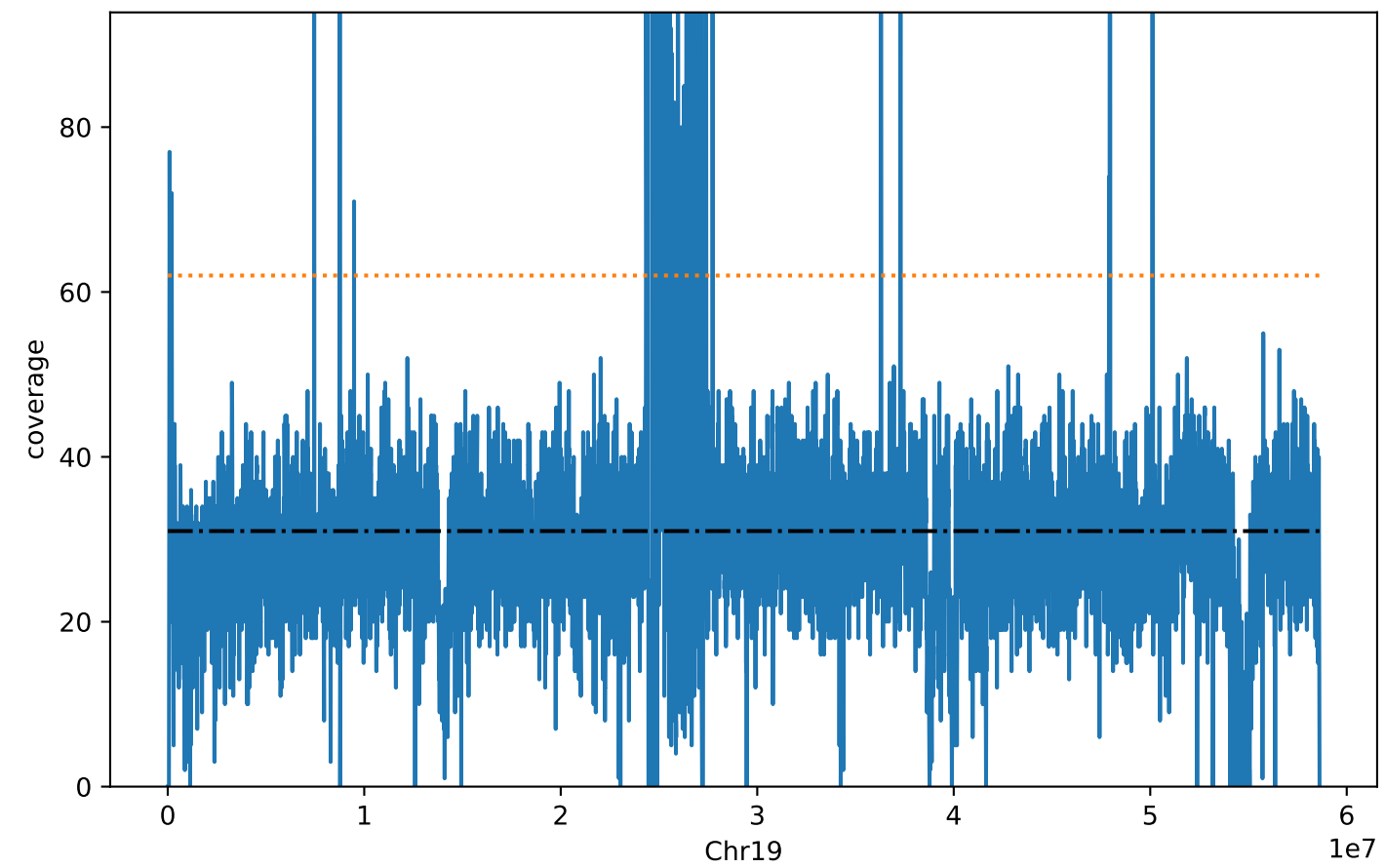
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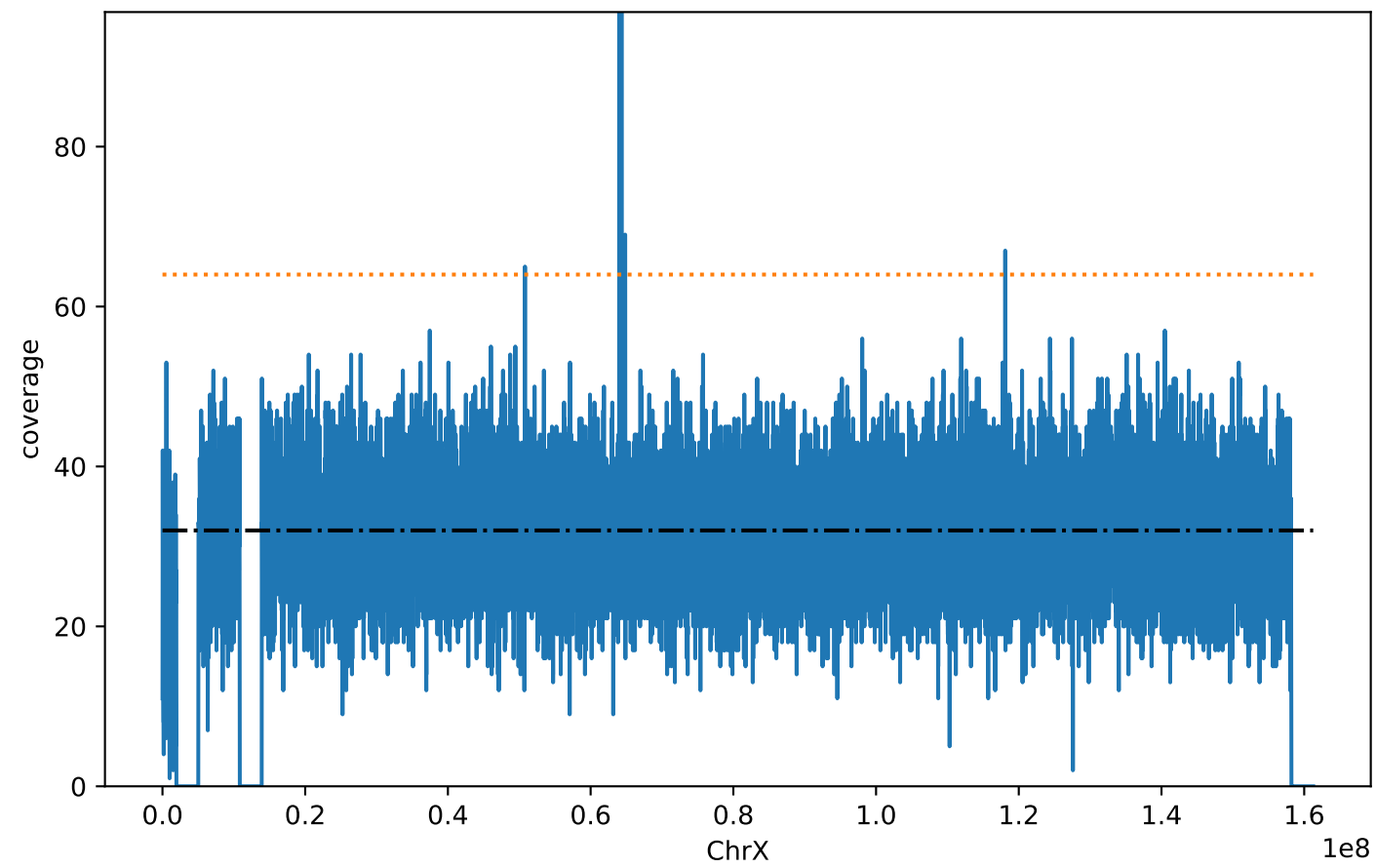
T2T V.1.0



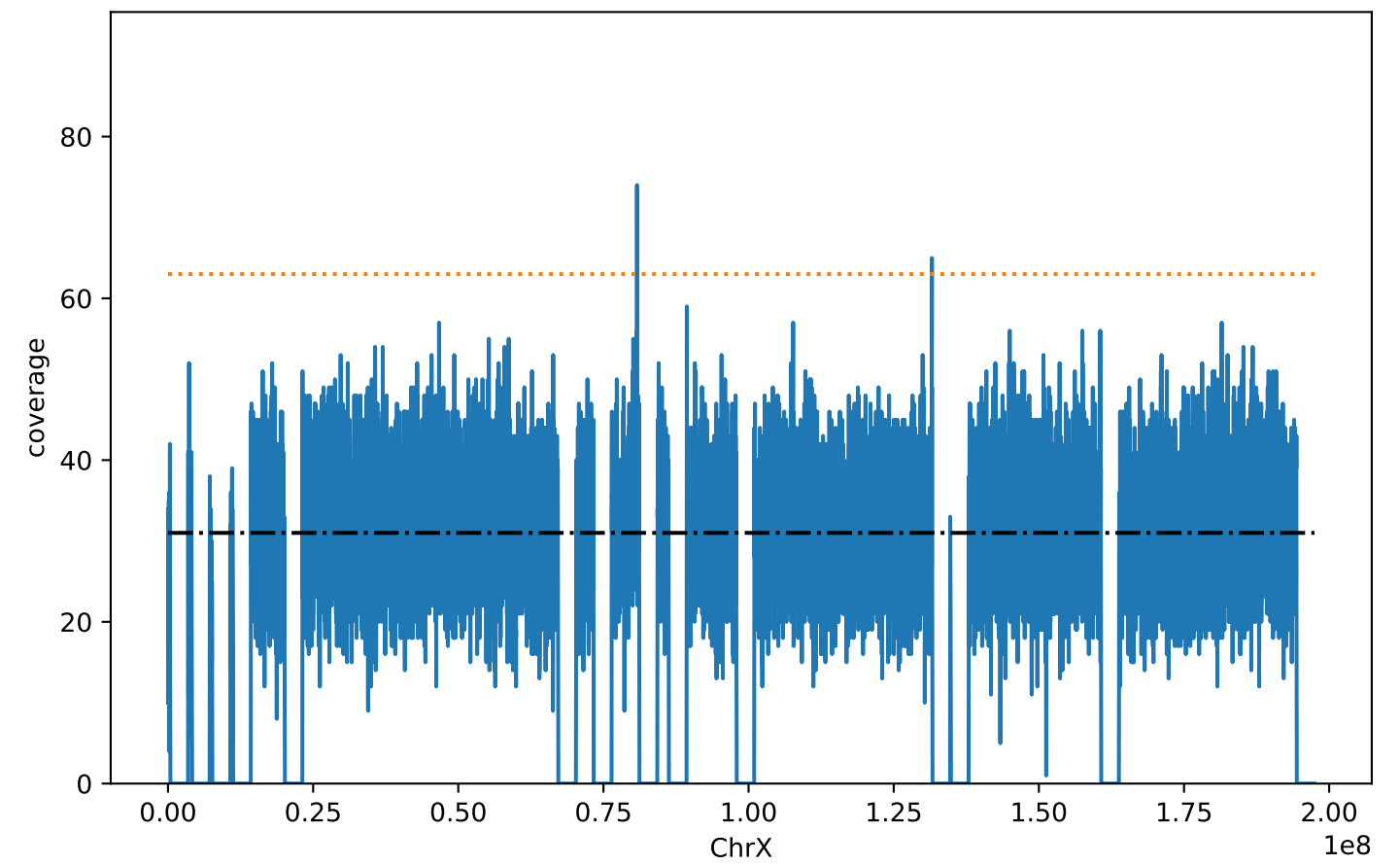
GRCh38 Reference Genome



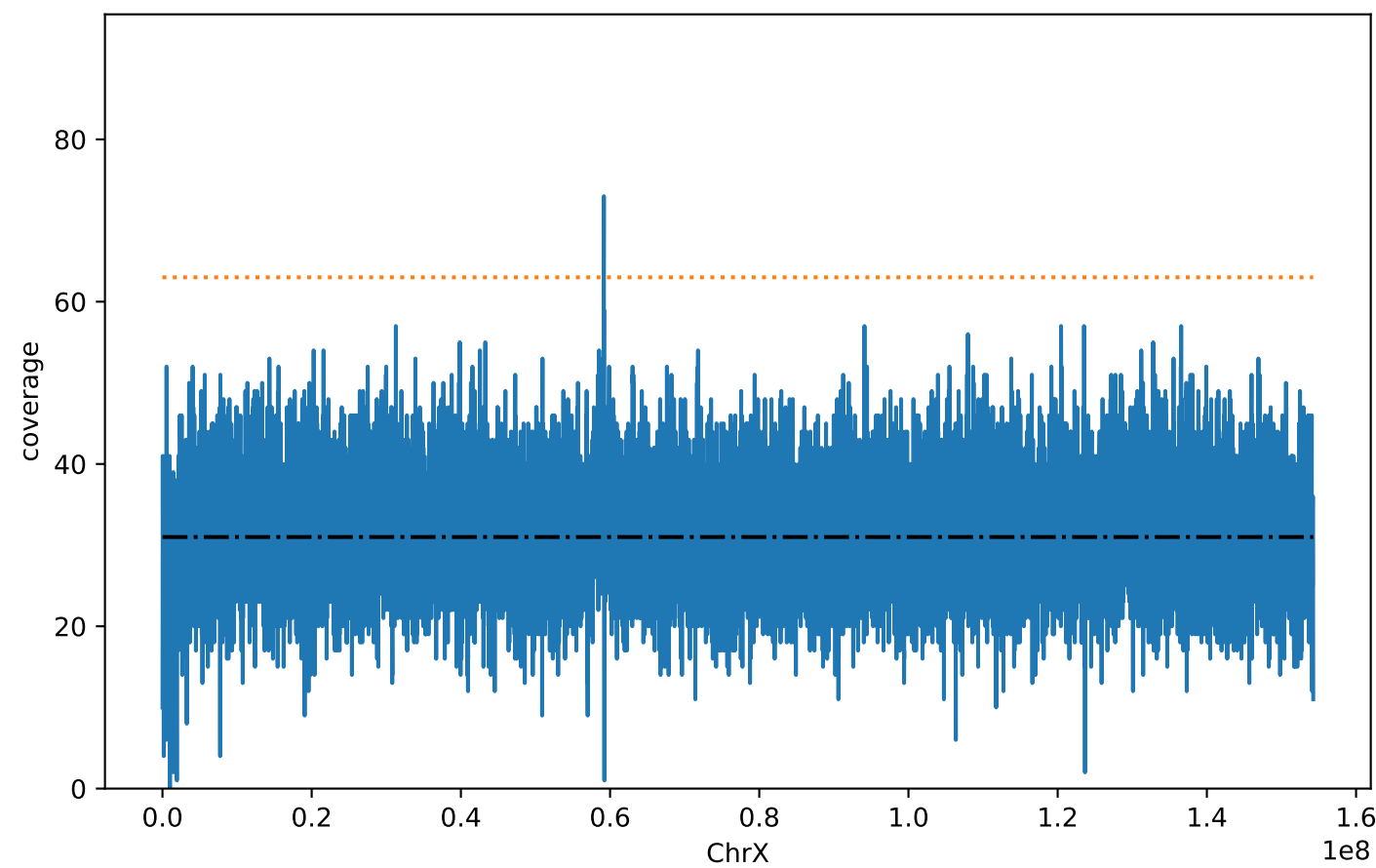
GALA



Canu



T2T V.1.0



GRCh38 Reference Genome

