

GEDmatch® Autosomal One-to-one Comparison - V1.0

Software Version Nov 12 2019 14:54:30

[Here](#) is a link to a useful YouTube video on using the One to One DNA comparison tool.

Individual marker indications:

Base Pairs with Full Match	
Base Pairs with Half Match	
Match with Phased data	
Base Pairs with No Match	

Validity of segments:

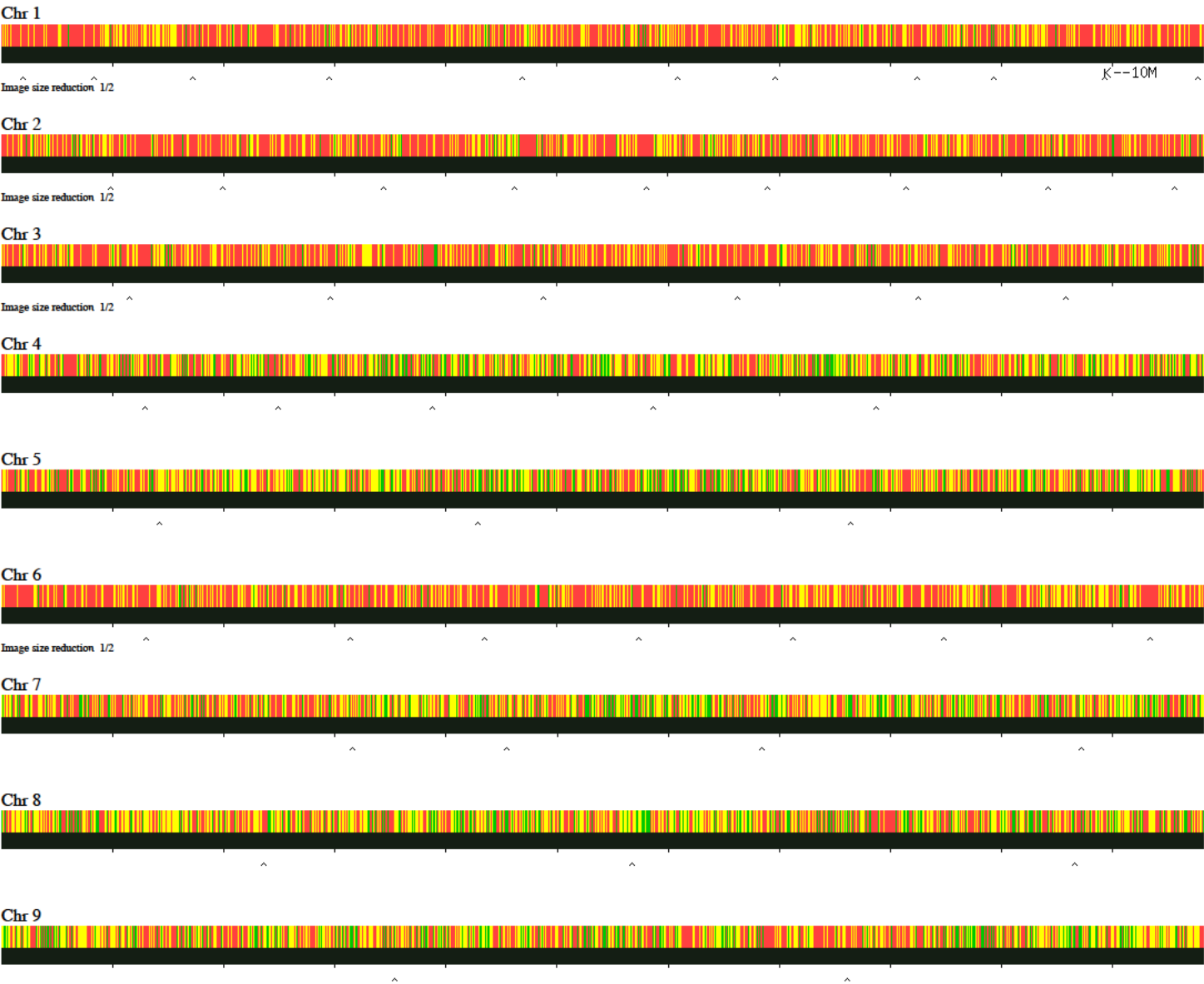
Significant	
Moderate	
Low	
Insignificant	
Large gap between adjacent SNPs	
No Match	
Centromere	

Comparing Ki [REDACTED] (*target.stripy.genome.3) [23andMe] and [REDACTED] (*22stripy.genome.1) [23andMe]

Minimum segment threshold size to be included in total = 25 SNPs

Minimum segment cM to be included in total = 0.7 cM

Mismatch-bunching Limit = 25 SNPs



Chr 10



Chr 11



Chr 12



Chr 13



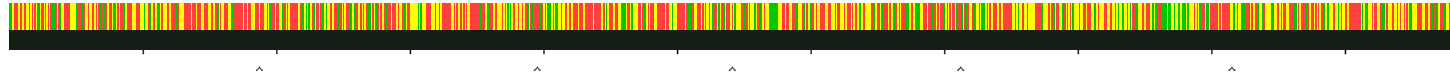
Chr 14



Chr 15



Chr 16



Chr 17



Chr 18



Chr 19



Chr 20



Chr 21



Chr 22



No shared DNA segments found

514763 SNPs used for this comparison.

Comparison took 0.345 seconds.

CPU time used: 0.156 cpu seconds.

