

Supplemental Table 2: Comparison of MP sequencing data from the average of 8 unamplified genomic DNA samples and 8 WGA samples.

	Total Number of MPs	% Replication	Bridged Coverage	Base Pair Coverage	Discordant MPs No Filter	Discordant MPs + Filter	% Discordant MPs + F	% MP noFilter NO Associates	% MP + Filter NO Associates
gDNA	96,772,473	19%	33 x	5 x	7,405,699	493,046	6.89%	82.88%	98.81%
WGA	91,023,609	21%	30 x	4 x	4,949,214	434,917	8.85%	91.13%	98.76%

