

Table S2 Summary of SNP/indel detection with PGM and MiSeq

	All reads				Reads with mapping quality above 40			
	Total SNPs and indels	Indels	% indels of total calls	Indels/1kb/sample	Total SNPs and indels	Indels	% indels of total calls	Indels/1kb/sample
PGM	9685	5544	57.2%	1.340	7574	5288	69.8%	1.270
MiSeq	3818	395	10.3%	0.096	3553	386	10.9%	0.093