

Table S1

<b>Fibroblast (GM02639) sequencing details (data run through Porechop)</b>			
<b>Sample Name</b>	<b>37°C</b>	<b>40°C - 24h</b>	<b>Rec</b>
Read count	597103	1528595	2349238
Base count	8003216090	3889197457	7184368217
Mean read length	13403.41	2544.30	3058.17
Reads mapped by bowtie2	572621	1285361	2010421
Aligned reads (%)	95.90	84.09	85.58