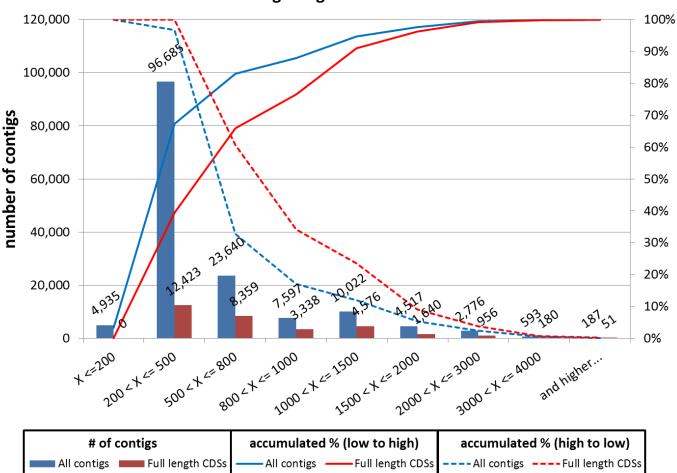
Figure S1: Distribution of contigs size obtained from the genome reference (hybrid assembly).



Contigs length distribution