Supplemental Fig. S4. GC content (a) and CpG frequency (b) of pig genomes. The x-axis indicates GC content (%) or CpG frequency and the y-axis indicates the proportion of the bin number divided by total windows. We used 10 kb, non-overlapping sliding windows across the assemblies. The data shows that the GC content of nine pig assemblies is around 41.88%, similar to that of the publicly available Tibetan wild boar (41.83%) and reference genomes (41.70%).