



Figure S5. Distribution of the number of SNPs in 1 Mb sliding windows in cattle (A) and sheep (B). Bars in A and B indicate the frequency of sliding windows containing various number of SNPs out of the genome-wide distribution, i.e., 38,610 SNPs of cattle and 47,502 SNPs of sheep data, respectively (details in Table 1, S3). The bars in red (black) colours show the mean \approx median (mode) numbers as 17 (18) and 19 (20) of SNPs for cattle and sheep data, respectively.