





Supplementary Figure S3. Manhattan plots for genomic distribution of zF_{ST} values estimated between the Cornish White (CRW), Plymouth Rock White (PRW) and Ushanka (USH) breeds. Values for the X-axis are chicken autosomes (breadth of autosomes corresponds to their length) and those for the Y-axis are zF_{ST} values. SNPs were plotted relative to their positions within each autosome. The threshold, which was estimated as the top 0.1% for zF_{ST} values, is indicated by a horizontal line. Breeds: (a) CW vs. PRW; (b) CW vs. USH; (c) PRW vs. USH.