**Suppl. Fig. 4. Fst analysis of individual virulence genes to identify genetic variants that are significantly more commonly in the Americas than in the rest of the world.**
The X axis indicates the nucleotide position in the gene, whereas the Y axis shows the Fst value for each position. a) cagA, b) vacA, c) babA, and d) ppiC. Known domains for each protein were taken from: for CagA, [41]; for VacA, [62]; for BabA, [63]; and for PpiC, [64].