

Figure S1 The distribution of the number of SNPs with large $|iHS|$ values in a 51-SNPs window across the genome. X-axis represents the number of SNPs with $|iHS| > 2$ in a 51-SNPs window. Y-axis represents top ratio across the genome. Top 1% was set as the threshold.

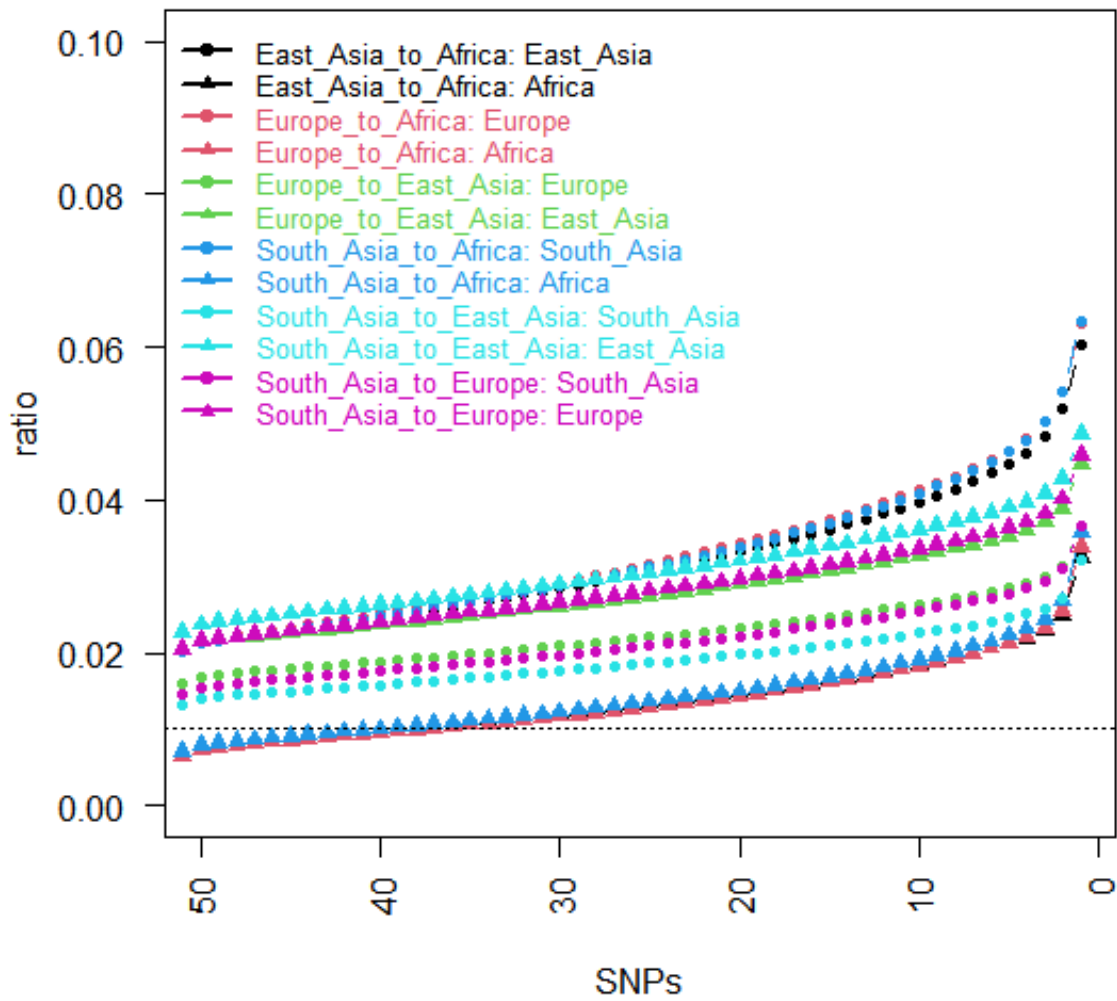


Figure S2 The distribution of the number of SNPs with large $|xpEHH|$ values in a 51-SNPs window across the genome. X-axis represents the number of SNPs with $|xpEHH| > 2$ in a 51-SNPs window. Y-axis represents top ratio across the genome. Top 1% was set as the threshold.