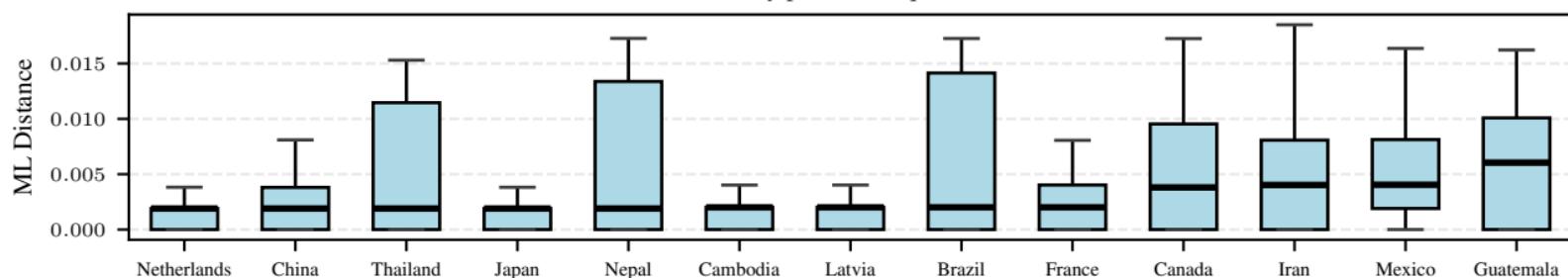


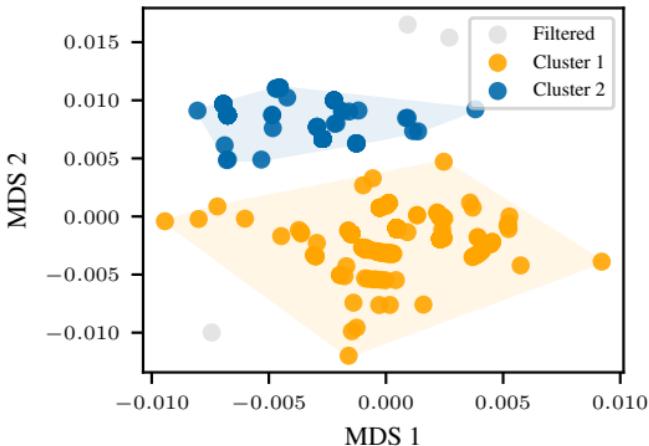
A

## Within-country pairwise sequence distances



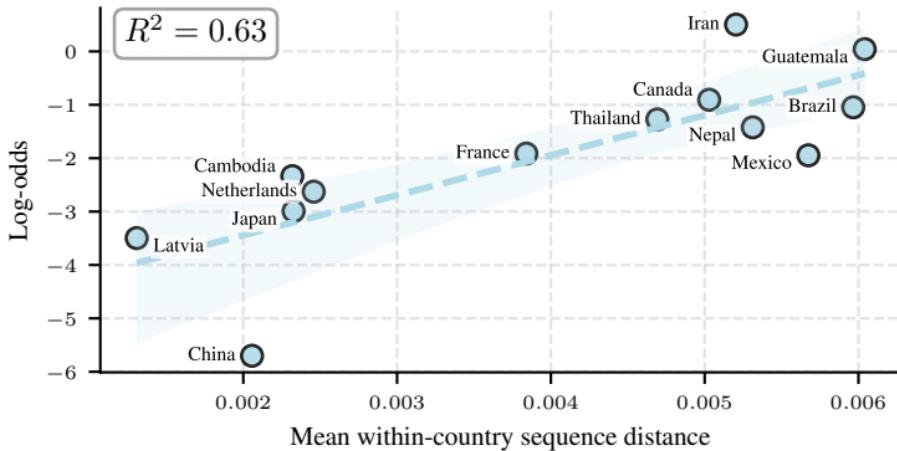
B

## MDS projection



C

## Diversity vs Log-odds



D

## Cluster sample counts over time

