

Supplementary Information

Patterns of Genomic Interrelatedness of Publicly Available Samples in the TB Portals Database.

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and Alex Rosenthal

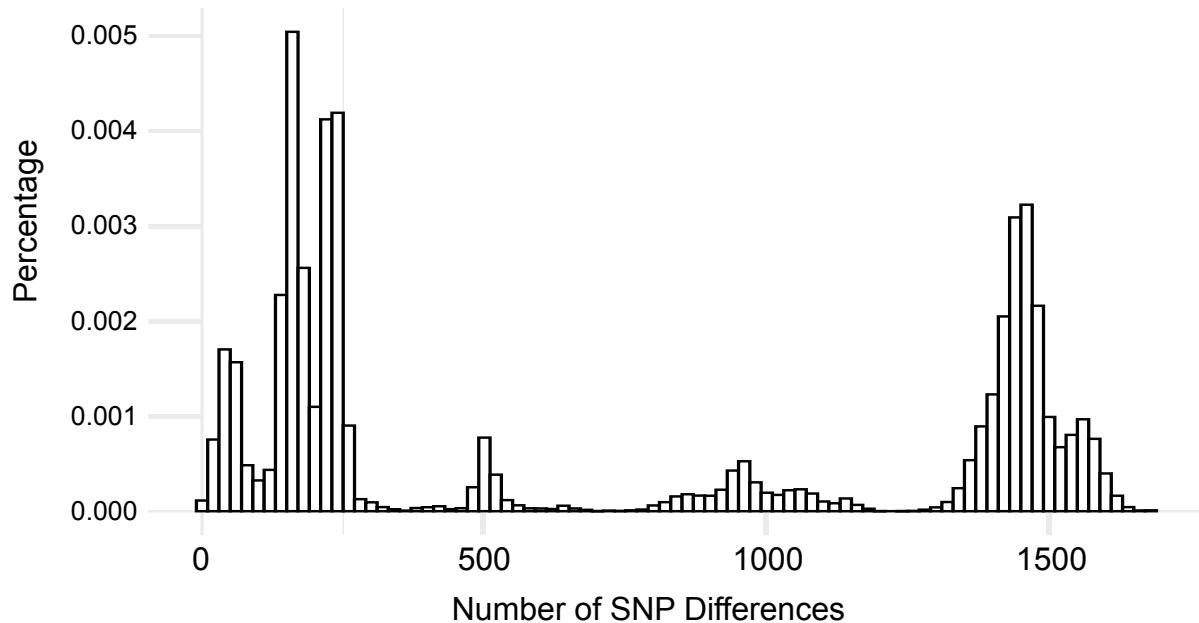
Office of Cyber Infrastructure and Computational Biology, National Institute of Allergy and
Infectious Disease, National Institutes of Health, Bethesda, Maryland, USA

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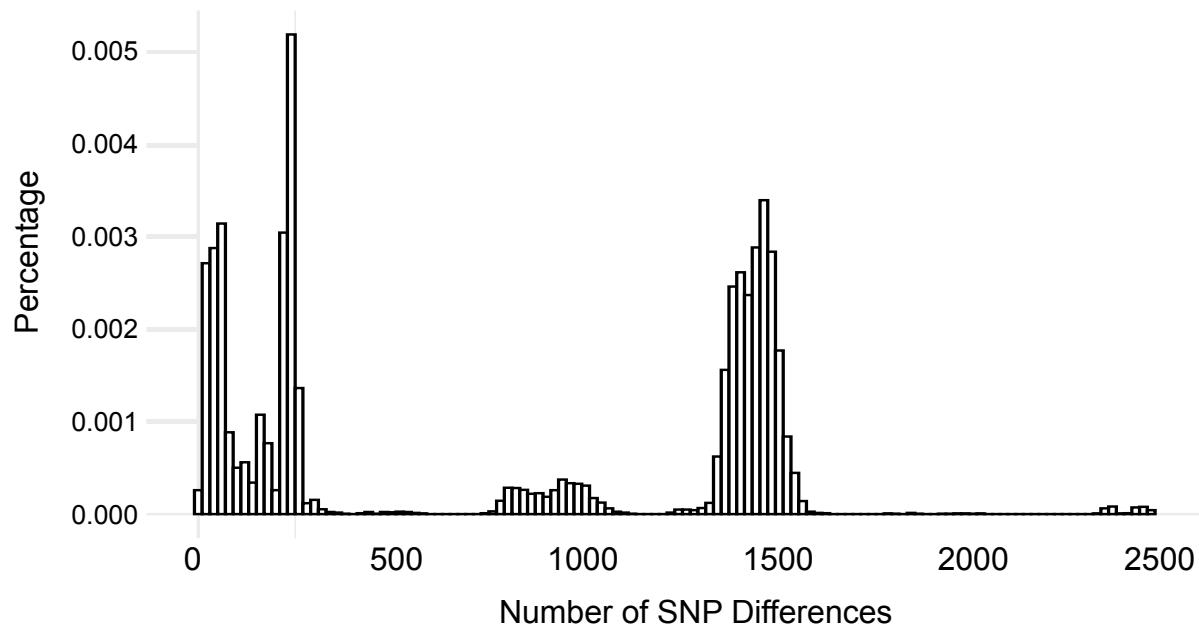
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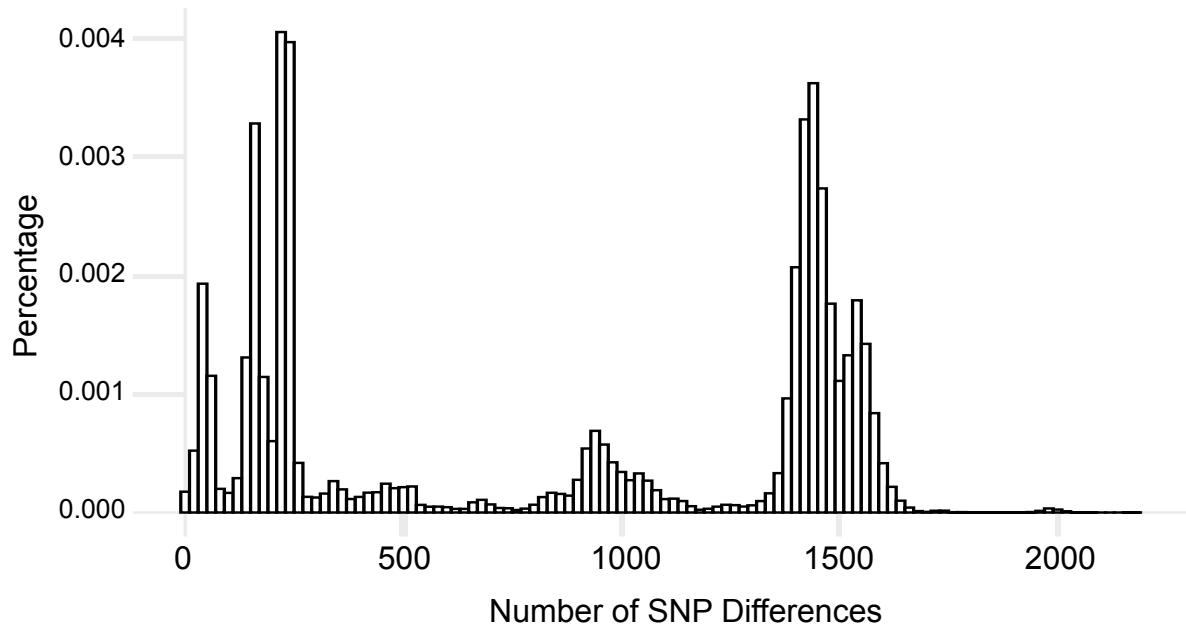
Azerbaijan



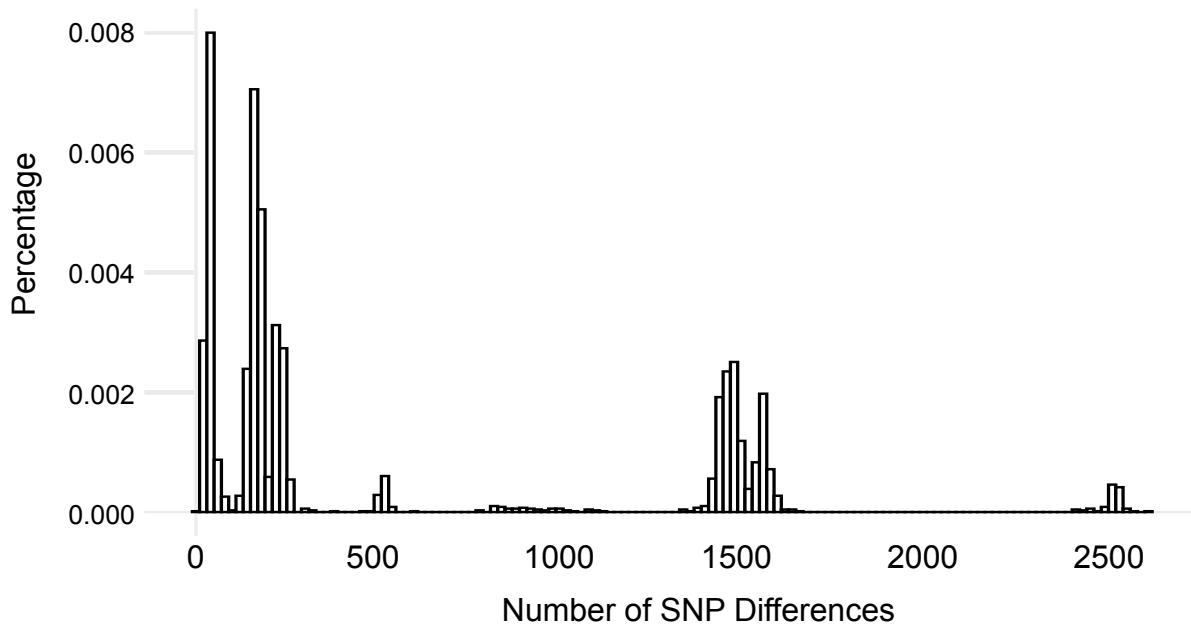
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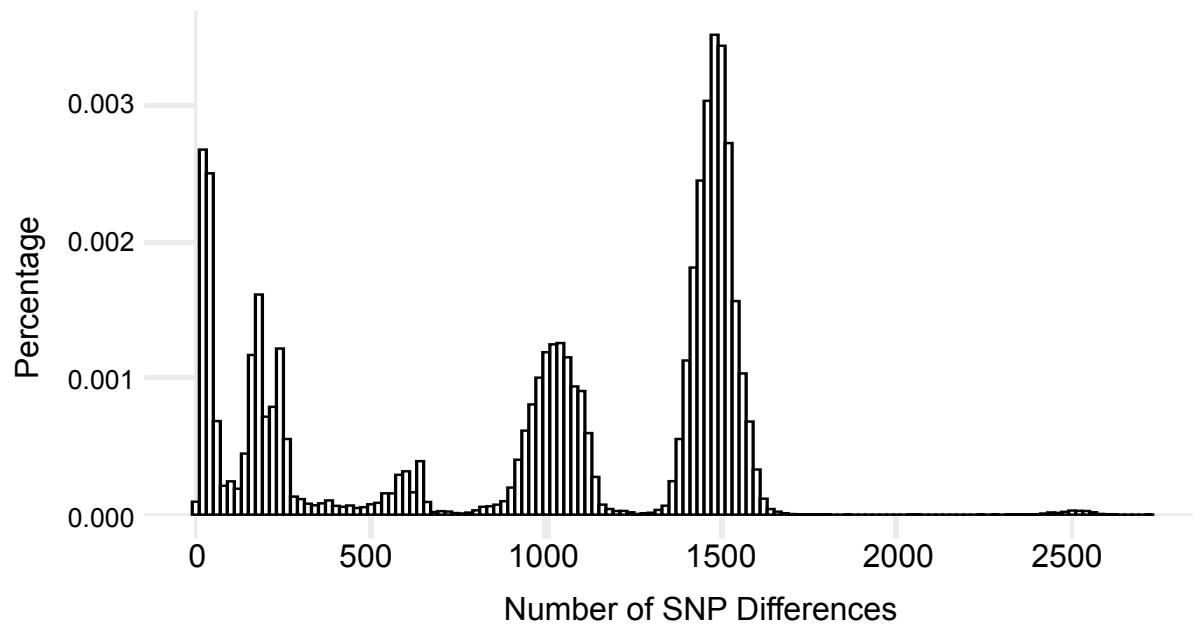
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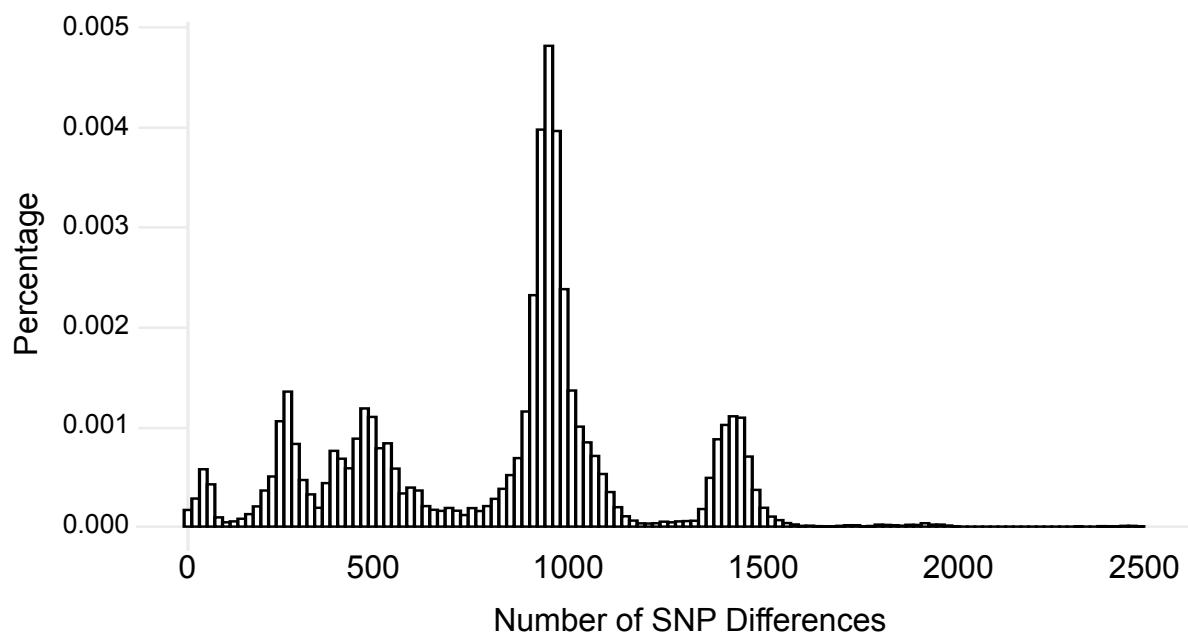
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Moldova

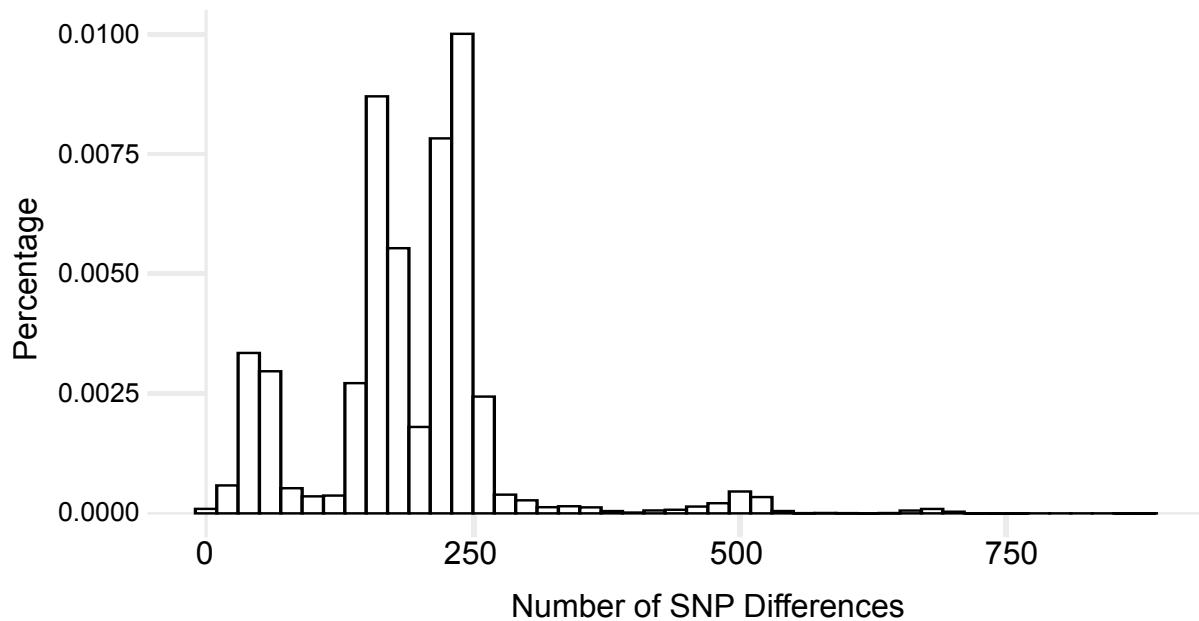


Romania

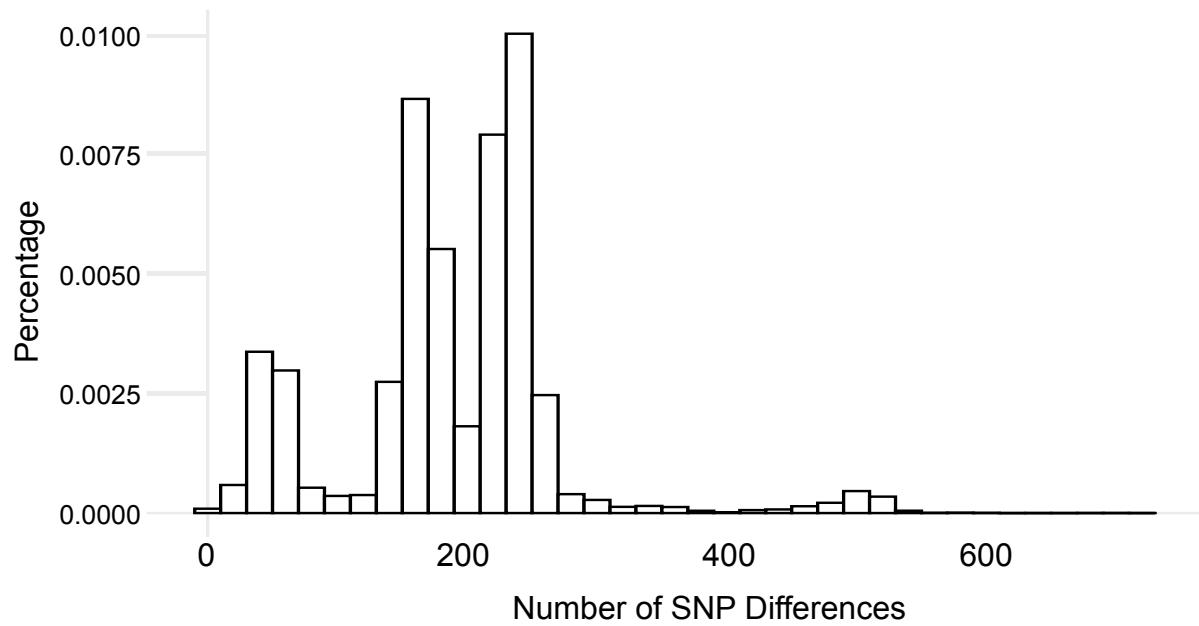


Supplementary Figure 1. Frequency histograms of pairwise distances within countries.

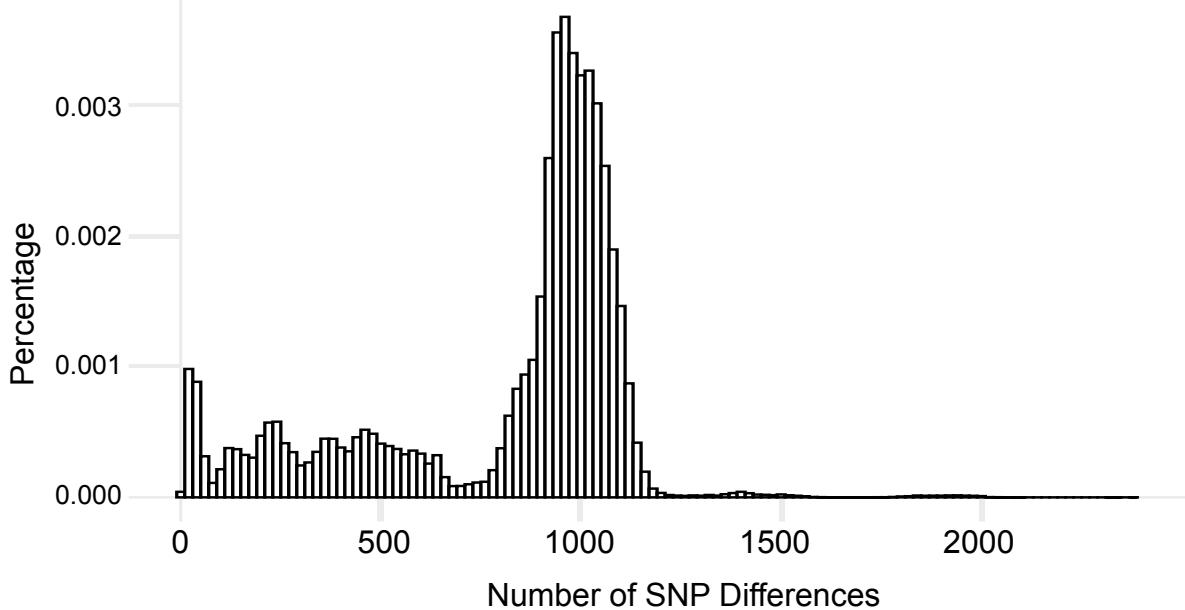
Lineage 2



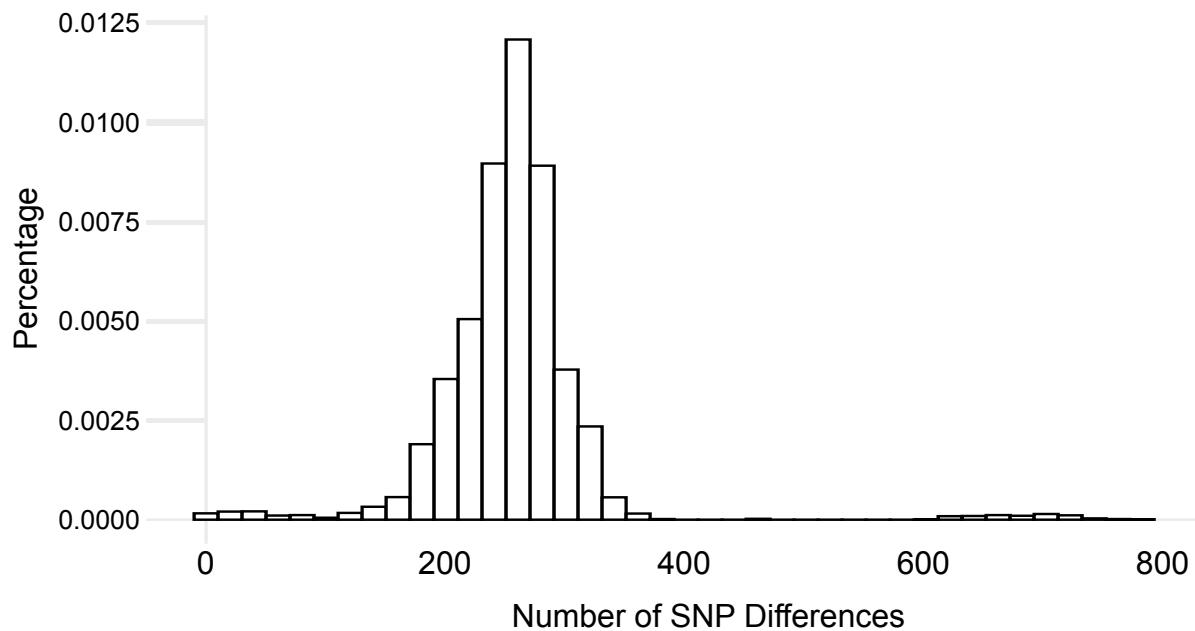
Lineage 2.2.1



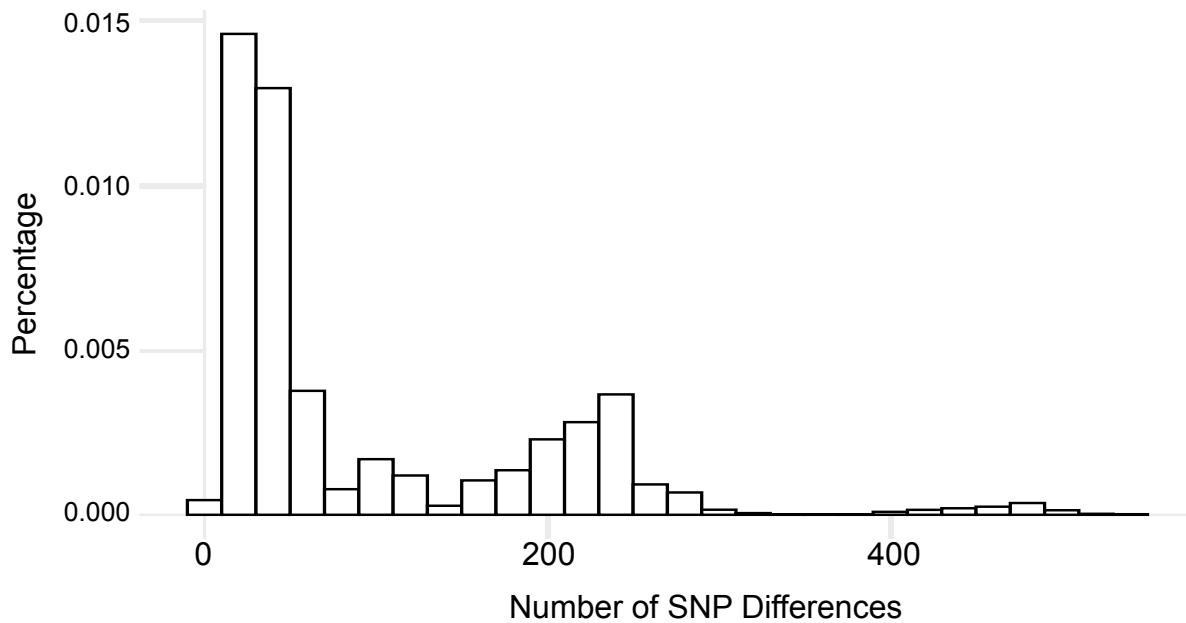
Lineage 4



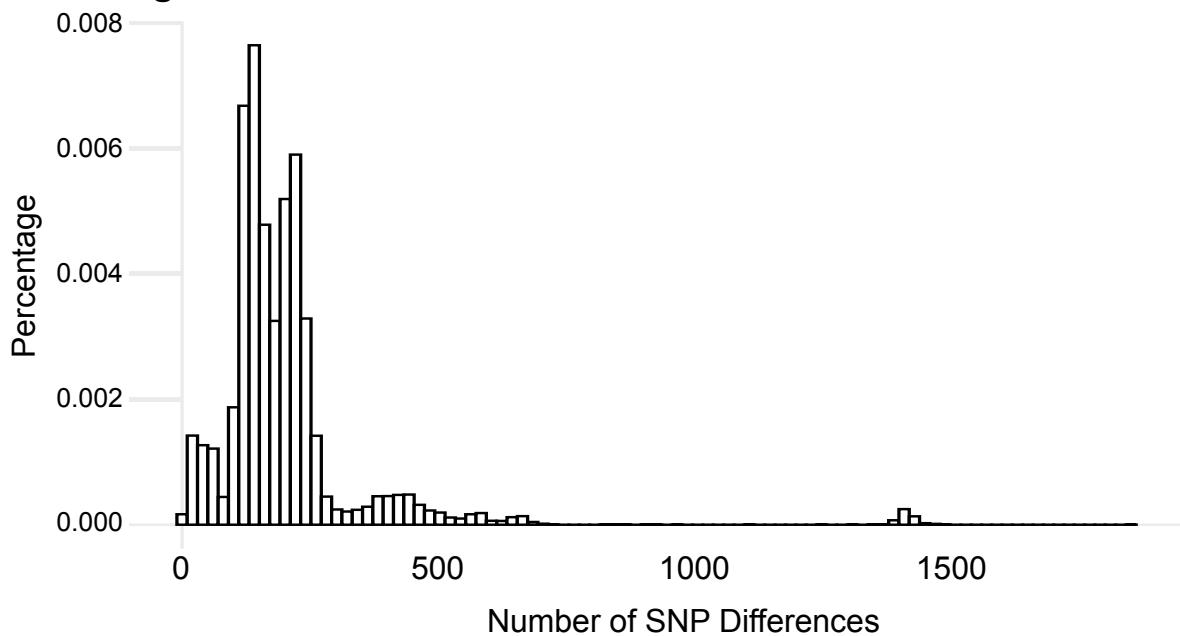
Lineage 4.1.2.1



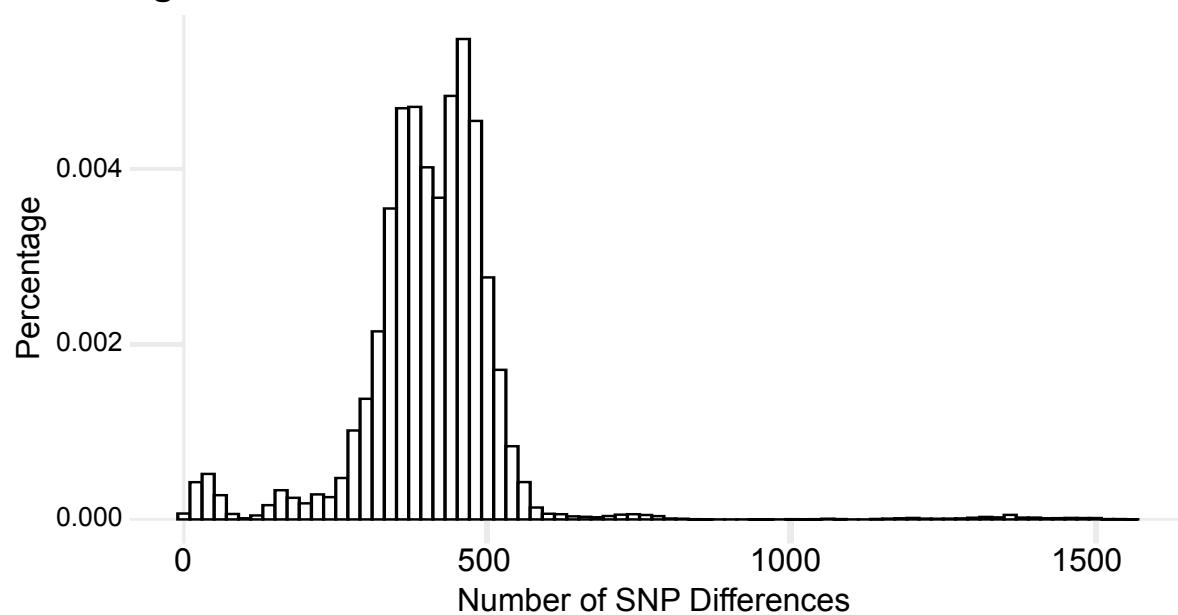
Lineage 4.2.1



Lineage 4.3.3



Lineage 4.8



Supplementary Figure 2. Frequency histograms of pairwise distances within phylogenomic lineages.