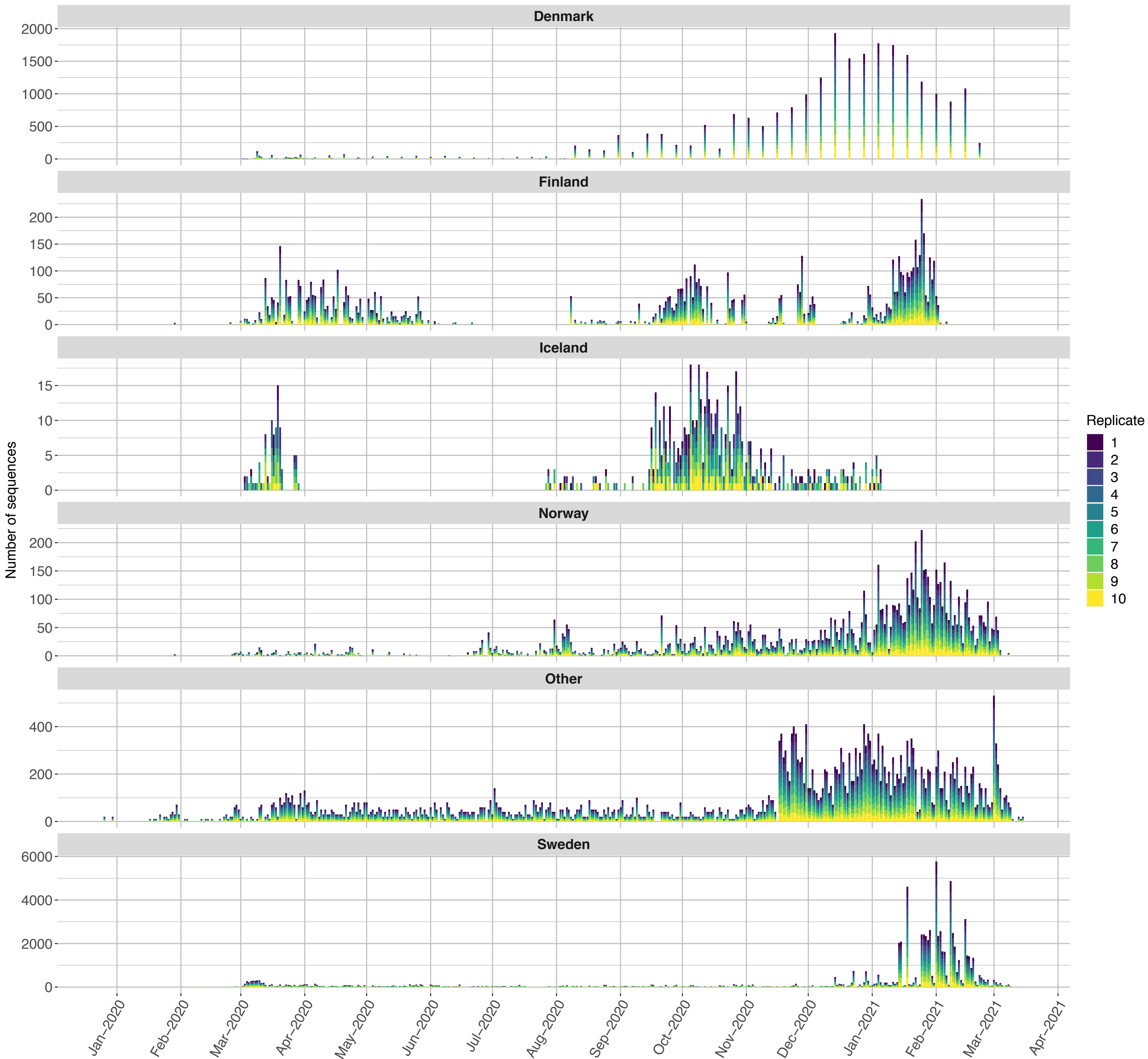


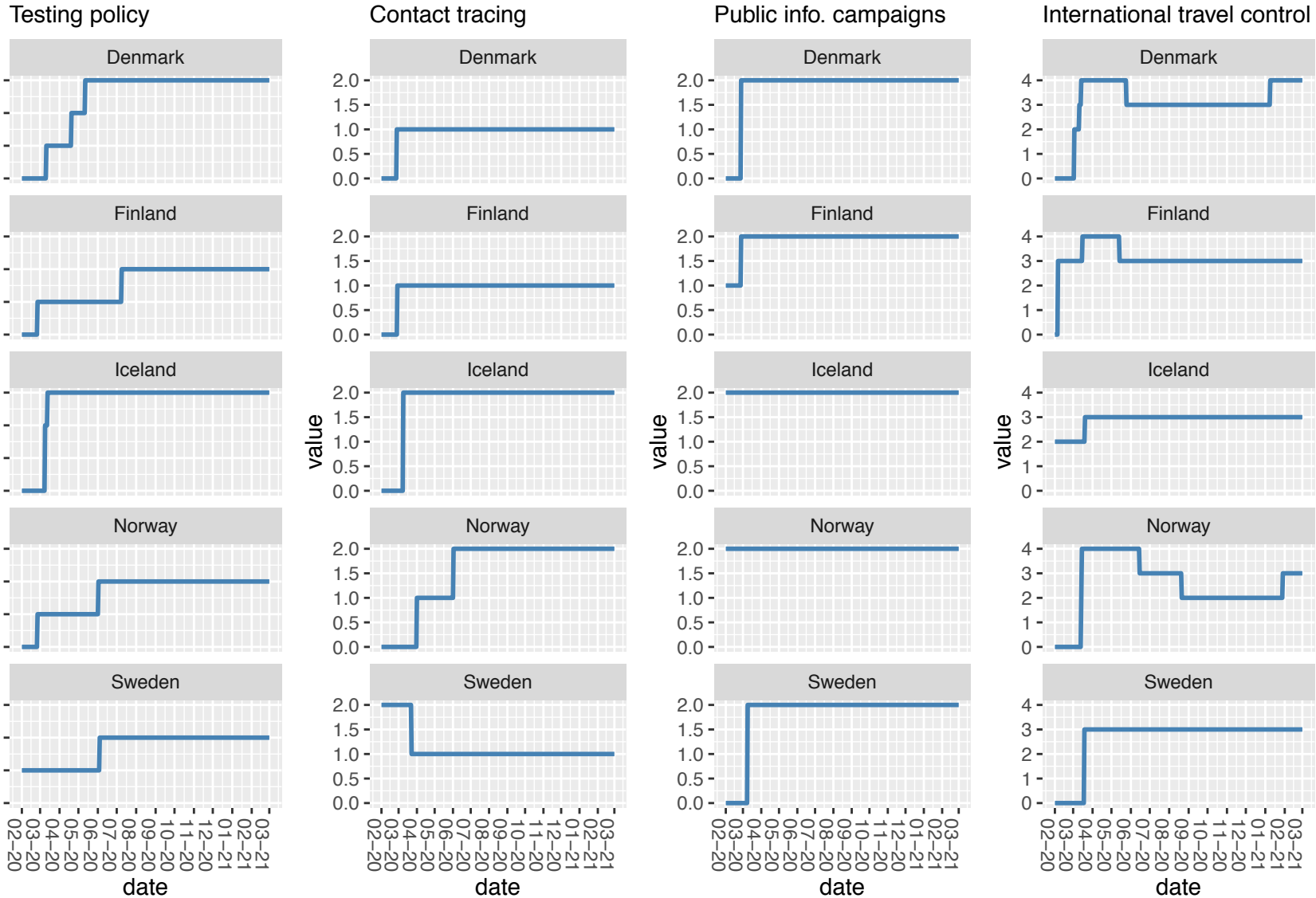
Supplementary Figure S1. Number of SARS-CoV-2 genomes sampled with time for all Nordic countries and global samples (i.e. 'other') included in the final alignment.

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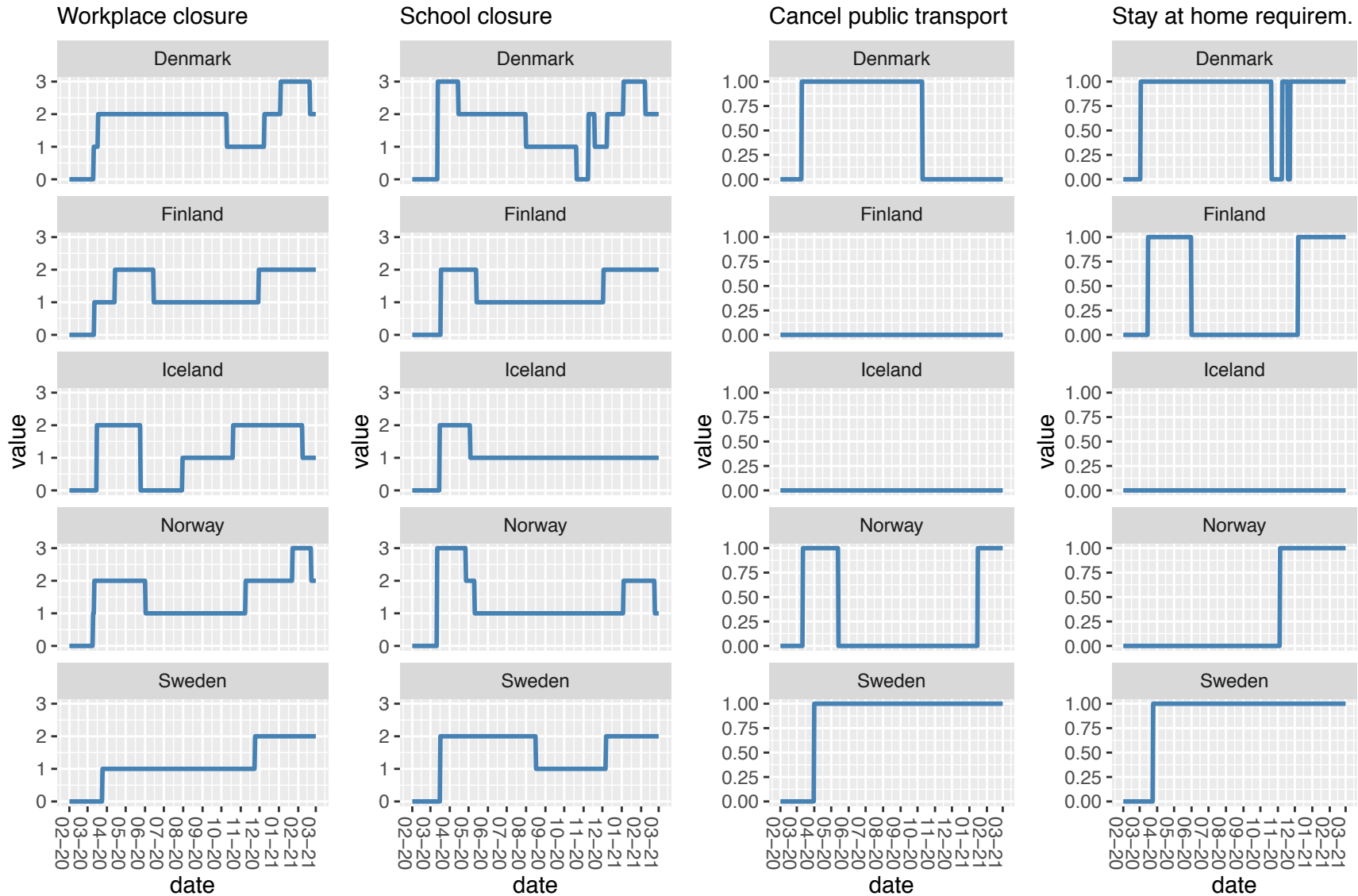
Supplementary Figure S2. Information on indicators of government response between February 2020 through March 2021 by country regarding (i) testing policy, (ii) contact tracing, (iii) public information campaigns, and (iv) international travel control.

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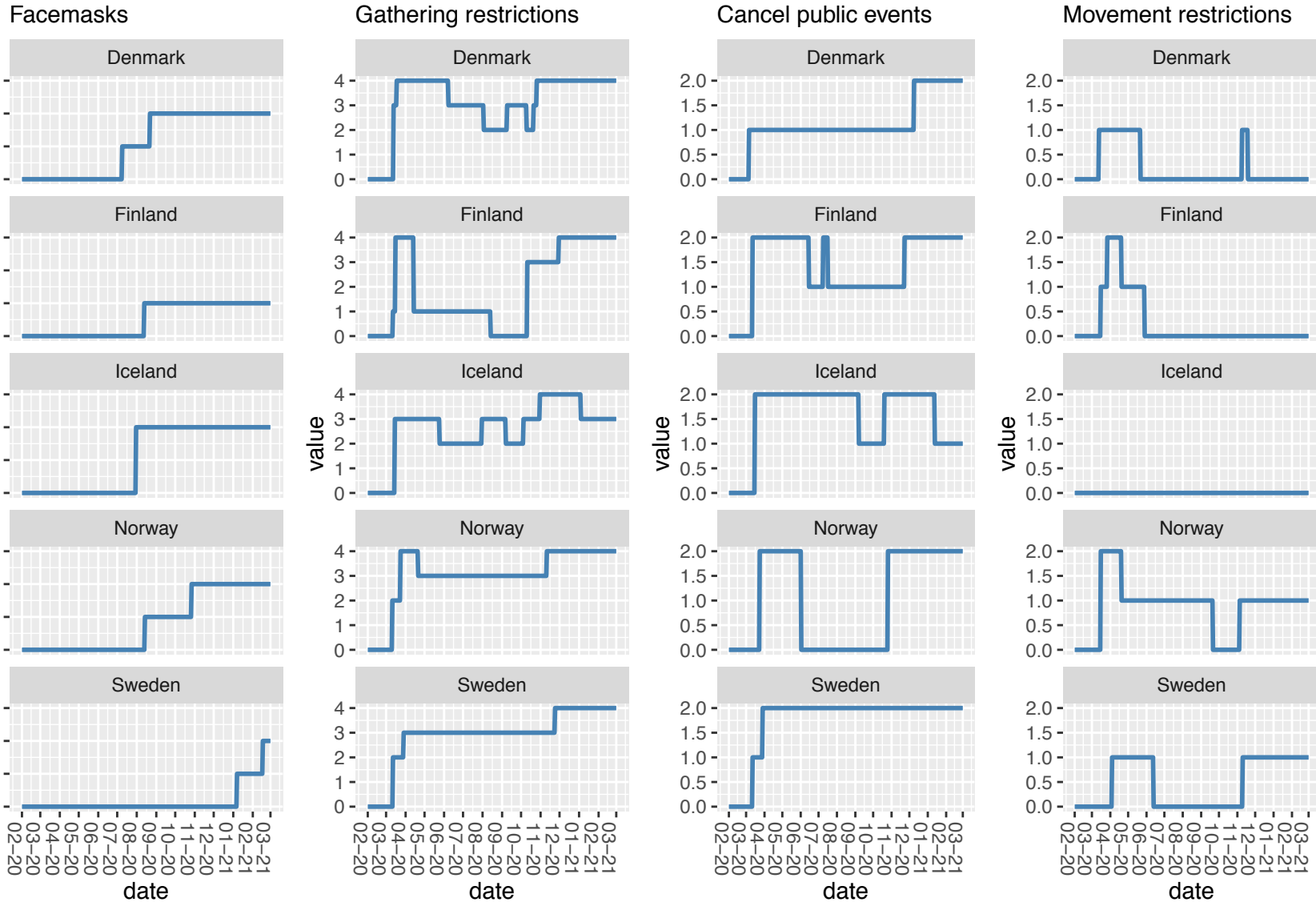
Supplementary Figure S3. Information on indicators of government response between February 2020 through March 2021 by country regarding (i) workplace closure, (ii) school closure, (iii) cancelation of public transportation, and (iv) stay at home requirements.

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Supplementary Figure S4. Information on indicators of government response between February 2020 through March 2021 by country regarding (i) facemasks, (ii) gathering restrictions, (iii) cancelation of public events, and (iv) restrictions on movement.

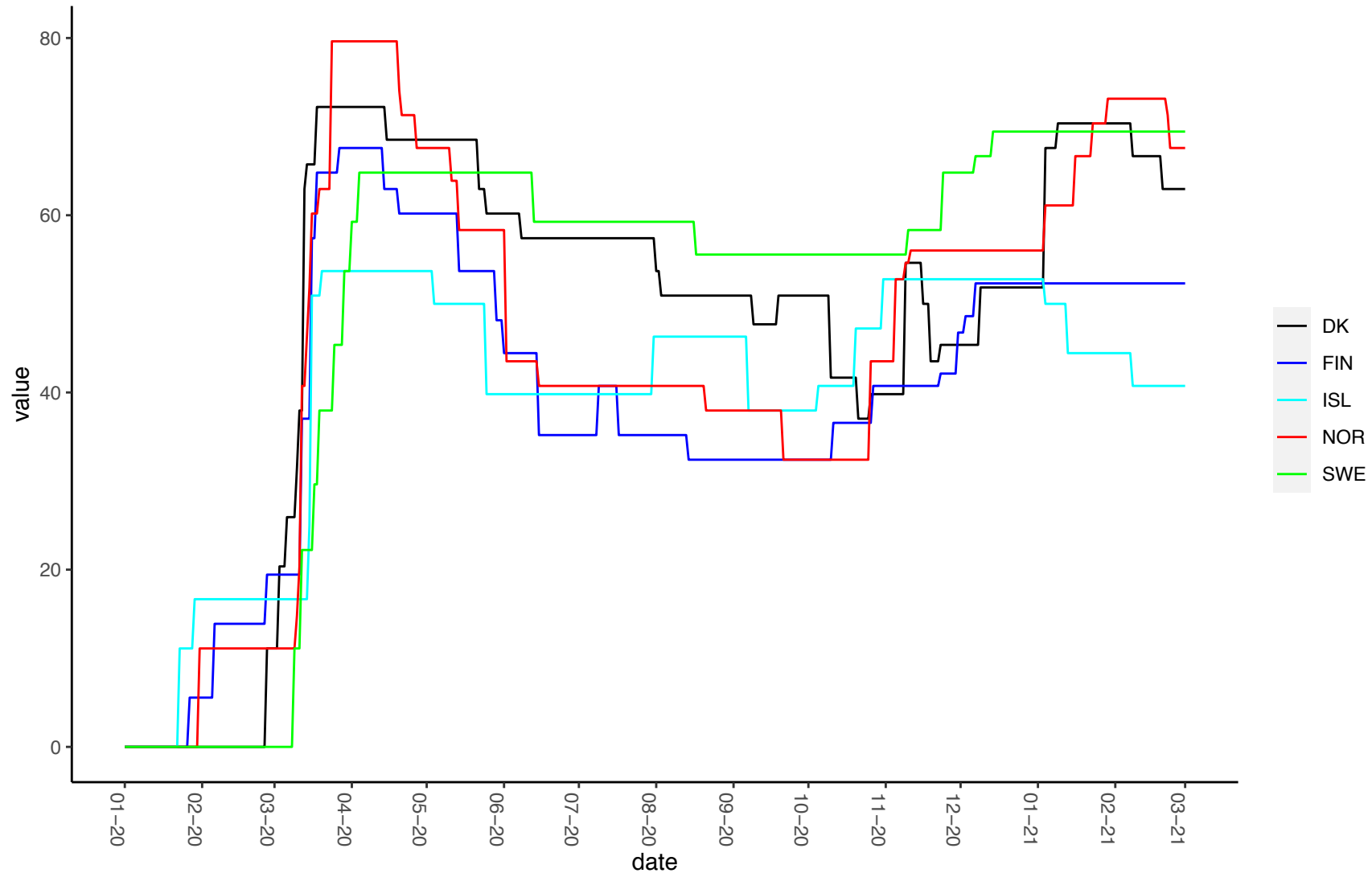
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Supplementary Figure S5. Stringency index of government response between February 2020 through March 2021 by country.

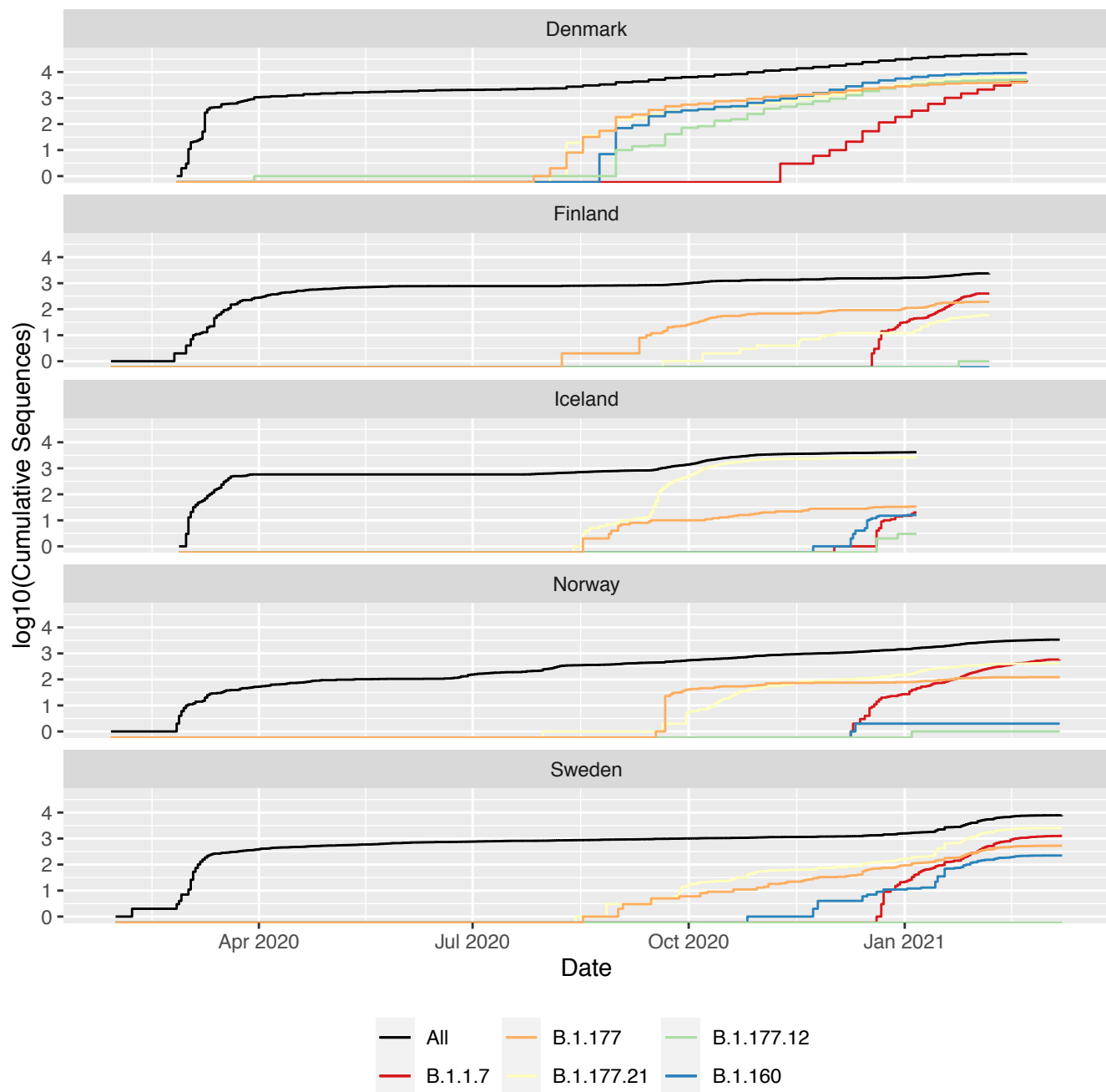
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Stringency index



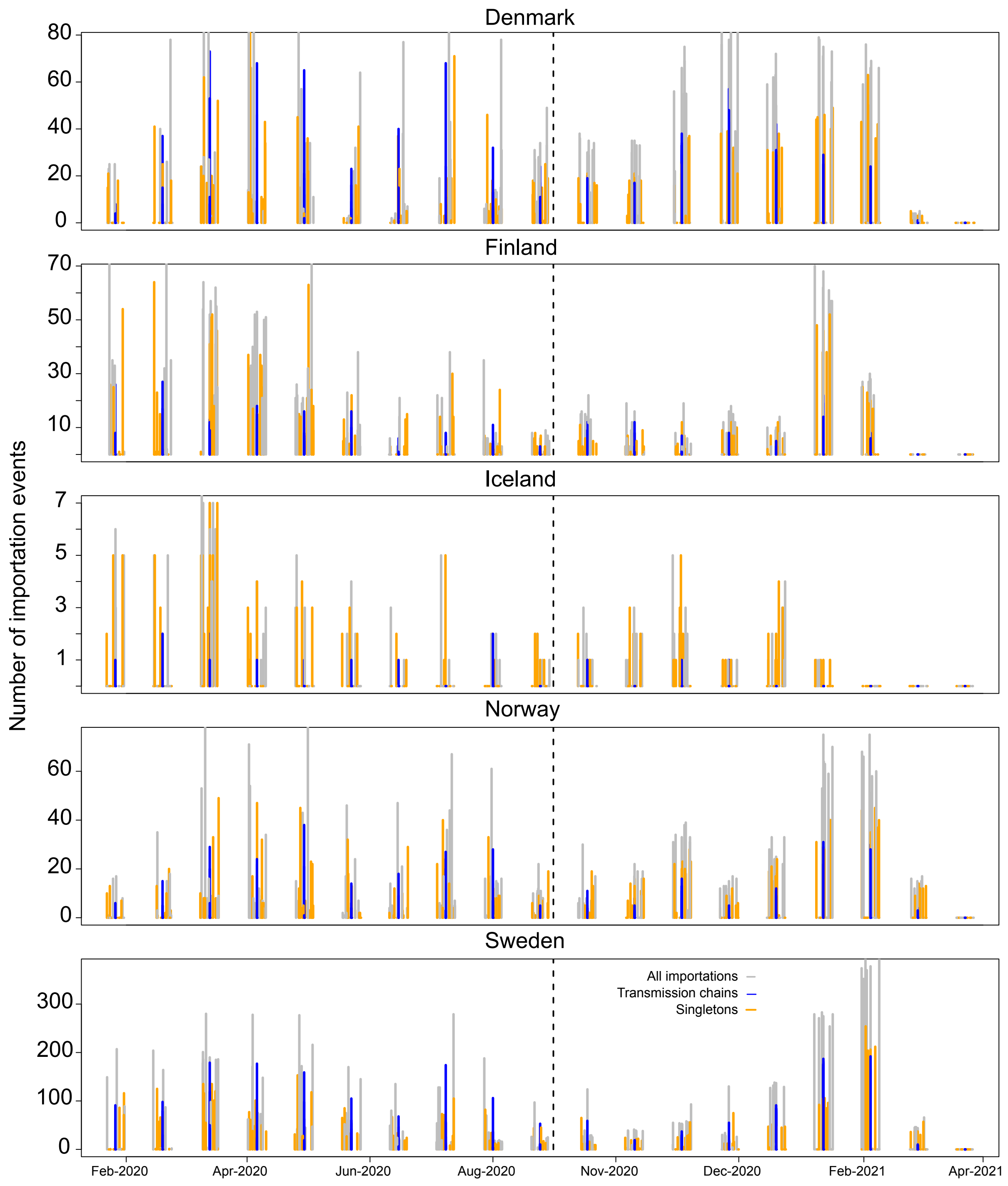
Supplementary Figure S6. Log number of genomes in each of the five most common pangolin lineages and B1.1.7 over time in the complete dataset.

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Supplementary Figure S7. Number of putative introductions identified by detecting transmission chains (monophyletic groups of at least 2 genomes from a Nordic country) or singletons (genomes from a Nordic country sitting outside a transmission chains).

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Supplementary Figure S8. Jittered markov rewards for each Nordic country and for the global diversity in the data (labelled as “other”).

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