

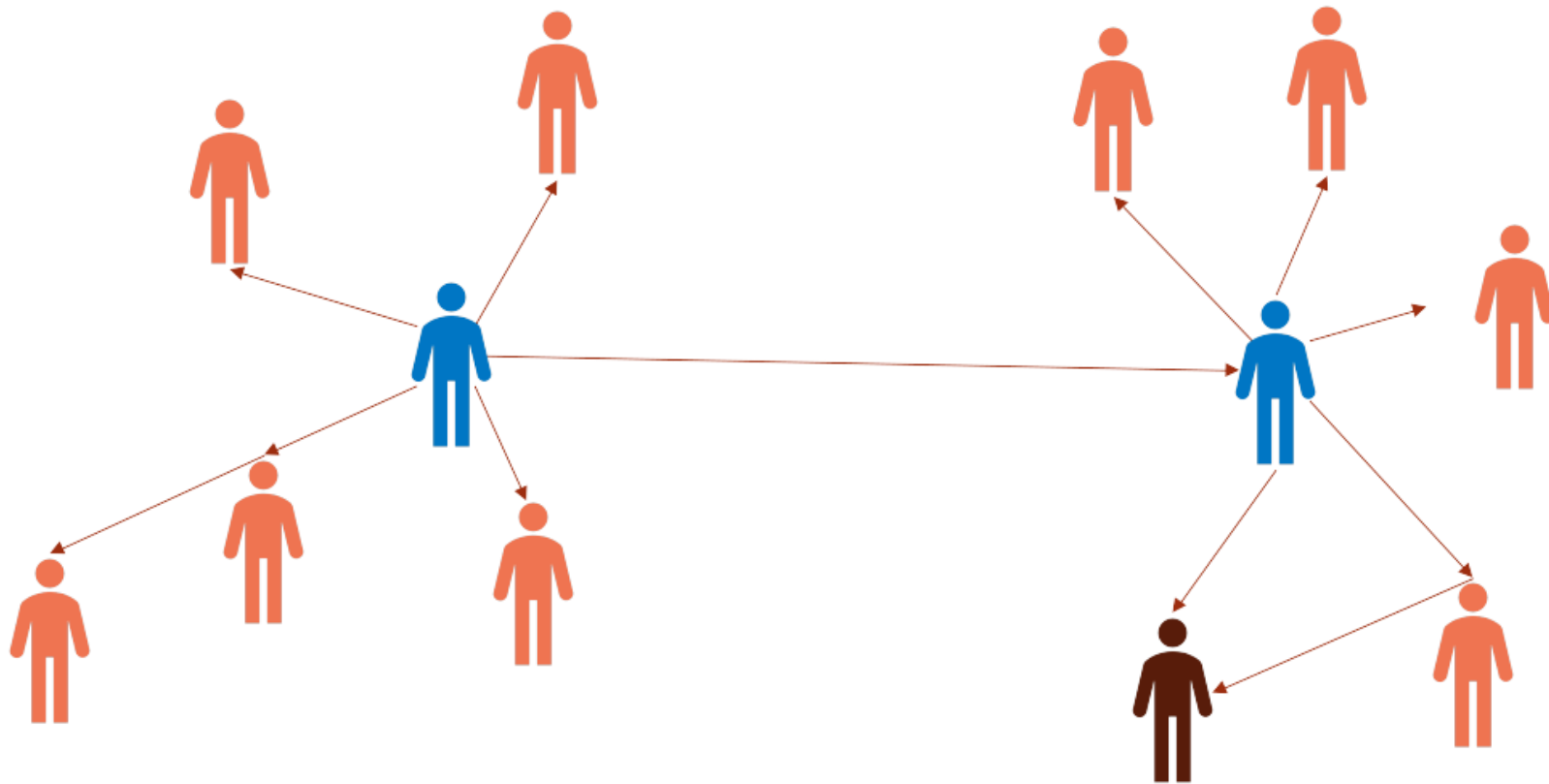
INFERRING THE GLOBAL COVID-19 TRANSMISSION NETWORK USING TNET

Saurav Dhar, Ion Mandoiu, Mukul S. Bansal

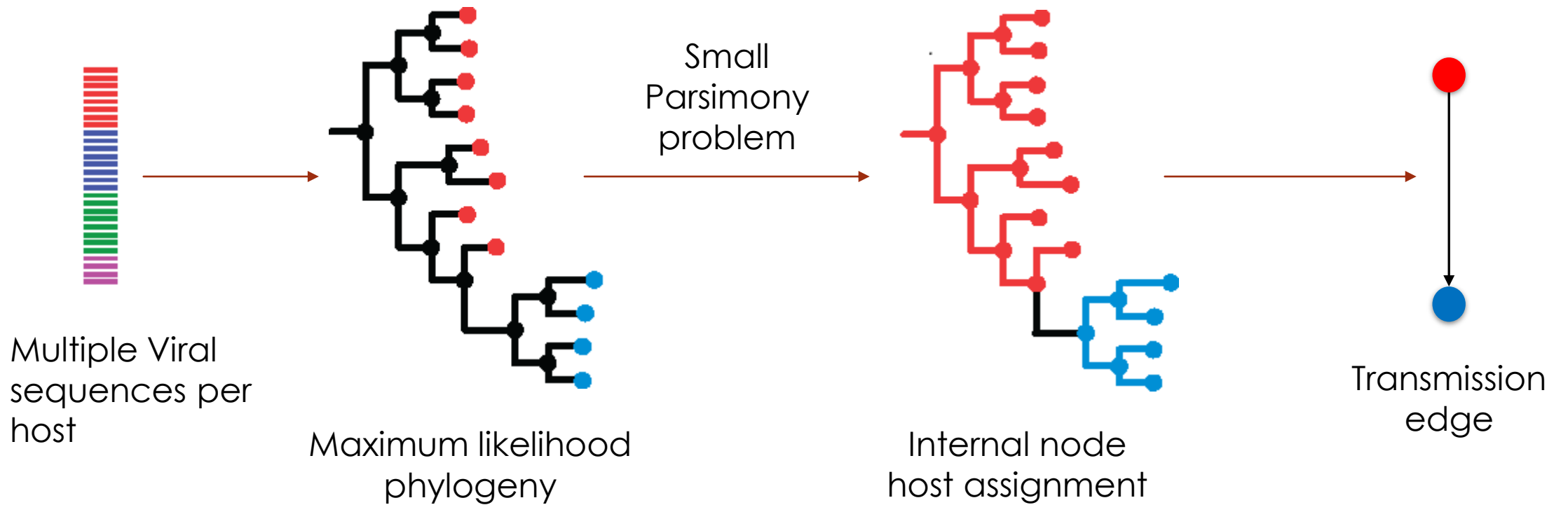
TRANSMISSION NETWORK

Who infected whom?

Multiple viral sequences sampled from the hosts.



TNET WORKFLOW



DEALING WITH MULTIPLE OPTIMAL HOST ASSIGNMENT

IDEA 1: RANDOM SAMPLING

| | | | | |
|-------|----|----|----|----|
| Score | 6 | 5 | 5 | 10 |
| Count | 20 | 10 | 30 | 5 |

H_2

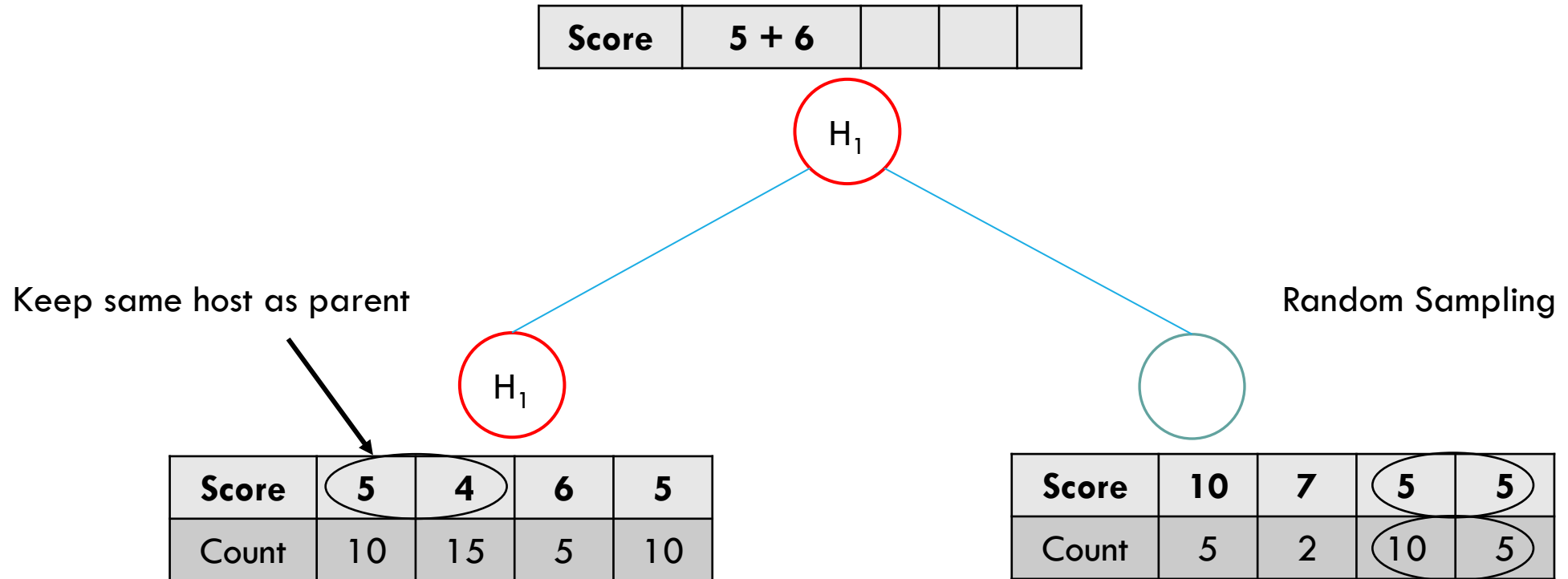
H_3

$$P_r(H_2) = \frac{10}{10+30} = \frac{1}{4}$$

$$P_r(H_3) = \frac{30}{10+30} = \frac{3}{4}$$

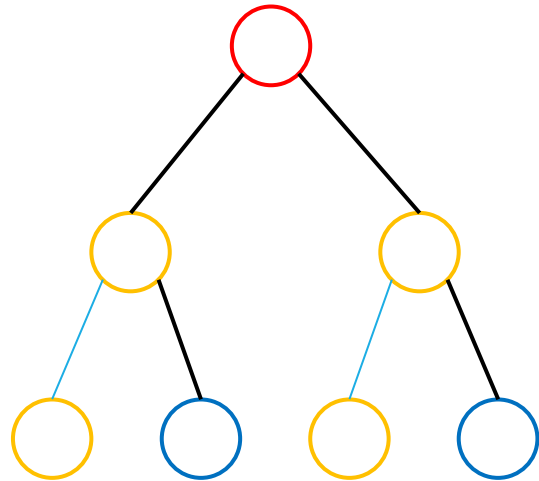
DEALING WITH MULTIPLE OPTIMAL HOST ASSIGNMENT

IDEA 2: BIASED SAMPLING

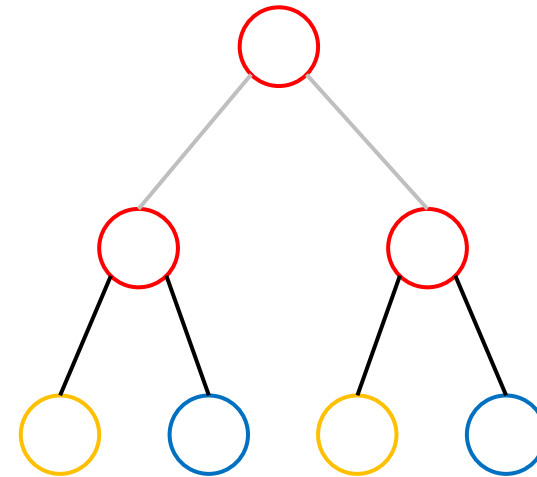


TNET-RANDOM VS. TNET-BIASED

Parsimony score = 4

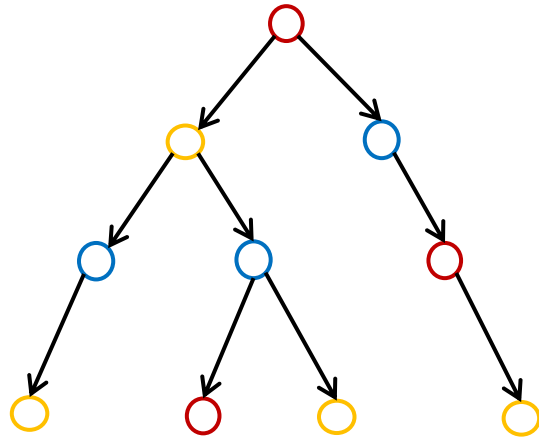


Parsimony score = 4

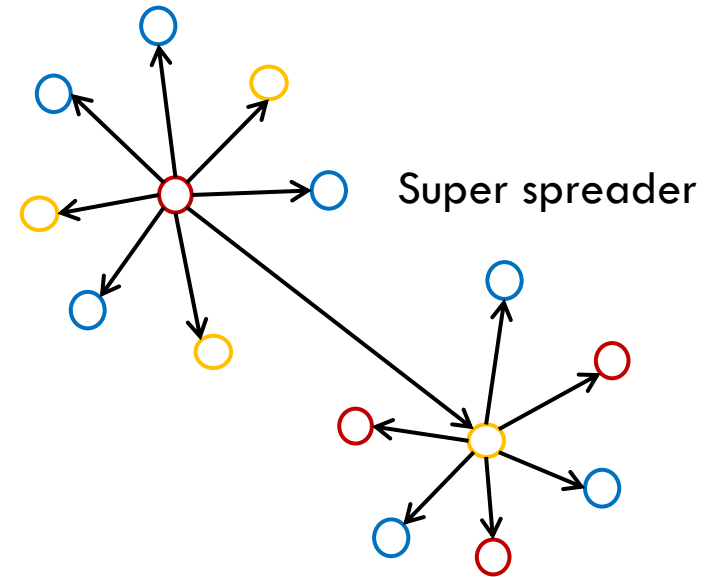


Changes are pushed further down the tree keeping the same parsimony score

TNET-RANDOM VS. TNET-BIASED: RELATIVE ACCURACY DEPENDS ON NETWORK



Infector ratio ~ 2



Infector ratio ~ 7

Based on simulated data, we see that TNet-Biased works better for networks with higher infector ratio.

TNET ON COVID-19 GLOBAL COUNTRY LEVEL ANALYSIS

Dataset

- 59 countries
- 2123 total complete sequences
- 10 to 100 sequences per country
- Up to June, 2020

Measure

- Country of exposure
- Top 5 spreaders and receivers in each month
- Top 5 spreaders and receivers of USA

Compared against NextStrain

- True country of exposure known

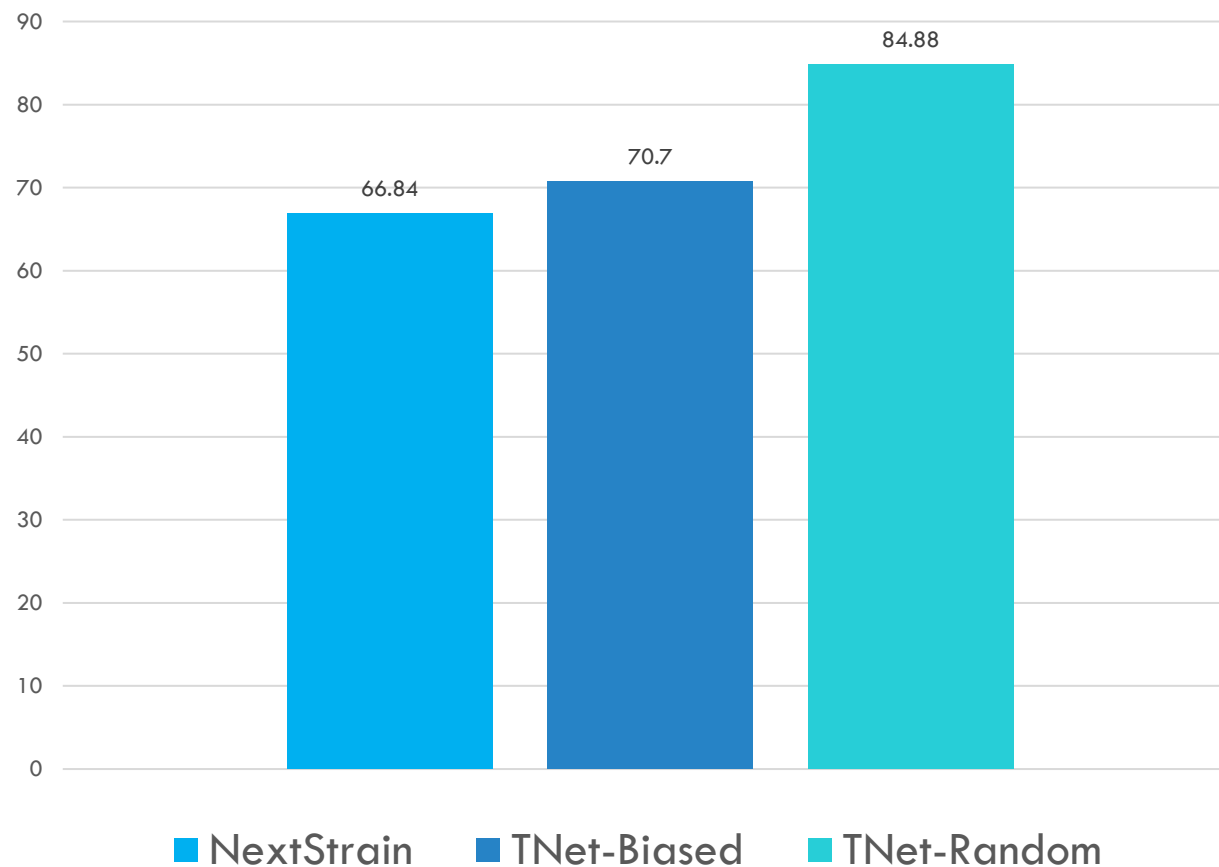
Handling topological uncertainty

- 10 bootstrap trees

Handling sampling uncertainty

- Used 100 samples per bootstrap tree

ACCURACY: KNOWN COUNTRY OF EXPOSURE FOR EACH SEQUENCE

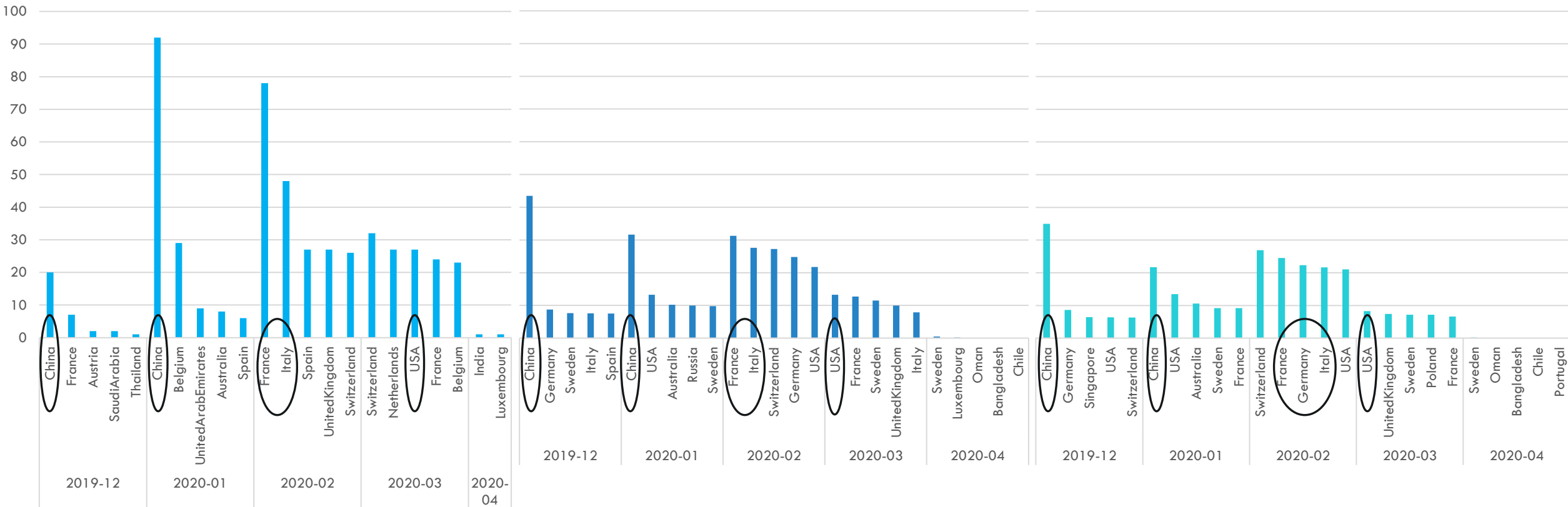


TOP 5 SPREADERS OF EACH MONTH (AVERAGED)

NextStrain

TNet-Biased

TNet-Random

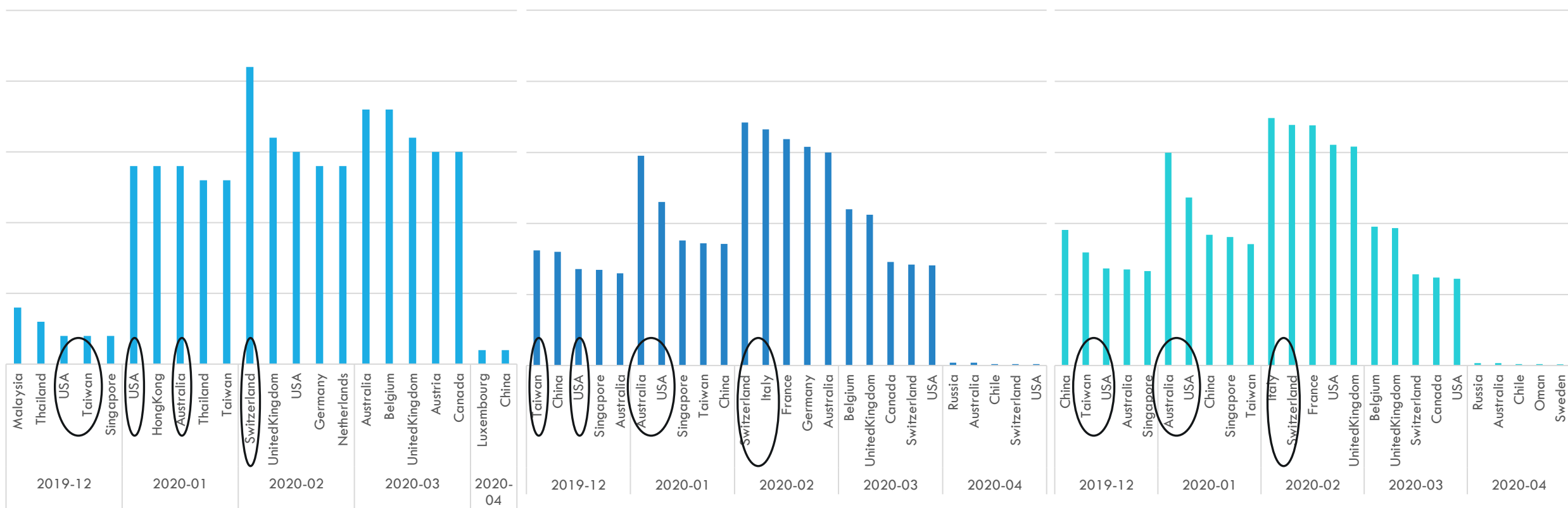


TOP 5 RECEIVERS OF EACH MONTH (AVERAGED)

NextStrain

TNet-Biased

TNet-Random

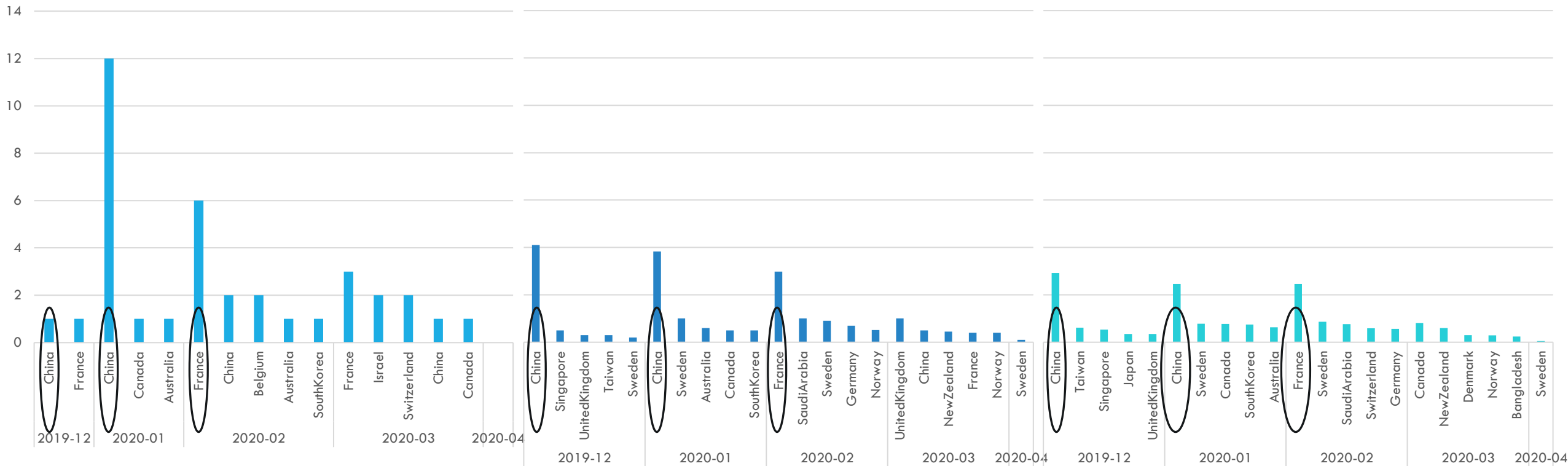


SPREAD TO USA: TOP 5 COUNTRIES

NextStrain

TNet-Biased

TNet-Random

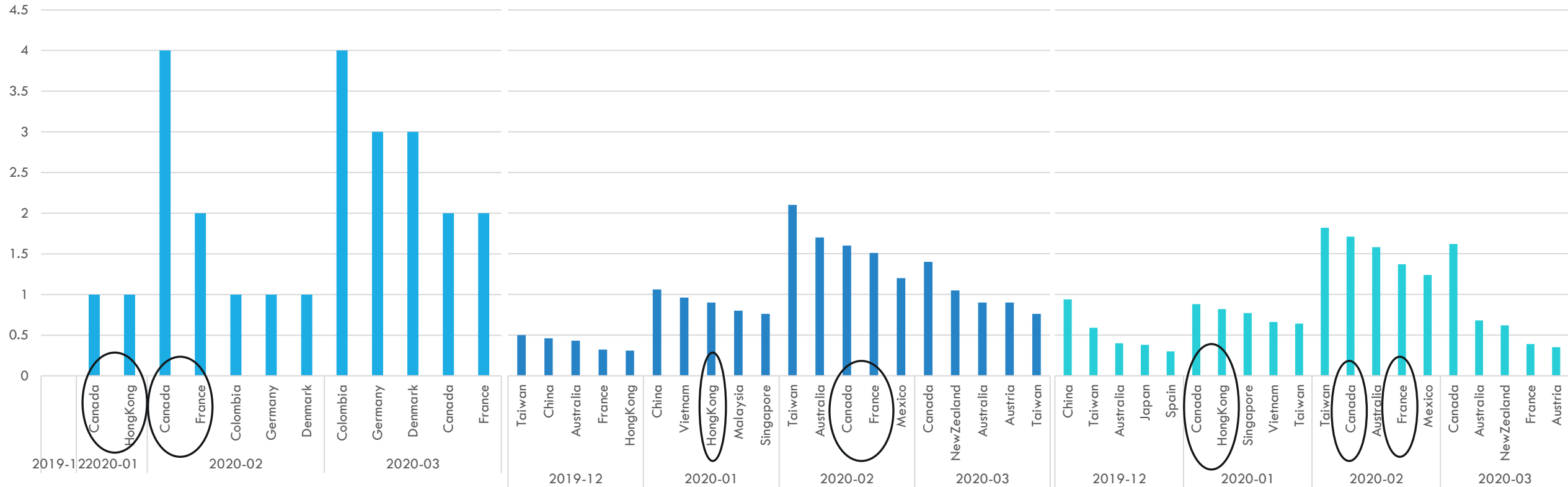


SPREAD FROM USA: TOP 5 COUNTRIES

NextStrain

TNet-Biased

TNet-Random



TNET ON COVID-19 STATE LEVEL ANALYSIS FOR USA

Dataset

- 30 states
- 1801 total complete sequences
- 10 to 100 sequences per state
- Up to July, 2020

Measure

- State of exposure
- Top 5 spreaders and receivers in each month
- Top 5 spreaders and receivers of New York

Compared against NextStrain

- True state of exposure known

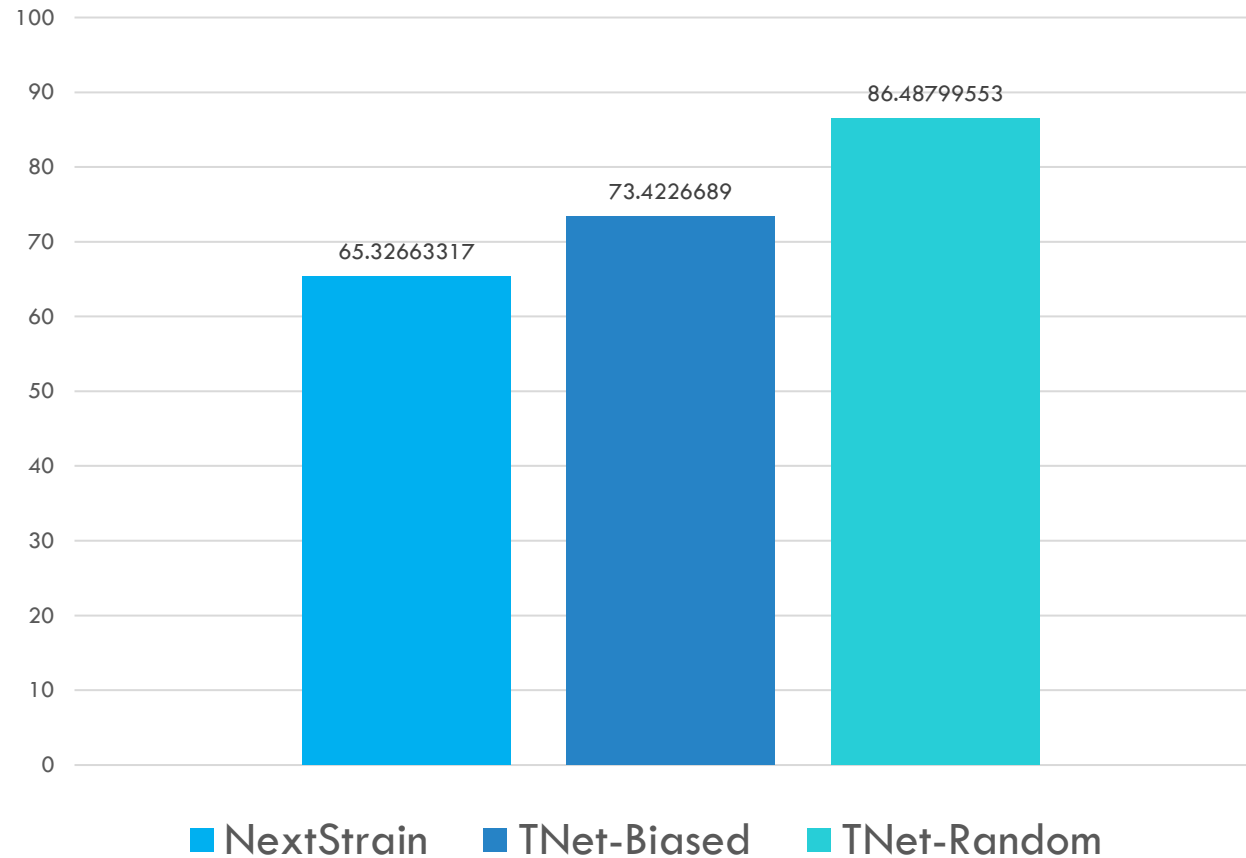
Handling topological uncertainty

- 10 bootstrap trees

Handling sampling uncertainty

- Used 100 samples per bootstrap tree

ACCURACY : KNOWN STATE OF EXPOSURE FOR EACH SEQUENCE

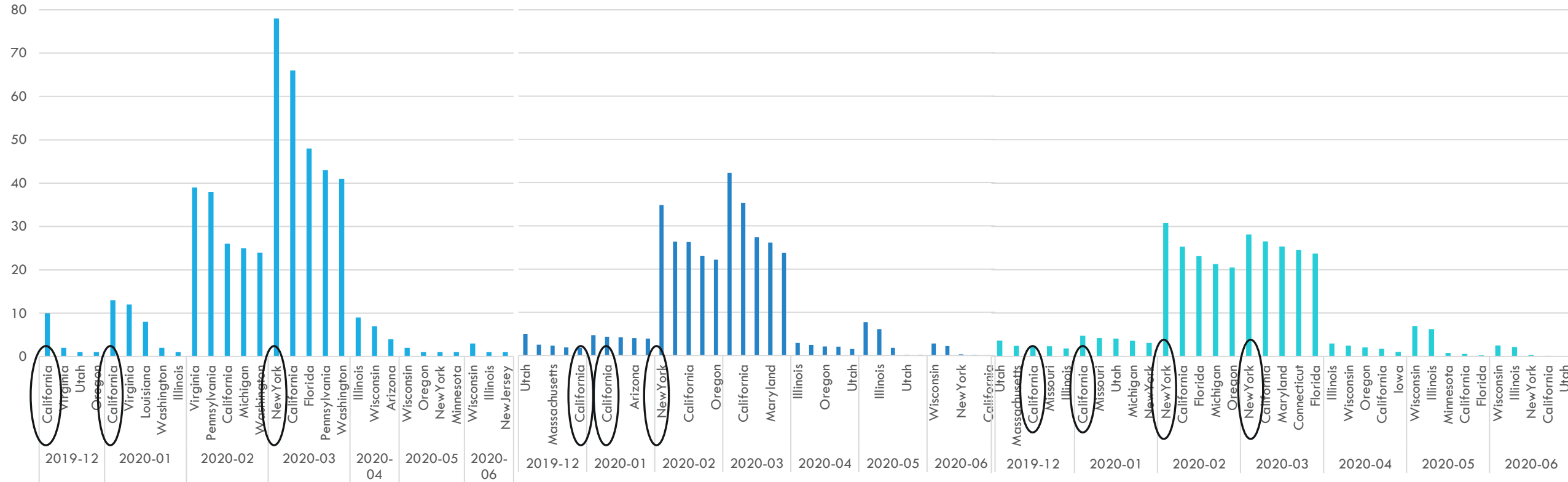


TOP 5 SPREADERS OVER TIME (AVERAGE)

NextStrain

TNet-Biased

TNet-Random

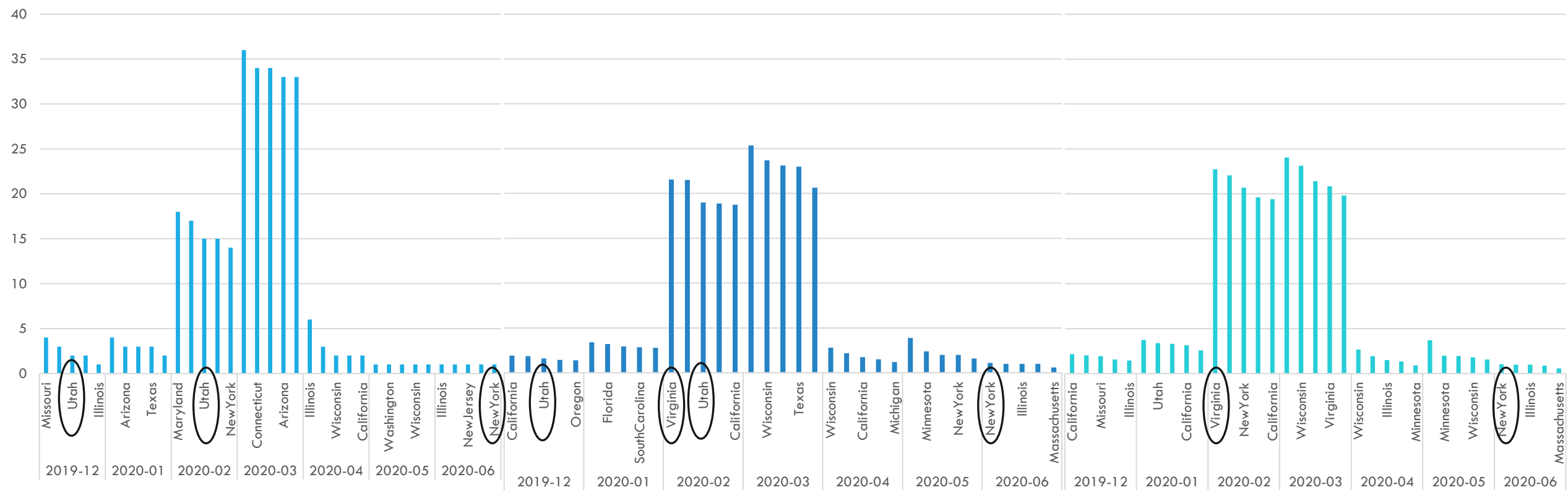


TOP 5 RECEIVERS OVER TIME (AVERAGE)

NextStrain

TNet-Biased

TNet-Random

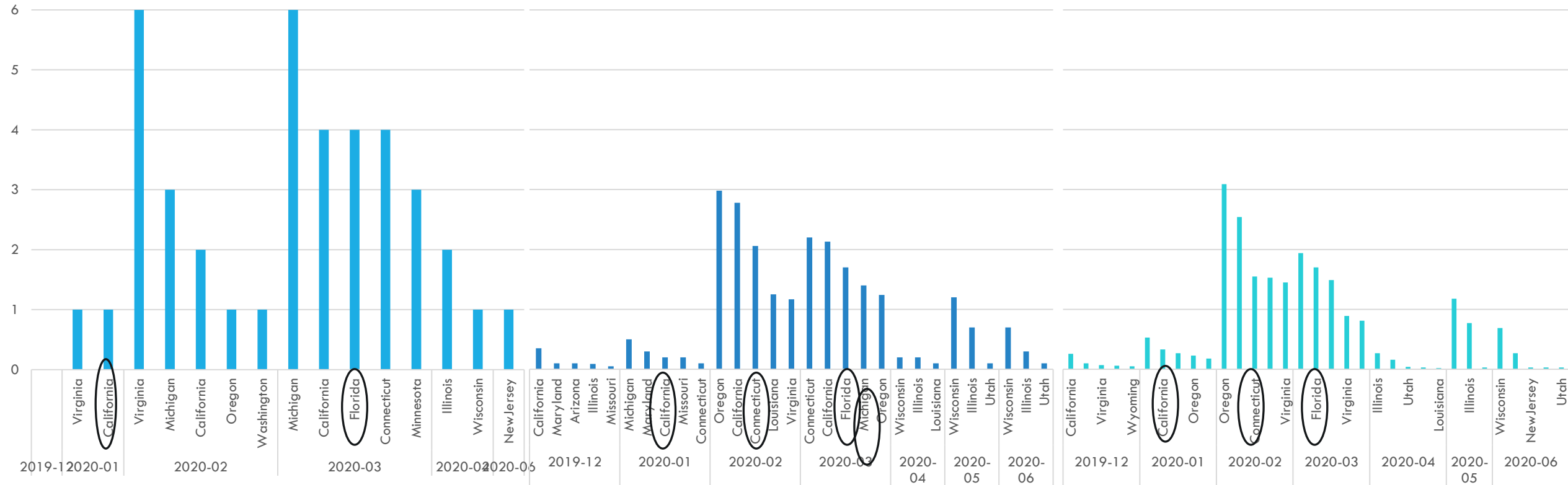


TOP 5 STATES NEW YORK RECEIVED FROM (AVERAGE)

NextStrain

TNet-Biased

TNet-Random

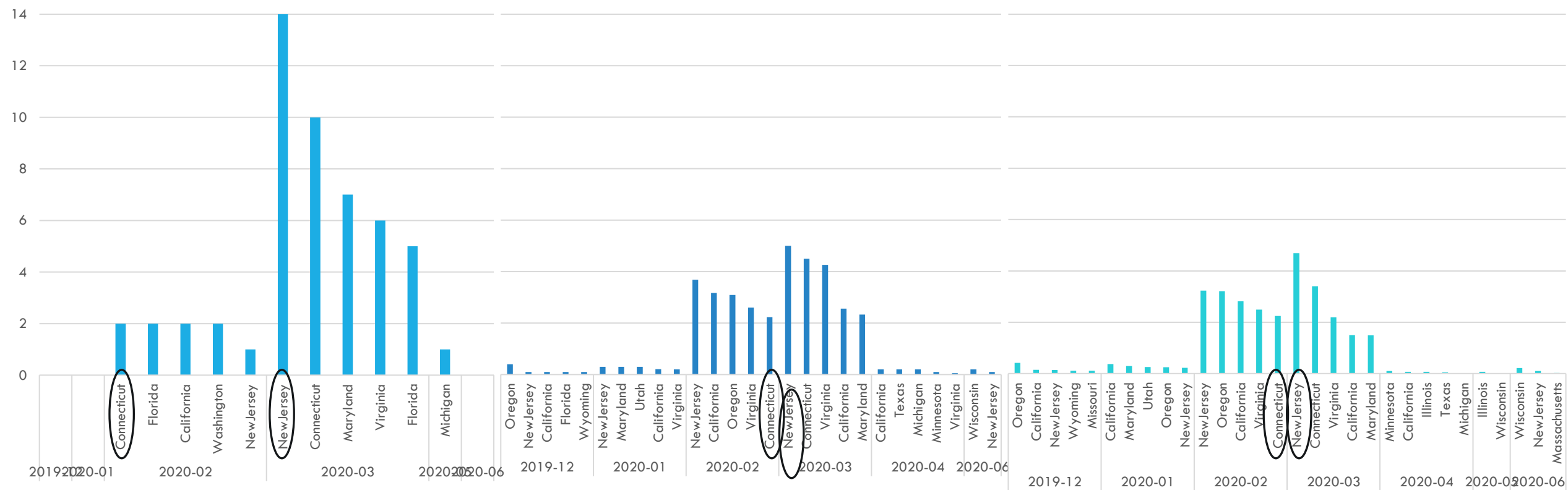


TOP 5 STATES NEW YORK SPREAD TO (AVERAGE)

NextStrain

TNet-Biased

TNet-Random



CONCLUSION

- Simple and scalable model like TNet-Random appears to work very well.
- Using countries and states as hosts may be a viable approach for such geographic spread analyses.



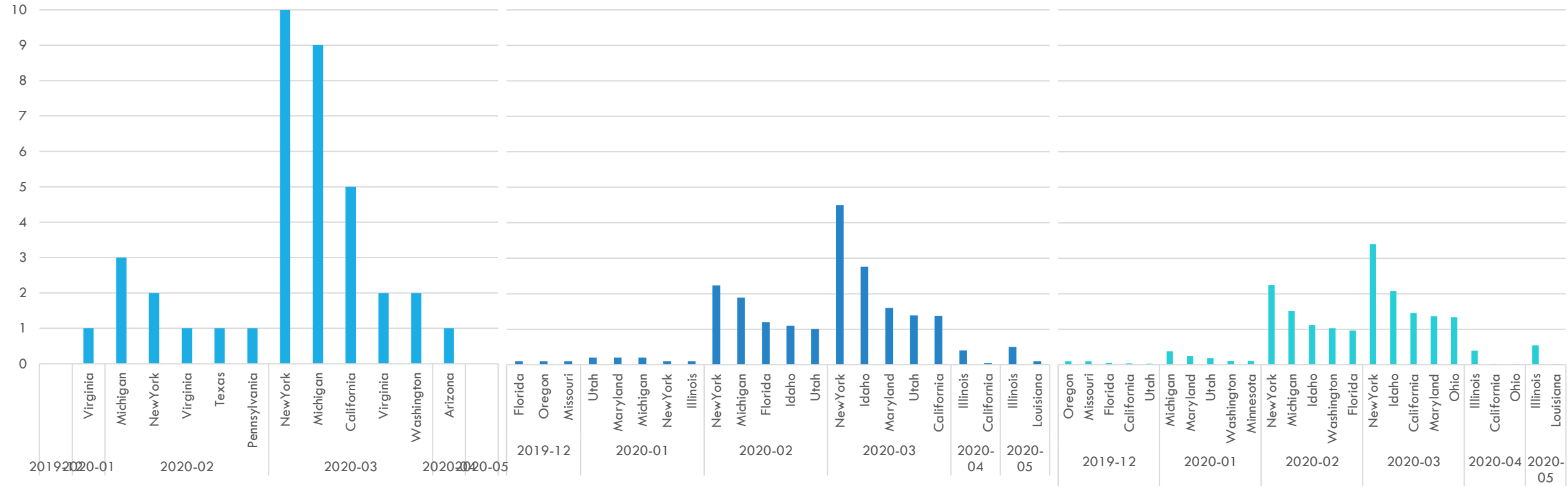
THANK YOU

TOP 5 STATES CONNECTICUT RECEIVED FROM (AVERAGE)

NextStrain

TNet-Biased

TNet-Random

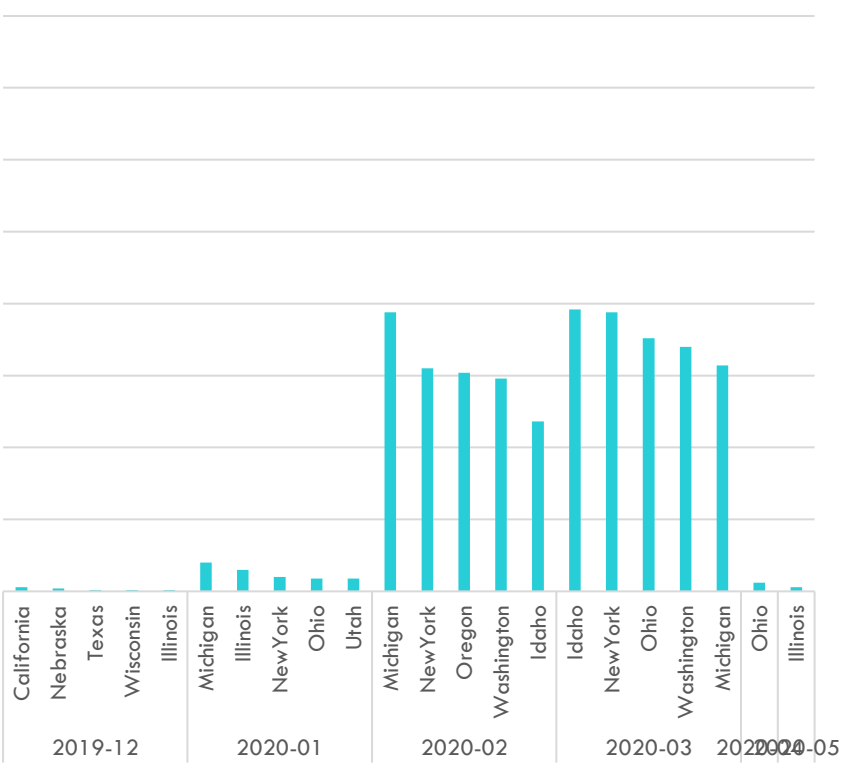
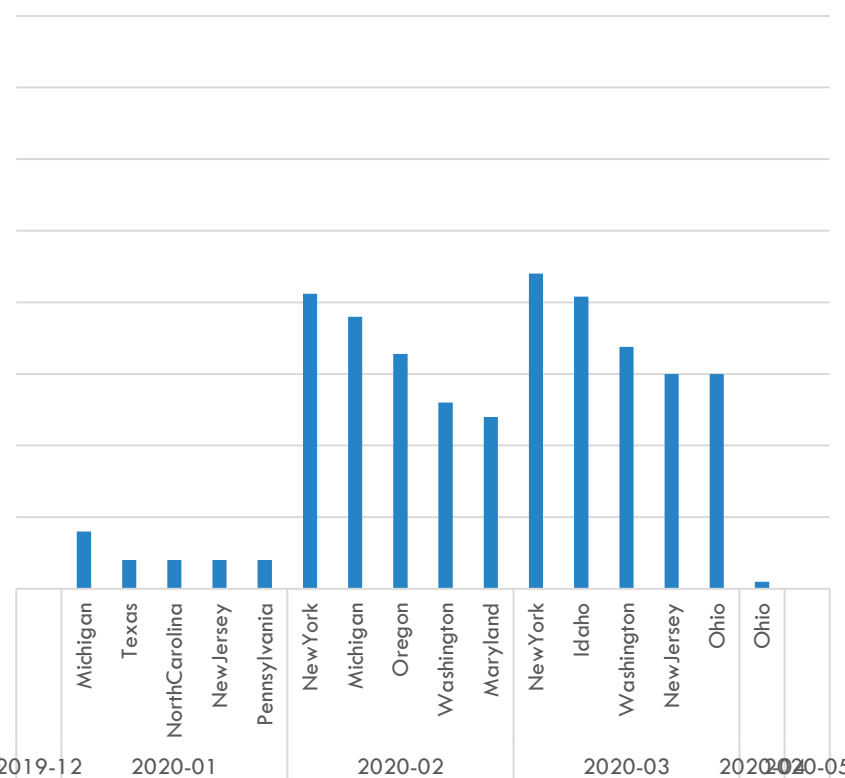
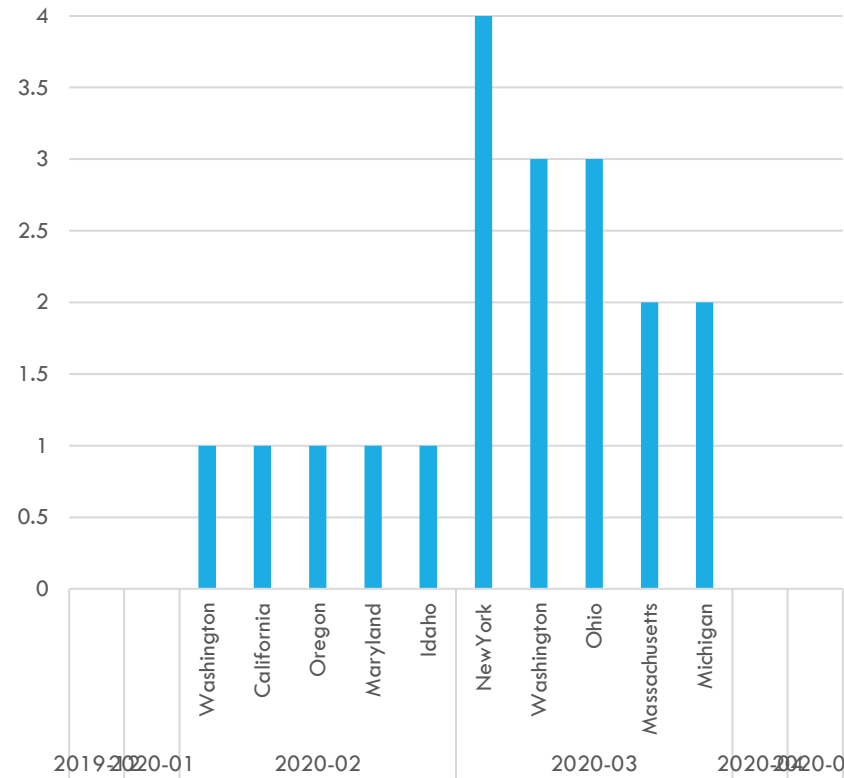


TOP 5 STATES CONNECTICUT SPREAD TO (AVERAGE)

NextStrain

TNet-Biased

TNet-Random

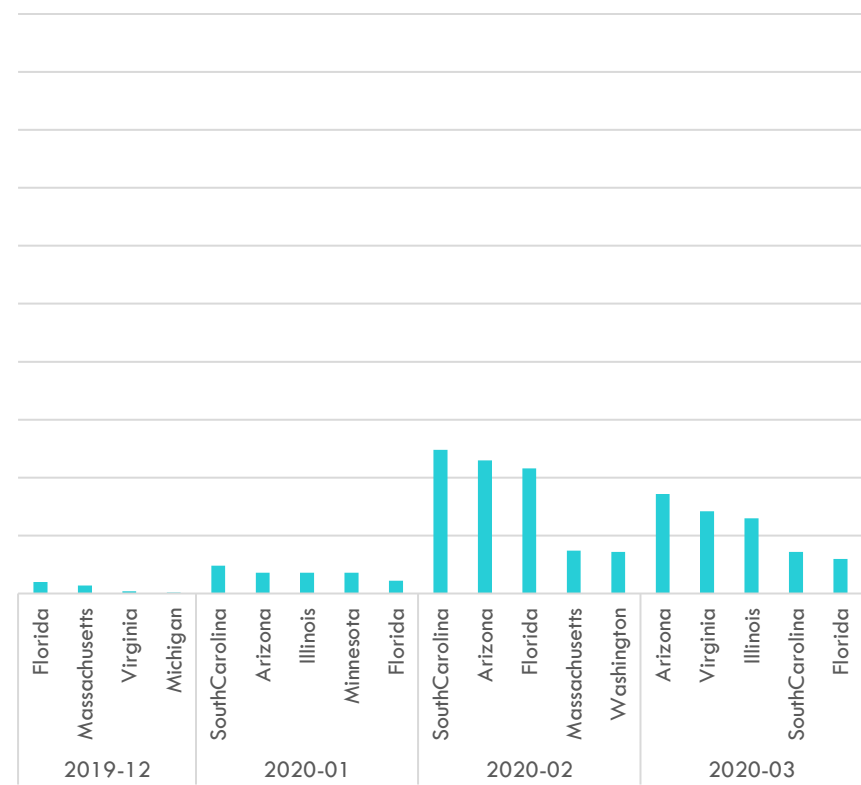
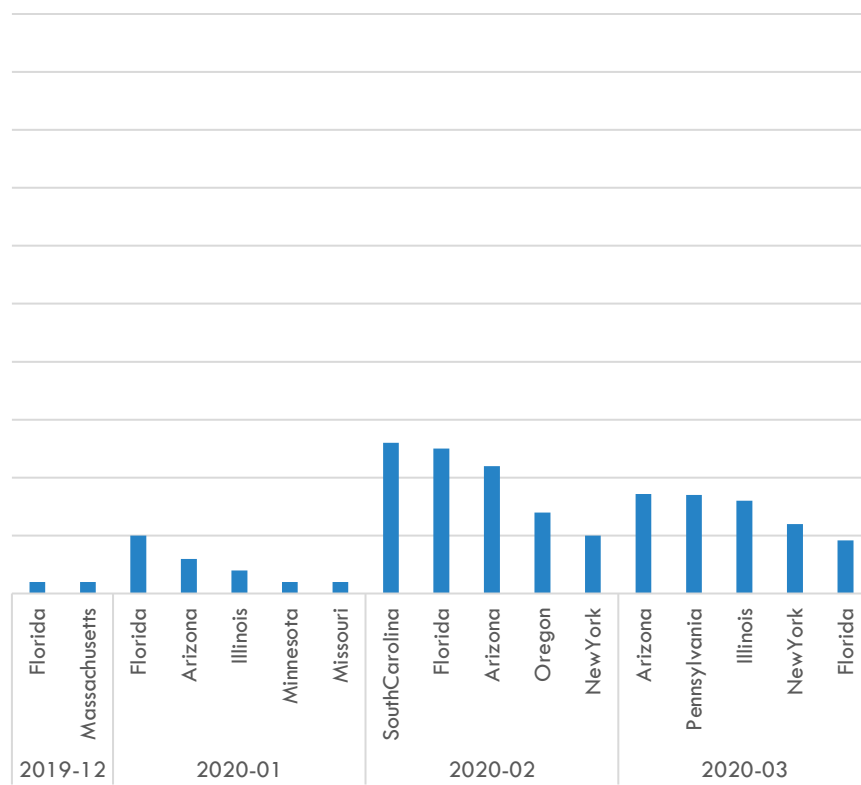
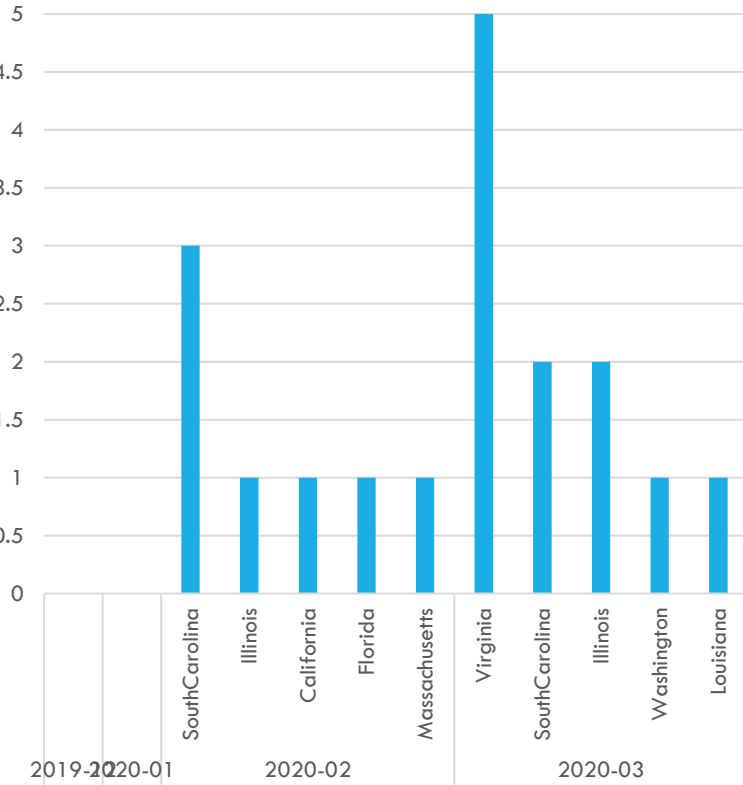


TOP 5 STATES GEORGIA RECEIVED FROM (AVERAGE)

NextStrain

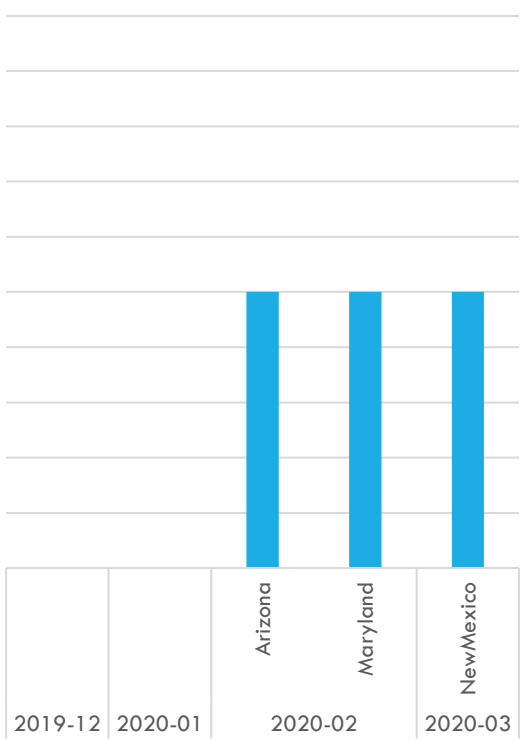
TNet-Biased

TNet-Random

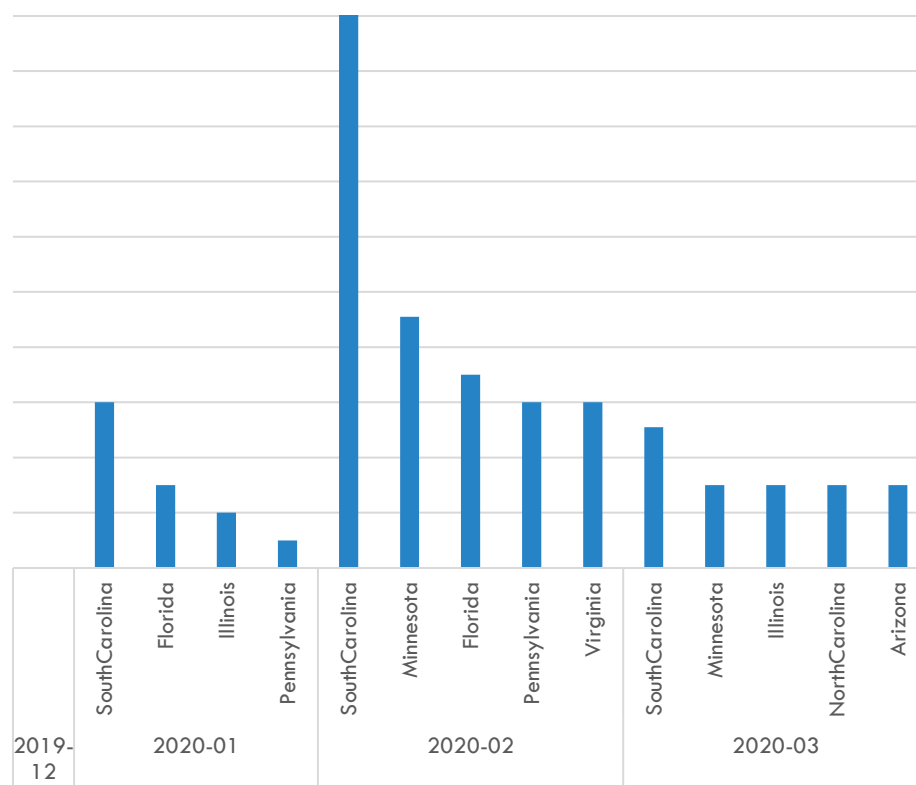


TOP 5 STATES GEORGIA SPREAD TO (AVERAGE)

NextStrain



TNet-Biased



TNet-Random

