Statistical Challenges in Tracking the Evolution of SARS-CoV-2

Lorenzo Cappello, Jaehee Kim, Sifan Liu and Julia A. Palacios

SUPPLEMENTARY MATERIAL

Table S1
Web URLs of phylodynamic method implementations for SARS-CoV-2 studies in Table 1.

Method	Implementation	Web URL
SMC	annealedSMC	https://github.com/liangliangwangsfu/annealedSMC
Online Bayesian phylodynamics	BEAST 1.10	https://beast.community/online_inference
VI for phylogenetics	bito	https://github.com/phylovi/bito
BDSP	BDSKY in BEAST2	https://github.com/BEAST2-Dev/bdsky
Structured BDSP	bdmm in BEAST2	https://github.com/denisekuehnert/bdmm
Coalescent	PhyDyn in BEAST2	https://github.com/mrc-ide/PhyDyn
Coalescent	LNAphyloDyn	https://github.com/MingweiWilliamTang/LNAphyloDyn
Parametric N_e , λ	NA	NA .
Nonparametric $N_e, \lambda = \exp(\beta_0) N_e^{\beta_1}$	phylodyn in R	https://github.com/mdkarcher/phylodyn
Epoch Skyline plot (Nonparametric N_e , λ)	BESP in BEAST2	https://github.com/laduplessis/besp
AdaPref (Nonparametric $N_e, \lambda = \beta N_e$)	adaPref in R	https://github.com/lorenzocapp/adapref
Nonparametric N_e , $\lambda = \exp(\beta_0) N_e^{\beta_1} + (\beta' \mathbf{X}(t))_{t \ge 0}$	BEAST	https://beast.community/