

Supplementary Information

Transmission dynamics of HIV-1 subtype B strains in

Indonesia

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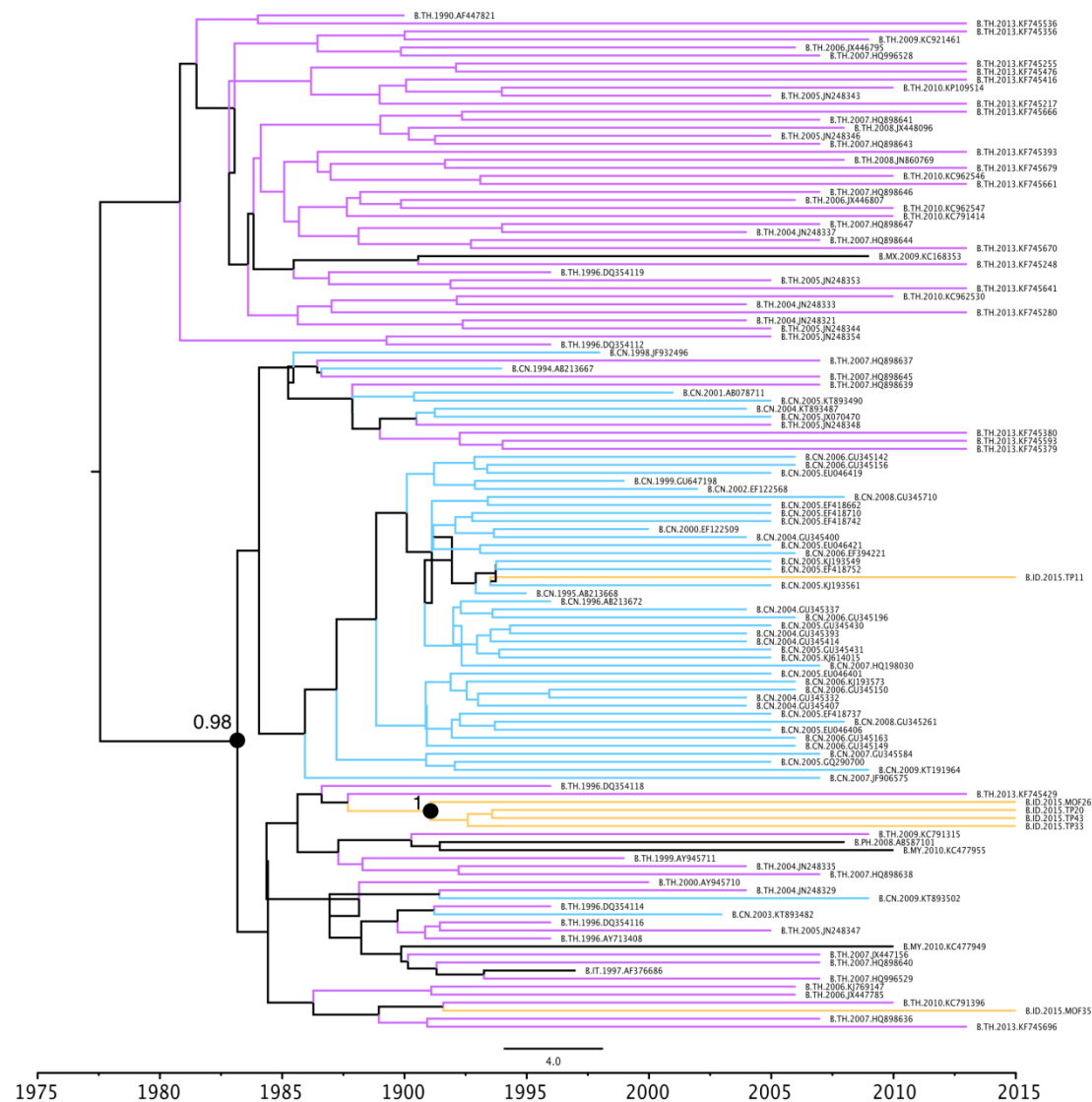
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Supplementary Figure S1. MCC phylogenetic tree of *pol* regions based on the CN clade after the addition of all available Thai strains.

All available Thai strains were added to CN clade strains. The MCC tree of HIV-1B was estimated by a Bayesian MCMC approach. Greater than or equal to 0.9 of posteriori probability was shown in the tree. The geographic origins of the sequences are represented by colors or two-letter country codes as follows: CN, China; ID, Indonesia; IT, Italy; MY, Malaysia; MX, Mexico; PH, Philippines; TH, Thailand.