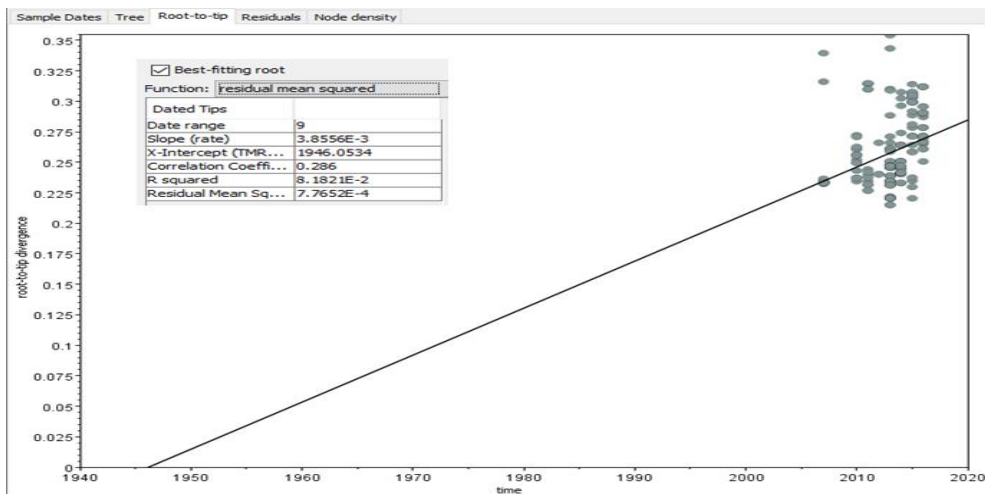


A



B

SupplementaryFigure 1. Root to tip regression analysis plots of the genetic distances against the year of isolation for the Bt-CoV sequences analyzed in this study. Figure 2a represents Afr-Bt-CoV, while 2b represents sequences of African, Asian and European origins.