

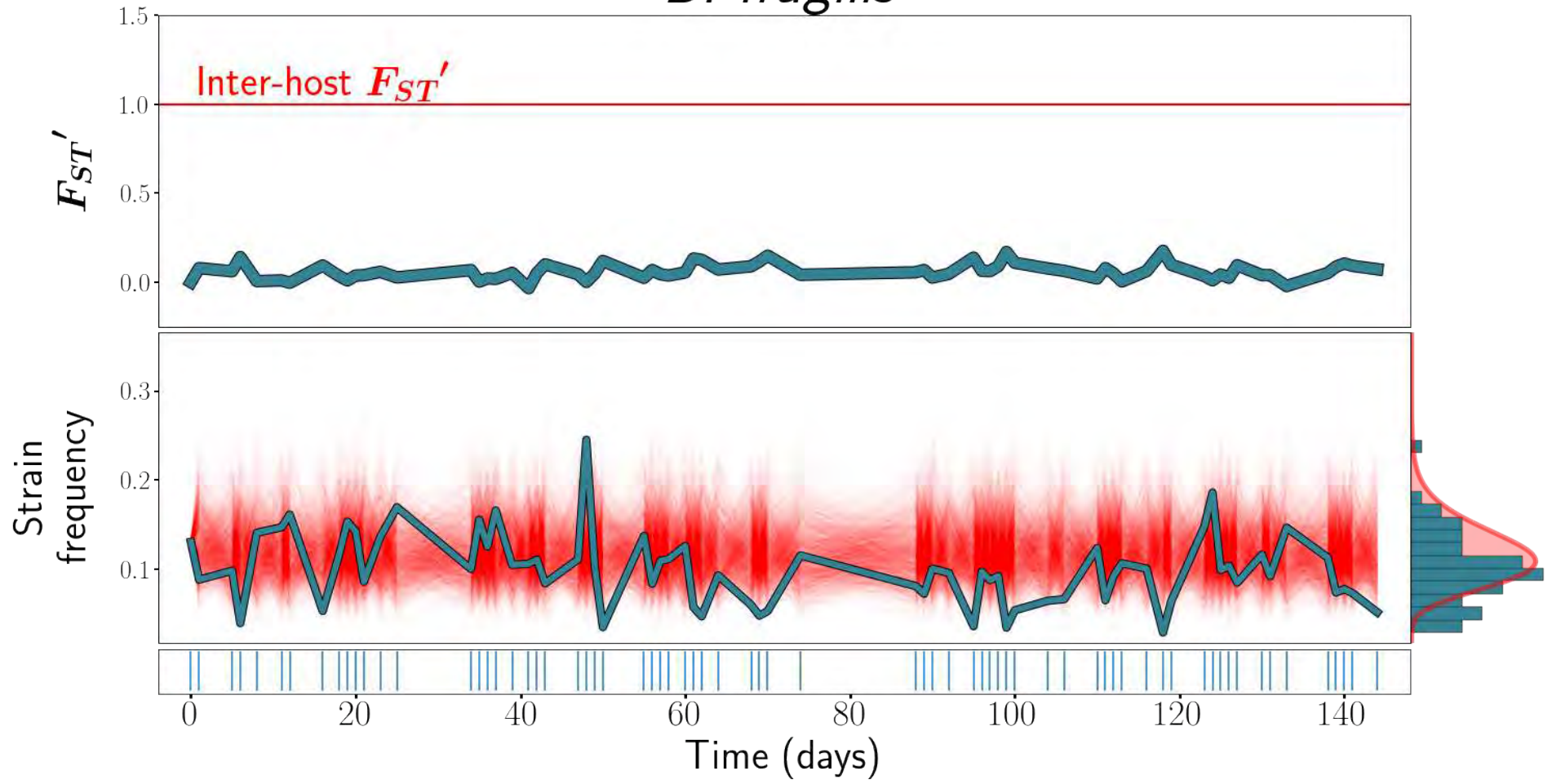
# S3 Text

$F_{ST}$ , strain frequency, and strain abundance dynamics plots for all species analyzed, for host *ao*. These plots are analogous to Main Text Figure 1. When only a single strain was detected, only  $F_{ST}$  and strain abundance dynamics plots, but no strain frequency plot, is included.

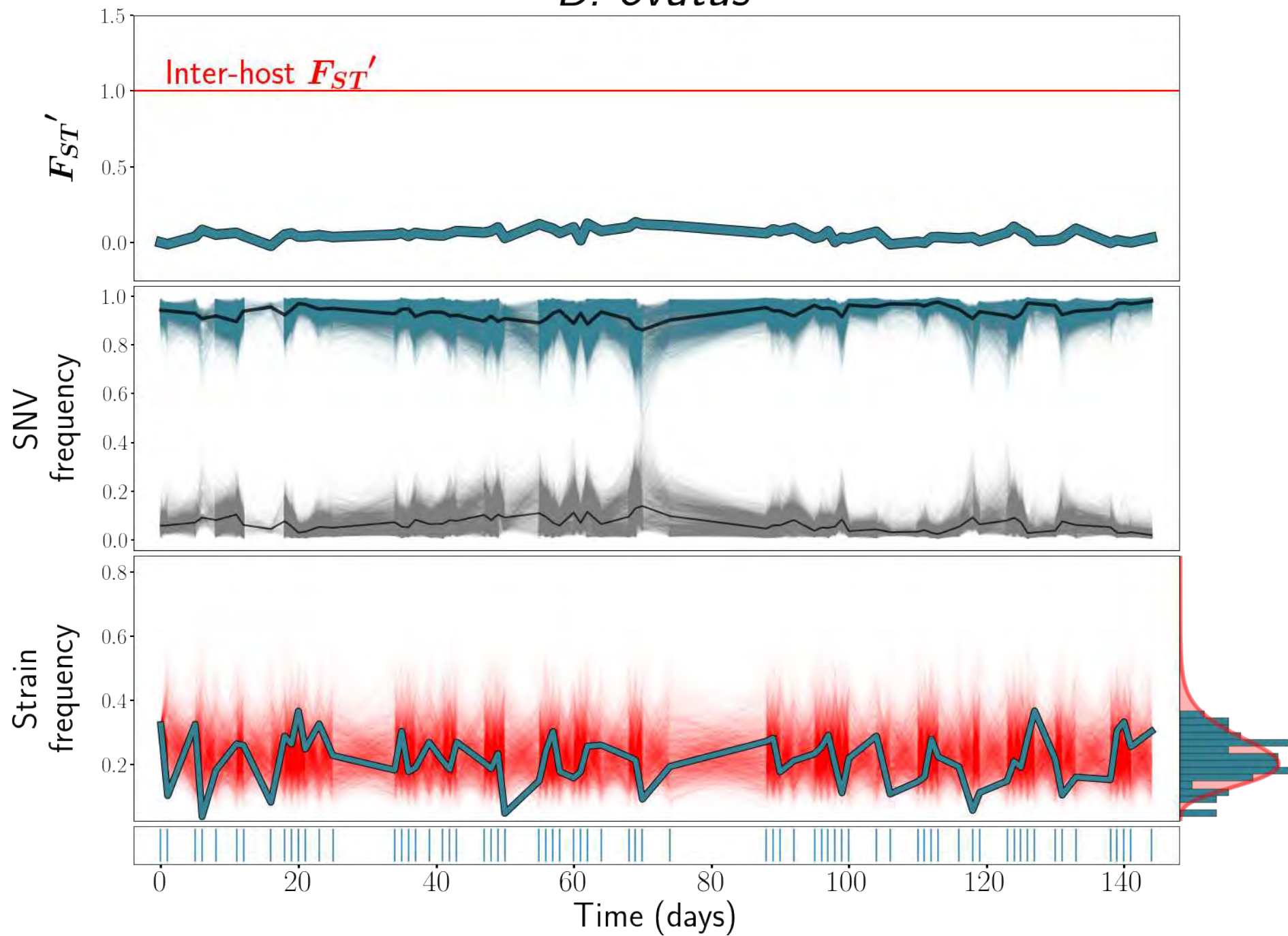
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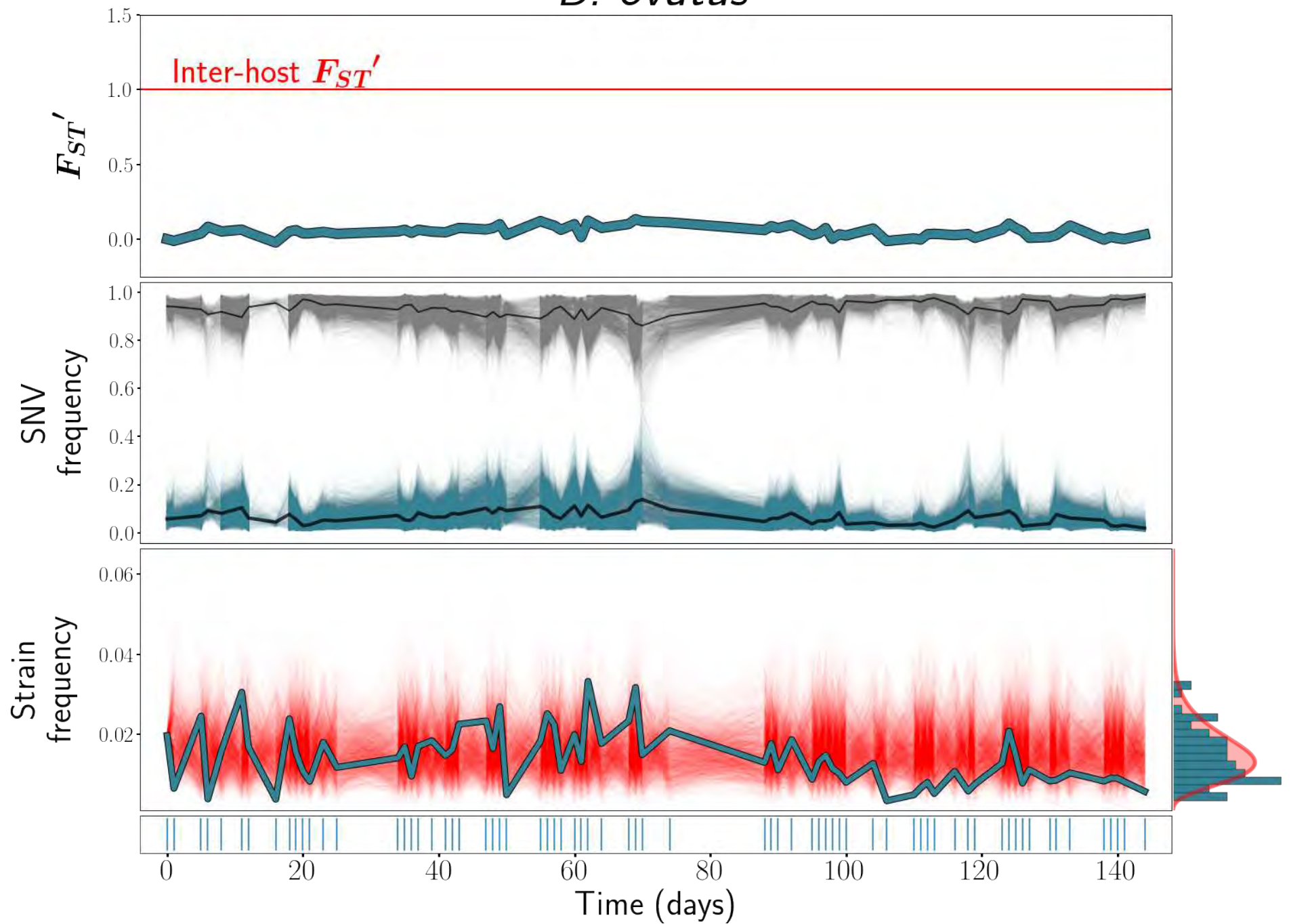
# *B. fragilis*



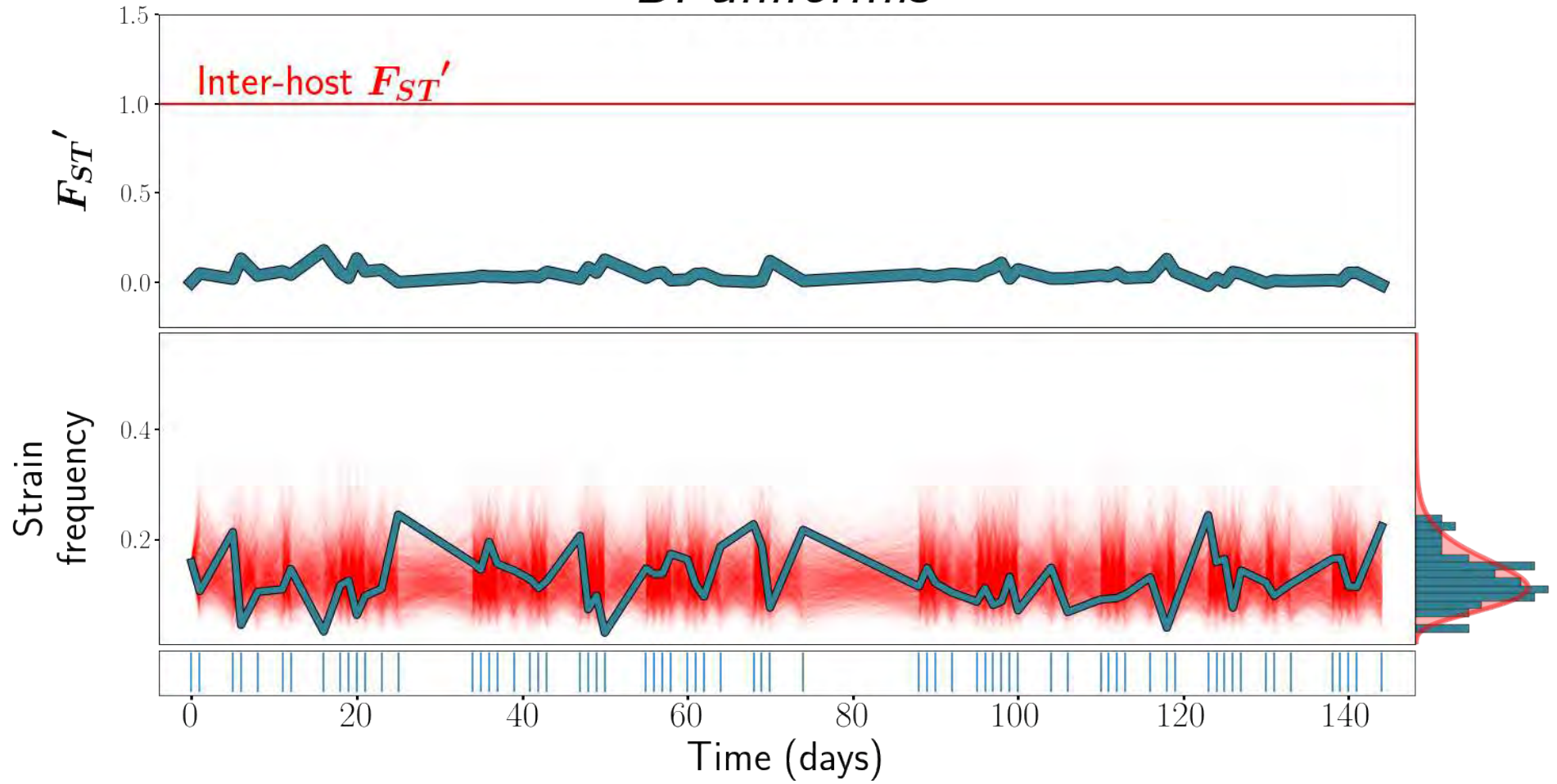
*B. ovatus*



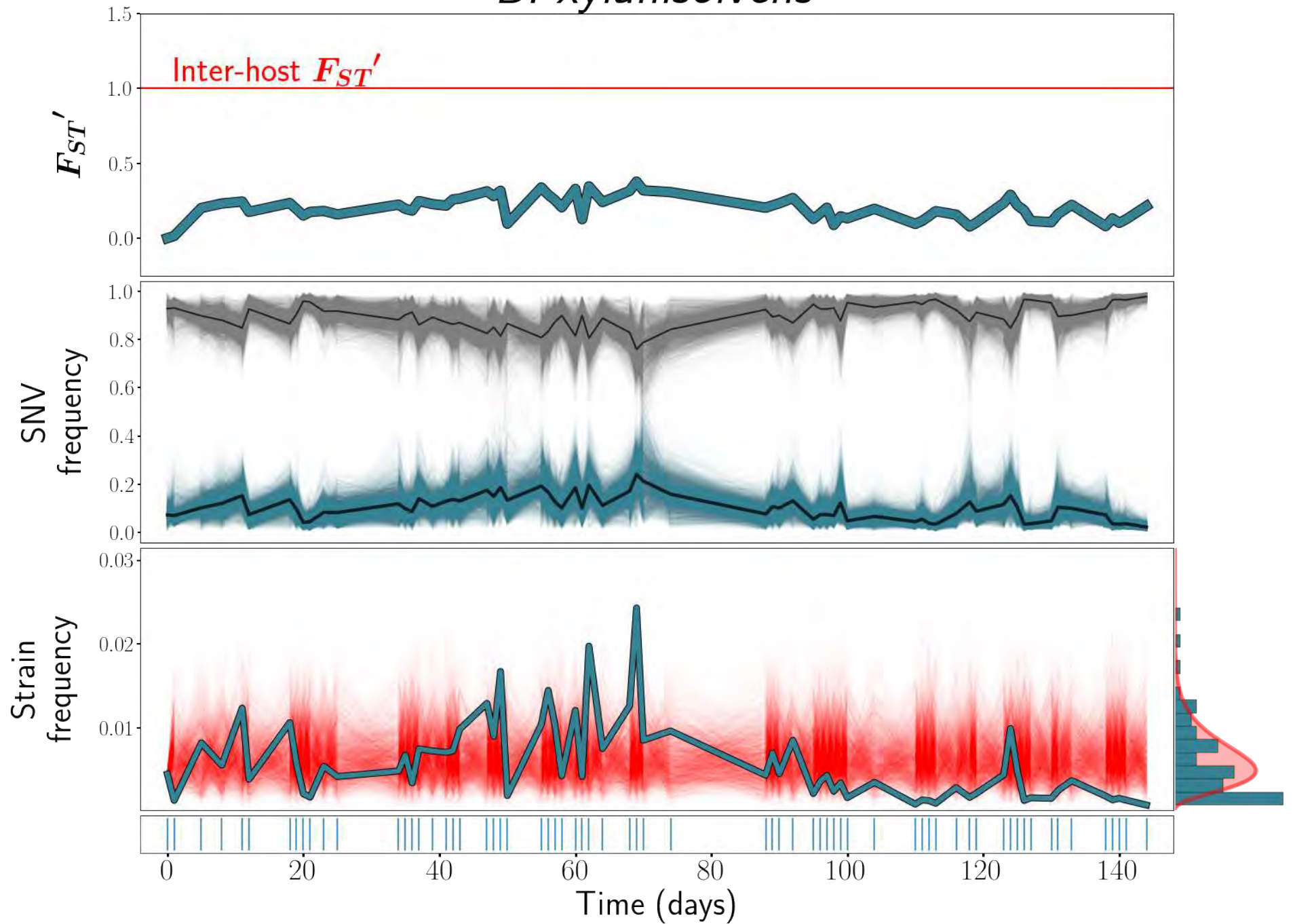
# *B. ovatus*



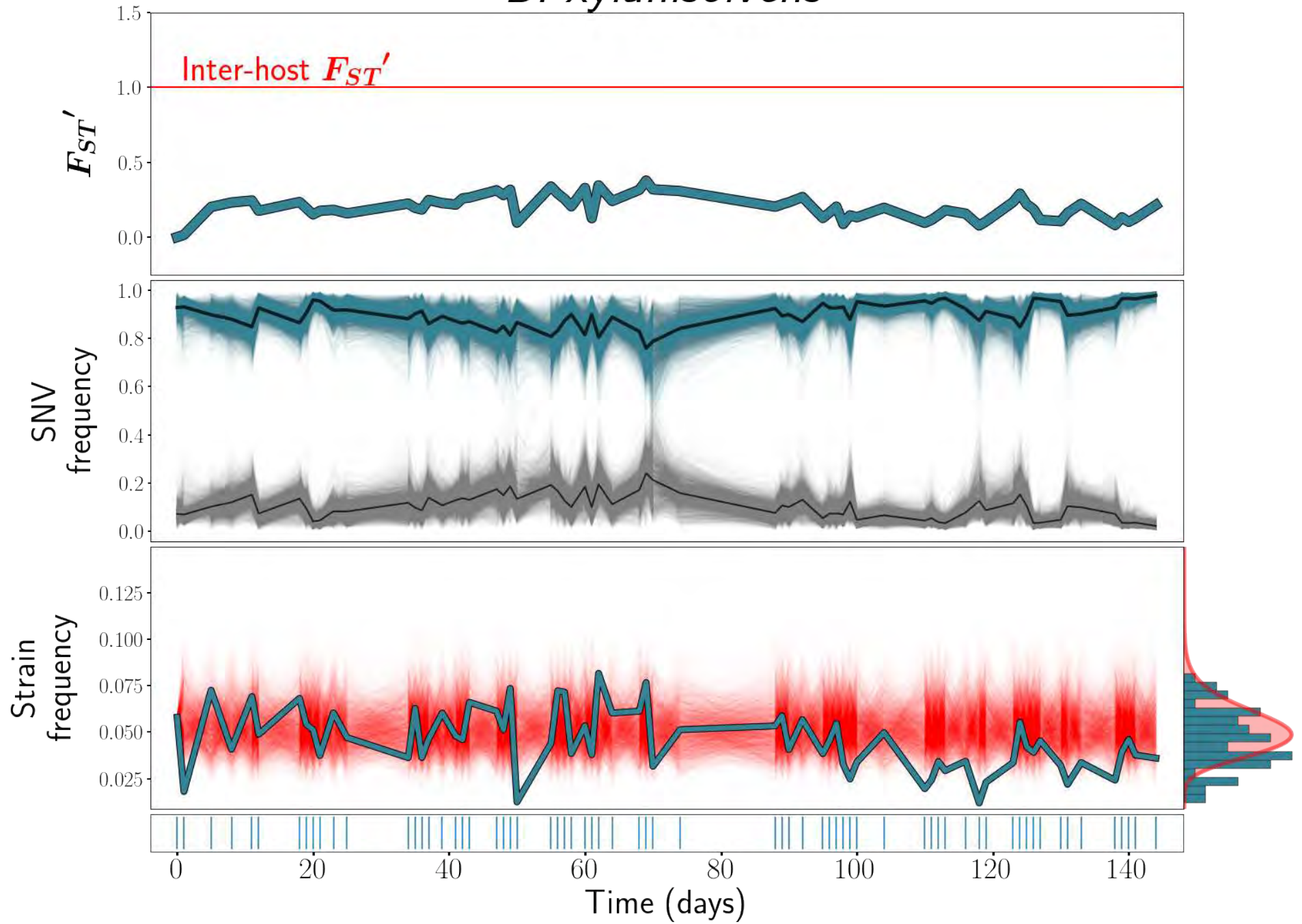
*B. uniformis*



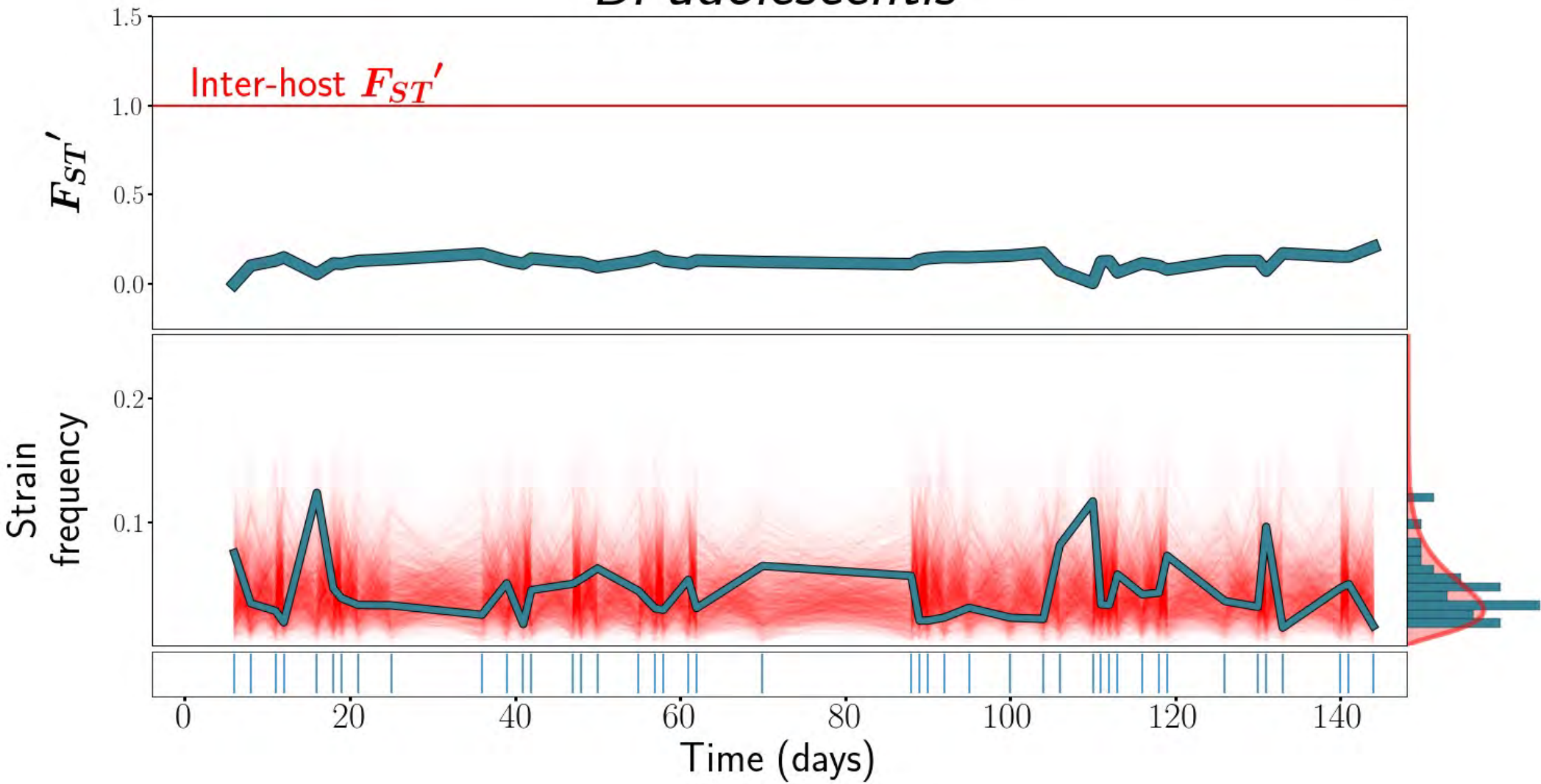
*B. xylanisolvens*



# *B. xylanisolvens*

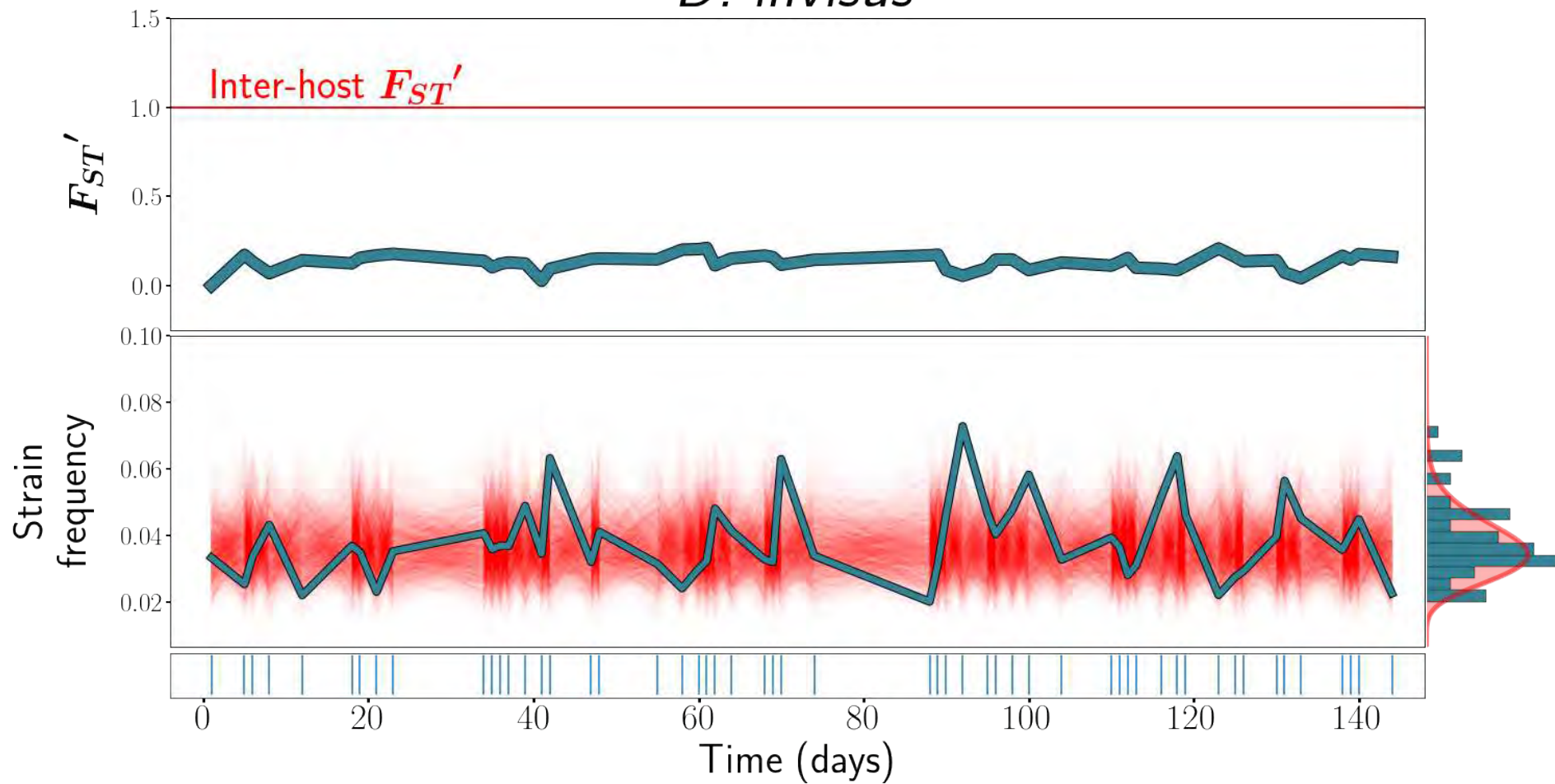


# *B. adolescentis*

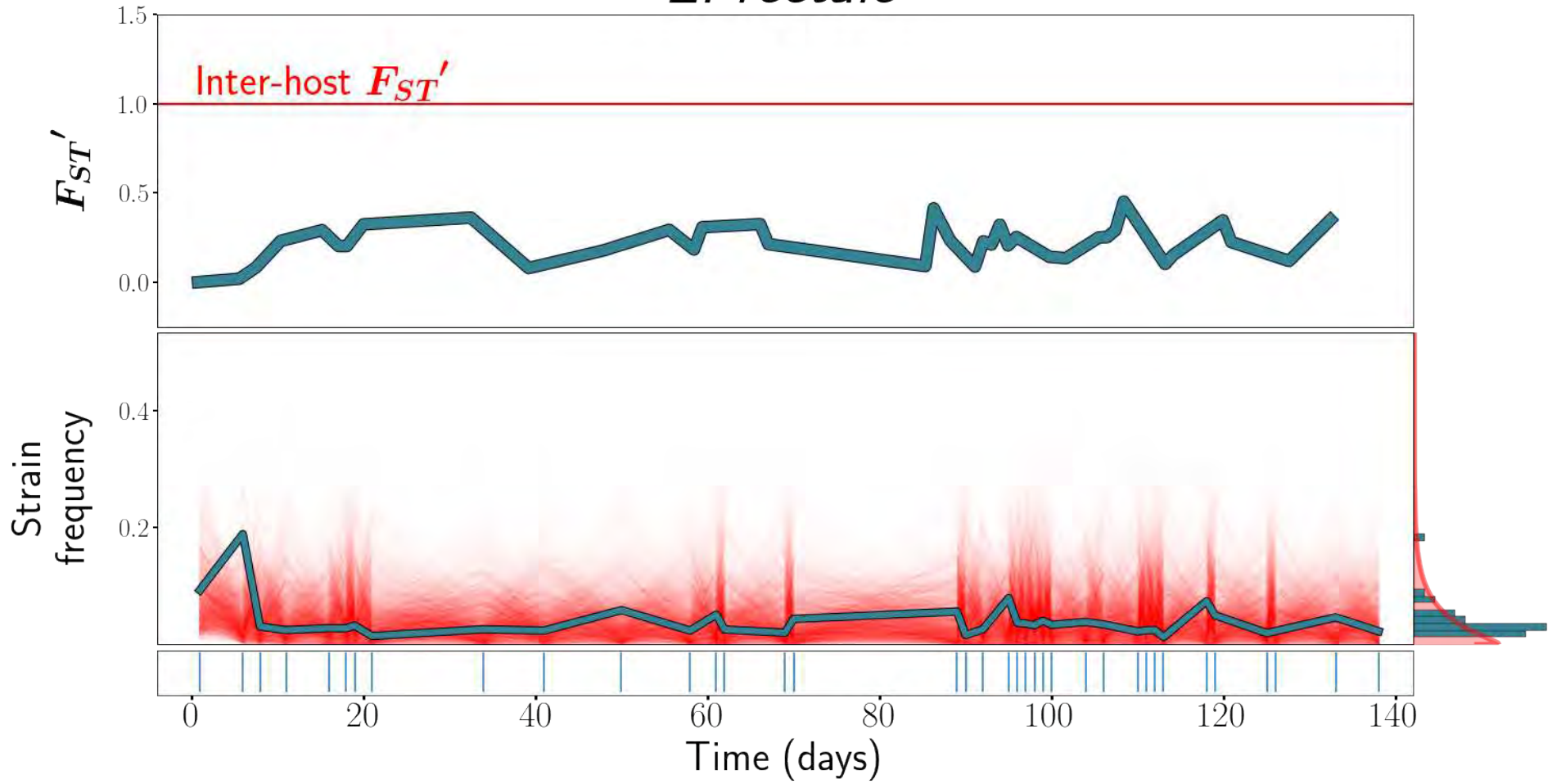




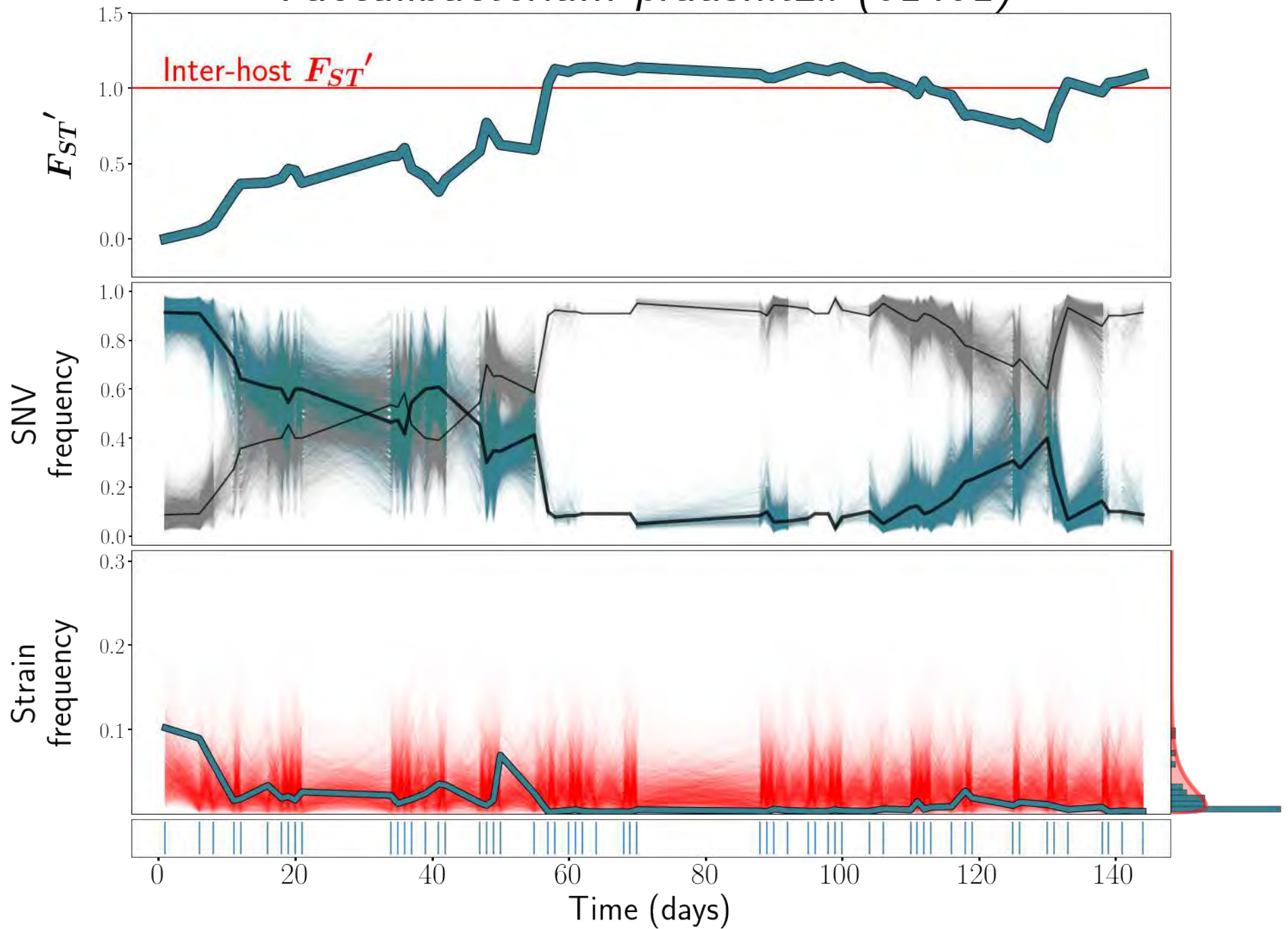
*D. invisus*



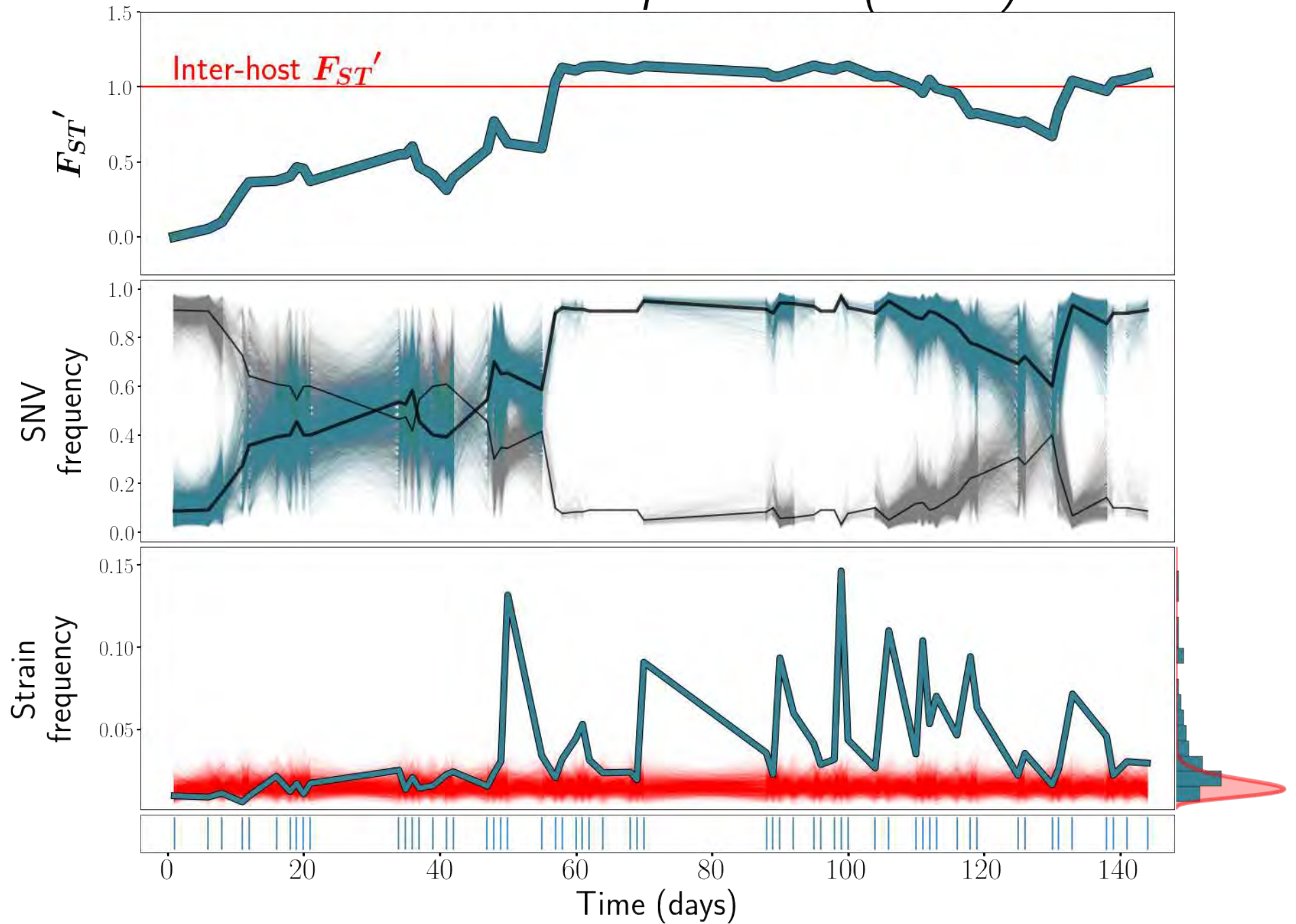
# *E. rectale*



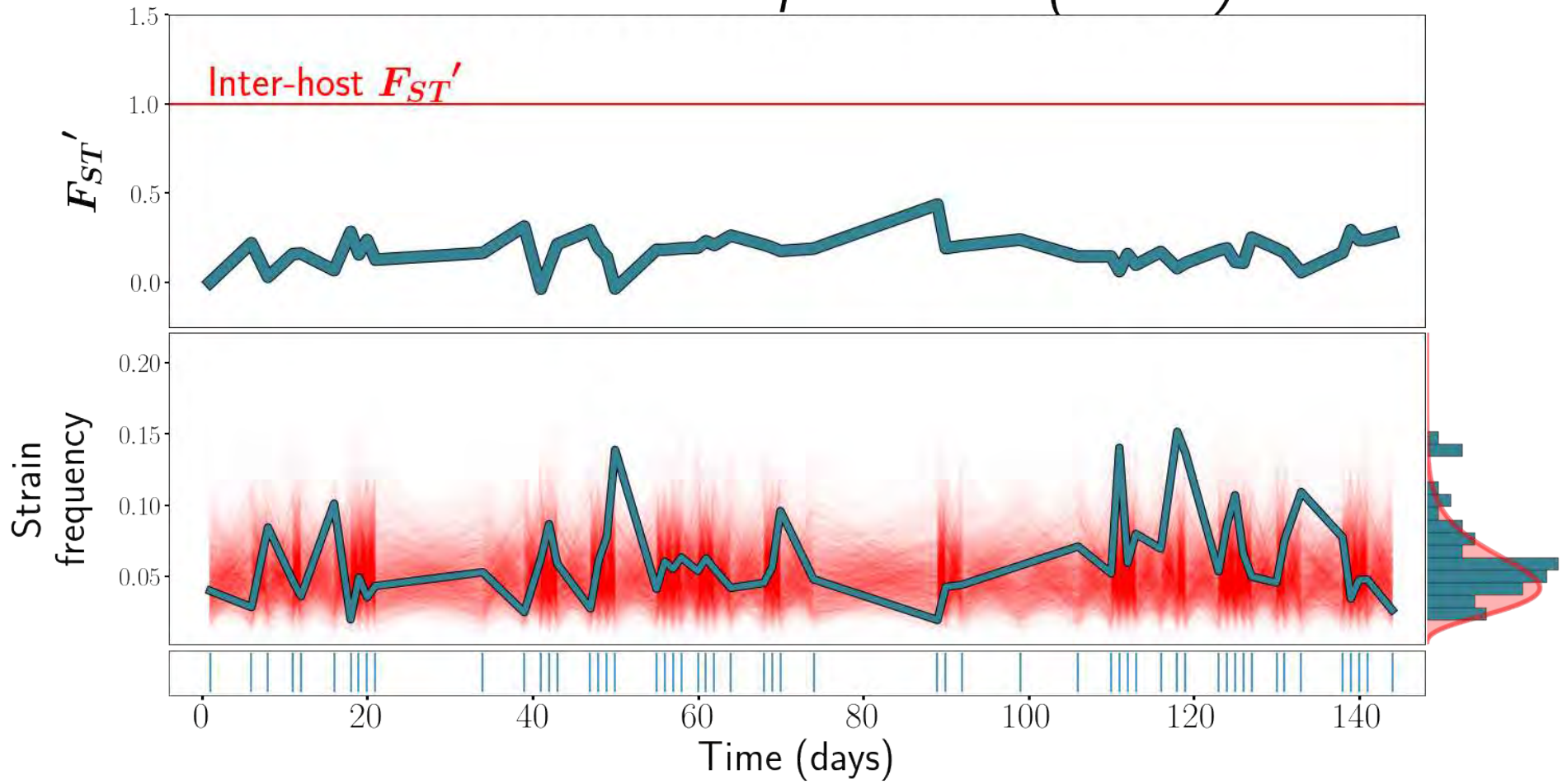
# *Faecalibacterium prausnitzii* (61481)



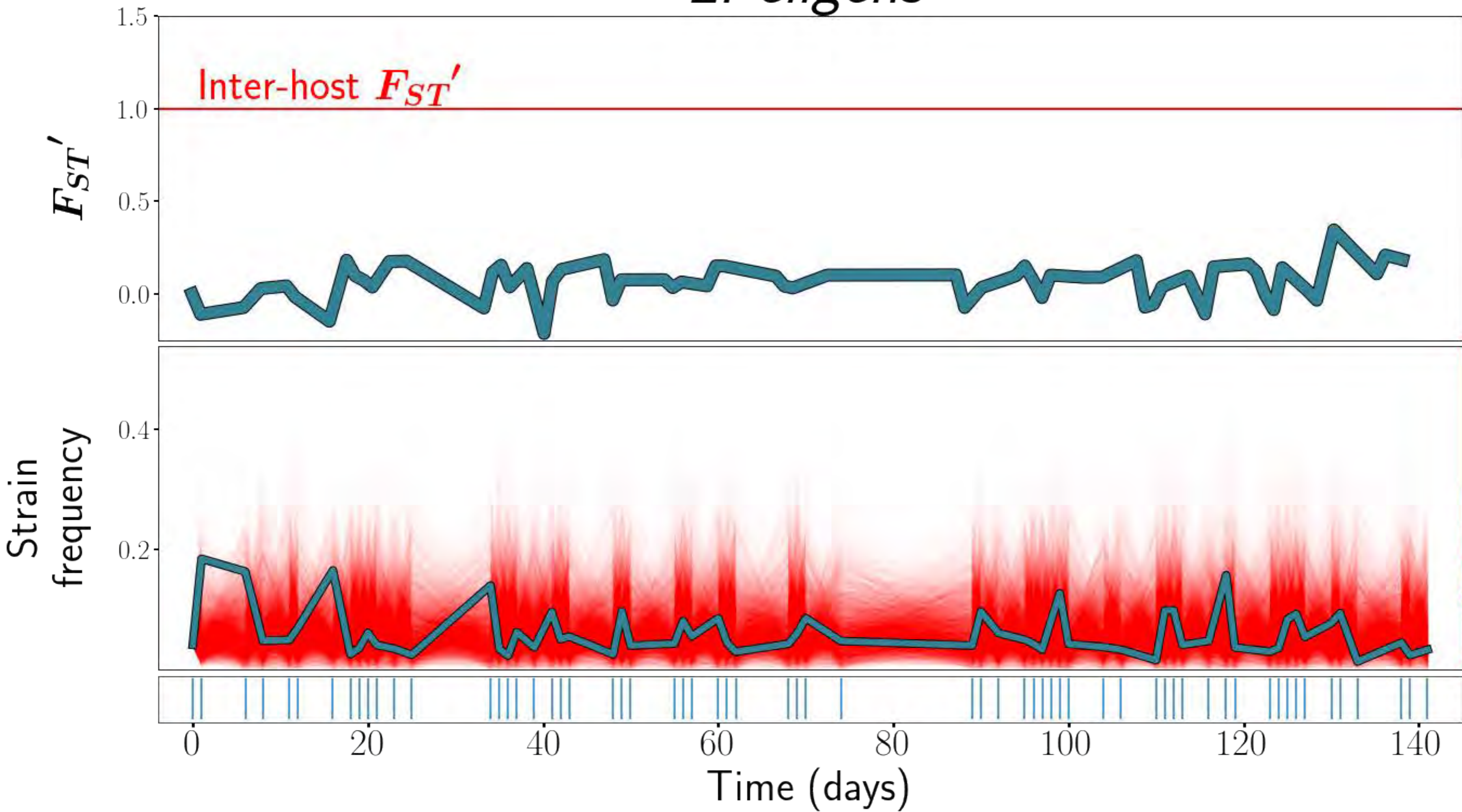
# *Faecalibacterium prausnitzii* (61481)



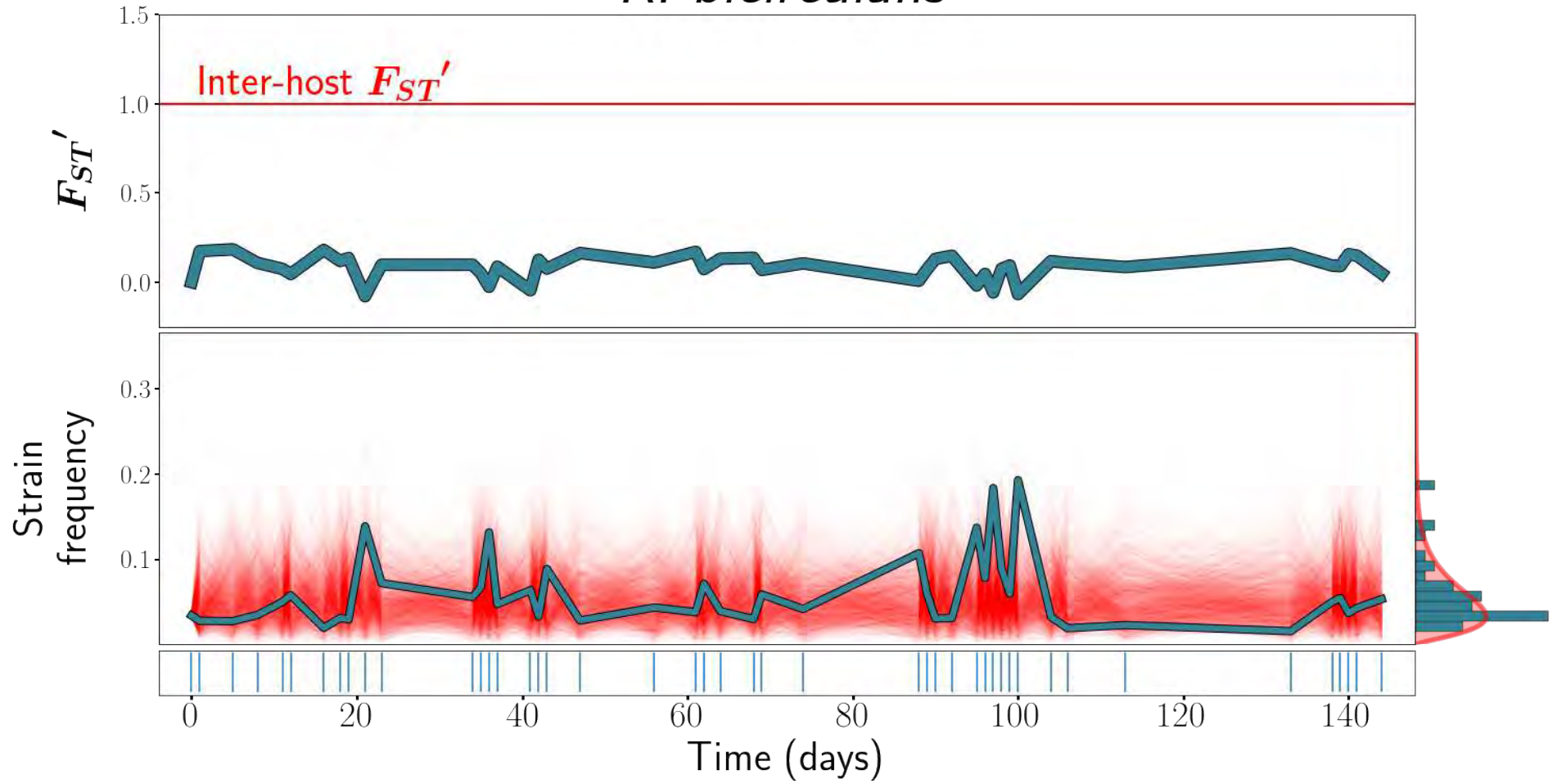
# *Faecalibacterium prausnitzii* (62201)



# *L. eligens*



# *R. bicirculans*



*P. distasonis*

