

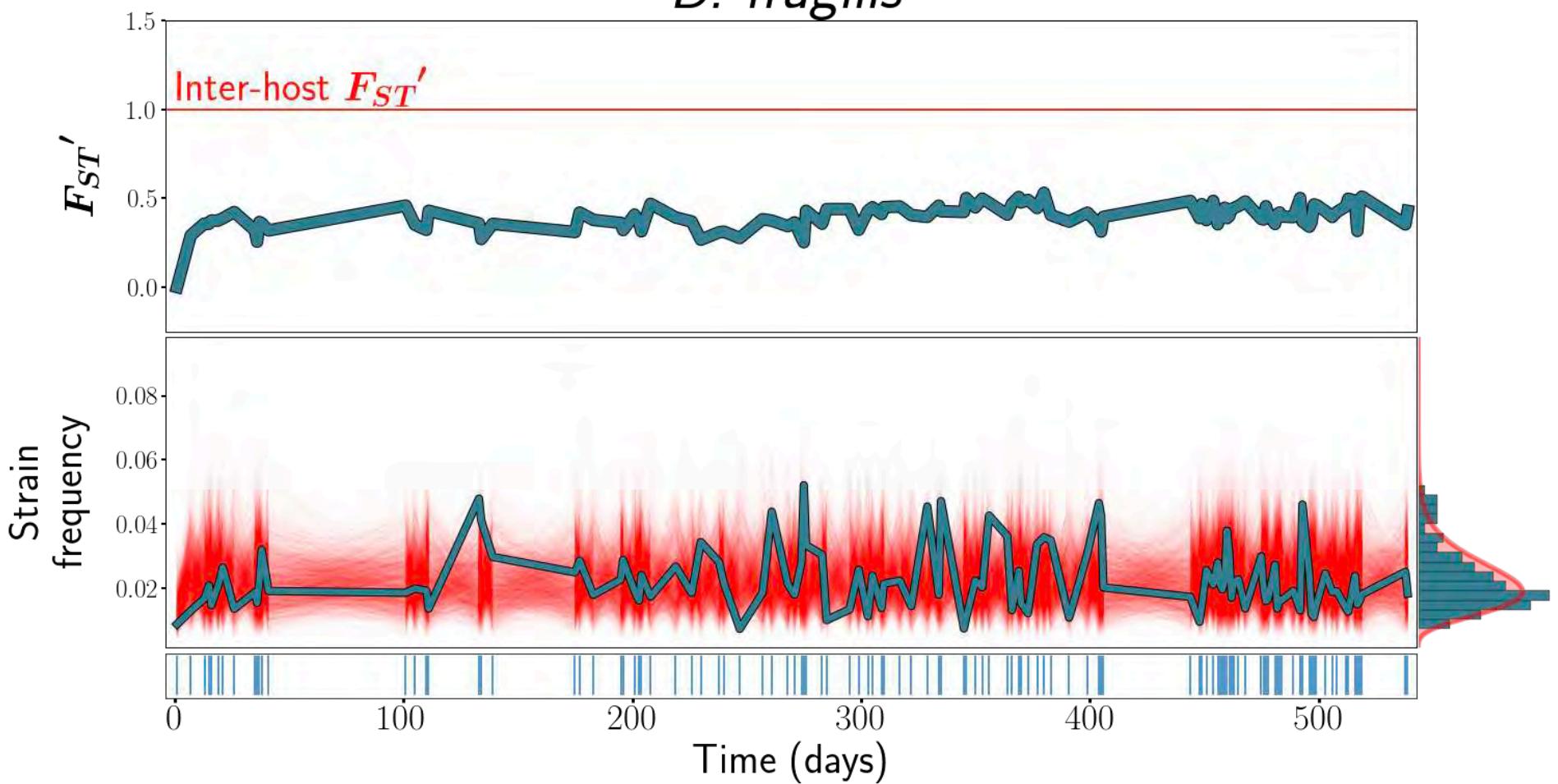
S2 Text

F_{ST}' , strain frequency, and strain abundance dynamics plots for all species analyzed, for host *am*. 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.

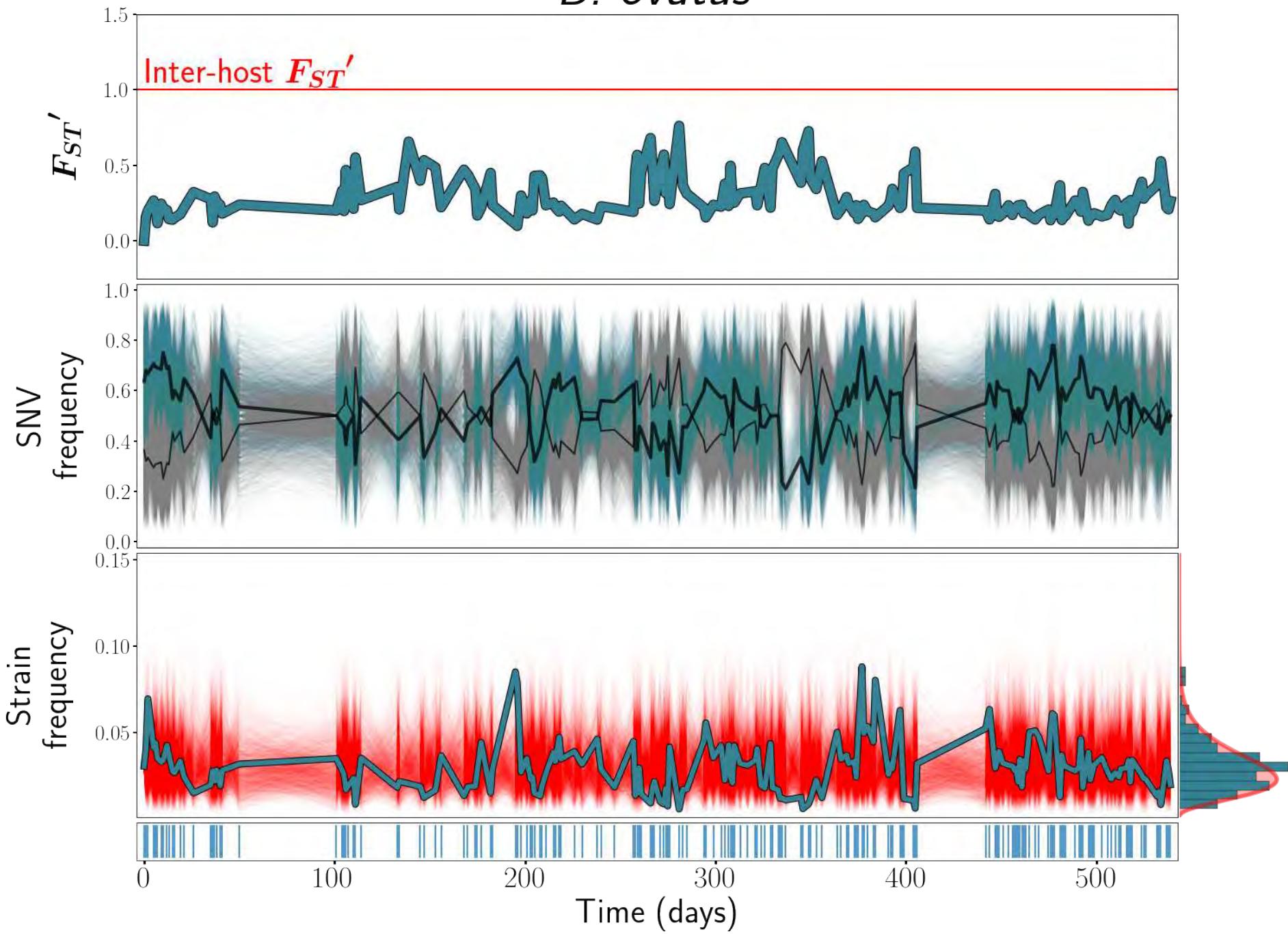
Table of Contents

<i>Bacteroides fragilis A</i>	1
<i>Bacteroides ovatus A</i>	2
<i>Bacteroides ovatus B</i>	3
<i>Bacteroides stercoris A</i>	4
<i>Bacteroides uniformis A</i>	5
<i>Phocaeicola vulgatus A</i>	6
<i>Phocaeicola vulgatus B</i>	7
<i>Phocaeicola vulgatus C</i>	8
<i>Bacteroides xylanisolvans A</i>	9
<i>Bacteroides xylanisolvans B</i>	10
<i>Barnesiella intenstinihominis A</i>	11
<i>Eubacterium rectale A</i>	12
<i>Eubacterium rectale B</i>	13
<i>Faecalibacterium prausnitzii (57453) A</i>	14
<i>Phascolarctobacterium sp. A</i>	15
<i>Parabacteroides merdae A</i>	16
<i>Ruminococcus bicirculans A</i>	17
<i>Ruminococcus bromii A</i>	18

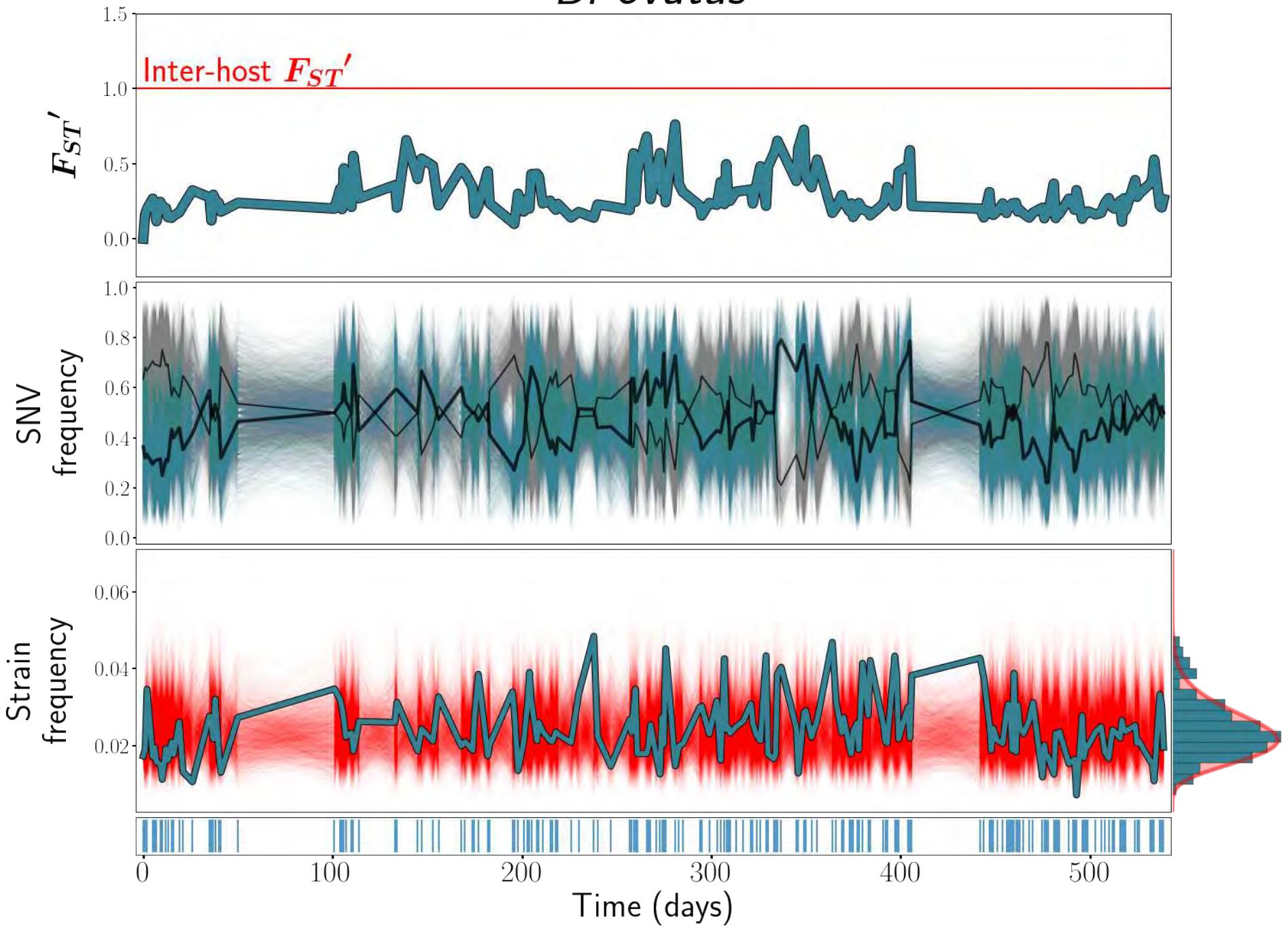
B. fragilis



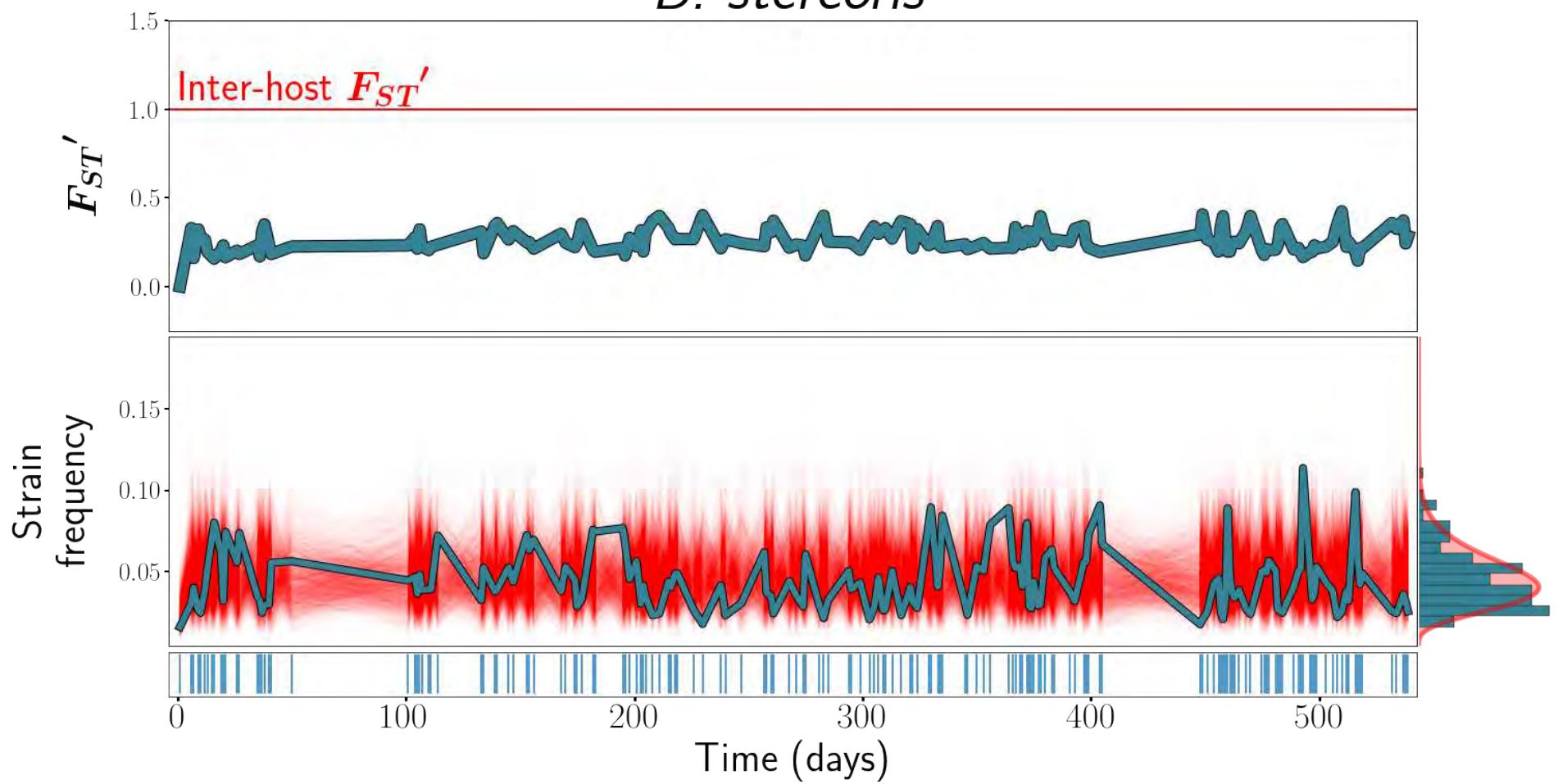
B. ovatus



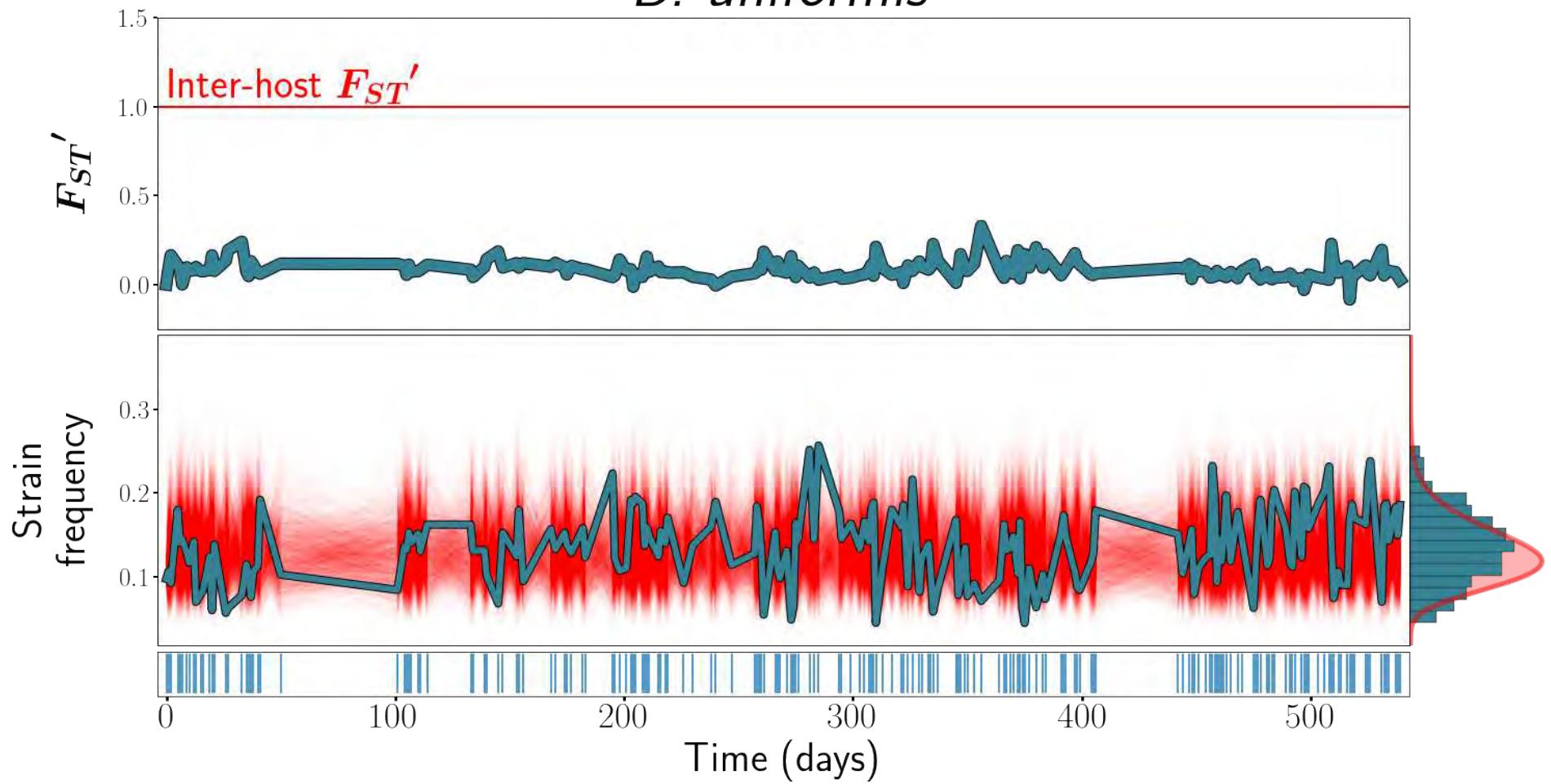
B. ovatus



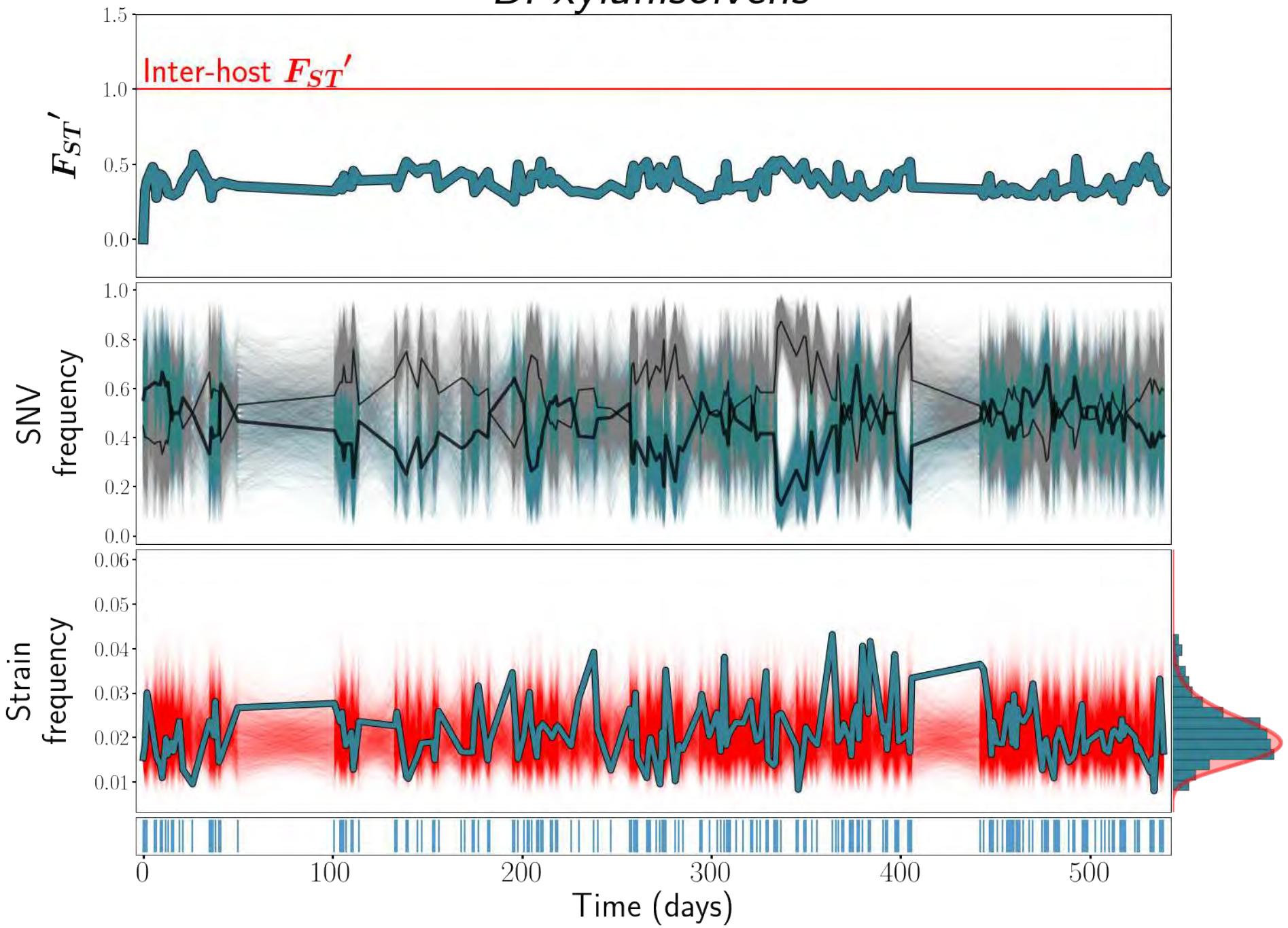
B. stercoris



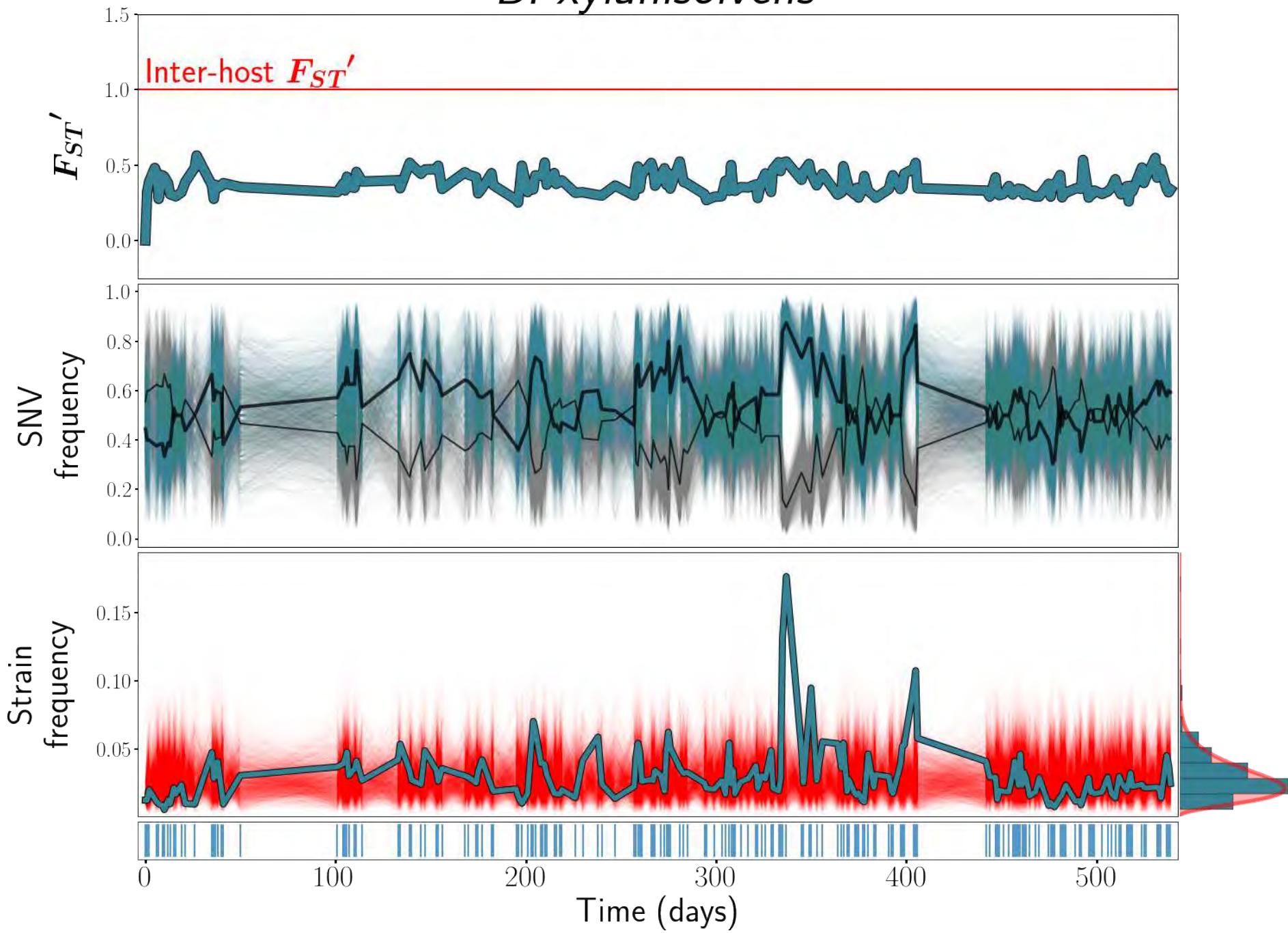
B. uniformis



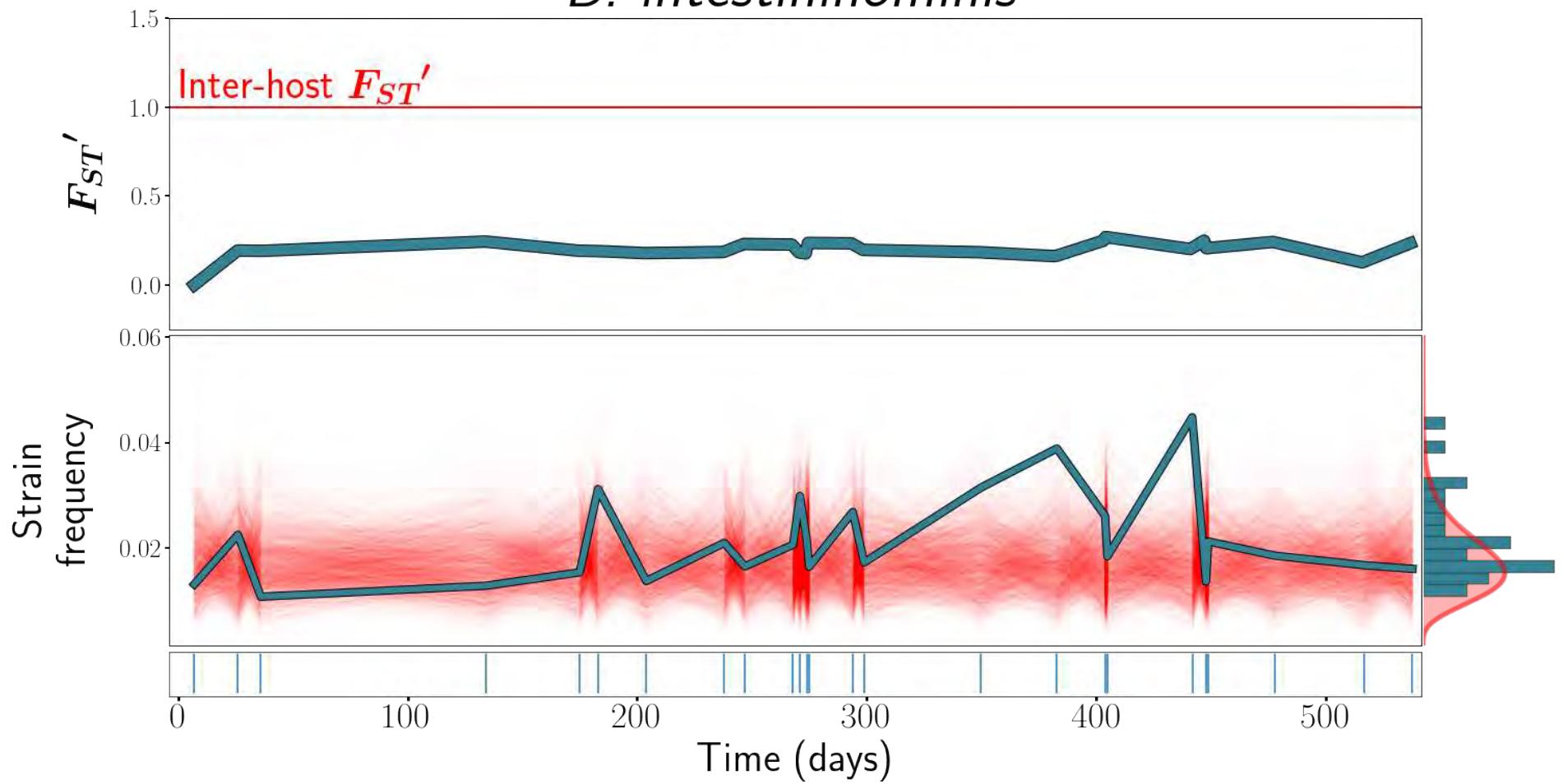
B. xylanisolvans



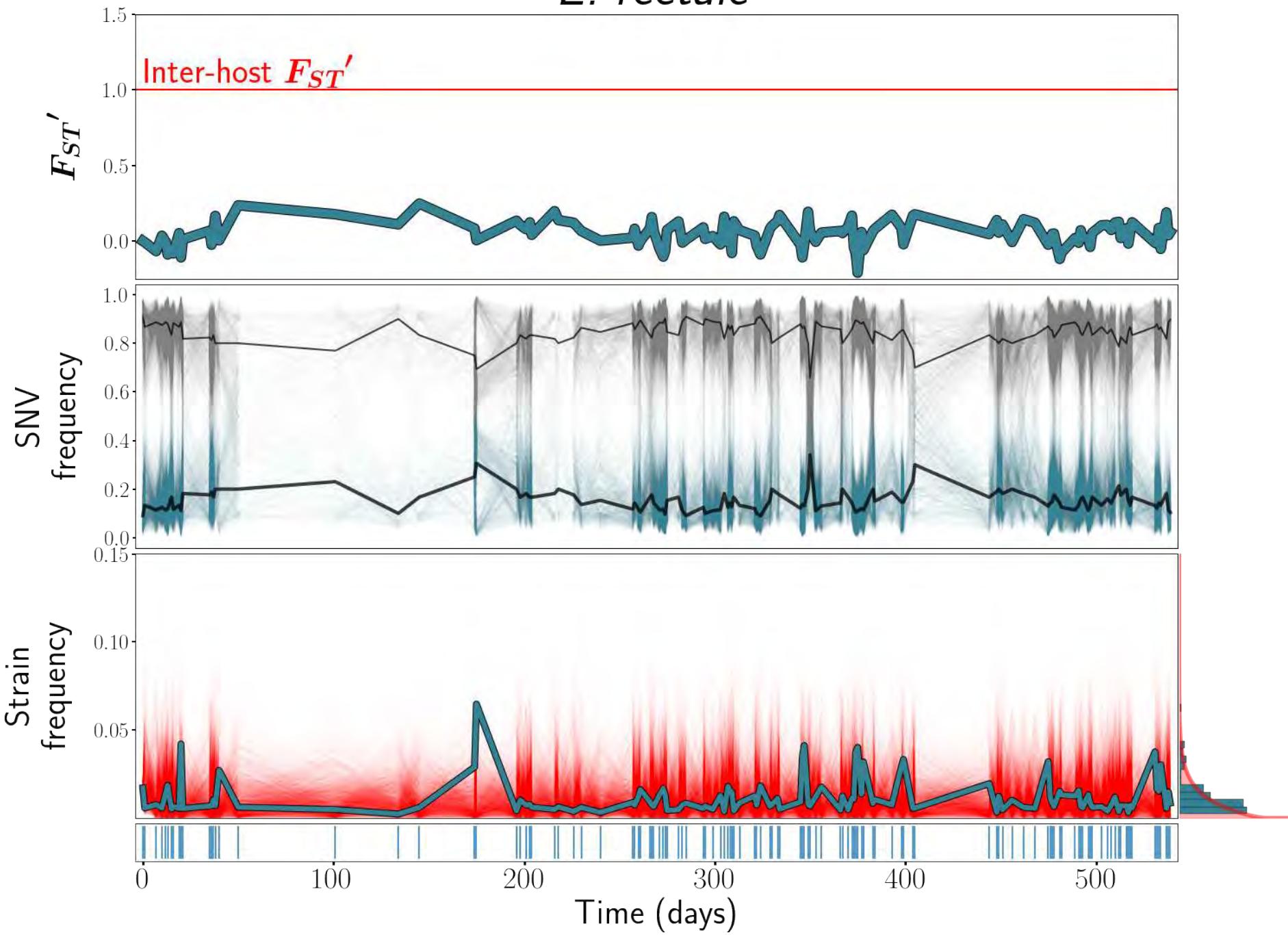
B. xylanisolvans



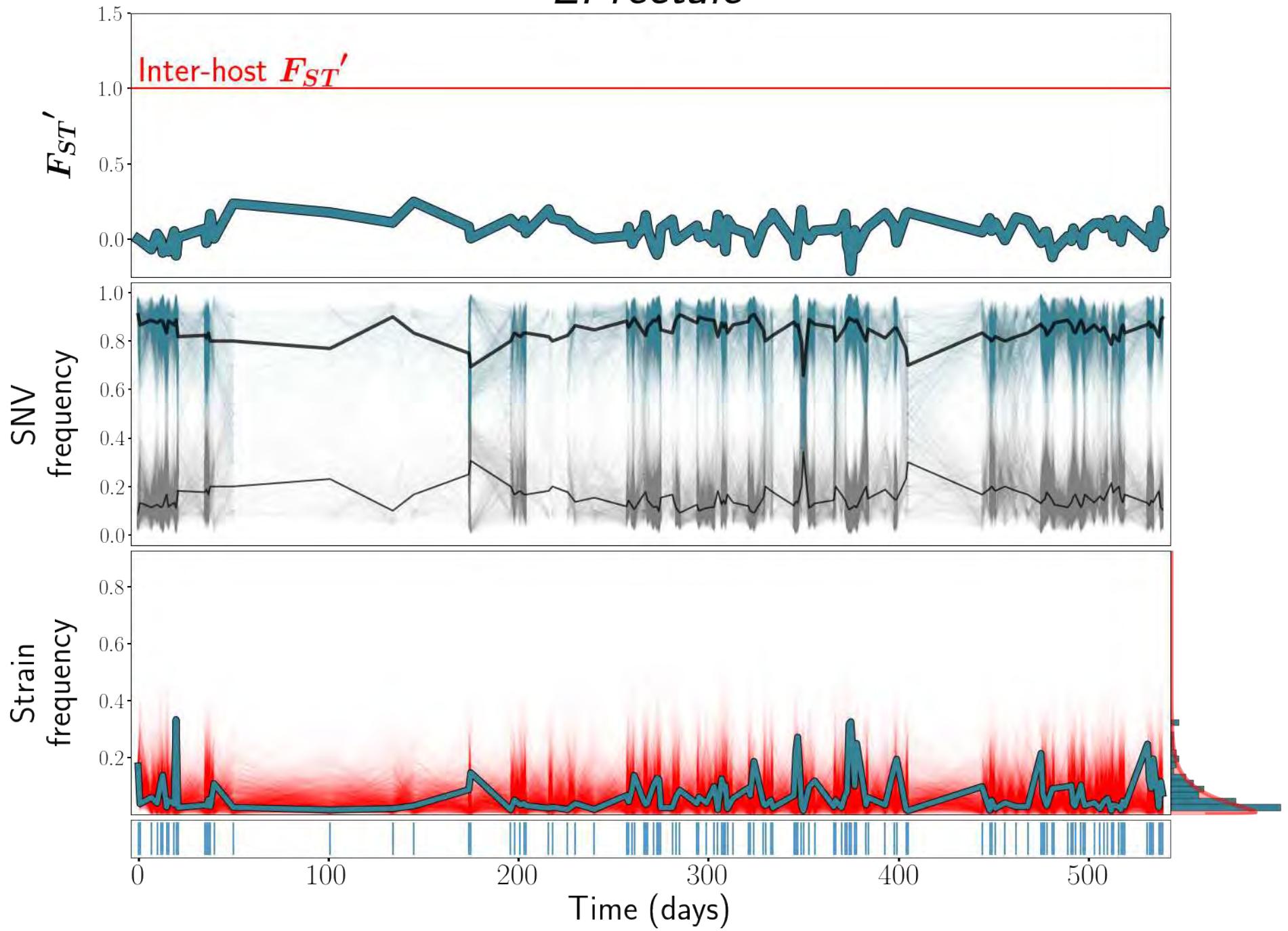
B. intestinihominis



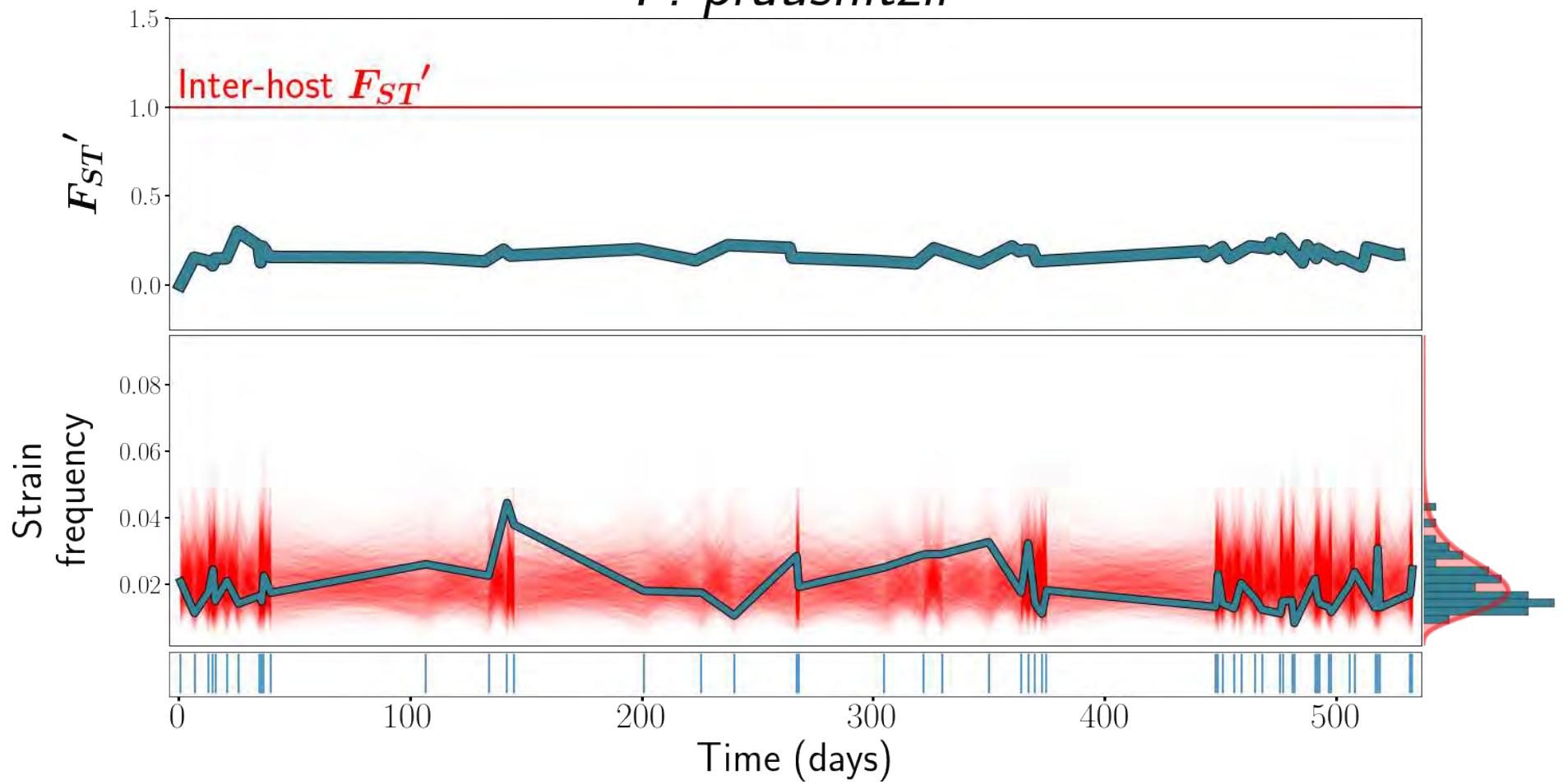
E. rectale



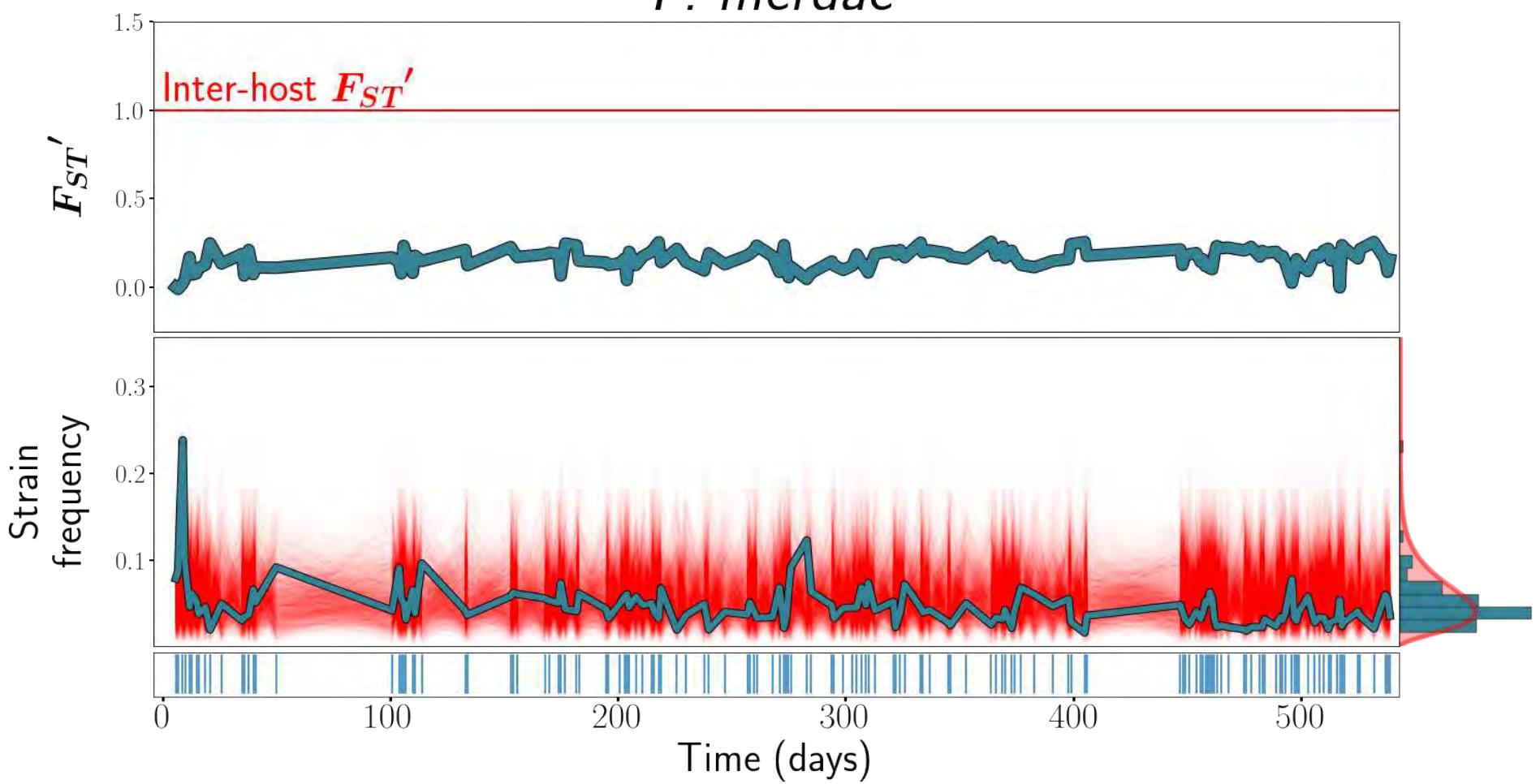
E. rectale



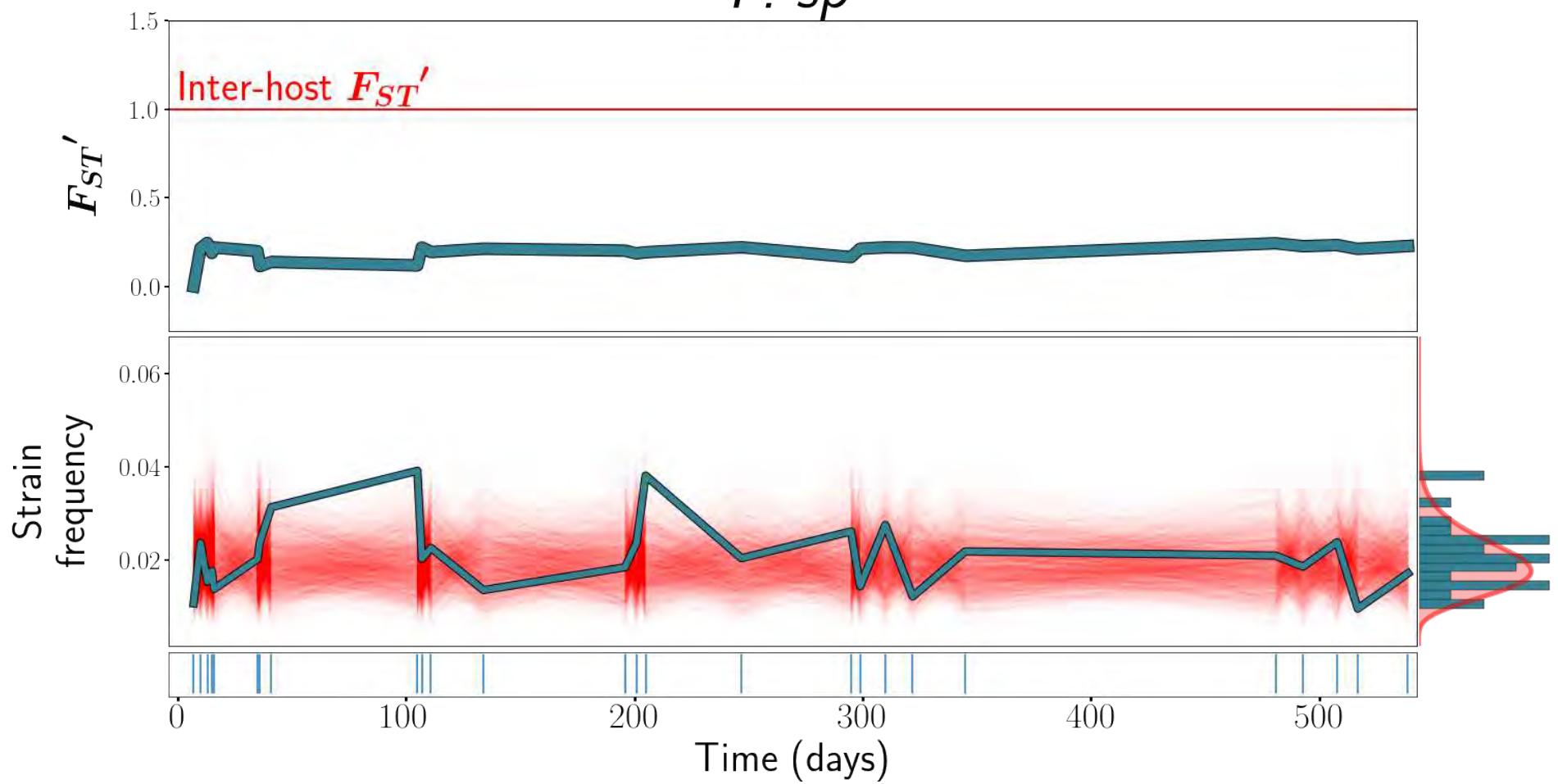
F. prausnitzii



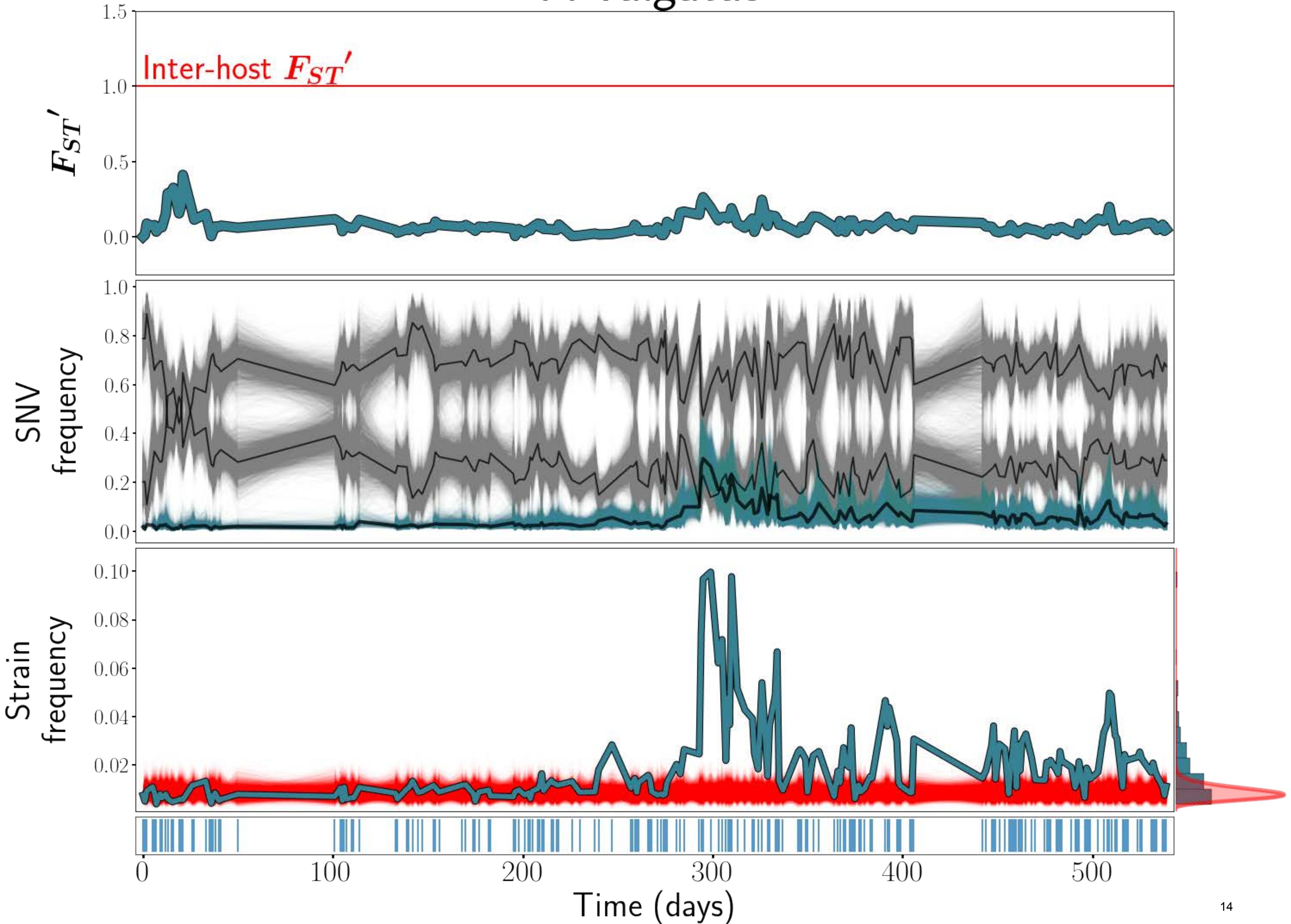
P. merdae



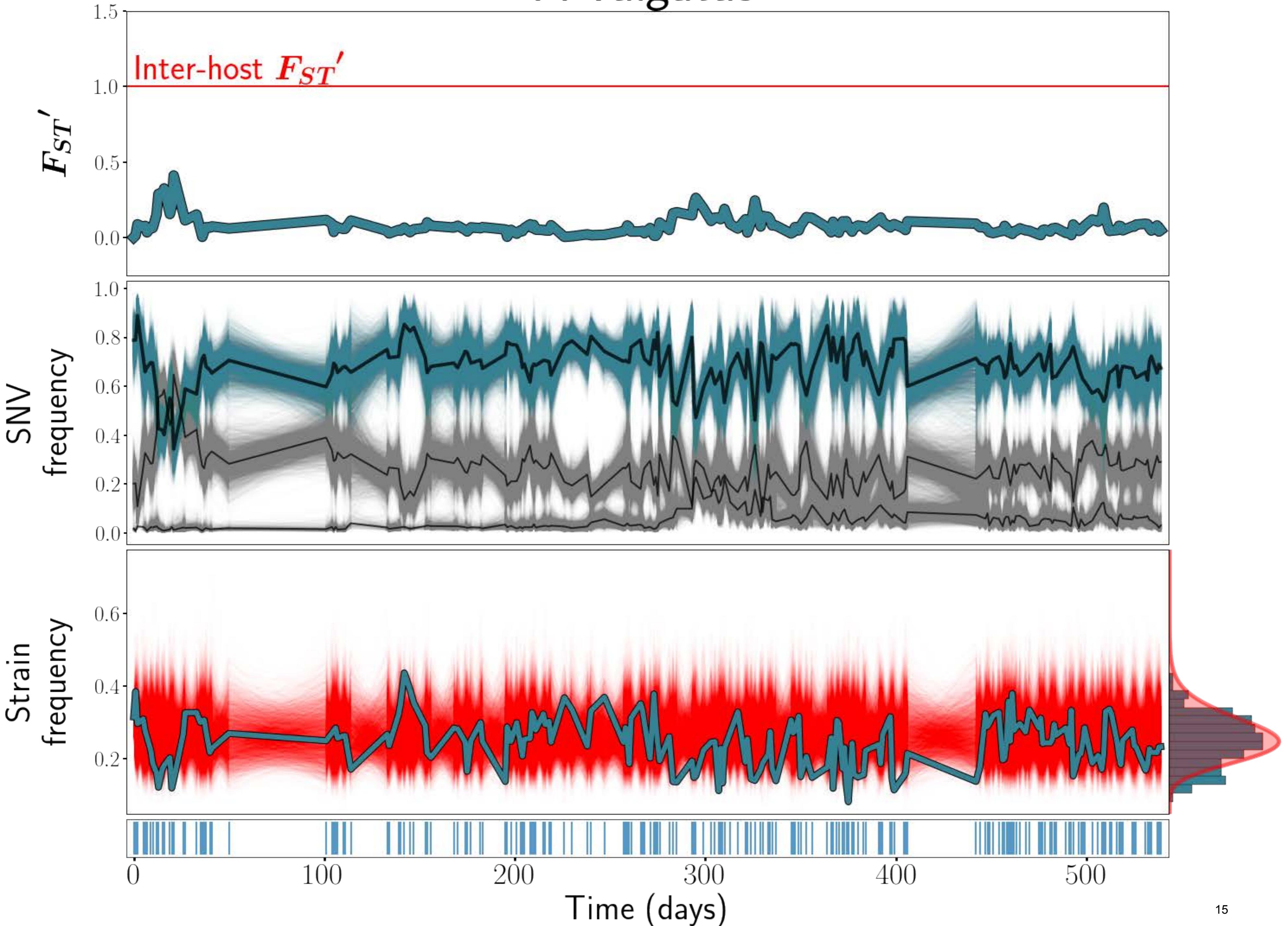
P. sp



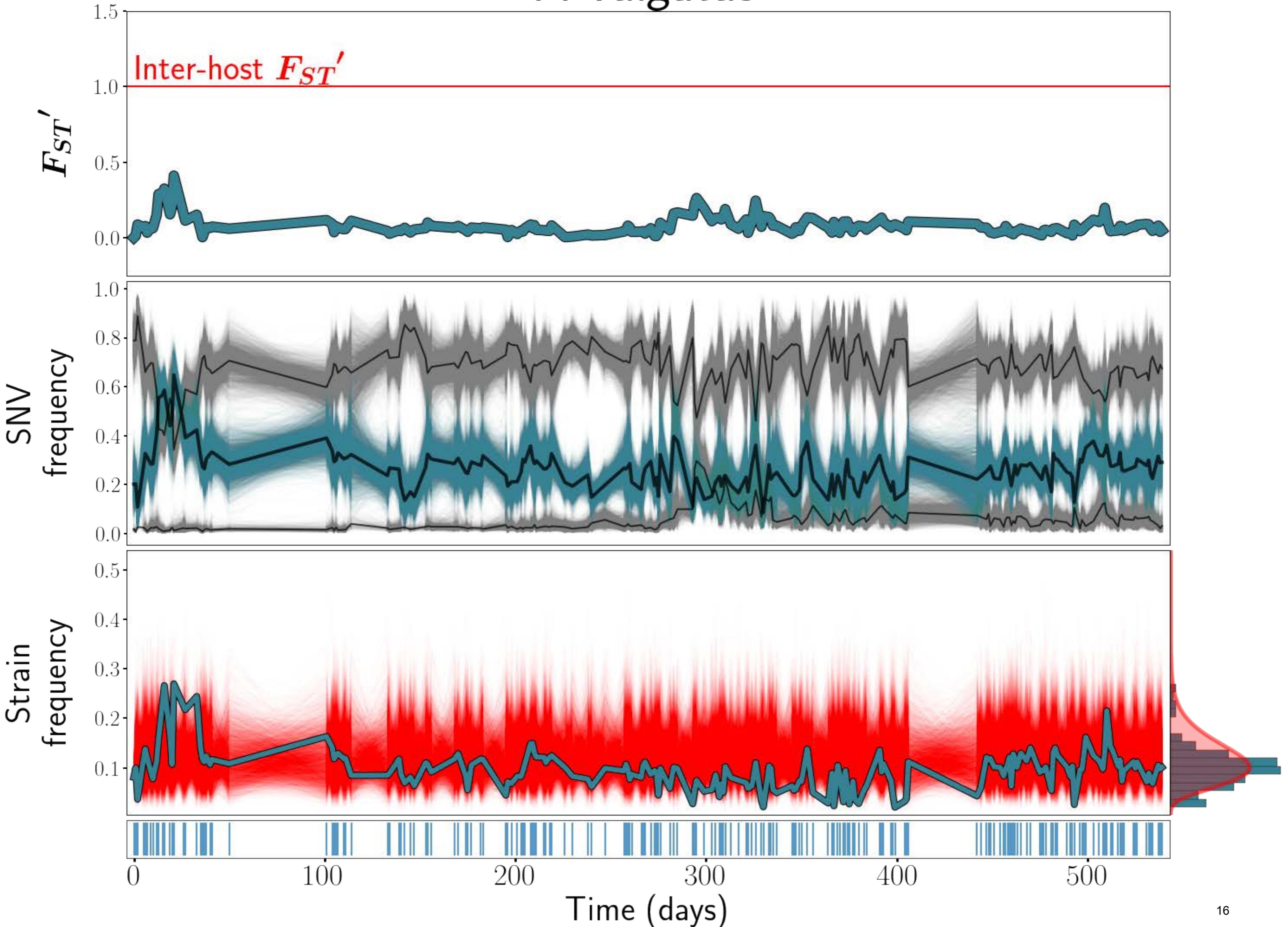
P. vulgatus



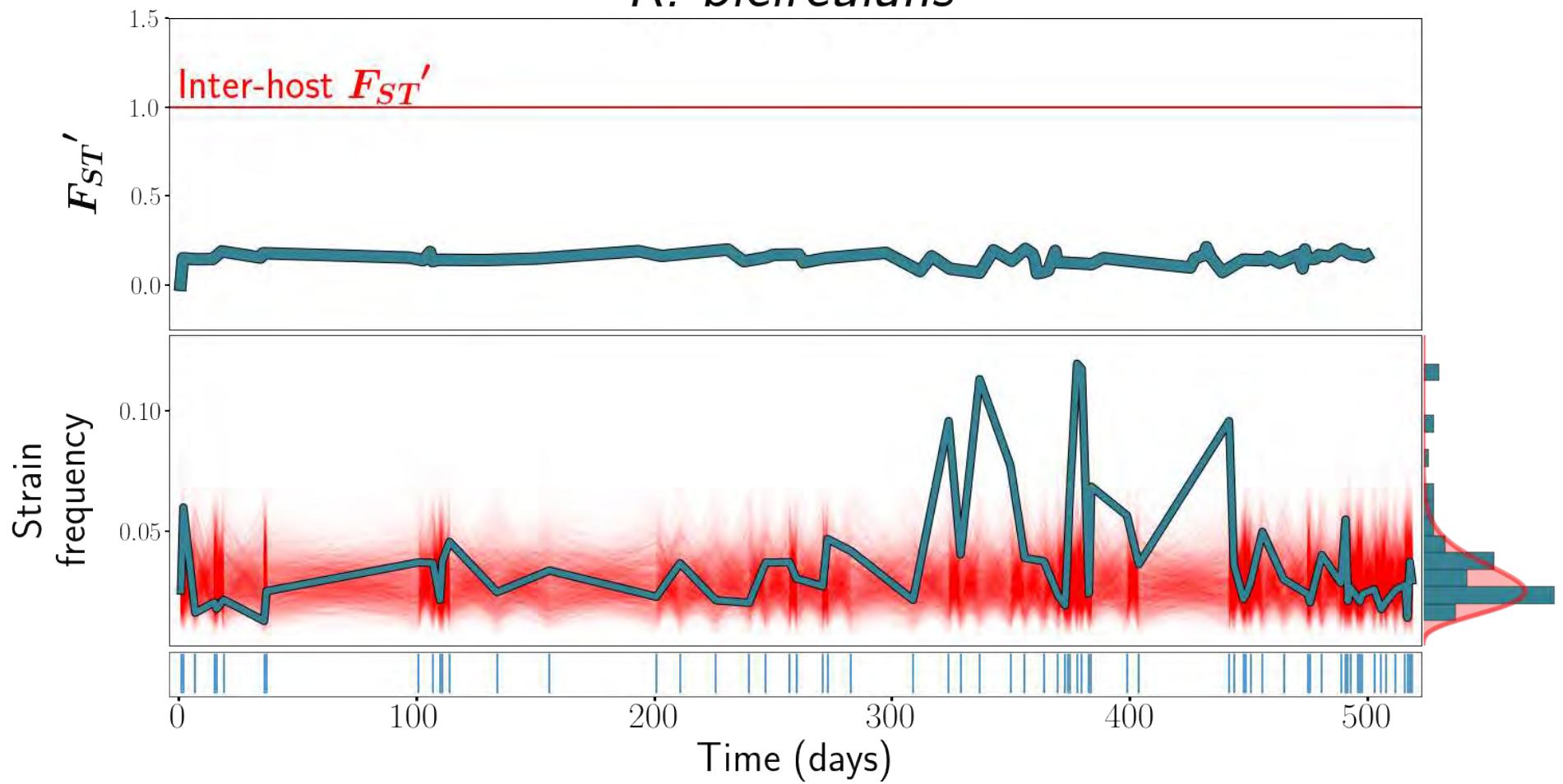
P. vulgatus



P. vulgatus



R. bicirculans



R. bromii

