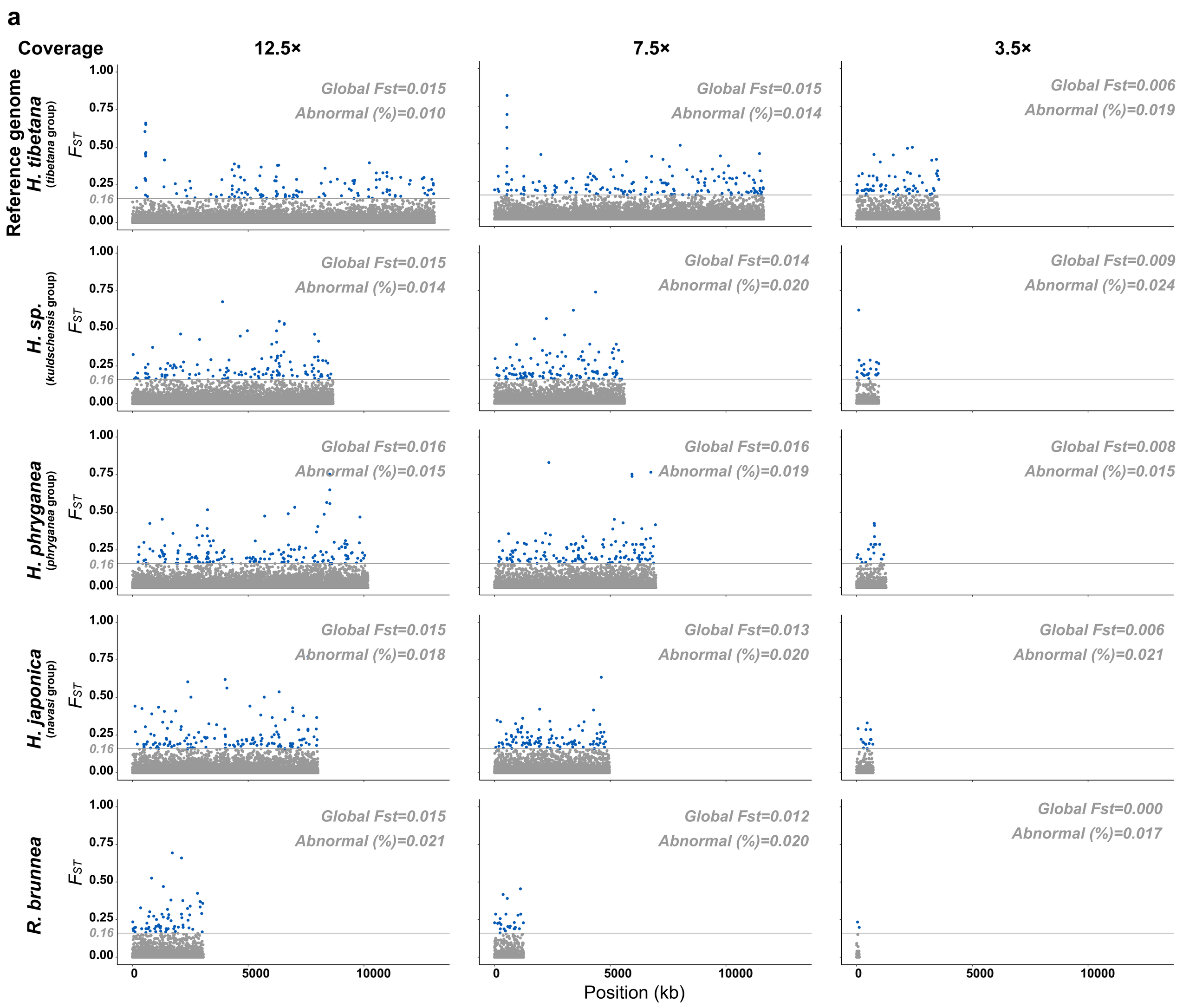
a
Coverage
12.5×
7.5×
3.5×
Reference genome
H. tibetana (tibetana group)
H. sp. (kuldschensis group)
H. phryganea (phryganea group)
H. japonica (navasi group)
R. brunnea
FST
1.00
0.75
0.50
0.25
0.16
0.00
Global Fst=0.015
Abnormal (%)=0.010
Global Fst=0.015
Abnormal (%)=0.014
Global Fst=0.006
Abnormal (%)=0.019
Global Fst=0.015
Abnormal (%)=0.014
Global Fst=0.014
Abnormal (%)=0.020
Global Fst=0.009
Abnormal (%)=0.024
Global Fst=0.016
Abnormal (%)=0.015
Global Fst=0.016
Abnormal (%)=0.019
Global Fst=0.008
Abnormal (%)=0.015
Global Fst=0.015
Abnormal (%)=0.018
Global Fst=0.013
Abnormal (%)=0.020
Global Fst=0.006
Abnormal (%)=0.021
Global Fst=0.015
Abnormal (%)=0.021
Global Fst=0.012
Abnormal (%)=0.020
Global Fst=0.000
Abnormal (%)=0.017
0
5000
10000
Position (kb)