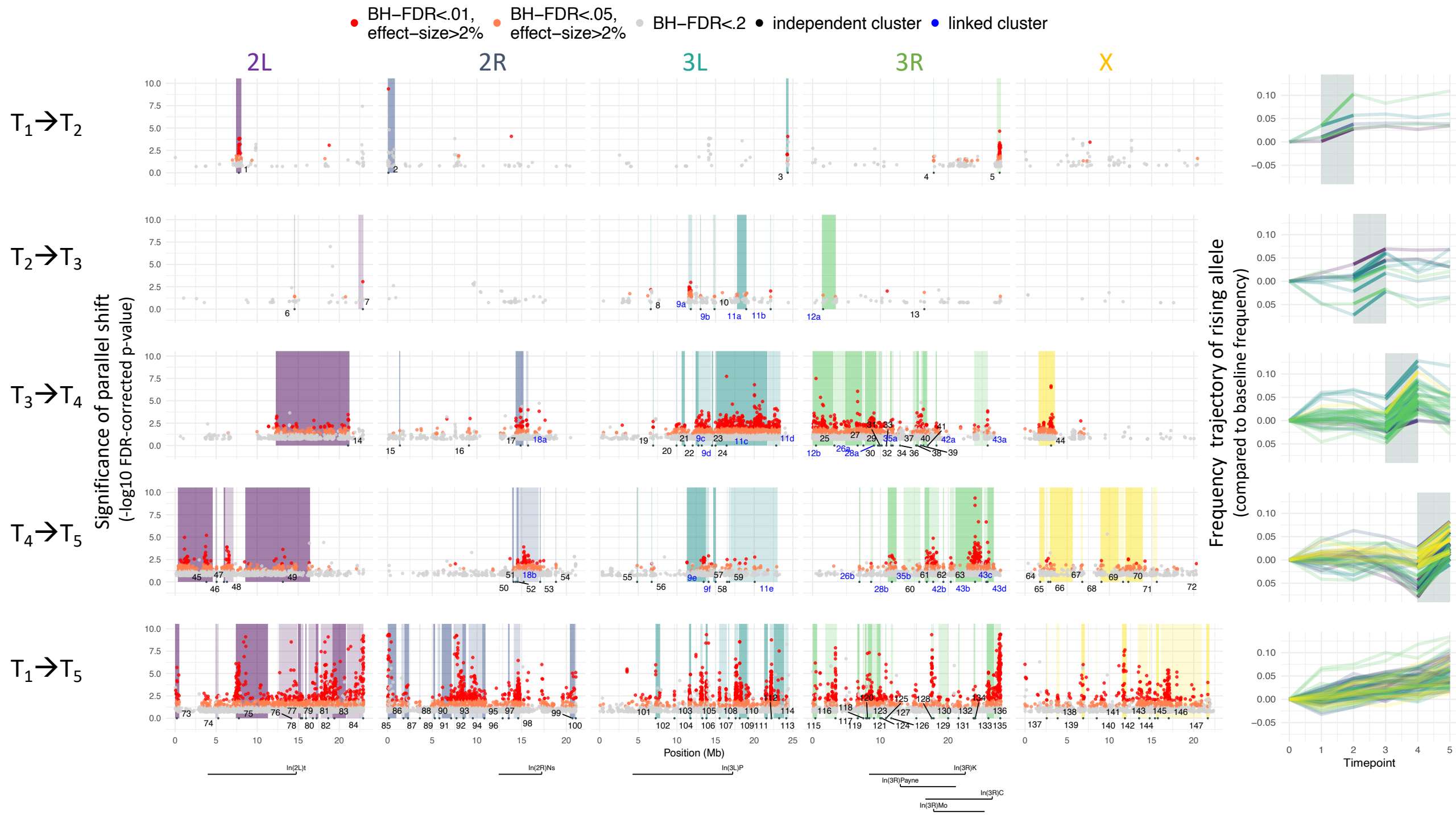
BH−FDR<.01, effect−size>2%
BH−FDR<.05, effect−size>2%
BH−FDR<.2
independent cluster
linked cluster
2L
2R
3L
3R
X
T1→T2
T2→T3
T3→T4
T4→T5
T1→T5
Significance of parallel shift
(-log10 FDR-corrected p-value)
Frequency trajectory of rising allele
(compared to baseline frequency)
Position (Mb)
Timepoint
In(2L)t
In(2R)Ns
In(3L)P
In(3R)K
In(3R)Payne
In(3R)C
In(3R)Mo