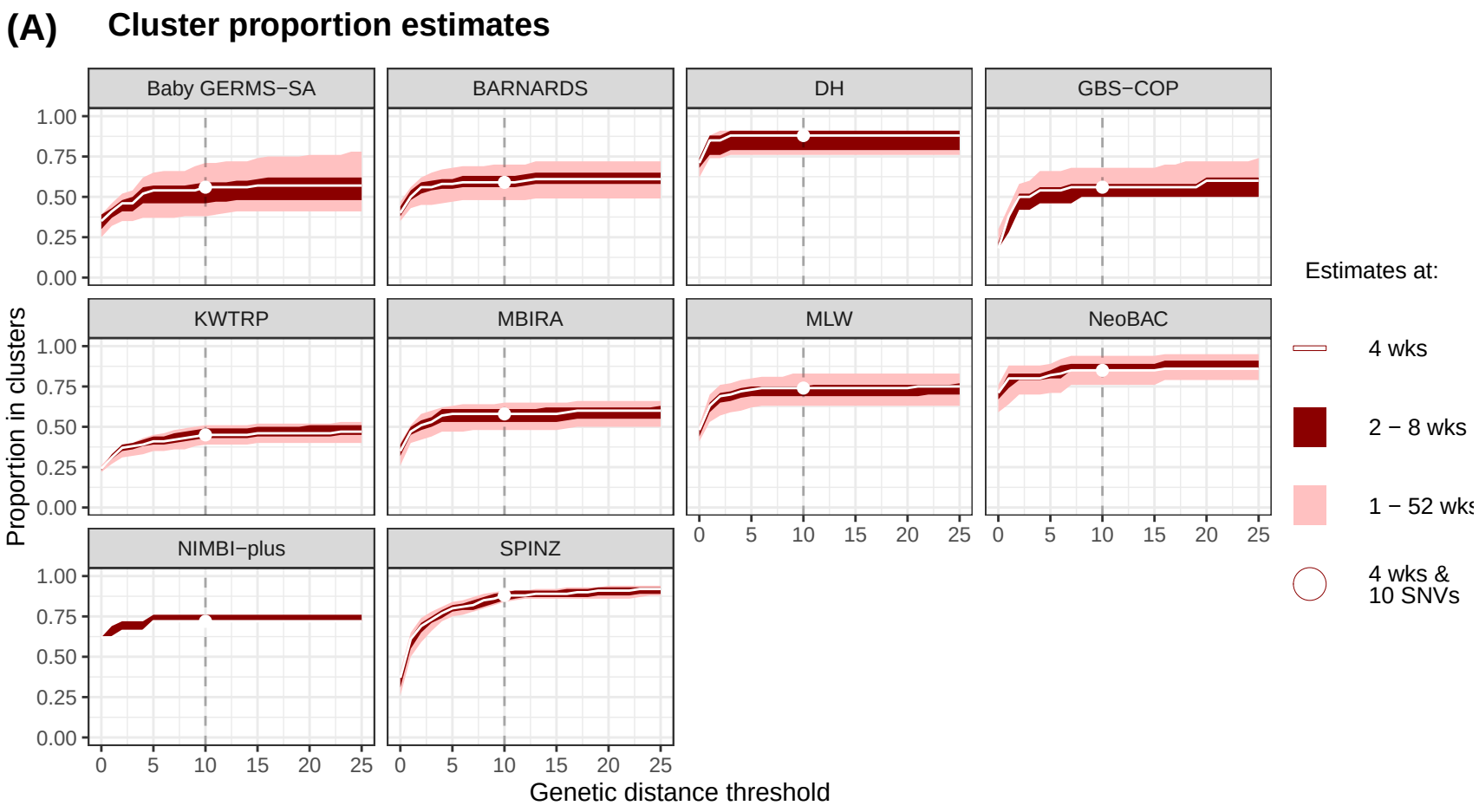
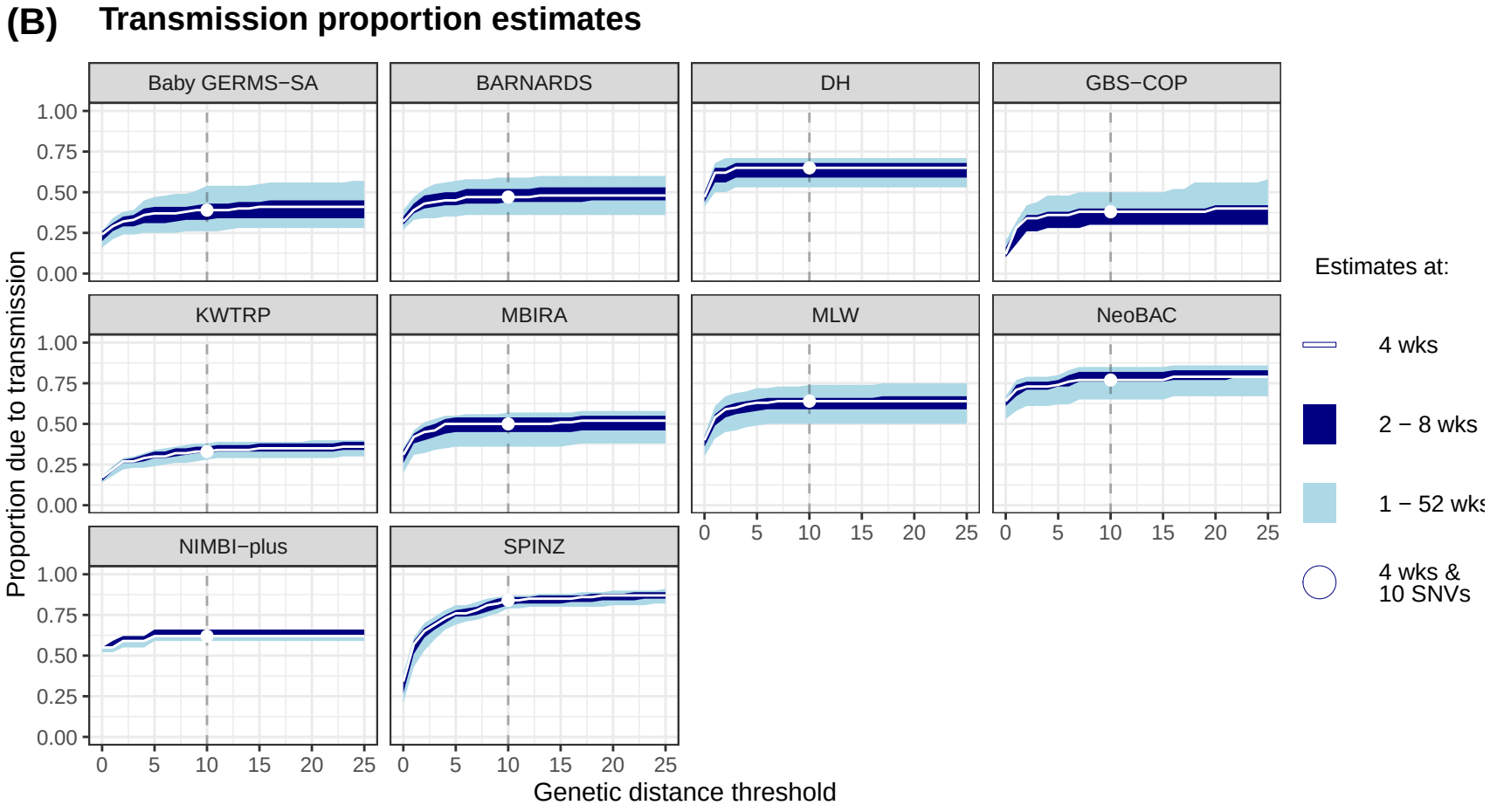

(A)
Cluster proportion estimates
Baby GERMS−SA
BARNARDS
DH
GBS−COP
KWTRP
MBIRA
MLW
NeoBAC
NIMBI−plus
SPINZ
Proportion in clusters
0.00
0.25
0.50
0.75
1.00
0
5
10
15
20
25
Genetic distance threshold
Estimates at:
4 wks
2 − 8 wks
1 − 52 wks
4 wks &
10 SNVs


(B)
Transmission proportion estimates
Baby GERMS−SA
BARNARDS
DH
GBS−COP
KWTRP
MBIRA
MLW
NeoBAC
NIMBI−plus
SPINZ
Proportion due to transmission
0.00
0.25
0.50
0.75
1.00
0
5
10
15
20
25
Genetic distance threshold
Estimates at:
4 wks
2 − 8 wks
1 − 52 wks
4 wks &
10 SNVs