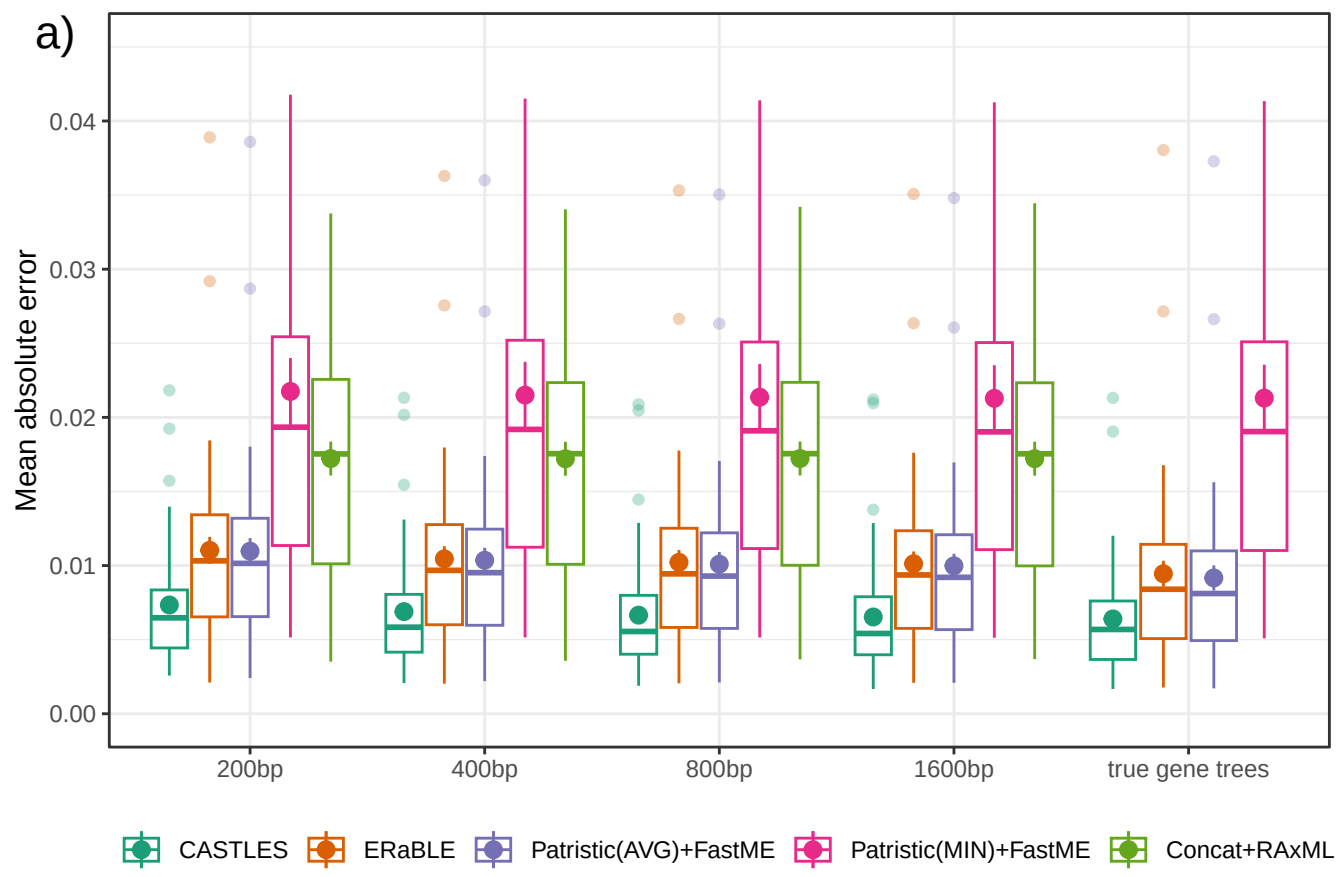
a)
Mean absolute error
0.04
0.03
0.02
0.01
0.00
200bp
400bp
800bp
1600bp
true gene trees
CASTLES
ERaBLE
Patristic(AVG)+FastME
Patristic(MIN)+FastME
Concat+RAxML