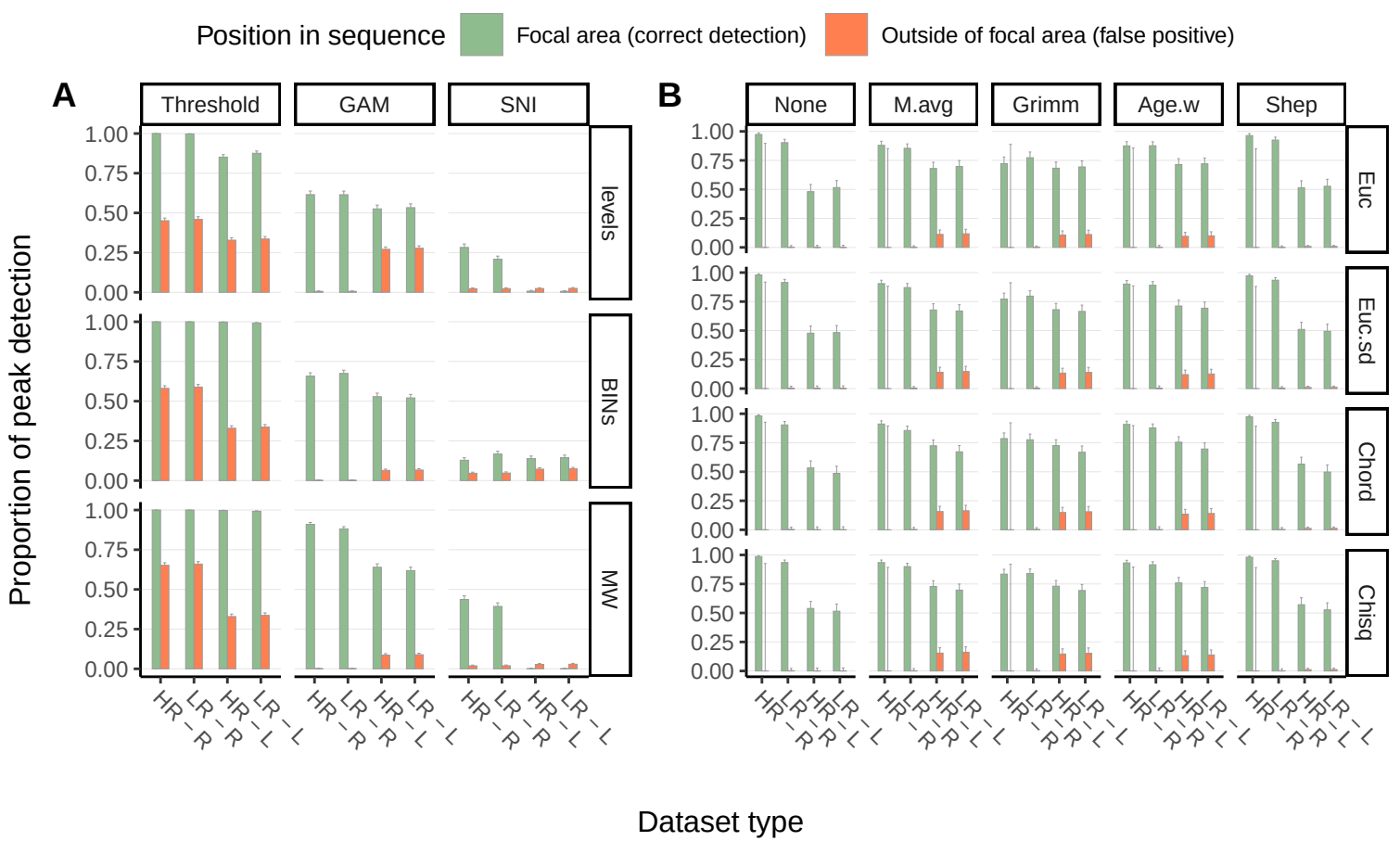
Position in sequence
Focal area (correct detection)
Outside of focal area (false positive)
A
B
Threshold
GAM
SNI
None
M.avg
Grimm
Age.w
Shep
levels
BINs
MW
Euc
Euc.sd
Chord
Chisq
Proportion of peak detection
0.00
0.25
0.50
0.75
1.00
HR_R
LR_R
HR_R
LR_L
Dataset type