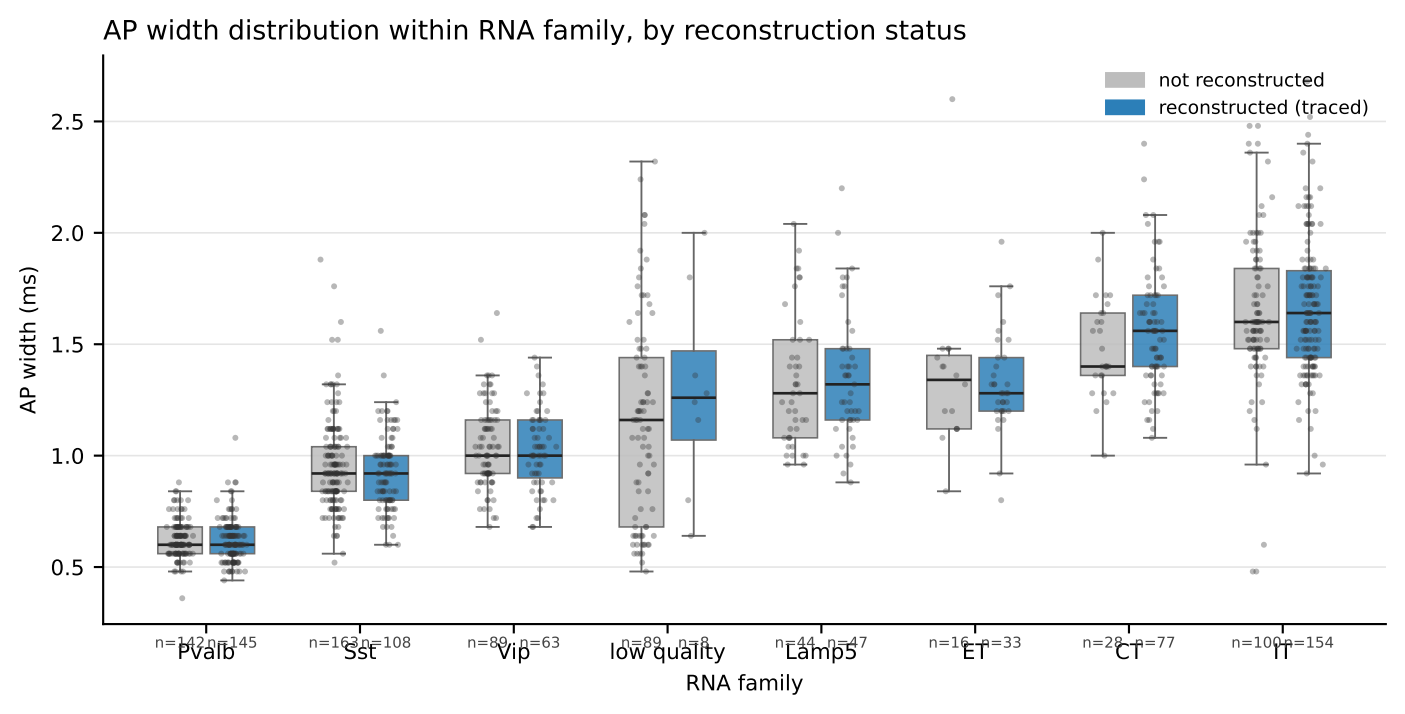

AP width distribution within RNA family, by reconstruction status
not reconstructed
reconstructed (traced)
AP width (ms)
2.5
2.0
1.5
1.0
0.5
n=142
n=145
n=163
n=108
n=89
n=63
n=89
n=8
n=44
n=47
n=16
n=33
n=28
n=77
n=100
n=154
Pvalb
Sst
Vip
low quality
Lamp5
ET
CT
IT
RNA family