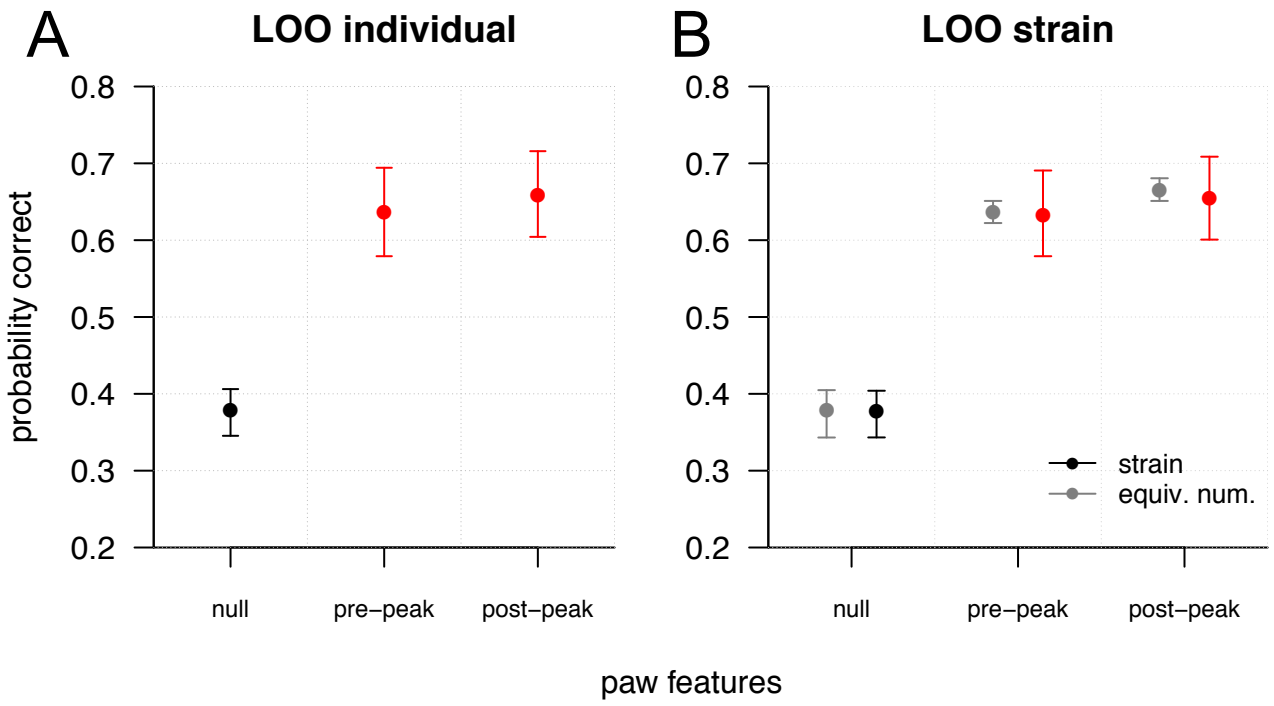
A
LOO individual
B
LOO strain
probability correct
0.8
0.7
0.6
0.5
0.4
0.3
0.2
null
pre−peak
post−peak
strain
equiv. num.
paw features