

Compartmental breakdown

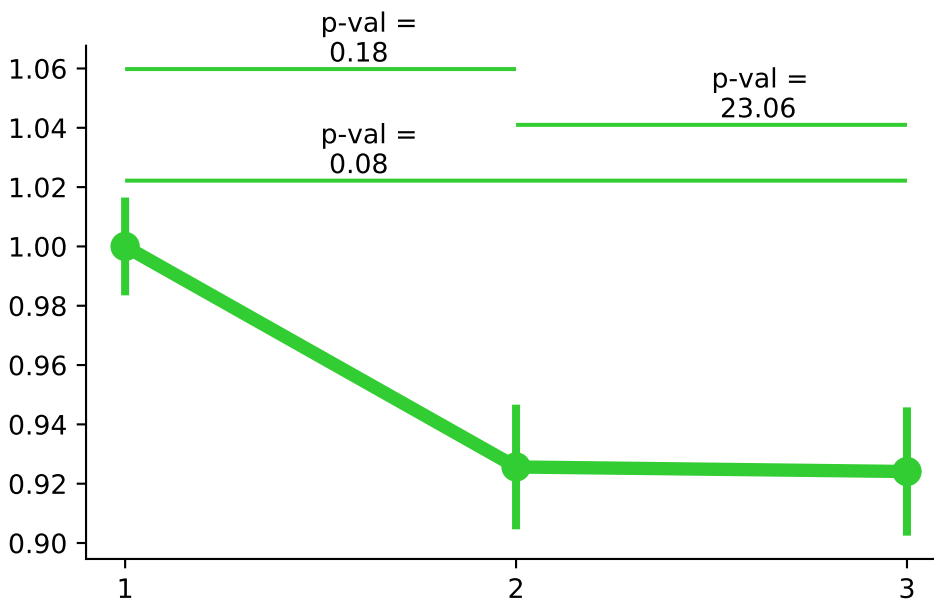
Frames ['brk']

Compartment amalgam
Frames ['brk']

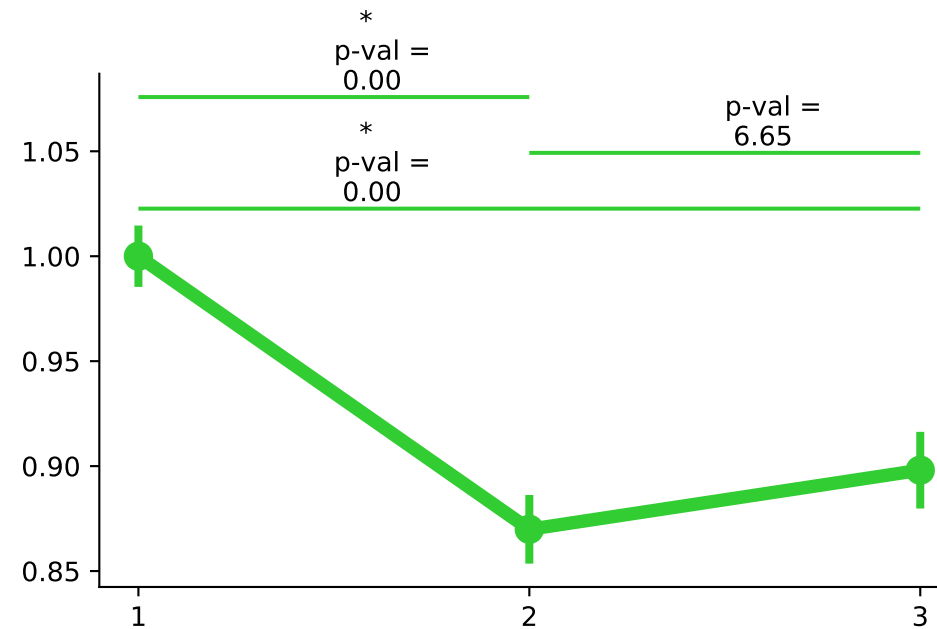
p-values are 2-tailed

Top row: Expected sequences
Bottom row: Unexpected sequences
p-values are 2-tailed

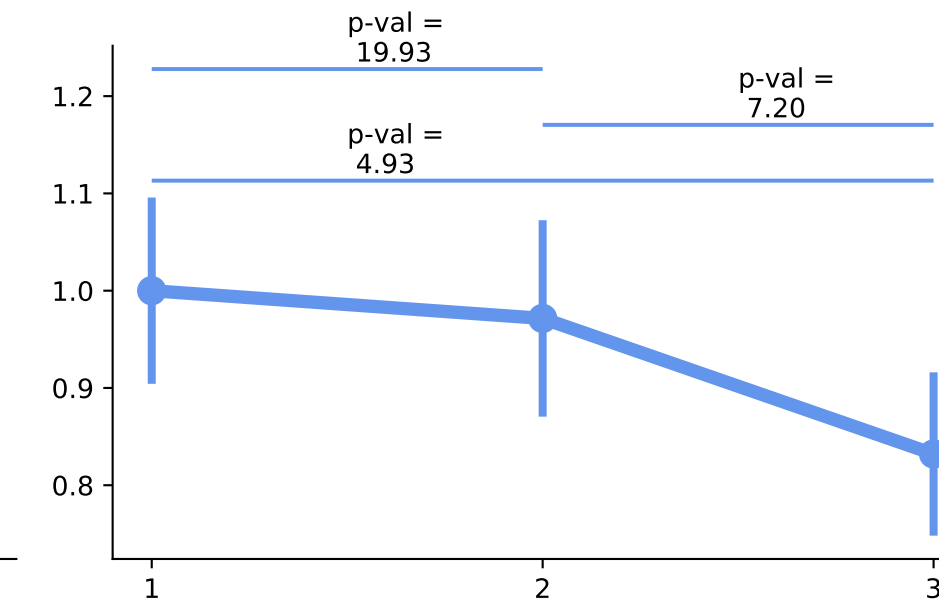
Expected sequences
L2/3 dend
Mice: [6 9]



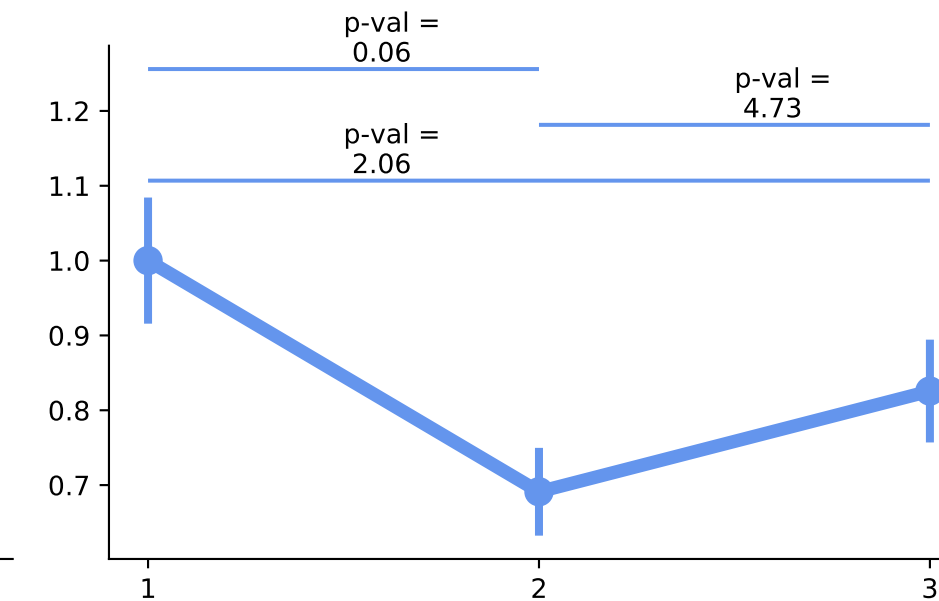
Expected sequences
L5 dend
Mice: [8 10 11]



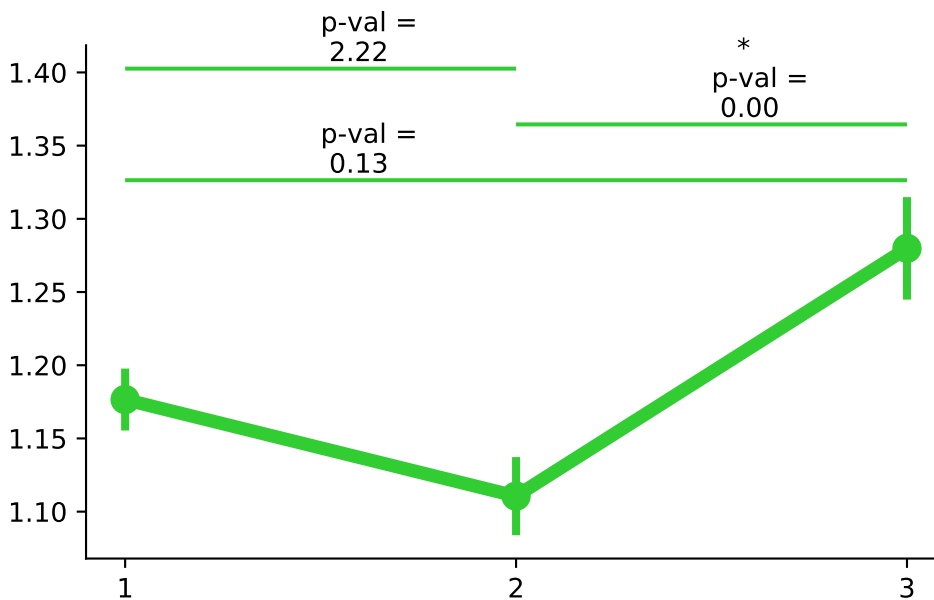
Expected sequences
L2/3 soma
Mice: [1 3 13]



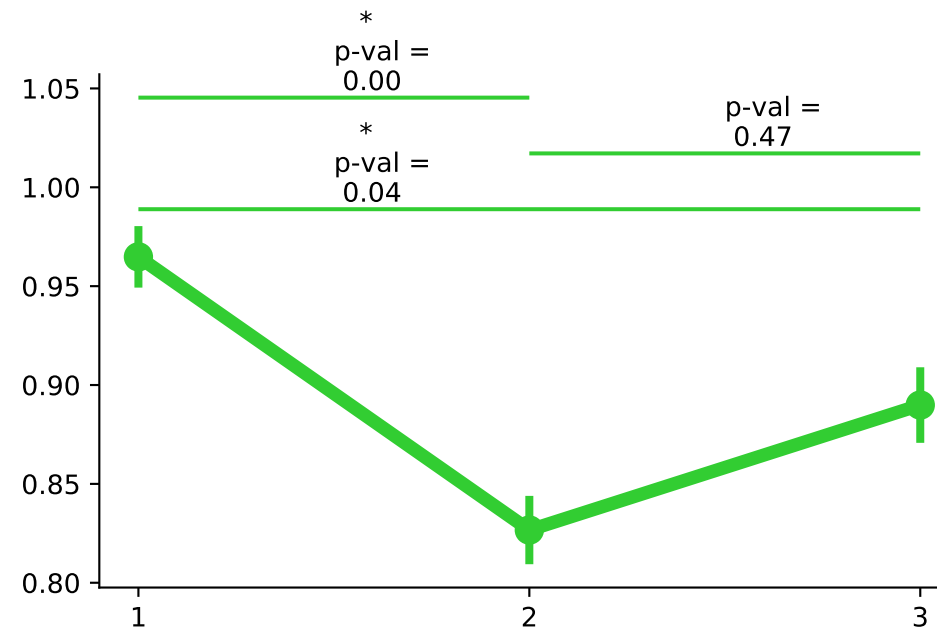
Expected sequences
L5 soma
Mice: [4 7 12]



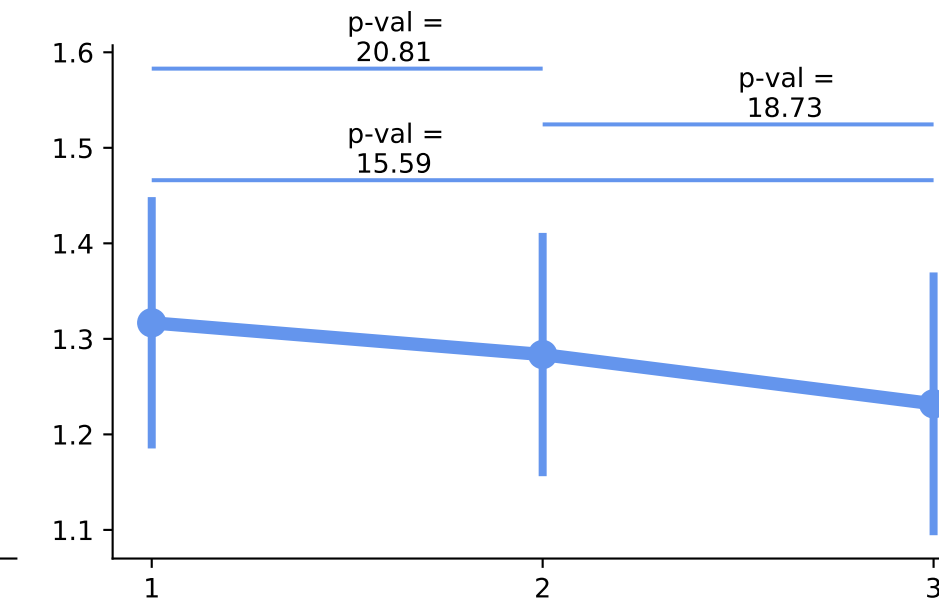
Unexpected sequences
L2/3 dend
Mice: [6 9]



Unexpected sequences
L5 dend
Mice: [8 10 11]



Unexpected sequences
L2/3 soma
Mice: [1 3 13]



Unexpected sequences
L5 soma
Mice: [4 7 12]

