

*Compartmental breakdown*

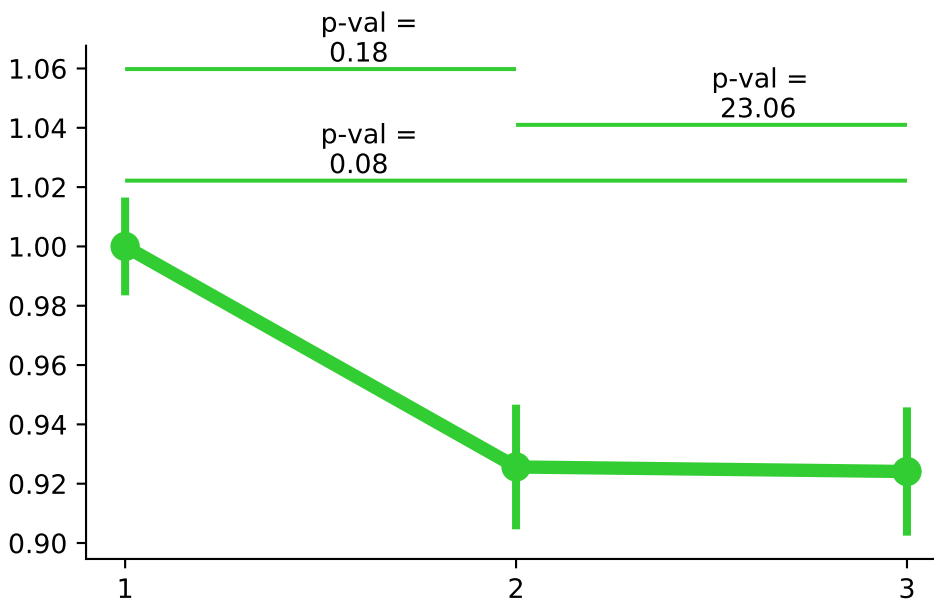
*Frames ['brk']*

Compartment amalgam  
Frames ['brk']

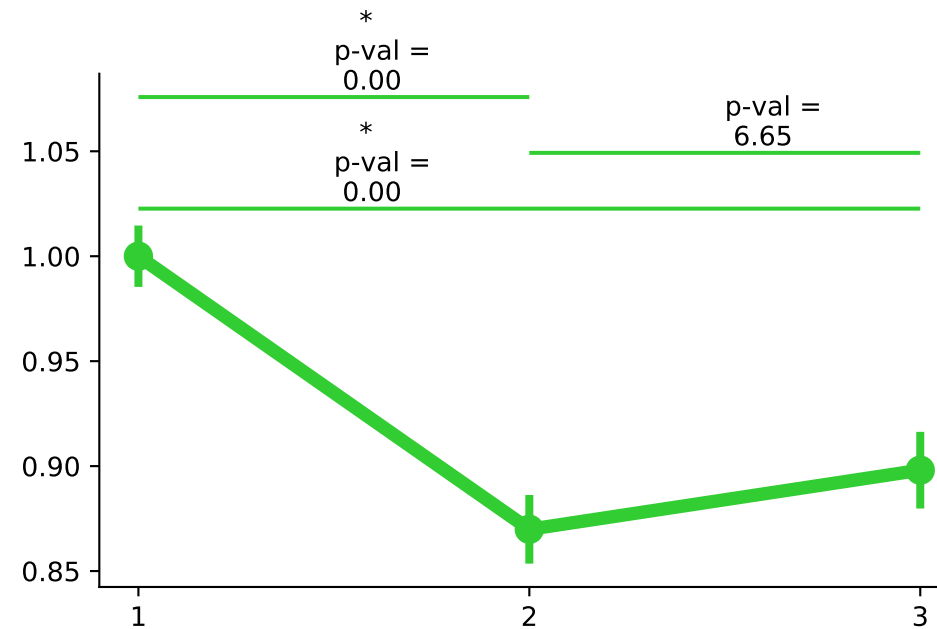
p-values are 2-tailed

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

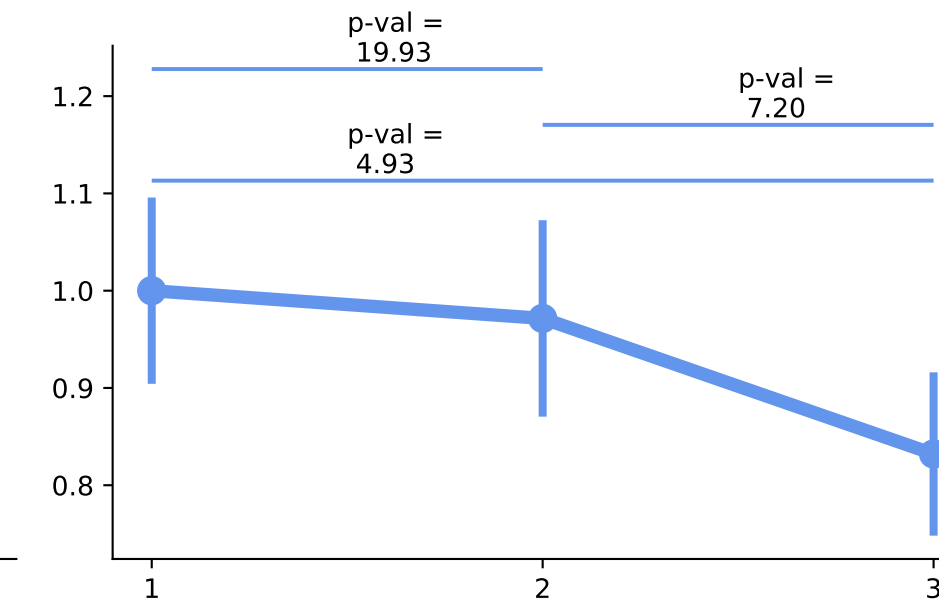
Expected sequences  
L2/3 dend  
Mice: [ 6 9]



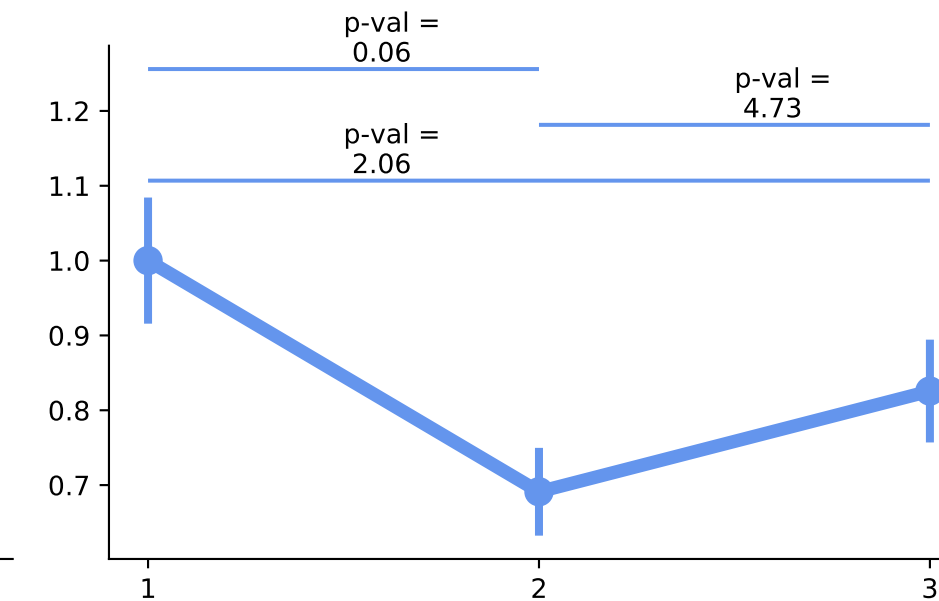
Expected sequences  
L5 dend  
Mice: [ 8 10 11]



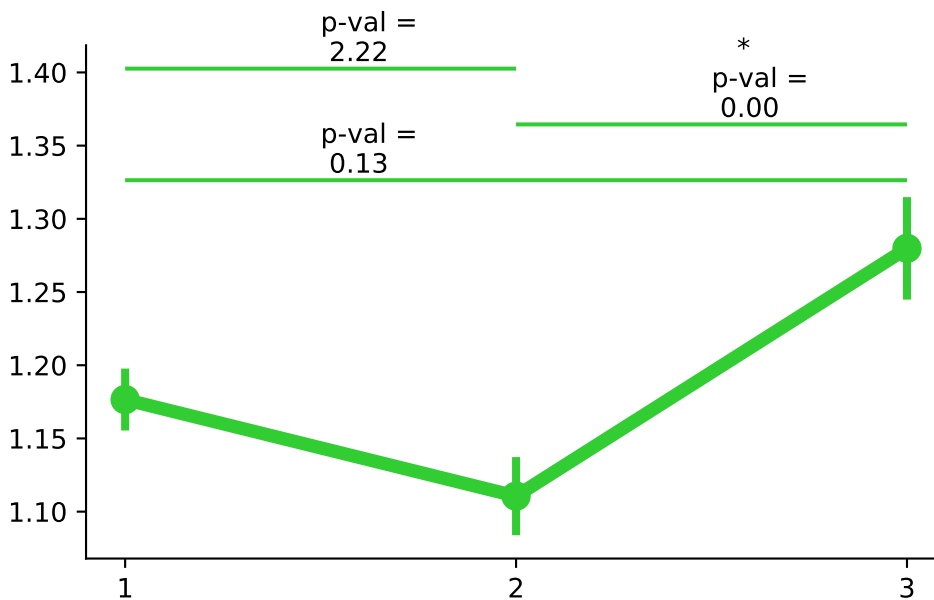
Expected sequences  
L2/3 soma  
Mice: [ 1 3 13]



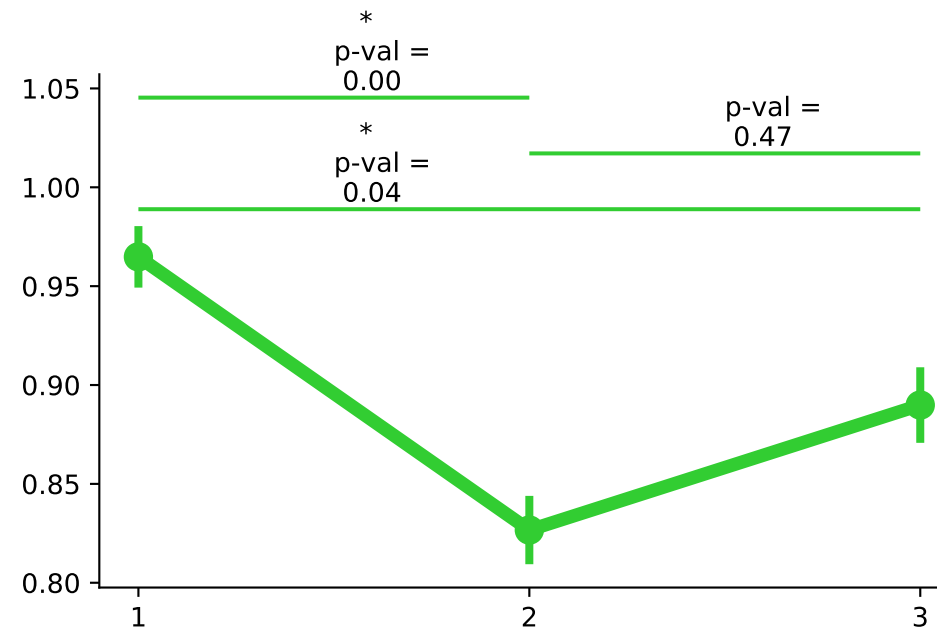
Expected sequences  
L5 soma  
Mice: [ 4 7 12]



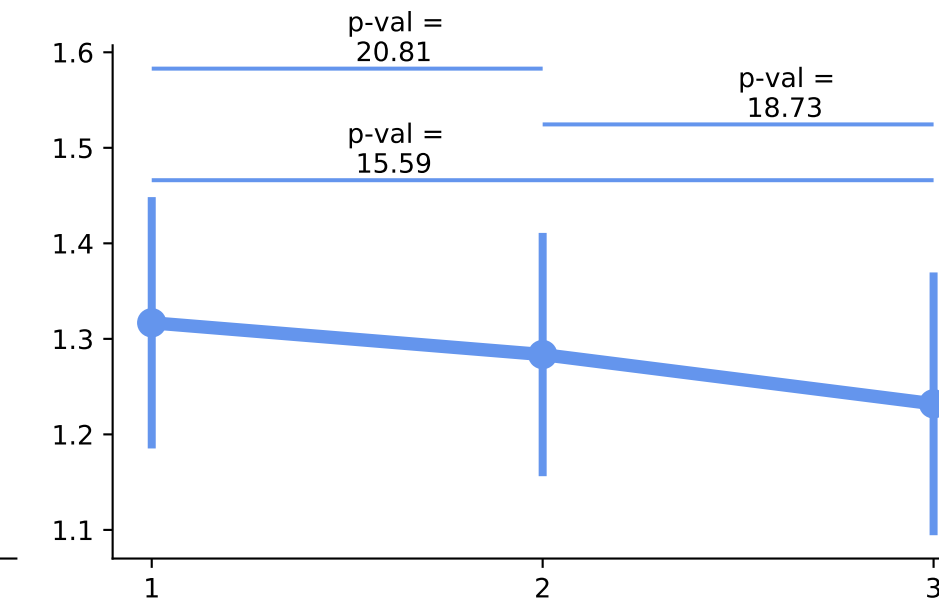
Unexpected sequences  
L2/3 dend  
Mice: [ 6 9]



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]

