

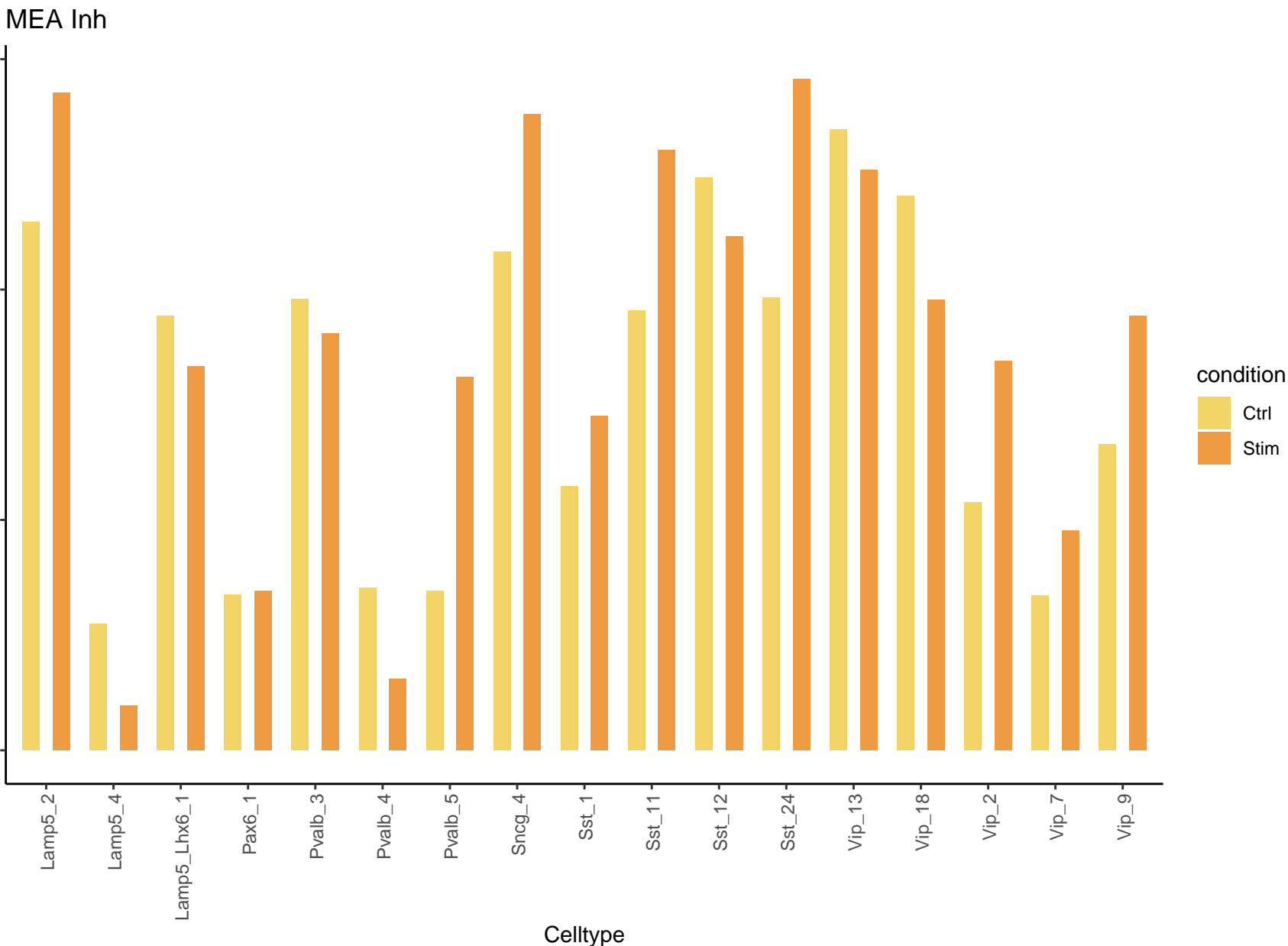
# MEA Inh

Number of significant gene-peak pairs

600  
400  
200  
0

Lamp5\_2 Lamp5\_4 Lamp5\_Lhx6\_1 Pax6\_1 Pvalb\_3 Pvalb\_4 Pvalb\_5 Sncg\_4 Sst\_1 Sst\_11 Sst\_12 Sst\_24 Vip\_13 Vip\_18 Vip\_2 Vip\_7 Vip\_9

condition  
Ctrl  
Stim



Celltype