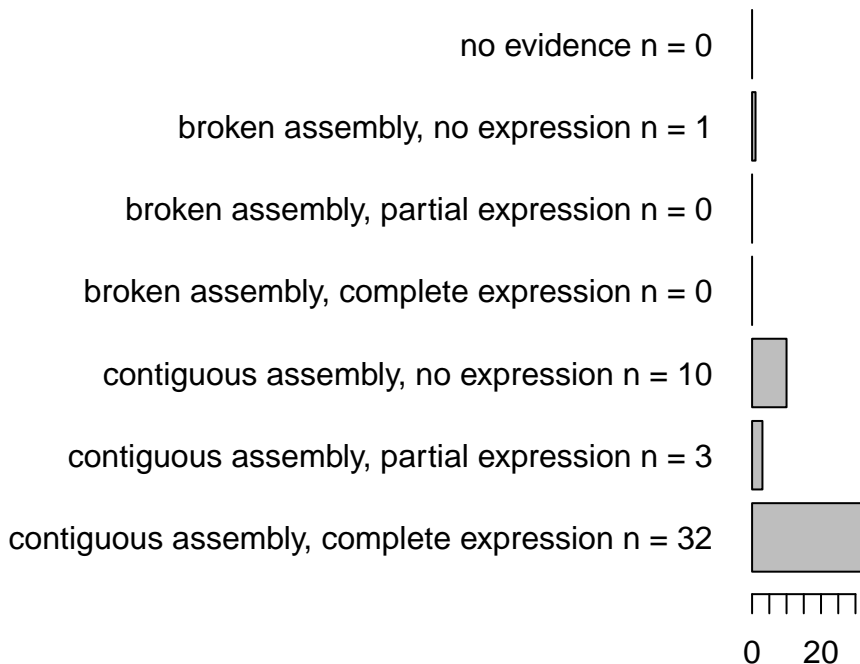


**histone fusions**  
**n=46**



# histone fusions

n=46

