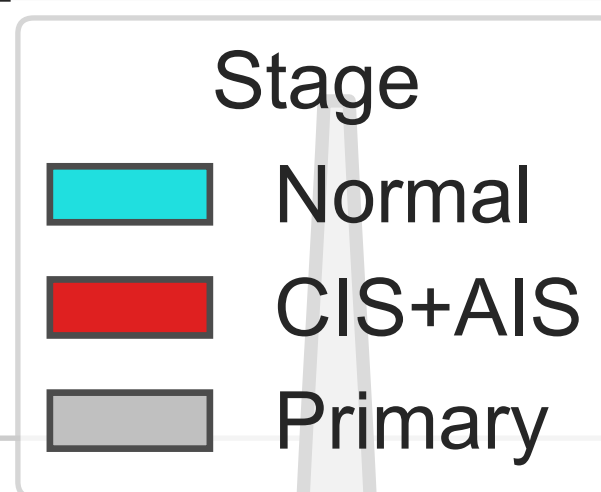


FOXR1: Kruskal-Wallis p=0.019

FOXR1 expression



1.0  
0.8  
0.6  
0.4  
0.2  
0.0  
-0.2

Lower Higher  
Mutation Shared Count (SYN) per TMB

