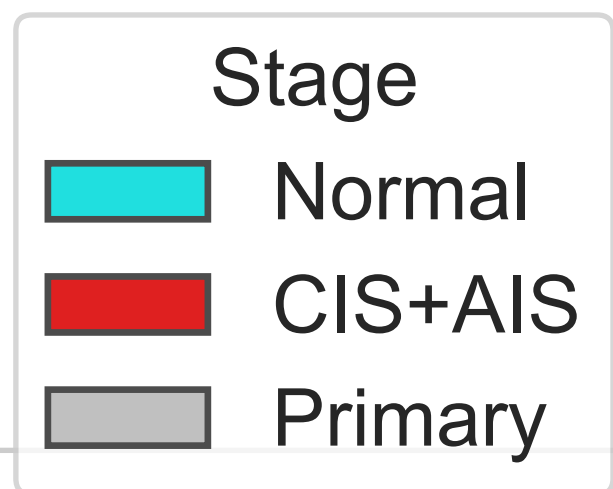


DGCR8: Kruskal-Wallis p=0.036

DGCR8 expression



M.W.W. $p \leq 0.05$

Lower Higher
Mutation Shared Count per TMB

