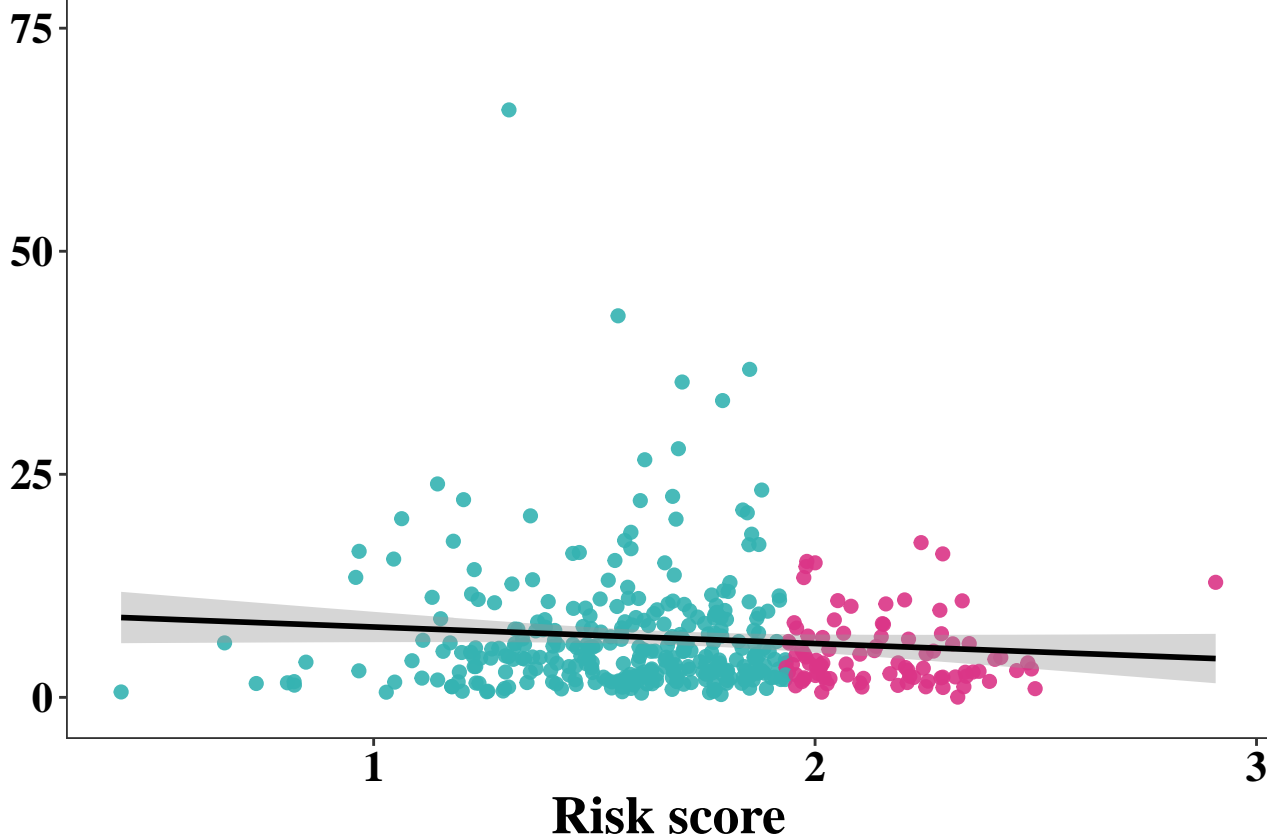


Tumor Burden Mutation

$R = -0.059, p = 0.24$



Risk    ● High risk    ● Low risk