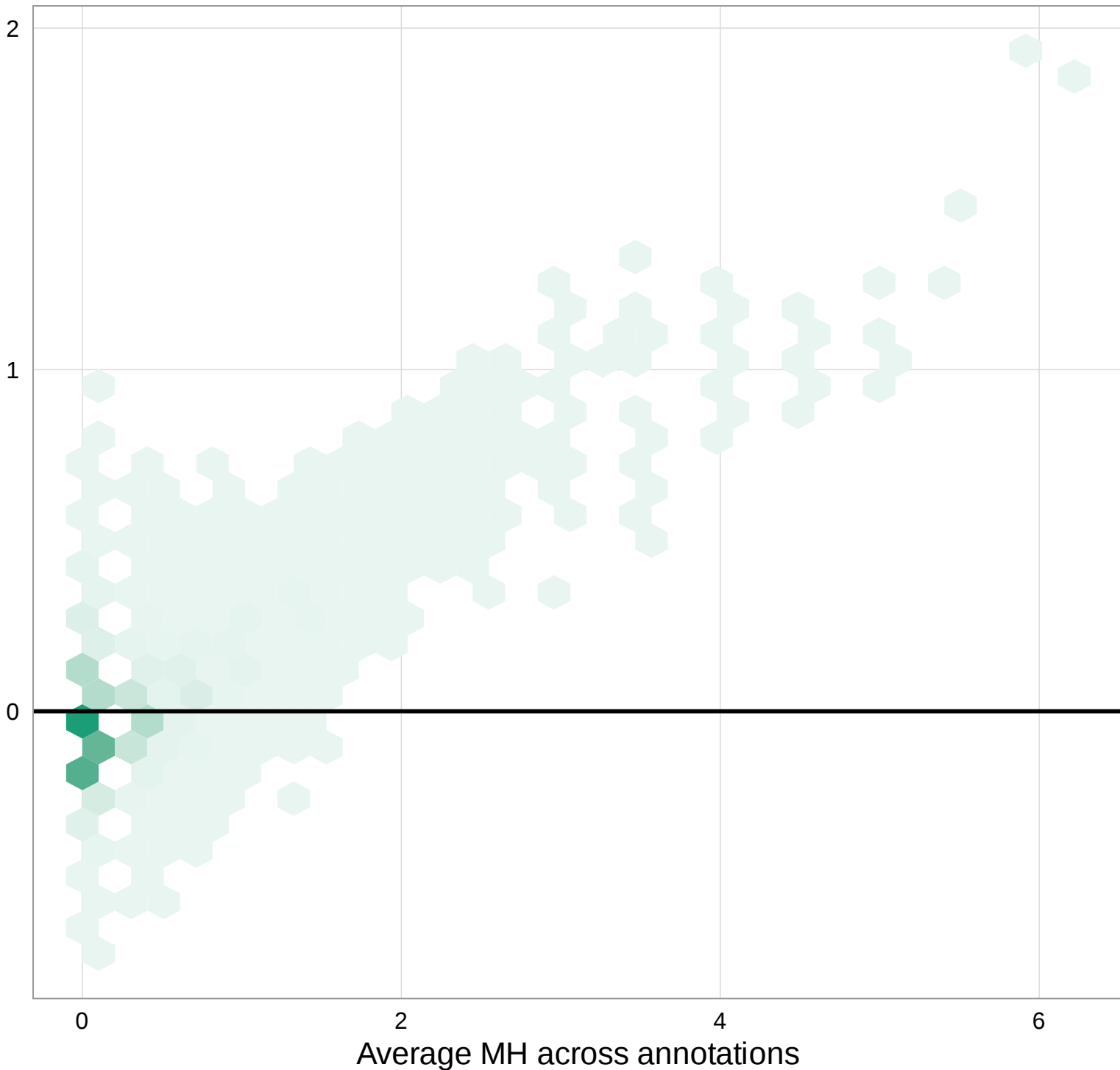


log10 difference in sequence generation probability
(MH vs no-MH model)



count

1e+05

2e+05

3e+05