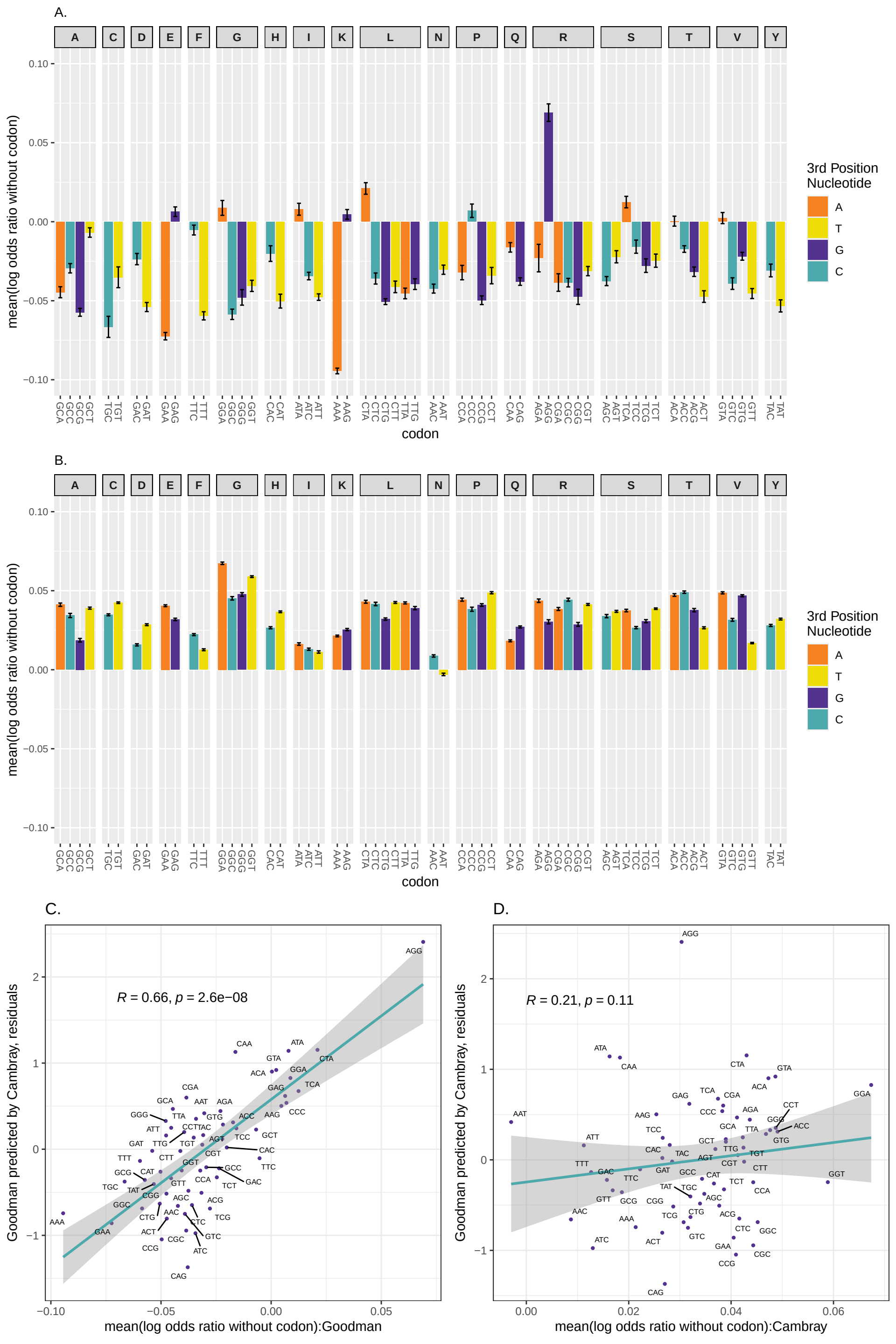
A.
B.
C.
D.
mean(log odds ratio without codon)
codon
3rd Position Nucleotide
A
T
G
C
R = 0.66, p = 2.6e−08
R = 0.21, p = 0.11
Goodman predicted by Cambray, residuals
mean(log odds ratio without codon):Goodman
mean(log odds ratio without codon):Cambray