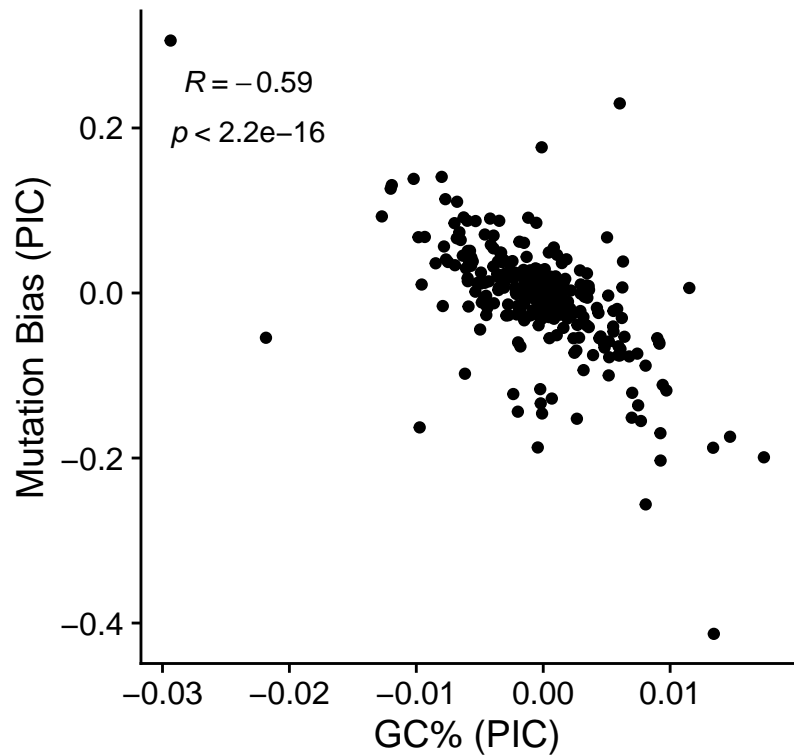


**A** Mutation bias vs. GC%  
across species (codon CAG)



**B** Shifts in mutation bias reflect changes  
to genome-wide GC%

