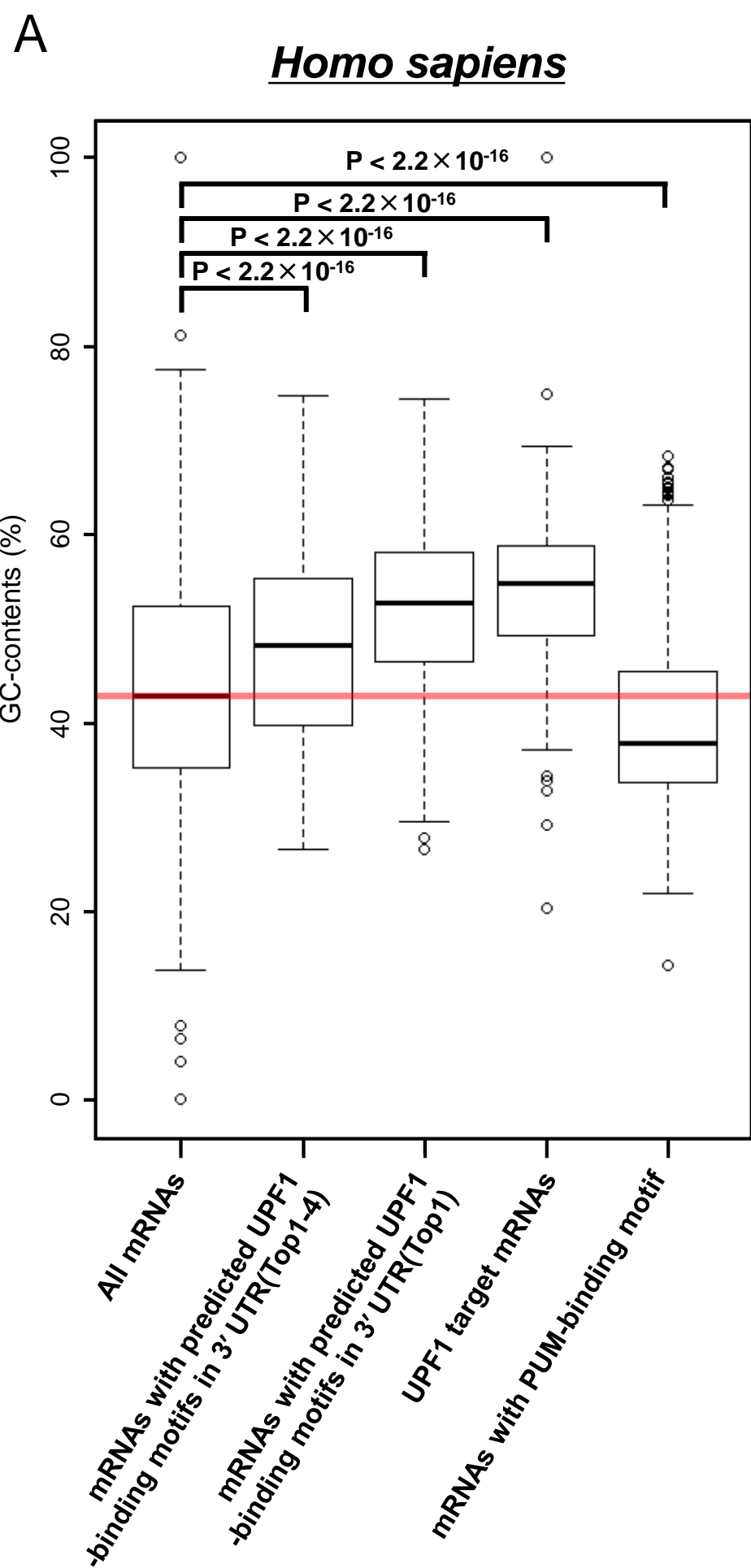


Supplemental Fig S6.



GC-contents (3' UTR): Median / Mean

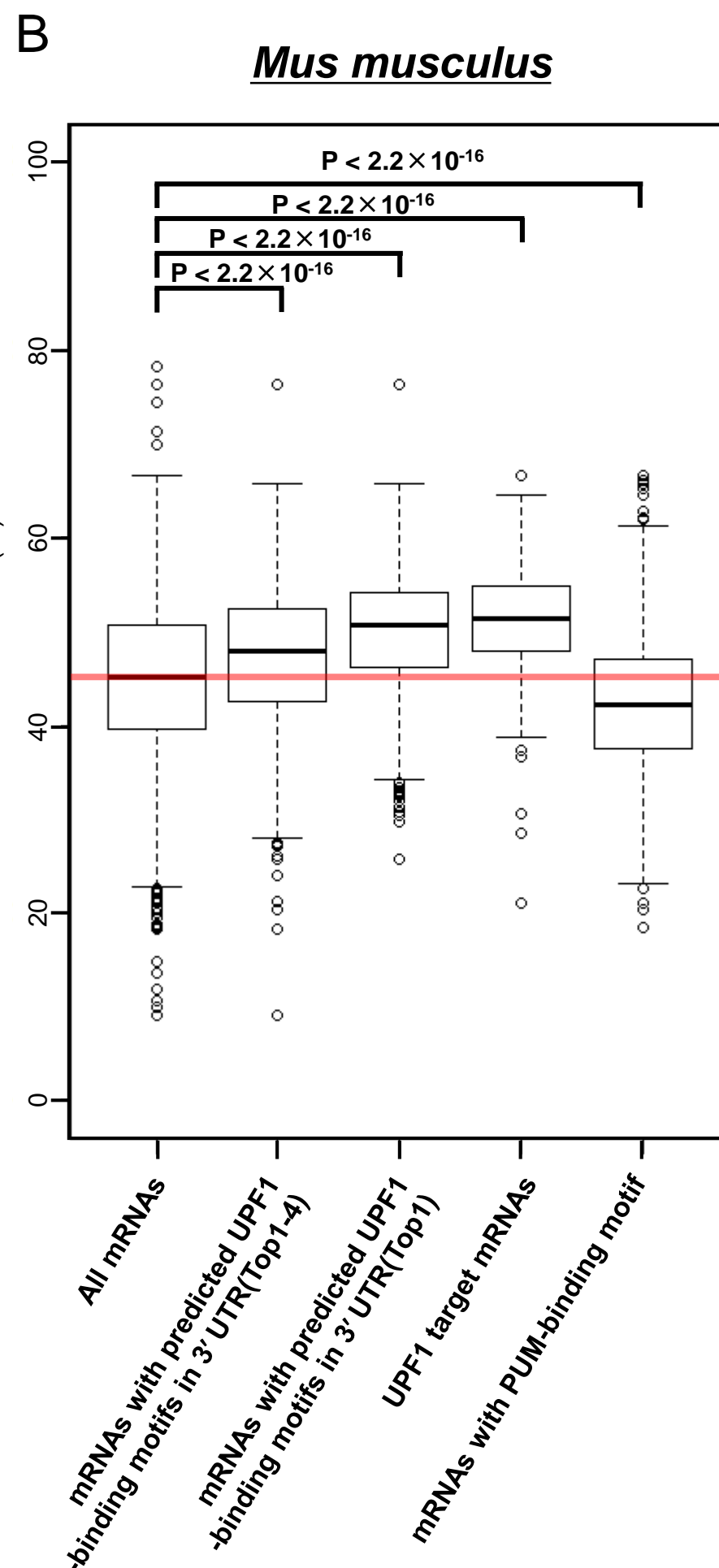
All mRNAs: 42.9 / 43.9

mRNAs with predicted UPF1-binding motifs
in 3' UTR (Top1-4): 48.2 / 47.8

mRNAs with predicted UPF1-binding motifs
in 3' UTR (Top1): 52.7 / 51.8

UPF1 target mRNAs: 54.8 / 53.9

mRNAs with PUM-binding motif in 3'UTR: 37.9 / 40.2



GC-contents (3' UTR): Median / Mean

All mRNAs: 45.2 / 45.0

mRNAs with predicted UPF1-binding motifs
in 3' UTR (Top1-4): 47.9 / 47.3

mRNAs with predicted UPF1-binding motifs
in 3' UTR (Top1): 50.6 / 49.9

UPF1 target mRNAs: 51.3 / 51.1

mRNAs with PUM-binding motif in 3'UTR: 42.1 / 42.6