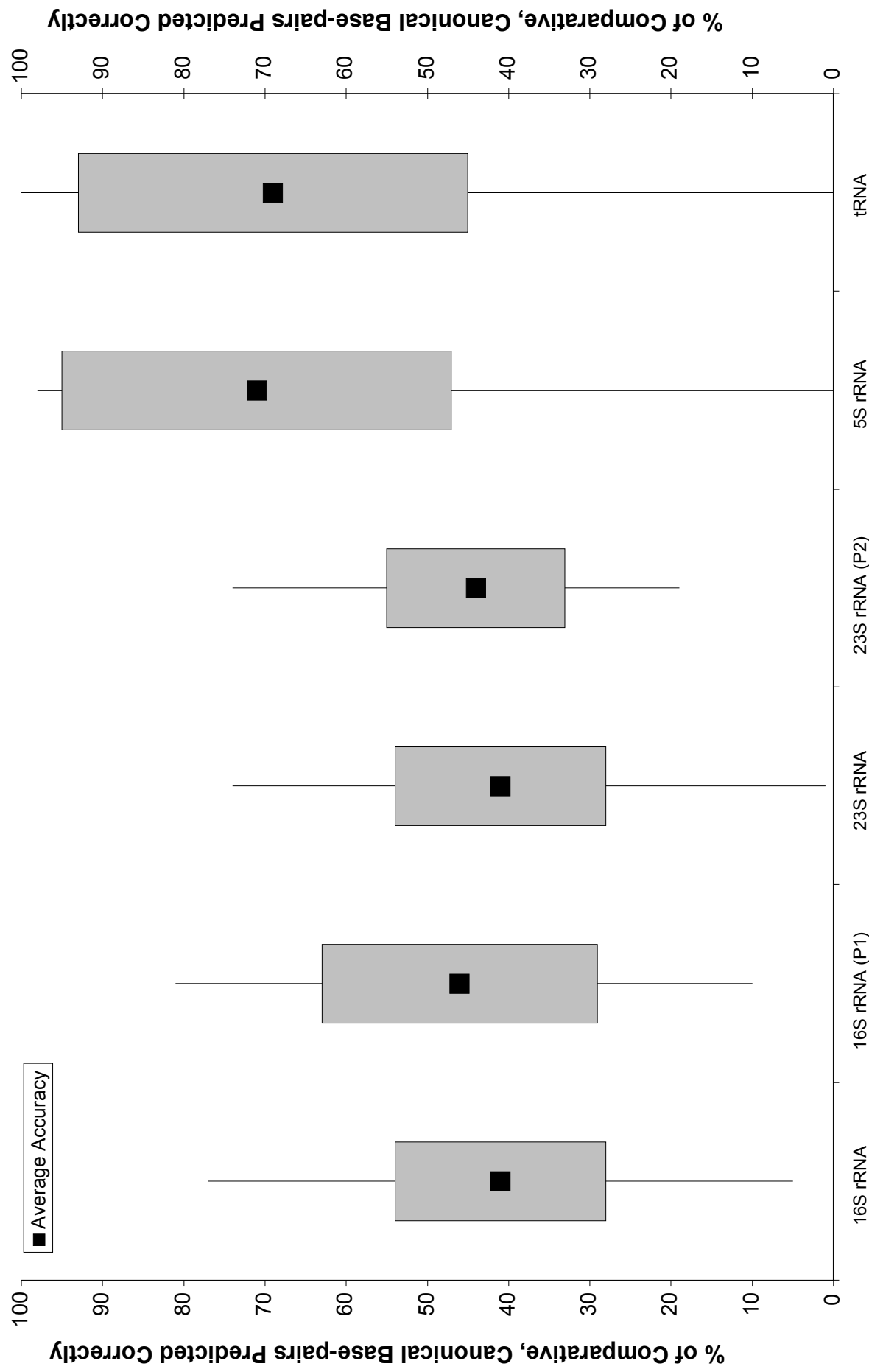


# Average, Minimum and Maximum Accuracy of MFold w/EFN2 for RNA Secondary Structure Prediction



Shaded boxes represent one standard deviation around the average folding accuracy. Vertical lines represent the extent of the range of folding accuracies observed in a particular category. Endpoints of the vertical lines represent the minimum and maximum observed folding accuracies. Folding parameters for Mfold 3.1 were window size (W)=1, max suboptimals (MAX)=750, percent suboptimality (P)=5%, and efn2 re-evaluation and re-ordering of the predicted folds.

(P1) Results from Konings DA & Gutell RR. RNA 1995 1:559-574

(P2) Results from Fields DS & Gutell RR. Folding and Design 1996 1:419-430