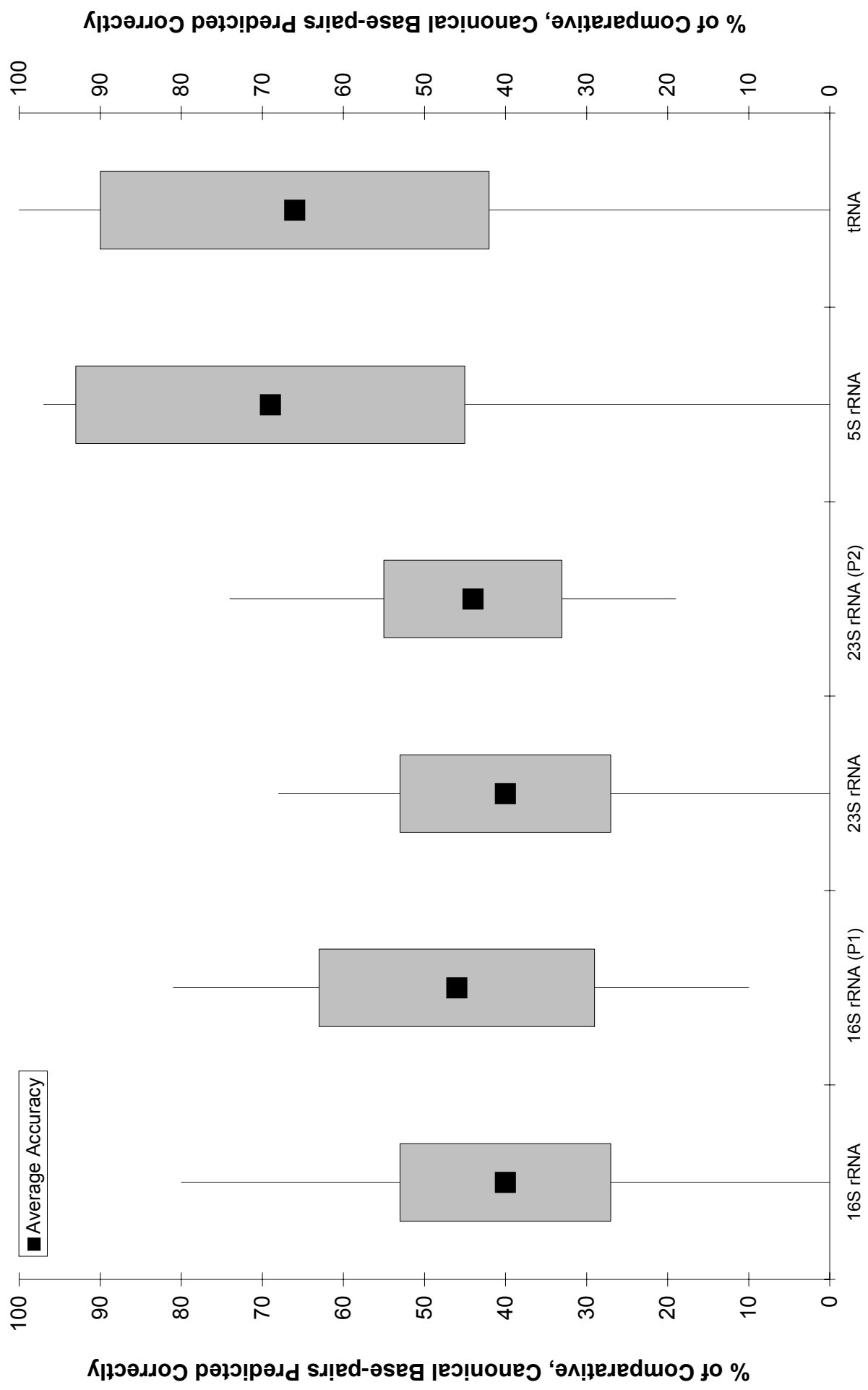


Average, Minimum and Maximum Accuracy of MFold for RNA Secondary Structure Prediction



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 with Mfold 3.1 were window size (W)=1, max suboptimals (MAX)=750, and percent suboptimality (P)=5%

(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