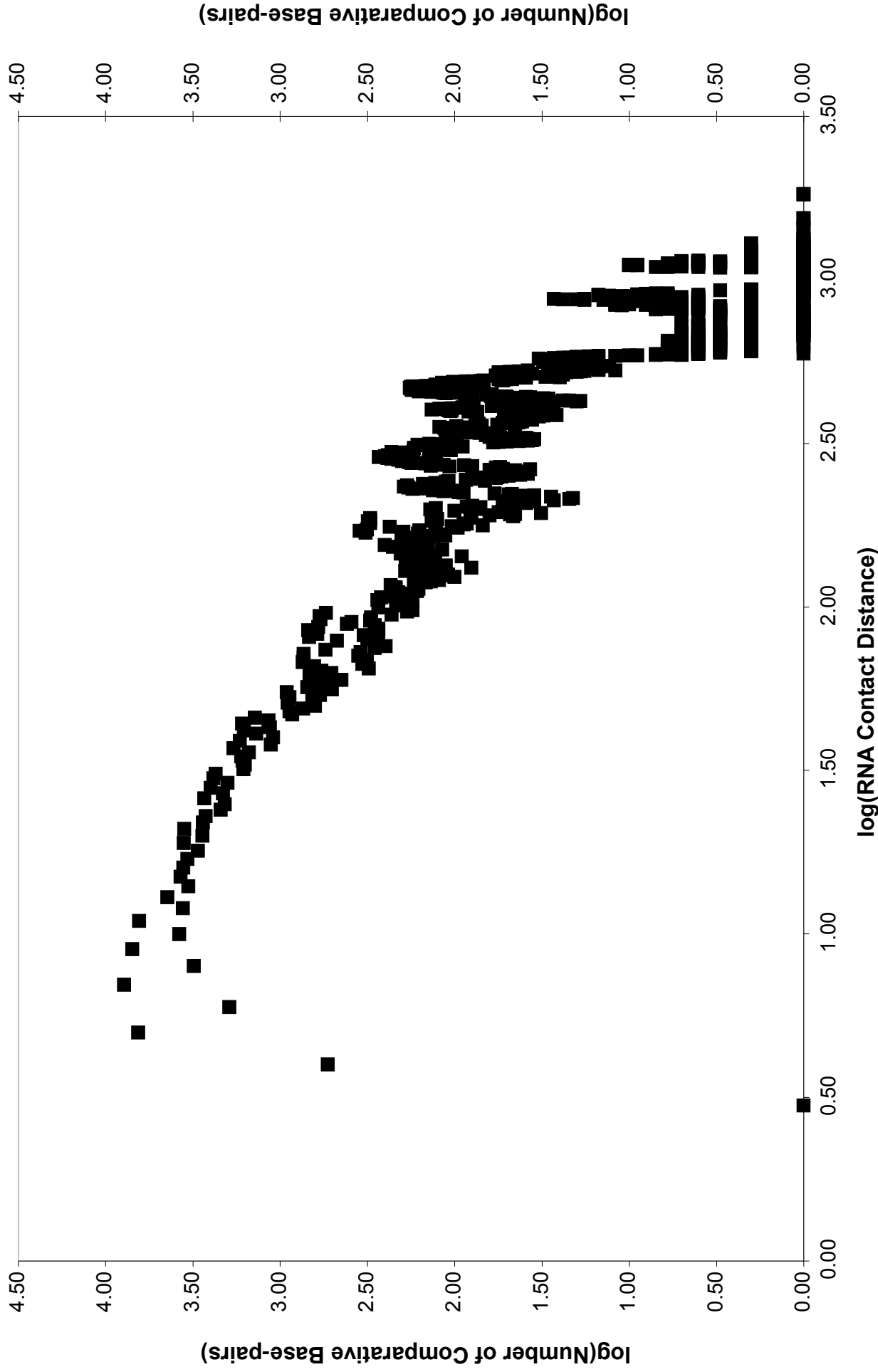


Distribution of 191,994 Comparative Canonical Base-pairs(from 496 16S rRNA Structures) by RNA Contact Distance



This chart shows a log-log plot of the distribution of 191,994 comparative, canonical base-pairs by contact distance. The base-pairs are taken from the 496 16S rRNA secondary structures in our dataset. the base-pairs range in contact distance from a minimum of 3 to a maximum of 1833. 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 predicted folds.