

## Input Mutational Profiling Data

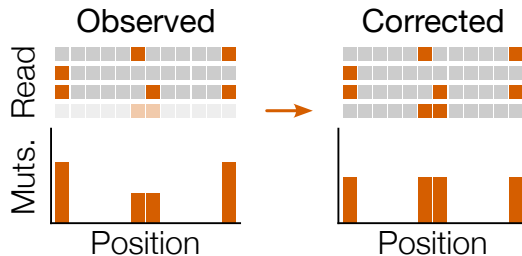
DMS-MaPseq    RNA editing  
SHAPE-MaP    Bisulfite-seq  
ETC-MaP    SNPs



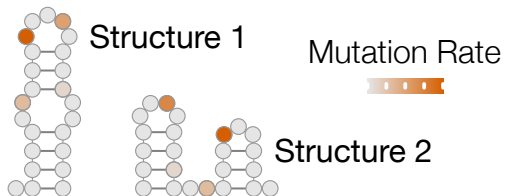
**Install and Run SEISMIC-RNA  
with One Command each**

```
> conda install seismic-rna  
> seismic wf refs.fa fqs/  
> _
```

## Fix Bias Against Nearby Mutations

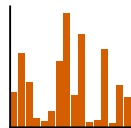


## Resolve RNA Structure Ensembles

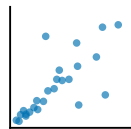


## Analyze in Depth

Profiles



Correlations



ROC Curves

