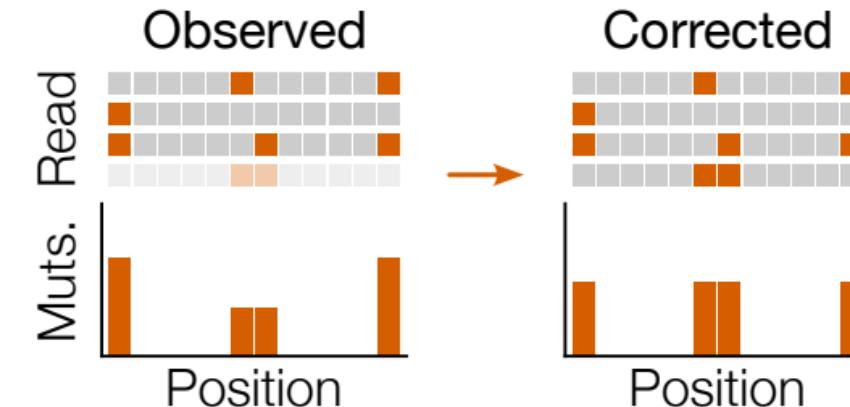


Input Mutational Profiling Data

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

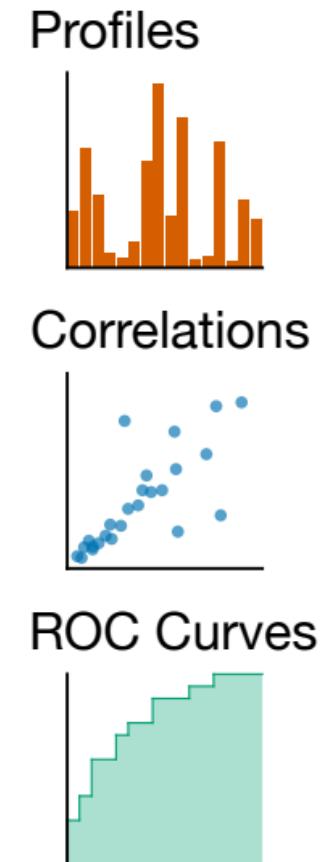
Fix Bias Against Nearby Mutations



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

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

Analyze in Depth



Resolve RNA Structure Ensembles

