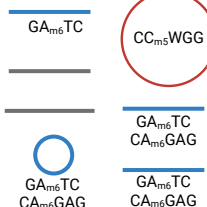
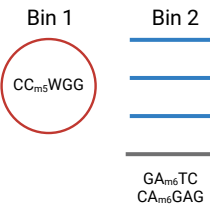


## Motif discovery

### Find motifs

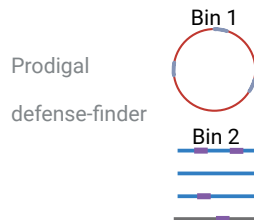


### bin-consensus



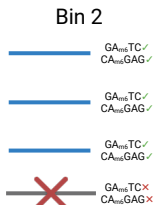
## MTase-Motif linking

### Predict MTase genes

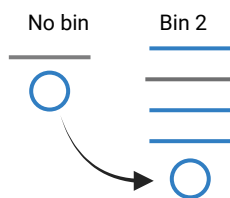


## Bin improvement

### Remove contaminants



### Associate unbinned



### Link Motifs

