

predict_genes
genome: *Vibrio.cholerae*

predict_genes
genome: *Vibrio.natriegens*

target

```
graph TD; A["predict_genes<br/>genome: Vibrio.cholerae"] --> C(target); B["predict_genes<br/>genome: Vibrio.natriegens"] --> C;
```

The diagram illustrates a workflow where two separate gene prediction tasks are performed. The first task uses the *Vibrio.cholerae* genome, and the second uses the *Vibrio.natriegens* genome. Both tasks output a 'target' value, which is represented by a red-bordered box at the bottom of the diagram.