

# Gene Prediction

2024-11-22

# Slide 1

Wow

## Slide 2

```
from Bio.Seq import Seq
from Bio import motifs

seqs = [Seq("GCCAT"), Seq("GCCAT"), Seq("GCCAT"), Seq("GCCA
        Seq("GCCAT"), Seq("GCCAT"), Seq("GCCAT"), Seq("GCCA
        Seq("GCCAT"), Seq("GCCAT"), Seq("GCCAT"), Seq("GCCA
        Seq("CGCAT"), Seq("CGATT"), Seq("CCAAT"), Seq("CCTA

motif = motifs.create(seqs)

print(motif.consensus)  #
print(motif.counts)    #
```