

Alineamiento de secuencias

```
#Alineamiento Ginger Yereni Cedeño
from Bio.Seq import Seq
from Bio import pairwise2
seq1 = Seq("TGTGACTA")
seq2 = Seq("CATGGTCA")
alignments = pairwise2.align.globalxx(seq1, seq2)

for match in alignments:
    print(match)
```

ModuleNotFoundError Traceback (most recent call last)

<ipython-input-24-4251102cf121> in <cell line: 1>()

----> 1 from Bio.Seq import Seq

2 from Bio import pairwise2

3 seq1 = Seq("TGTGACTA")

4 seq2 = Seq("CATGGTCA")

5 alignments = pairwise2.align.globalxx(seq1, seq2)

ModuleNotFoundError: No module named 'Bio'

NOTE: If your import is failing due to a missing package, you can manually install dependencies using either !pip or !apt.

To view examples of installing some common dependencies, click the "Open Examples" button below.

OPEN EXAMPLES

link de bioPython:

https://colab.research.google.com/drive/1vEbTZmQZpkLg0YVw_Sp1J4tgK3T4Ly2Y?usp=sharing