Trabalho 2: Filogenia

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a) Descubra o organismo correspondente de cada uma das 10 sequência (dica: usar o Blast para esta tarefa, https://blast.ncbi.nlm.nih.gov/Blast.cgi).

```
Formato da Resposta:
```

```
seq_1: Streptococcus agalactiae
seq_2: Neisseria gonorrhoeae
seq_3: Mycobacterium tuberculosis
seq_4: Staphylococcus aureus
seq_5: Treponema pallidum
seq_6: Bacillus anthracis
seq_7: Yersinia pestis
seq_8: Acinetobacter baumannii
seq_9: Pseudomonas aeruginosa
```

b) Realize o alinhamento global par-a-par de todas as sequências entre si. Para esta tarefa deve ser utilizado o algoritmo Needleman e Wunsch implementado no Trabalho I da disciplina. Você deve reportar o score de cada alinhamento considerando o seguinte esquema de pontuação: Match = +1; Mismatch = -1; Gap = -2.

Formato da Resposta:

seq 10: Helicobacter pylori

```
seq_1 vs seq_2: xxx seq_1 vs seq_3: xxx ... seq_10 vs seq_9: xxx
```

```
In [1]: import numpy as np
    sequences_ids = list(range(1, 11))
    matrix_size = len(sequences_ids)

distance_matrix = np.zeros((matrix_size, matrix_size))
In [21: # b)
```

```
In [2]: # b)
import global_alignment
import numpy as np
```

```
dir = "./2 phylogenetics"
format = "fasta"
id1 = "seq 1"
id2 = "seq 2"
scoring_file = "scoring"
sequences ids = list(range(1, 11))
num of seqs = len(sequences ids)
distance_matrix = np.zeros((num_of_seqs, num_of_seqs))
for target id in sequences ids:
   print("Alvo:", target_id)
   candidates ids = sequences ids.copy()
   candidates_ids.remove(target_id)
   print("Candidatos:", candidates ids)
   for candidate id in candidates ids:
        print(f"\nseq {target id} vs seq {candidate id}\n-----
        result = global_alignment.main(dir, format, f"seq_{target_id}", f"se
       print(result)
       # c)
        distance matrix[target id - 1, candidate id - 1] = result["score"]
       distance matrix[candidate id - 1, target id - 1] = result["score"]
   print("\n=====
   distance matrix
```

```
Alvo: 1
Candidatos: [2, 3, 4, 5, 6, 7, 8, 9, 10]
seq 1 vs seq 2
                    -----
{'seq 1': 'AAAC-ACCTCCAGTCATAAT-A-TTCGTAAACCAATCAAAAACTCATGTTTTAAATCAATAAAAA
ATACTTAG-', 'seq 2': 'CCCCTGCCGATTTTCGGAGTCAGACCGT-GCGTAATATAAAACGC-CG--GCCC
GCCGATGTATTTGCCGTGGC', 'score': -26, 'identity': 0.3783783783783784, 'matc
h': 28}
seq 1 vs seq 3
{'seq 1': '-AAACAC-CTCCAGTCATAATATTCGTAAACCAATCAAAAACTCA-TGTTTTAAATCAATAAAAA
AT-ACTTA-G', 'seq 2': 'TCACCACGCCCCCACGAGGAGCTCGT-GGGC-ACCCAGCATTCACTG-CTT-
-ACCACTACGATCTCGCTCACG', 'score': -15, 'identity': 0.466666666666667, 'matc
h': 35}
seq 1 vs seq 4
{'seq 1': 'AAACACCTCCAGTCATAATATTCGT-AAACCAATCAAAAACTCAT-GTTTTAAATCAATA-AAAA
ATACTTAG', 'seq 2': 'TTATTTTGAAAGTAACAAT-GGCATCATTATTATAAATGGT-ATAGGTTT-CTT
GTTTACTGGTATGCTTCA', 'score': -17, 'identity': 0.4246575342465753, 'match':
31}
seq 1 vs seq 5
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{'seq 1': '--AAACACCTCCAGTCATAATATTCGTAAACCAATCA-AAAACTCATGTTTTAAATCAATAAAAA
ATACTTAG', 'seq 2': 'TGATATACTTCAAG-CA-CAT-CACGCAGCCCAACCACTTTTCCCGCGGACGAAG
ACATCTTCCCAGGCAAAC', 'score': -17, 'identity': 0.4246575342465753, 'match':
31}
seq 1 vs seq 6
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{'seq 1': 'AAACACCTCCAGTCATAATA-TTCGTAAACCAAT-CAAAAACTCATGTTTTAAATCAATAAAAAA
TACTTA-G', 'seq 2': '-CTTGCAT-AAGT-ACAACACCGCATAAAATAATACCCGACATAACTAATTCCTT
CATTGGGAGTTTTTTATG', 'score': -13, 'identity': 0.4520547945205479, 'match':
33}
seq 1 vs seq 7
{'seq 1': '-AAACACCTCCAGTCATAATATTCG--TAAACCAATCAAAAACTCATGTTTTAAATCAATAAAAA
\verb|ATACTTAG', 'seq_2': 'GGTTGACAGCGAAT-ACAATACTCGCCGCCAACAGTGTGAAGCTGCCG-CCCGTT| \\
TCTTTGGCGTCAAAGCA-', 'score': -25, 'identity': 0.3698630136986301, 'match':
27}
seq 1 vs seq 8
{'seq 1': 'AAACACCTCCA-GTCATAATATTC-GTAAACCAATCAAAAACTCATGTTTTAAATCAATAAAAAA
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TATCATTTAAAGGGGT', 'score': -20, 'identity': 0.388888888888888, 'match': 2
8}
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{'seq_1': 'AAACACCTCCAGTCATAATATTCG-TAAACCAATCAAAAACTCATGTTTTAAATCAATAAAAAAT ACTTAG--', 'seq 2': '--TCGCCGAACGTCCAGGCCCTCGCCCAGCGCCCACCGCCCG-TACCGCCC

seq 1 vs seq 9

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TGGAACGCCTGCCCAGCC', 'score': -29, 'identity': 0.3424657534246575, 'match':
seq 1 vs seq 10
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{'seq 1': 'AAACACCTCCAGTCATAATATTCGTAA-ACCA--ATCAAAAACTCAT-GTTTTA-AATCAATAAA
AAATACTTAG--', 'seq 2': '---CA-CT-GAATCA-ATCCTTCTTAACTTTAGGATCACTTA-TTATGGGG
CTAGGATCAATAAAGGCTATCAAGCA', 'score': -11, 'identity': 0.5194805194805194,
'match': 40}
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Alvo: 2
Candidatos: [1, 3, 4, 5, 6, 7, 8, 9, 10]
seq 2 vs seq 1
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{'seq 1': 'CCCCTGCCGATTTTCGGAGTCAGACCGT-GCGTAATATAAAACGC-CG--GCCCGCCGATGTATT
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ATCAATAAAAAATACTTAG-', 'score': -26, 'identity': 0.3783783783783784, 'matc
h': 28}
seq 2 vs seq 3
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{'seq 1': '-C-CC-CTGCCGATTTTCGGAG-TCAGACCGTGCGTAATATAAAACGCCGGCC-CGCCGATGTAT
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ACTACGA--T-CTCGCTCACG-', 'score': -19, 'identity': 0.44, 'match': 33}
seq 2 vs seq 4
{'seq 1': 'CCCCTGCCGATTTTCGGAGTCAGACCGTGCGTAATATAAAACG-CCGGCCCGCCGATGTATTTG-
CCG-TGGC', 'seq 2': '-TTATTTTGAAAGTAACAATGGCATCAT-TATTAT-TAAATGGTATAGGTTTCTT
GTTTACTGGTATGCTTCA', 'score': -27, 'identity': 0.3561643835616438, 'match':
26}
seq 2 vs seq 5
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{'seq 1': 'CCCCTGCCGAT-TTTCGGAGTCAGA-C-CG-TGCGTAATATAAAACGCCGGCCCGCCGATG-TAT
-TTGCC--G-TGGC', 'seq 2': '----TG-ATATACTTC-AAG-CACATCACGCAGCCCAA-CCACTTTTCC
CG-CGGACGAAGACATCTTCCCAGGCAAAC', 'score': -23, 'identity': 0.468354430379746
83, 'match': 37}
seq 2 vs seq 6
         {'seq 1': 'C--CCCT--GCCGATTTTCGGAGTCAGACCGTGCGTAATATAA--AACGCCGGCCGCCGATGTA
TTTGCCGTGGC', 'seq 2': 'CTTGCATAAGTACAACACCGCA-TAAAATAACCGGACATAACTAATTCCT
TCATTGGGA-GT-TTT-TTAT-G-', 'score': -24, 'identity': 0.42105263157894735, 'm
atch': 32}
seq 2 vs seq 7
{'seq 1': 'CCCCTG-C--CGATTTTCGGAGTCAGACCGTGCGTAATATAAAACGCCGGCCCGCCGATGTATTT
GCCGT---GGC-', 'seq 2': '-GGTTGACAGCGAATACAATACTC-G-CCGCCAACAGTGT-GAA-G-CTG-
CCGCCCGTTTCTTTGGCGTCAAAGCA', 'score': -15, 'identity': 0.4935064935064935,
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'match': 38}

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seq 2 vs seq 8
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{'seq 1': 'CCCCTGCCGATTTTCGGAGTCAGACCGTGC-GTAATATAAAACGCCGGCCGCCGATGT-ATTTG CCGTGGC', 'seq 2': '-AGAGACTAATATGTTTTTTCACACTTTTTCTGCTTTTTTACA-TCCTGAAATATAT

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seq 2 vs seq 9

{'seq 1': '-CCCCTGCCGATTTTCGGAGTCAGACCGTGCGTAATATAAAACG-CCGGCCCGCCGATGTA--TT TGCCGTGGC', 'seq 2': 'TCGCCGAACG-TCCAGGCCGTC-G-CCCTGC-CCAGCGCCACCGCCCGTACCGC CCTGGAACGCCTGCCCAGCC', 'score': -10, 'identity': 0.4864864864864865, 'matc h': 36}

seq 2 vs seq 10

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Alvo: 3

Candidatos: [1, 2, 4, 5, 6, 7, 8, 9, 10]

seq 3 vs seq 1

{'seq 1': 'TCACCACGCCCCCACGACGGAGCTCGT-GGGC-ACCCAGCATTCACTG-CTT--ACCACTACGAT CTCGCTCACG', 'seq 2': '-AAACAC-CTCCAGTCATAATATTCGTAAACCAATCAAAAACTCA-TGTTTTA AATCAATAAAAAT-ACTTA-G', 'score': -15, 'identity': 0.46666666666667, 'matc h': 35}

seq_3 vs seq_2

{'seq 1': 'TCACCACGCCCCCACGAGGAGCTC-GTGGGCACCCAGCAT-TCACTGCTTACC-ACTACG-ATC T-CGCTCACG', 'seq 2': '-C-CC-CTGCCGATTTTCGGAG-TCAGACCGTGCGTAATATAAAACGCCGGCC CGCCGATGTATTTGCCGTGGC-', 'score': -19, 'identity': 0.44, 'match': 33}

seq 3 vs seq 4

{'seq 1': 'TCACCACG-CCCCCACGACGG-AGC-TCGTGGGCACCCAGCATTCACTGCTT-ACCACTACGATC TCGCTCACG', 'seq 2': 'TTATTTTGAAAGTAACAATGGCATCATTATTATAAATGGTATAGGTTTCTTGT TTACT--GGTAT-GCT-TCA', 'score': -22, 'identity': 0.40540540540540543, 'matc h': 30}

seq_3 vs seq_5

{'seq 1': 'TCACCACGCCCCCACGAGGAGCTCG-TGGGCACCCAGCATTCACTGCTTACCACTACGATC-TC GC--TC-ACG', 'seq 2': 'TGA-TA-TACTTCAAG-CACATCACGCAGCCCAACCA-CTTTTCCCGCGGACG AAGAC-ATCTTCCCAGGCAAAC', 'score': -11, 'identity': 0.493333333333335, 'mat ch': 37}

seq 3 vs seq 6

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{'seq 1': '-TCAC-CACGCCCCACGACGGAGCTCGTGGGCACCCAGCATTCACTGCTTAC--CACTACGATC
TCGCTCACG', 'seq 2': 'CTTGCATAAG-TACAACACCGCATAAAAT-AATACCCGACA-TAACTAATTCCT
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h': 33}
seq 3 vs seq 7
{'seq 1': 'TCACCAC-GCCCCCACGACGGAGCTCGTGGGCACCCAGCATTCA-CTGCTTACCACTACGATCTC
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GTTTCTTTGGCGTCAAAGCA', 'score': -14, 'identity': 0.4594594594594595, 'matc
h': 34}
seq 3 vs seq 8
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{'seq 1': '--TCACCACGCCCCCACGACGGAGCTCGTGGGC-ACCCAGCATTCACTG-CTTACCACTACGATC
TCGCTCACG', 'seq 2': 'AGAGACTAATATGTTTTTTCACA-CTTTTTCTGCTTTTTTACA-TC-CTGAAATA
-TATTATCATTTAAAGGGGT', 'score': -32, 'identity': 0.33783783783783783, 'matc
h': 25}
seq 3 vs seq 9
{'seq 1': 'TCACCACGCCCCCACGACGGAGCTCGTGGGCA-C-CCAGCATTCACTGCTTACC-ACTACGATCT
CGCTCA-CG', 'seq 2': 'TCGCCGAACGTCCAGGCCGTCGC-CCTGCCCAGCGCCCA-CCGCCCGTACCGCCC
TGGAACG-CCT-GCCCAGCC', 'score': -6, 'identity': 0.5135135135135135, 'match':
38}
seq 3 vs seq 10
{'seq 1': 'TCAC-CACGCCCCCACGACGAGCTCGTGG-GCACCCAGCAT-TCACTGCTTACCACTACGATCT
CGCTCACG', 'seq 2': '-CACTGAATCAATC-CTTCTTAACTTTAGGATCACTTATTATGGGGCT-AGGATC
AATAAAGGCTATCAAGCA', 'score': -19, 'identity': 0.410958904109589, 'match': 3
0}
______
_____
Alvo: 4
Candidatos: [1, 2, 3, 5, 6, 7, 8, 9, 10]
seq 4 vs seq 1
{'seq 1': 'TTATTTTGAAAGTAACAAT-GGCATCATTATTAAATGGT-ATAGGTTT-CTTGTTTACTGGT
CAATA-AAAAATACTTAG', 'score': -17, 'identity': 0.4246575342465753, 'match':
31}
seq 4 vs seq 2
{'seq 1': '-TTATTTTGAAAGTAACAATGGCATCAT-TATTAT-TAAATGGTATAGGTTTCTTGTTTACTGGT
ATGCTTCA', 'seq 2': 'CCCCTGCCGATTTTCGGAGTCAGACCGTGCGTAATATAAAACG-CCGGCCCGCCG
ATGTATTTG-CCG-TGGC', 'score': -27, 'identity': 0.3561643835616438, 'match':
26}
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seq 4 vs seq_3

```
{'seq 1': 'TTATTTTGAAAGTAACAATGGCATCATTATTATTAAATGGTATAGGTTTCTT--GTTTACTGGTA
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CACTAC-GATCTCGCTCACG', 'score': -22, 'identity': 0.40540540540543, 'matc
h': 30}
seq 4 vs seq 5
{'seq 1': 'T--TATTTTGAAAGTA-A-CA-ATGGCATCATTATTATTAAATGGTATAGGTTTCTTGTTTACTG
GTATGCTTCA', 'seq 2': 'TGATATACTTCAAGCACATCACGCAGCCCAACCACTTTTCCCGCGGA-CGAAG
ACATCTTCCCAGGCA---AAC-', 'score': -23, 'identity': 0.413333333333333, 'mat
ch': 31}
seq 4 vs seq 6
{'seq 1': 'TTATTTTGAAAGTA-ACAATGGCATCATTATTATTAAATGGTATAGGTTTCTTGTTTACTGGTA-
TGCTTCA--', 'seq 2': '--CTTGCATAAGTACACCCGCAT-A-AAATAATACCCGACATAACTAATTCC
TTCATTGGGAGTTTTTTATG', 'score': -10, 'identity': 0.4864864864864865, 'matc
h': 36}
seq 4 vs seq 7
{'seq 1': '--TT-ATTTTGAAAGTAACAAT-GGCATCATTATTATAA-ATGGTATAGGTTTCTTGTTTACTG
GTATGCTTCA', 'seq 2': 'GGTTGACAGCGAATACAATACTCGCCGCCAACAGTGTGAAGCTGCCGCCCGTT
TC-T-TTGGC--GT-CAAAGCA', 'score': -17, 'identity': 0.4533333333333333, 'matc
h': 34}
seq 4 vs seq 8
------
{'seq 1': 'TTATTTTGA-AAGTAACAATGGCA-TCAT-T--ATTATTAAATGGTATAGGTTTCTTGTTTACTG
GTATGCTTCA', 'seq_2': 'AGAGACTAATATGT-TTTTTCACACTTTTTTTACATCCT-GAAATA
TATT-ATCATTTAAAGGGGT--', 'score': -23, 'identity': 0.4133333333333333, 'mat
ch': 31}
seq 4 vs seq 9
{'seq 1': 'TTATTTTGAAAGTAACAATGGCATCATTATTATTAAATGGTATAGGTTTCTTGTTTACTGGTATG
-CT--TCA---', 'seq 2': '-T-CGCCGAACGT--CCAGGCCGTC-GCCCTGCCCAGCGCCA-CCGCCCGTA
CCGCCCTGGAACGCCTGCCCAGCC', 'score': -32, 'identity': 0.3684210526315789, 'ma
tch': 28}
seq 4 vs seq 10
{'seq 1': 'TTATTTTGAAAGTAACAATGGCATCATTATTATTAAATGGTAT-AGG-TTTCTTGTTTACTGG-T
ATGCTTCA', 'seq 2': '-CA--CTGAATCATCCTTCTTAACTTTAGGATCACTTATTATGGGGCTAGGATC
AATAAAGGCTATCAAGCA', 'score': -13, 'identity': 0.4520547945205479, 'match':
33}
Alvo: 5
```

Candidatos: [1, 2, 3, 4, 6, 7, 8, 9, 10]

seq_5 vs seq_1

{'seq 1': 'TGATATACTTCAAG-CA-CAT-CACGCAGCCCAACCACTTTTCCCGGGACGAAGACATCTTCCC

AGGCAAAC', 'seq 2': '--AAACACCTCCAGTCATAATATTCGTAAACCCAATCA-AAAACTCATGTTTTAAA TCAATAAAAATACTTAG', 'score': -17, 'identity': 0.4246575342465753, 'match': 31} seq_5 vs seq_2 {'seq 1': '----TG-ATATACTTC-AAG-CACATCACGCAGCCCAA-CCACTTTTCCCG-CGGACGAAGACAT CTTCCCAGGCAAAC', 'seq 2': 'CCCCTGCCGAT-TTTCGGAGTCAGA-C-CG-TGCGTAATATAAAACGCC GGCCCGCCGATG-TAT-TTGCC--G-TGGC', 'score': -23, 'identity': 0.468354430379746 83, 'match': 37} seq 5 vs seq 3 {'seq 1': 'TGA-TA-TACTTCAAG-CACATCACGCAGCCCAACCA-CTTTTCCCGCGGACGAAGAC-ATCTTC CCAGGCAAAC', 'seq 2': 'TCACCACGCCCCCACGACGGAGCTCG-TGGGCACCCAGCATTCACTGCTTACC ACTACGATC-TCGC--TC-ACG', 'score': -11, 'identity': 0.493333333333335, 'mat ch': 37} seq 5 vs seq 4 ______ {'seq 1': 'TGATATACTTCAAGCACATCACGCAGCCCAACCACTTTTCCCGCGGA-CGAAGACATCTTCCCAG GCA---AAC-', 'seq 2': 'T--TATTTTGAAAGTA-A-CA-ATGGCATCATTATTATATAAATGGTATAGGTT TCTTGTTTACTGGTATGCTTCA', 'score': -23, 'identity': 0.4133333333333333, 'mat ch': 31} seq 5 vs seq 6 {'seq 1': 'TGATATACTTCAAGCACATCACGCAGCCCAACCACTTTTCCCG-C-GGACGAA-GACATC-TTCC CAG---GCAAAC', 'seq 2': '--CT-TGCAT-AAGTACAACAC-C-GCATAA-AATAACACGACATAACT AATTCCTTCATTGGGAGTTTTTTATG', 'score': -17, 'identity': 0.4805194805194805, 'match': 37} seq 5 vs seq_7 {'seq 1': 'TGATATACTTCAAGCAC-AT-CACGCAGCCCAACCACTTTTCCCGCGGACGAAGACATCTTCCCA GGCAAA-C-', 'seq 2': '-GGTTGACAGCGAATACAATACTCGCCG-CCAA-CA-GTGTGAAGCTGCCGCCC GTTTCTTTGGCGTCAAAGCA', 'score': -6, 'identity': 0.5135135135135135, 'match': 38} seq 5 vs seq 8 {'seq 1': 'TGATA-TACTTCAAG-CACATCACGCAGCCCAACCACTTTTCCCGCGGACGAAGA-CATC-TTCC CAGGCAAAC', 'seq 2': 'AGAGACTA-AT-ATGTTTTTTCACACTTTTTCTGCTTTTTTACATCCTGAAATAT ATTATCATTTAAAGG--GGT', 'score': -18, 'identity': 0.43243243243243246, 'matc h': 32} seq 5 vs seq 9 ______ {'seq 1': 'T-GATATACTTCAAGCACATCACGCAGCCCAACCACTTTTCCCG--CGGACGAAG-ACATCTTCC CAGGCAAAC', 'seq 2': 'TCGCCGAACGTCCAGGCCGTCGCCCTGCCCAGCGCCACCGCCCGTACCGCCCTG GAACGCCTGCCCA-GC---C', 'score': -4, 'identity': 0.527027027027027, 'match': 39} seq 5 vs seq 10

{'seq_1': 'TGATATACTTCAAGCACATC-ACGCAGCCCAACCACTTTTCCCGCGGACGAAGA-C-ATCTTCCC

AGGCAAAC-', 'seq_2': '-CA-CTGAATCAATC-CTTCTTAACTTTAGGATCACTTATTATG-GGGCTAGGA TCAATAAAGGCTATCAAGCA', 'score': -16, 'identity': 0.44594594594594594, 'matc h': 33}

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Alvo: 6

Candidatos: [1, 2, 3, 4, 5, 7, 8, 9, 10]

seq_6 vs seq_1

{'seq_1': '-CTTGCAT-AAGT-ACAACACCGCATAAAATAATACCCGACATAACTAATTCCTTCATTGGGAGT TTTTTATG', 'seq_2': 'AAACACCTCCAGTCATAATA-TTCGTAAACCCAAT-CAAAAACTCATGTTTTAAAT CAATAAAAAAATACTTA-G', 'score': -13, 'identity': 0.4520547945205479, 'match': 33}

seq_6 vs seq_2

{'seq_1': 'CTTGCATAAGTACACCCGCA-TAAAATAATACCCGACATAACTAATTCCTTCATTGGGA-GT-TTT-TTAT-G-', 'seq_2': 'C--CCCT--GCCGATTTTCGGAGTCAGACCGTGCGTAATATAA--AACGCCGGCCCGCCGATGTATTTGCCGTGGC', 'score': -24, 'identity': 0.42105263157894735, 'm atch': 32}

seq 6 vs seq 3

{'seq_1': 'CTTGCATAAG-TACAACACCGCATAAAAT-AATACCCGACA-TAACTAATTCCTTCATTGGGA-G
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h': 33}

seq 6 vs seq 4

{'seq_1': '--CTTGCATAAGT-ACAA-CACCGCATAAAATAATACCCGACATAACTAATTCCTTCATTGGGAG TTTTTTTATG', 'seq_2': 'TTATTTTGAAAGTAACAATGGCATCAT-TATTATTAAATGGTATAGGTTTCTTG TTTACTGGTA-TGCTTCA--', 'score': -10, 'identity': 0.4864864864864865, 'matc h': 36}

seq 6 vs seq 5

{'seq_1': '--CT-TGCAT-AAGTACAACAC-C-GCATAA-AATAATACCCGACATAACTAATTCCTTCATTGG GAGTTTTTTATG', 'seq_2': 'TGATATACTTCAAGCACATCACGCAGCCCAACCACTTTTCCCG-C-GGACG AA-GACATC-TTCCCAG---GCAAAC', 'score': -17, 'identity': 0.4805194805194805, 'match': 37}

seq 6 vs seq 7

{'seq_1': '-CTTG-CA-TAAGTACAACAC-CGCATAAAATAATACCCGACATAACTAATTCCTTCATTGGGAG TTTTTTTATG', 'seq_2': 'GGTTGACAGCGAATACAATACTCGCCGCCAACAGT--GTGA-AGCTGCCGCCCG TTTCTTTGGCG-TCAAAGCA', 'score': -18, 'identity': 0.43243243243243243246, 'matc h': 32}

seq 6 vs seq 8

{'seq_1': 'CTTG-C--ATAAG-TACAACACCGCATAAAATAATACCCGACATAACT-AATTCCTTCATTGGGAGTTTTTTATG-', 'seq_2': 'AGAGACTAATATGTTTTTTCA-CAC-TTTTCTGCTTTTTTACAT-CCTGAAA

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atch': 31}
seq 6 vs seq 9
______
{'seq 1': 'CTTGCATAA-GTACAACACCGCATAAAATAATACCCGACATAACTAATTCCTTCATTGGGA-G-T
TTTTTA-TG', 'seq 2': '-TCGCCGAACGT-CCAGGCCG-TCGCCCTGCCCAGCGCCCCCGTACC-GC
CCTGGAACGCCTGCCCAGCC', 'score': -24, 'identity': 0.3918918918919, 'matc
h': 29}
seq 6 vs seq 10
{'seq_1': 'C-TTGCATAAGTACAACACCGCATAAAATAATACCCGACATAACTAATTCCTTCATTGGGAGTTT
TTTATG--', 'seq 2': 'CACTGAATCATCCTTCTTAAC-TTTAGGATCACTTATTAT-GGGGCTAGGATCA
AT-AAAGGCTATCAAGCA', 'score': -23, 'identity': 0.3835616438356164, 'match':
28}
Alvo: 7
Candidatos: [1, 2, 3, 4, 5, 6, 8, 9, 10]
seq 7 vs seq 1
               -----
{'seq 1': 'GGTTGACAGCGAAT-ACAATACTCGCCGCCAACAGTGTGAAGCTGCCG-CCCGTTTCTTTGGCGT
CAAAGCA-', 'seq 2': '-AAACACCTCCAGTCATAATATTCG--TAAACCAATCAAAAACTCATGTTTTAAA
TCAATAAAAATACTTAG', 'score': -25, 'identity': 0.3698630136986301, 'match':
27}
seq 7 vs seq 2
{'seq 1': '-GGTTGACAGCGAATACATACTC-G-CCGCCAACAGTGT-GAA-G-CTG-CCGCCCGTTTCTTT
GGCGTCAAAGCA', 'seq 2': 'CCCCTG-C--CGATTTTCGGAGTCAGACCGTGCGTAATATAAAACGCCGGC
CCGCCGATGTATTTGCCGT---GGC-', 'score': -15, 'identity': 0.4935064935064935,
'match': 38}
seq 7 vs seq 3
{'seq 1': 'GGTTGACAGCGAATAC-A-ATA-CTCGCCGCCA-ACAGTGTGAAGCTGCCGCCCGTTTCTTTGGC
GTCAAAGCA', 'seq 2': 'TCACCAC-GCCCCCACGACGGAGCTCGTGGGCACCCAGCATTCA-CTGCTTACC
ACTACGATCTCG-CTCA-CG', 'score': -14, 'identity': 0.4594594594595, 'matc
h': 34}
seq 7 vs seq 4
{'seq 1': 'GGTTGACAGCGAATACAATACTCGCCGCCAACAGTGTGAAGCTGCCGCCCGTTTC-T-TTGGC--
GT-CAAAGCA', 'seq 2': '--TT-ATTTTGAAAGTAACAAT-GGCATCATTATTATTAA-ATGGTATAGGTT
TCTTGTTTACTGGTATGCTTCA', 'score': -17, 'identity': 0.453333333333333, 'matc
h': 34}
seq 7 vs seq 5
______
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{'seq_1': '-GGTTGACAGCGAATACATACTCGCCG-CCAA-CA-GTGTGAAGCTGCCGCCCGTTTCTTTGGC GTCAAAGCA', 'seq_2': 'TGATATACTTCAAGCAC-AT-CACGCAGCCCAACCACTTTTCCCGCGGACGAAG ACATCTTCCCAGGCAAA-C-', 'score': -6, 'identity': 0.5135135135135135, 'match':

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seq 7 vs seq 6
```

{'seq_1': 'GGTTGACAGCGAATACAATACTCGCCGCCAACAGT--GTGA-AGCTGCCGCCCGTTTCTTTGGCG
-TCAAAGCA', 'seq_2': '-CTTG-CA-TAAGTACAACAC-CGCATAAAATAATACCCGACATAACTAATTCC
TTCATTGGGAGTTTTTTATG', 'score': -18, 'identity': 0.43243243243243243246, 'matc
h': 32}

seq_7 vs seq_8

{'seq_1': 'GGTTGACAGCGAATA-CAATACTCGCCGCCAACAG-TGTGAAGCTGCCGCCCGTTTCTT-TGGCG
TCAA-AGCA', 'seq_2': '---AGAGA-CTAATATGTTTTTTCACACTTTTTTCACACTTTTTTACATCCTGAAAT
ATATTATCATTTAAAGGGGT', 'score': -26, 'identity': 0.3783783783783784, 'matc
h': 28}

seq_7 vs seq_9

{'seq_1': 'GGTTGAC-AGCGAATACATAC-TCGCCGCCAACAGTGTGAAGCTGCCG-CCCG-TTTCTTTGGC
GT-CAAAGCA', 'seq_2': '--TCGCCGAACG--TCCAGGCCGTCGCCCTGCCCAGCG-CCACCGCCCGTACC
GCCCTGGAACGCCTGCCCAGCC', 'score': -15, 'identity': 0.4666666666666667, 'matc
h': 35}

seq_7 vs seq_10

Alvo: 8

Candidatos: [1, 2, 3, 4, 5, 6, 7, 9, 10]

seq 8 vs seq 1

{'seq_1': 'AGAGACTAATATGTTTTTTCACAC-TTTTCTGCTTTTTTTACATCCTGAAATATAT-TATCATTTA AAGGGGT', 'seq_2': 'AAACACCTCCA-GTCATAATATTCGTAAACCAATCAAAAAC-TCATGTTTTAAATC AATAAAAAAATACTTAG', 'score': -20, 'identity': 0.388888888888888, 'match': 2 8}

seq 8 vs seq 2

{'seq_1': '-AGAGACTAATATGTTTTTTCACACTTTTC-TGCTTTTTTACATCCTGAAATATATTATCATTTA AAGGGGT', 'seq_2': 'CCCCTGCCGATTTTCGGAGTCAGACCGTGCGTAATATAAAAC-GCCGGCCCGA TGT-ATTTGCCGTGGC', 'score': -20, 'identity': 0.388888888888888, 'match': 2 8}

seq 8 vs seq 3

{'seq_1': 'AGAGACTAATATGTTTTTTCACA-CTTTTCTGCTTTTTTACA-TC-CTGAAATA-TATTATCATT TAAAGGGGT', 'seq_2': '--TCACCACGCCCCCACGACGGAGCTCGTGGGC-ACCCAGCATTCACTG-CTTA CCACTACGATCTCGCTCACG', 'score': -32, 'identity': 0.33783783783783783783, 'matc h': 25}

.....

seq_8 vs seq_5

{'seq_1': 'AGAGACTA-AT-ATGTTTTTTCACACTTTTCTGCTTTTTTACATCCTGAAATATATTATCATTTA AAGG--GGT', 'seq_2': 'TGATA-TACTTCAAG-CACATCACGCAGCCCAACCACCTTTTCCCGCGGACGAAG A-CATC-TTCCCAGGCAAAC', 'score': -18, 'identity': 0.43243243243243246, 'matc h': 32}

seq_8 vs seq_6

{'seq_1': 'AGAGACTAATATGTTTTTTCACACTTTTCTGCTTTTTTACAT-CCTGAAAT--AT-ATTATCATT TAAAGGGGT', 'seq_2': 'CTTGCATAA-GTACAACACCGCA-TAAAATAATACCCGACATAACT-AATTCCT TCATTGGGAGTTTTTTATG-', 'score': -26, 'identity': 0.3783783783783784, 'matc h': 28}

seq_8 vs seq_7

{'seq_1': '---AGAGA-CTAATATGTTTTTTCACACTTTTTCTGCTTTTTTACATCCTGAAATATATTATCATT TAAAGGGGT', 'seq_2': 'GGTTGACAGCGAATA-CAATACTCGCCGCCAACAG-TGTGAAGCTGCCGCCCGT TTCTT-TGGCGTCAA-AGCA', 'score': -26, 'identity': 0.3783783783783784, 'matc h': 28}

seq_8 vs seq_9

<u>-</u>

{'seq_1': 'AGAGACTAATATGTTTTTTCACACTTTTCTGCTTTTTTACATCCTG-AAATATAT-TATCATTTA AAGGGGT', 'seq_2': '-TCGCCGAACGTCCAGGCCGTCGCCCTGC-CCAGCGCCCACCGCCCGTACCGCCCTG GAACGCCTGCCCAGCC', 'score': -34, 'identity': 0.2916666666666667, 'match': 2 1}

seq 8 vs seq 10

{'seq_1': '-AGAGACTAATATGTTTTTTCACACTTTTCTGCTTTTTTA-CATCCTG-AAATAT-ATTATCATT TAAAGGGGT', 'seq_2': 'CACTGAATCA-ATCCTTCTT-A-AC-TTTAGGATCACTTATTATGGGGCTAGGA TCAATAAAGGCTATCAAGCA', 'score': -14, 'identity': 0.4594594594594595, 'matc h': 34}

......

Alvo: 9

Candidatos: [1, 2, 3, 4, 5, 6, 7, 8, 10]

seq_9 vs seq_1

{'seq_1': '--TCGCCGAACGTCCAGGCCGTCGCCCTGCCCAGCGCCCACCGCCCG-TACCGCCCTGGAACGCCTGCCCAGCC', 'seq_2': 'AAACACCTCCAGTCATAATATTCG-TAAACCAATCAAAAAACTCATGTTTTAAATCAATAAAAAAATACTTAG--', 'score': -29, 'identity': 0.3424657534246575, 'match': 25}

{'seq_1': 'TCGCCGAACG-TCCAGGCCGTC-G-CCCTGC-CCAGCGCCCACCGCCCGTACCGCCCTGGAACGCC
TGCCCAGCC', 'seq_2': '-CCCCTGCCGATTTTCGGAGTCAGACCGTGCGTAATATAAAACG-CCGGCCCGC
CGATGTA--TTTGCCGTGGC', 'score': -10, 'identity': 0.4864864864864865, 'matc
h': 36}

seq_9 vs seq_3

{'seq_1': 'TCGCCGAACGTCCAGGCCGTCGC-CCTGCCCA-GC-GCCACCGC-CCGTACCGCCCTGGAACGCC
TGCCCAGCC', 'seq_2': 'TCACCACGCCCCCACGACGGAGCTCGTGGGCACCCAGCATTCACTGCTTACC-A
CTACGATC-TC-GCTCA-CG', 'score': -6, 'identity': 0.5135135135135135, 'match':
38}

seq_9 vs seq_4

{'seq_1': '-T-CGCCGAACGT--CCAGGCCGTC-GCCCTGCCCAGCGCCA-CCGCCCGTACCGCCCTGGAACG CCTGCCCAGCC', 'seq_2': 'TTATTTTGAAAGTAACAATGGCATCATTATTATTAAATGGTATAGGTTTCTT GTTTACTGGTATG-CT--TCA---', 'score': -32, 'identity': 0.3684210526315789, 'ma tch': 28}

seq_9 vs seq_5

{'seq_1': 'TCGCCGAACGTCCAGGCCGTCGCCCTGCCCAGCGCCCACCGCCCGTACCGCCCTGGAACGCCTGCC
CA-GC---C', 'seq_2': 'T-GATATACTTCAAGCACATCACGCAGCCCAACCACTTTTCCCG--CGGACGAA
G-ACATCTTCCCAGGCAAAC', 'score': -4, 'identity': 0.527027027027027, 'match':
39}

seq 9 vs seq 6

{'seq_1': '-TCGCCGAACGT-CCAGGCCG-TCGCCCTGCCCAGCGCCCCGTACC-GCCCTGGAACGCC TGCCCAGCC', 'seq_2': 'CTTGCATAA-GTACAACACCGCATAAAATAACCCGACATAACTAATTCCTTC ATTGGGA-G-TTTTTTA-TG', 'score': -24, 'identity': 0.3918918918918919, 'matc h': 29}

seq 9 vs seq 7

{'seq_1': '--TCGCCGAACG--TCCAGGCCGTCGCCCTGCCCAGCG-CCACCGCCCGTACCGCCCTGGAACGC
CTGCCCAGCC', 'seq_2': 'GGTTGAC-AGCGAATACAATAC-TCGCCGCCAACAGTGTGAAGCTGCCG-CCC
G-TTTCTTTGGCGT-CAAAGCA', 'score': -15, 'identity': 0.4666666666666667, 'matc
h': 35}

seq 9 vs seq 8

{'seq_1': '-TCGCCGAACGTCCAGGCCGTCGCCCTGC-CCAGCGCCACCGCCCGTACCGCCCTGGAACGCCTG
CCCAGCC', 'seq_2': 'AGAGACTAATATGTTTTTTCACACTTTTTCTGCTTTTTTACATCCTG-AAATATATTATCATTTAAAAGGGGT', 'score': -34, 'identity': 0.291666666666667, 'match': 2
1}

seq_9 vs seq_10

{'seq_1': 'TCGCCGAACGTCCAGGCCGTC--GCCCTGCCCAGCGC-CA--CCGCCCGTACCGCCCTGGAACGC
CTGCCCAGCC', 'seq_2': '-CACTGAA--T-CAATCCTTCTTAACTTTAGGATCACTTATTATGGGGCTA-G
GATCAATAAAGGCTATCAAGCA', 'score': -21, 'identity': 0.4266666666666667, 'matc
h': 32}

Alvo: 10

Candidatos: [1, 2, 3, 4, 5, 6, 7, 8, 9]

seq_10 vs seq_1

{'seq_1': '---CA-CT-GAATCA-ATCCTTCTTAACTTTAGGATCACTTA-TTATGGGGCTAGGATCAATAAA GGCTATCAAGCA', 'seq_2': 'AAACACCTCCAGTCATAATATTCGTAA-ACCA--ATCAAAAAACTCAT-GTT TTA-AATCAATAAAAAAATACTTAG--', 'score': -11, 'identity': 0.5194805194805194, 'match': 40}

seq_10 vs seq_2

seq 10 vs seq 3

{'seq_1': '-CACTGAATCAATC-CTTCTTAACTTTAGGATCACTTATTATGGGGCT-AGGATCAATAAAGGCT ATCAAGCA', 'seq_2': 'TCAC-CACGCCCCCACGACGGAGCTCGTGG-GCACCCAGCAT-TCACTGCTTACC ACTACGATCTCGCTCACG', 'score': -19, 'identity': 0.410958904109589, 'match': 3 0}

seq_10 vs seq_4

{'seq_1': '-CA--CTGAATCAATCCTTCTTAACTTTAGGATCACTTATTATGGGGCTAGGATCAATAAAGGCT ATCAAGCA', 'seq_2': 'TTATTTTGAAAGTAACAATGGCATCATTATTATTAAATGGTAT-AGG-TTTCTTG TTTACTGG-TATGCTTCA', 'score': -13, 'identity': 0.4520547945205479, 'match': 33}

seq 10 vs seq 5

{'seq_1': '-CA-CTGAATCAATC-CTTCTTAACTTTAGGATCACTTATTATG-GGGCTAGGATCAATAAAGGC TATCAAGCA', 'seq_2': 'TGATATACTTCAAGCACATC-ACGCAGCCCAACCACTTTTCCCGCGGACGAAGA-C-ATCTTCCCAGGCAAAC-', 'score': -16, 'identity': 0.44594594594594594, 'matc h': 33}

seq_10 vs seq_6

seq_10 vs seq_7

{'seq_1': 'CACTGA-ATC-AAT-C-CTTCTTAACTTTAGGATCACTTATTATGGGGCTAGGATCAATAAAGGC TATC-AAGCA', 'seq_2': 'GGTTGACAGCGAATACAATACTCGCCGCCAACA-GTGTGAAGCTGCCGCCCGT TTC--T-TTGGC-GTCAAAGCA', 'score': -15, 'identity': 0.4666666666666667, 'matc h': 35}

seq 10 vs seq 8

```
{'seq 1': 'CACTGAATCA-ATCCTTCTT-A-AC-TTTAGGATCACTTATTATGGGGCTAGGATCAATAAAGGC
TATCAAGCA', 'seq_2': '-AGAGACTAATATGTTTTTTCACACTTTTCTGCTTTTTTA-CATCCTG-AAATA
T-ATTATCATTTAAAGGGGT', 'score': -14, 'identity': 0.4594594594595, 'matc
h': 34}
seq_10 vs seq_9
{'seq 1': '-CACTGAA--T-CAATCCTTCTTAACTTTAGGATCACTTATTATGGGGCTA-GGATCAATAAAGG
CTATCAAGCA', 'seq 2': 'TCGCCGAACGTCCAGGCCGTC--GCCCTGCCCAGCGC-CA--CCGCCCGTACC
GCCCTGGAACGCCTGCCCAGCC', 'score': -21, 'identity': 0.426666666666667, 'matc
h': 32}
______
 c) Utilize o score de cada alinhamento obtido na letra "b" e construir uma matriz
 de distância entre as 10 OTUs.
 Formato da Resposta:
 seq_1 seq_2 seq_3 seq_4 seq_5 seq_6 seq_7 seq_8 seq_9 seq_10
 seq 1
 seq 2
 seq 3
 seq 4
 seq 5
 seq 6
 seq 7
 seq 8
 seq 9
 seq 10
```

In [6]: print(distance matrix)

```
[[ 0. -26. -15. -17. -17. -13. -25. -20. -29. -11.]
        0. -19. -27. -23. -24. -15. -20. -10. -23.]
              0. -22. -11. -16. -14. -32. -6. -19.]
 [-15. -19.
                   0. -23. -10. -17. -23. -32. -13.]
 [-17. -27. -22.
 [-17. -23. -11. -23.
                        0. -17.
                                -6. -18.
                             0. -18. -26. -24. -23.]
 [-13. -24. -16. -10. -17.
 [-25. -15. -14. -17. -6. -18.
                                  0. -26. -15. -15.]
 [-20. -20. -32. -23. -18. -26. -26.
                                       0. -34. -14.]
 [-29. -10. -6. -32. -4. -24. -15. -34.
                                            0. -21.]
 [-11. -23. -19. -13. -16. -23. -15. -14. -21.
```

d) Implemente o algoritmo UPGMA (unweighted pair group method with arithmetic mean) para realizar a construção da árvore filogenética das 10 OTUs consideradas na letra "a"

Entrada do algoritmo: matriz de distâncias construída no item "c"

In [3]:

```
print(distance matrix)
         np.savetxt(f'{dir}/dist_matrix.txt', distance_matrix, fmt='%d')
        [[ 0. -26. -15. -17. -17. -13. -25. -20. -29. -11.]
         [-26.
                 0. -19. -27. -23. -24. -15. -20. -10. -23.]
                      0. -22. -11. -16. -14. -32. -6. -19.]
         [-15. -19.
         [-17. -27. -22.
                           0. -23. -10. -17. -23. -32. -13.]
         [-17. -23. -11. -23.
                                0. -17. -6. -18.
                                                   -4. -16.]
                                     0. -18. -26. -24. -23.]
         [-13. -24. -16. -10. -17.
         [-25. -15. -14. -17. -6. -18.
                                           0. -26. -15. -15.]
         [-20. -20. -32. -23. -18. -26. -26.
                                                0. -34. -14.]
         [-29. -10. -6. -32. -4. -24. -15. -34.
                                                     0. -21.1
         [-11. -23. -19. -13. -16. -23. -15. -14. -21.
         Saída: a árvore filogenética
         Os índices das sequências começam em 0.
         seq 1 -> 0,
         seq 2 -> 1,
In [10]:
         import upgma
         upgma.run(distance_matrix)
```

Out[10]: (4, ((6, ((3, (8, 7)), (1, 0))), (2, (9, 5))))

This notebook was converted with convert.ploomber.io