

Part 1:

Before either banded/unrestricted are called, there is a cost of slicing each string to make it the size of the align length, but since that is a constant, it can be dropped from the overall time and space complexity.

For the unrestricted algorithm, the space complexity is simply $O(nm)$ where n is the length of one string and m is the length of the other string. Since each cell of the table stores a tuple with an integer value and a short string dictating which direction that score came from, it does not affect the asymptotic space complexity of the algorithm.

There are two for loops in the unrestricted algorithm, the outer one being $O(n)$ where n is the length of the string or the align length, and the inner loop being $O(m)$ where m is the length of the other string or align length, whichever is shorter.

The backtrace time complexity for unrestricted is $O(n)$, since it simply constructs the alignments from the directions stored in the table. This would make the overall time complexity $O(nm + n)$, but asymptotically just $O(nm)$

The above adds another $O(n + m)$ to the space complexity, but again does not contribute asymptotically.

For the banded algorithm, the space complexity is $O(kn)$, where k is the bandwidth (in this case 7), and n is the length of the shorter string. The backtrace adds another $O(n + m)$ to the space complexity, but again does not contribute asymptotically.

The first for loop in the banded algorithm is $O(n)$ time complexity, where n is the minimum of the shorter string's length or the align length. The inner for loop contributes an effectively constant time, in this case $O(7)$ which is asymptotically $O(1)$, since the bandwidth is 7 and the inner loop only goes as long as the bandwidth.

The backtrace for unbanded then is also $O(n)$, as it traces the path to the start from the end and constructs the alignments while doing so. It is $O(n)$ because it is the length of the string (or the align length).

Part 2:

My algorithm uses the Needleman-Wunsch cost function, and goes through the shorter string character by character, comparing each one to every character of the longer (or horizontally-oriented) string. It formed a DAG because no higher row was dependent on a lower row, and no value was dependent on the value to its right. I stored the value and direction the value came from in tuples in each cell of the table, so that the backtrace could follow the directions in reverse to construct the alignment and arrive at the beginning of the string.

Part 3:

Gene Sequence Alignment

	sequence1	sequence2	sequence3	sequence4	sequence5	sequence6	sequence7	sequence8	sequence9	sequence10
sequence1	-30	-1	4956	4956	4956	4956	4956	4956	4956	4956
sequence2		-33	4948	4948	4948	4948	4948	4948	4948	4948
sequence3			-3000	-2996	-2956	-2944	-1431	-1448	-1399	-1448
sequence4				-3000	-2960	-2948	-1431	-1448	-1399	-1448
sequence5					-3000	-2988	-1423	-1452	-1391	-1448
sequence6						-3000	-1426	-1452	-1394	-1448
sequence7							-3000	-2771	-2814	-2767
sequence8								-3000	-2731	-2996
sequence9									-3000	-2727
sequence10										-3000

Label I:

Sequence I:

Sequence J:

Label J:

☐ Banded Align Length:

Done. Time taken: 39.534 seconds.

Gene Sequence Alignment

	sequence1	sequence2	sequence3	sequence4	sequence5	sequence6	sequence7	sequence8	sequence9	sequence10
sequence1	-30	-1	inf	inf	inf	inf	inf	inf	inf	inf
sequence2		-33	inf	inf	inf	inf	inf	inf	inf	inf
sequence3			-9000	-8984	-8888	-8848	-2735	-2743	-1429	-2735
sequence4				-9000	-8888	-8848	-2739	-2748	-1426	-2740
sequence5					-9000	-8960	-2711	-2739	-1426	-2727
sequence6						-9000	-2708	-2728	-1415	-2716
sequence7							-9000	-8103	-1256	-8099
sequence8								-9000	-1310	-8980
sequence9									-9000	-1315
sequence10										-9000

Label I:

Sequence I:

Sequence J:

Label J:

☒ Banded Align Length:

Done. Time taken: 1.096 seconds.

Part 4:

Unrestricted:

attgcgagcgatttgcgtgcgtgcatcccgcttc-actg--at-
ctcttgtagatcttttcataatctaaactttataaaaacatccactccctgta-g
ataa-gagtgattggcggtccgtacgtaccctttctactctcaaactcttgtagtttaaadc-
taatctaaactttataaa--cggc-acttcctgtgtg

sequences 3 and 10

```
attgcgagcgatttgcgtgcgtgcatcccgcttc-actg--at-ctcttgtagatcttttcataatctaaactttataaaaacatccactccctgta-g
ataa-gagtgattggcggtccgtacgtaccctttctactctcaaactcttgtagtttaaadc-taatctaaactttataaa--cggc-acttcctgtgtg
```

Banded:

gattgcgagcgatttgcgtgcgtgcat-ccc--gcttcact-
gatctcttgtagatcttttcataatctaaactttataaaaacatccactccctgt-a
-a-taagagtgattggcggtccgtacgtaccctttctactctcaaactcttgtagtttaaadc-
taatctaaactttat--aac-ggcacttcctgtgt

sequences 3 and 10 for banded

```
gattgcgagcgatttgcgtgcgtgcat-ccc--gcttcact-gatctcttgtagatcttttcataatctaaactttataaaaacatccactccctgt-a
-a-taagagtgattggcggtccgtacgtaccctttctactctcaaactcttgtagtttaaadc-taatctaaactttat--aac-ggcacttcctgtgt
```

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Part 5:

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for i in range(len(sequences)):
    jresults = []
    for j in range(len(sequences)):
        if j < i:
            s = {}
        else:
            a = "-" + sequences[i][:align_length]
            b = "-" + sequences[j][:align_length]
            if banded:
                if abs(len(a) - len(b)) > MAXINDELS:
                    score = math.inf
                    alignment1 = alignment2 = "No Alignment Possible"
                else:
                    if len(b) < len(a):
                        score, alignment1, alignment2 = self.banded_algorithm(b, a, align_length)
                    else:
                        score, alignment1, alignment2 = self.banded_algorithm(a, b, align_length)
            else:
                score, alignment1, alignment2 = self.unrestricted(a, b, align_length)

            s = {'align_cost':score, 'seqi_first100':alignment1, 'seqj_first100':alignment2}
            table.item(i,j).setText('{}'.format(int(score) if score != math.inf else score))
            table.repaint()
        jresults.append(s)
    results.append(jresults)
return results
```

```

59 | def unrestricted(self, a, b, align_length): # a is vertical, along the side and b is horizontal along the top
60 |     table = []
61 |     a_end = min(len(a), align_length + 1)
62 |     b_end = min(len(b), align_length + 1)
63 |     for i in range(a_end):
64 |         row = []
65 |         for j in range(b_end):
66 |             if i == 0:
67 |                 value = (j * INDEL, "left")
68 |             elif j == 0:
69 |                 value = (i * INDEL, "top")
70 |             else:
71 |                 diagonal = MATCH if a[i] == b[j] else SUB
72 |                 value = self.score_direction(table[i-1][j][0] + INDEL, row[j-1][0] + INDEL, table[i-1][j-1][0] + diagonal)
73 |             row.append(value)
74 |         table.append(row)
75 |     score = table[-1][-1][0]
76 |     i = a_end - 1
77 |     j = b_end - 1
78 |     alignment1 = ""
79 |     alignment2 = ""
80 |     while i != 0 and j != 0:
81 |         if table[i][j][1] == "top":
82 |             alignment1 = a[i] + alignment1
83 |             alignment2 = "-" + alignment2
84 |             i -= 1
85 |         elif table[i][j][1] == "left":
86 |             alignment1 = "-" + alignment1
87 |             alignment2 = b[j] + alignment2
88 |             j -= 1
89 |         else:
90 |             alignment1 = a[i] + alignment1
91 |             alignment2 = b[j] + alignment2
92 |             i -= 1
93 |             j -= 1
94 |
95 |     return score, alignment1, alignment2
96 |
97 | def score_direction(self, top, left, diagonal):
98 |     if top <= left and top <= diagonal:
99 |         return top, "top"
100 |     elif left <= top and left <= diagonal:
101 |         return left, "left"
102 |     else:
103 |         return diagonal, "diagonal"
104 |

```

```

105 def banded_algorithm(self, a, b, align_length):
106     table = []
107     z = (2 * MAXINDELS) + 1
108     for i in range(min(len(a), align_length + 1)):
109         row = [(math.inf, "")] * z
110         row_start = MAXINDELS - i
111         for j in range(max(0, row_start), z):
112             if i == 0:
113                 value = ((j - MAXINDELS) * INDEL, "left")
114             elif j == row_start and i < MAXINDELS + 1:
115                 value = (i * INDEL, "diagonal")
116             else:
117                 if i + j - MAXINDELS >= len(b):
118                     continue
119                 top = MATCH if a[i] == b[i + j - MAXINDELS] else SUB
120                 left = row[j-1][0] + INDEL if j > 0 else math.inf
121                 diagonal = table[i-1][j+1][0] + INDEL if j < (2 * MAXINDELS) else math.inf
122                 value = self.score_direction(table[i-1][j][0] + top, left, diagonal)
123             row[j] = value
124         table.append(row)
125     score_index = (-MAXINDELS-1) + (len(b) - len(a))
126     score = table[-1][score_index][0]
127     alignment1 = ""
128     alignment2 = ""
129     i = min(len(a), align_length - 1) - 1
130     j = z + score_index
131     while i != 0 and j != MAXINDELS:
132         if table[i][j][1] == "top":
133             alignment1 = a[i] + alignment1
134             alignment2 = b[i + j - MAXINDELS] + alignment2
135             i -= 1
136         elif table[i][j][1] == "left":
137             alignment1 = "-" + alignment1
138             alignment2 = b[i + j - MAXINDELS] + alignment2
139             j -= 1
140         else:
141             alignment1 = a[i] + alignment1
142             alignment2 = "-" + alignment2
143             i -= 1
144             j += 1
145
146     return score, alignment1, alignment2
147

```