Analysis – Unrestricted Algorithm:

String size n, string size m

The unrestricted algorithm starts by initializing an array of size n+1 * m+1, (which is O(n+1 * m+1), but as n and m get bigger and bigger the +1 becomes so small in comparison that we can simplify to O(nm)) which takes O(nm) time and O(nm) space. This is the only substantial space that we initialize in the algorithm, as all other parts of our algorithm only take O(1) space when making a single variable. Later, we concatenate the sequences as strings, which takes O(m+n) space, which simplifies to either O(n) or O(m) space depending on which input is longer as our ending strings will be length of the longest string. This is small in comparison to O(nm) space of the 2d array, so we can ignore this.

Apart from initializing the 2d array, we then loop through each element, each taking O(1) time to determine the minimum value from its neighbors, which leaves us with n*m elements each with O(1) time. This leaves us with another O(nm) time chunk, but when added together to get O(2nm), we simplify to O(nm). When concatenating the strings at the end, our time complexity is O(m+n), which simplifies to either O(n) or O(m) as we traverse all the way to the origin of our 2d matrix of size n*m. By comparison, these times are small when compared to O(nm), so we are left with only O(nm) time complexity and O(nm) space complexity.

See code for more analysis on each function's individual time and complexity

Analysis - Banded Algorithm:

String size n, string size m, size k bandwidth

The unrestricted algorithm starts by initializing an array of size k^*n , where n is the longest string which takes O(kn) time and O(kn) space. This is the only substantial space that we initialize in the algorithm, as all other parts of our algorithm only take O(1) space when making a single variable. Later, we concatenate the sequences as strings, which takes O(m+n) space, which simplifies to either O(n) or O(m) space depending on which input is longer as our ending strings will be length of the longest string. This is small in comparison to O(kn) space of the 2d array, so we can ignore this.

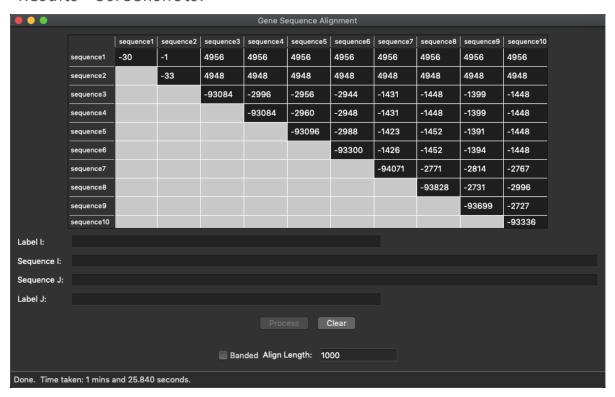
Apart from initializing the 2d array, we then loop through each element, each taking O(1) time to determine the minimum value from its neighbors, which leaves us with k*n elements each with O(1) time. Because we only care about the elements that are within d=3 elements from the diagonal, we only need to iterate through 7 elements on each row of the matrix, so even though we are comparing each element of n and m, we only do this when m is +- 3 from the diagonal, which leaves us iterating through k*n elements. This leaves us with another O(kn) time chunk, but when added together to get O(2kn), we simplify to O(kn). When concatenating the strings at the end, our time complexity is O(m+n), which simplifies to either O(n) or O(m) as we are traversing our 2d matrix of size k*n all the way back to the origin point. By comparison, these times are small when compared to O(kn), so we are left with only O(kn) time complexity and O(kn) space complexity.

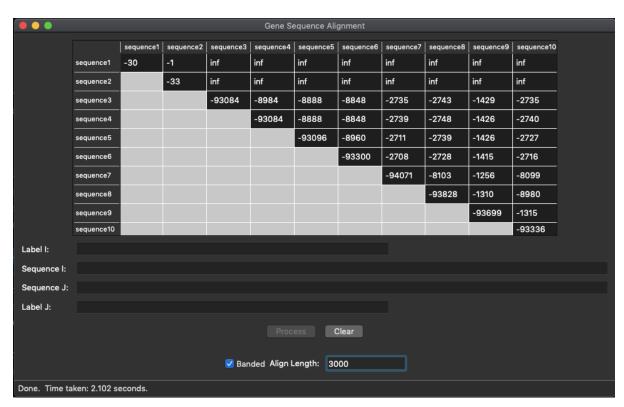
See code for more analysis on each function's individual time and complexity

How it works:

First we make a 2d matrix, size dependent on whether we want it banded or unrestricted (see analysis for difference). Then we loop through each element, and using its neighbors (top, left, and diagonal) we calculate the minimum value of the sequence if we have an indel, a match, or a mismatch. We would then update the current position with the current minimum value from my neighbors, and go to the next element in my sequences. In my matrix, I used a tuple to determine where I was coming from. If I came from a diagonal, my tuple would contain the current value and a value "D" so I knew when I would traverse back from the end of the matrix, I would see the "D" in my tuple and know that the next node I needed to visit was matrix[i-1][j-1]. I would store the same for top ("T") and left ("L"), with the origin point containing a "\$" so my algorithm knew when it was done and had reached the upper left hand corner. We would start from the bottom right hand corner of my cost matrix, and follow the back pointers until we reached the start, each time either adding a character from sequence1, sequence2, or both if a diagonal. If we came from the Left, we would add a – to the beginning of sequence1, and a character to the beginning of sequence2, if from the top, we would add a – to the beginning of sequence2 and a character to the beginning of sequence1. If we come from a diagonal, we add a character to both sequences' beginnings.

Results - Screenshots:





Results - Alignment:

K = 1000

K = 3000

 $\label{thm:condition} gattgcgacgatttgcgtgcgtcat-ccc--gcttcact-gatctcttgttagatcttttcataatctaaactttataaaacatccactccctgt-a\\ -a-taagagtgattggcgtccgtacgtaccctttctactctcaaactcttgttagtttaaatc-taatctaaactttat--aaac-ggcacttcctgtgt$

Commented Code:

Unbanded Algorithm:

```
# Unrestricted Algorithm
# Time: O(nm) time, to fill up the matrix, and to iterate through each element in the 2d matrix
# Space: O(nm) space, as we fill up a 2d matrix and alter those values, but it is the only major object we create

def sequenceal = sequence2:
    return MATCH * len(sequence1), sequence1, sequence2
else:
    sequence1 = sequence2[:align_length]
    sequence2 = sequence2[:align_length]

if(len(sequence2) > len(sequence1)):
    temp = sequence2
    sequence2 = sequence2
    sequence2 = sequence3
    sequence3 = sequence4
    sequence4 = sequence5
    sequence5 = sequence6

matrix[0][0] = (0, '$')

for i in range(len(sequence1) + 1): #O(nm) time

for j in range(len(sequence2) + 1):
    newTuple = self.nextValue(i, j, matrix, sequence2, sequence2)
    if newTuple != 0:
        matrix[i][j] = newTuple

bestfit = matrix[len(sequence3)][len(sequence2)][0]
alignment1, alignment2 = self.getAlignments(matrix, sequence1, sequence2)

return bestFit, alignment1, alignment2
```

```
tuple = self.getMin(diagonalValue, topValue, leftValue); #0(1) time and space
```

```
letterIndex1 = len(sequence1) - 1
letterIndex2 = len(sequence2) - 1
currentAlignment2 = ""
            currentAlignment2 = "-" + currentAlignment2
currentAlignment1 = sequence1[letterIndex1] + currentAlignment1
      elif currentTuple[1] = "L":
    currentAlignment1 = "-" + currentAlignment1
```

Banded Algorithm:

```
1f sequence1 == sequence2:
    return MATCH * len(sequence1), sequence1, sequence2
     sequence1 = sequence1[:align_length]
sequence2 = sequence2[:align_length]
    if abs(len(sequence1) - len(sequence2)) > MAXINDELS:
    return float("inf"), "No Alignment Possible", "No Alignment Possible"
     sequence1 = sequence1[:align_length]
sequence2 = sequence2[:align_length]
          temp = sequence2
sequence2 = sequence1
sequence1 = temp
      for 1 in range(len(sequencel) + 1): # Loops through k*n times, Q(kn) complexity
    if 1 - (len(sequencel) + 1) >= -3:
        pass
    elif 1 > 3:
                 matrix(1)(j) = newTuple
else;
           alignment1, alignment2 = self.getBandwidthAlignments(matrix, sequence1, sequence2, indexOfBestFit) #0(n) or 0(m) time and space return bestFit, alignment1, alignment2
```

```
topValue = float("inf")
validOption = False
topValid = False
shiftedI = 1 - 3
    validOption = True
    topValid = True
    topValue = matrix[i - 1][j + 1 - shiftedI][0]
    leftValid = True
if validOption:
        if sequence1[1 -1] = sequence2[j - 1]:
            diagonalValue = diagonalValue + MATCH
    if topValid:
        topValue = topValue + INDEL
    if leftValid:
        leftValue = leftValue + INDEL
    tuple = self.getMin(diagonalValue, topValue, leftValue); #0(1) time and space
    return tuple
```