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using System;
using System.Collections.Generic;
using System.Text;

namespace GeneticsLab
{
    class PairWiseAlign
    {
        int MaxCharactersToAlign;

        public PairWiseAlign()
        {
            // Default is to align only 5000 characters in each sequence.
            this.MaxCharactersToAlign = 5000;
        }

        public PairWiseAlign(int len)
        {
            // Alternatively, we can use an different length; typically used with the
            // banded option checked.
            this.MaxCharactersToAlign = len;
        }

        /// <summary>
        /// this is the function you implement.
        /// </summary>
        /// <param name="sequenceA">the first sequence</param>
        /// <param name="sequenceB">the second sequence, may have length not equal to
        /// the length of the first seq.</param>
        /// <param name="banded">true if alignment should be band limited.</param>
        /// <returns>the alignment score and the alignment (in a Result object) for
        /// sequenceA and sequenceB. The calling function places the result in the
        /// display appropriately.
        ///
        public ResultTable.Result Align_And_Extract(GeneSequence sequenceA,
            GeneSequence sequenceB, bool banded)
        {
            //
            // -----
            // Setup section. Also returns without calculation for banded analysis of
            // sequences which will be impossible
            // O(1)

            int maxlength = MaxCharactersToAlign;

            const int indel = 5;
            const int sub = 1;
            const int match = -3;

            //Console.WriteLine("Sequence a: " + sequenceA.Sequence);
            //Console.WriteLine("Sequence b: " + sequenceB.Sequence);
        }
    }
}

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int lengthA, lengthB;
if (sequenceA.Sequence.Length > maxlength) {
    lengthA = maxlength;
} else {
    lengthA = sequenceA.Sequence.Length;
}

if (sequenceB.Sequence.Length > maxlength)
{
    lengthB = maxlength;
}
else
{
    lengthB = sequenceB.Sequence.Length;
}

ResultTable.Result result = new ResultTable.Result();
int score; // place ↗
    your computed alignment score here
string[] alignment = new string[2]; // place ↗
    your two computed alignments here
// these "alignments" are just the two strings, with "-" added where an ↗
    insertion/deletion has occurred. simple enough.

// ***** these are placeholder assignments that you'll replace with ↗
    your code *****
score = 0;
alignment[0] = "";
alignment[1] = "";
//
    ***** ↗
    *****

// We will not be able to get a banded result if the lengths differ by ↗
    more than 3
if (banded && Math.Abs(lengthA - lengthB) > 3) {
    score = int.MaxValue;
    result.Update(score, "No Alignment Possible", "No Alignment ↗
        Possible");
    return result;
}

// Sequence length because it has to fit the string AND a space for the ↗
    "empty string" at the beginning
node[,] calcTable = new node[lengthA+1 , lengthB+1];

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// Go through the entire table to calculate the things.
int a, b = 0;

// do top left corner first
calcTable[0, 0] = new node(-1, -1, 0);

// END setup section
//
-----

if (banded)
{
    //-----
    // Banded table scores calculation.
    // It goes through the length of A, each time doing 7 calculations for B.
    //  $O(7n)$  - where n is the length of A
    // Also note that the length difference between A and B is MAX 4, so  $O(n)$ 

    // Do entire top row
    for (a = 1; a < 4; a++)
    {
        calcTable[a, 0] = new node(a - 1, 0, a * indel);
    }

    // And entire left row
    for (b = 1; b < 4; b++)
    {
        calcTable[0, b] = new node(0, b - 1, b * indel);
    }

    // and the rest
    for (a = 1; a < lengthA + 1; a++)
    {
        for (b = a - 3; b < a + 4; b++)
        { // Can only calculate in the band
            if (b < 1 || b > lengthB) {
                // Can't be having those System.IndexOutOfRangeException,
                can we?
                continue;
            }

            int topCost, leftCost, diagCost;

            // Calculate cost of coming from top
            if (calcTable[a, b - 1] == null)
            {

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        topCost = int.MaxValue;           // null is very bad.
    }
    else
    {
        topCost = calcTable[a, b - 1].score + indel;           // coming from top is an insert/delete
    }

    // Calculate cost of coming from left
    if (calcTable[a - 1, b] == null)
    {
        leftCost = int.MaxValue;           // again, null is very bad
    }
    else
    {
        leftCost = calcTable[a - 1, b].score + indel;           // coming from left is also an insert/delete
    }

    // Calculate cost of coming from the diagonal
    // We don't worry about nulls here, because they are impossible
    if (sequenceA.Sequence[a - 1] == sequenceB.Sequence[b - 1])
    {
        // If the two strings match at this character
        diagCost = calcTable[a - 1, b - 1].score + match;           // coming from diagonal on a match!
    }
    else
    {
        diagCost = calcTable[a - 1, b - 1].score + sub;           // coming from diagonal on a substitution
    }

    // Now to make our table entry
    if (diagCost <= leftCost && diagCost <= topCost)
    {
        // Diagonal is cheapest
        calcTable[a, b] = new node(a - 1, b - 1, diagCost);
    }
    else if (leftCost <= diagCost && leftCost <= topCost)
    {
        // Left is cheapest
        calcTable[a, b] = new node(a - 1, b, leftCost);
    }
    else
    {
        // Top is cheapest
        calcTable[a, b] = new node(a, b - 1, topCost);
    }
}
}

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// END Banded table scores calculation
//----->
-----
} else { // ends our if section for banded; starts our section to calculate unbanded

//
----->
-----
// Start unbanded table scores calculation
// Goes through the length of A, each time going through the length of B.
// Therefore  $O(n*m)$ , where n is the length of A, and m is the length of B.

// Do the entire top row first
for (a = 1; a < lengthA + 1; a++)
{
    calcTable[a, 0] = new node(a - 1, 0, a * indel);
}
// and the entire left row
for (b = 1; b < lengthB + 1; b++)
{
    // Skipping the first one which was already done
    calcTable[0, b] = new node(0, b - 1, b * indel);
}

// And the rest
for (a = 1; a < lengthA + 1; a++)
{
    for (b = 1; b < lengthB + 1; b++)
    {
        // Calculate cost of coming from top
        int topCost = calcTable[a, b - 1].score + indel; //
        coming from top is an insert/delete

        // Calculate cost of coming from left
        int leftCost = calcTable[a - 1, b].score + indel; //
        coming from left is also an insert/delete

        // Calculate cost of coming from the diagonal
        int diagCost;
        if (sequenceA.Sequence[a - 1] == sequenceB.Sequence[b - 1])
        {
            // If the two strings match at this character
            diagCost = calcTable[a - 1, b - 1].score + match; //
            coming from diagonal on a match!
        }
        else
        {
            diagCost = calcTable[a - 1, b - 1].score + sub; //

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        coming from diagonal on a substitution
    }

    // Now to make our table entry
    if (diagCost <= leftCost && diagCost <= topCost)
    {
        // Diagonal is cheapest
        calcTable[a, b] = new node(a - 1, b - 1, diagCost);
    }
    else if (leftCost <= diagCost && leftCost <= topCost)
    {
        // Left is cheapest
        calcTable[a, b] = new node(a - 1, b, leftCost);
    }
    else
    {
        // Top is
        cheapest
        calcTable[a, b] = new node(a, b - 1, topCost);
    }
}
//END unbanded calculation
//-----
} // This ends the difference between banded and unbanded calculation

//
//-----

// At this point our scores table should be complete. Now we just take the
// final node and walk back to the beginning with it
// The length of this string is the larger of the length of A and the
// length of B
// O(max(n,m)) where n is A's length and m is B's length
a = lengthA;
b = lengthB;

StringBuilder strA = new StringBuilder(maxlength);
StringBuilder strB = new StringBuilder(maxlength);

// Go until we hit either the top or left row/column
while (a != 0 && b != 0) {
    int parent_a = calcTable[a, b].parent_x;
    int parent_b = calcTable[a, b].parent_y;
    if (parent_a < a) {
        // To get here, we came from left
        //alignment[0].Insert(0,sequenceA.Sequence[a-1].ToString
        ());
        // Which means we used a char of
        sequence a
        strA.Insert(0, sequenceA.Sequence[a - 1].ToString());
    }
    if (parent_b < b) {
        // Also came from top = came from

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        diagonal
        //alignment[1].Insert(0, sequenceB.Sequence[b-1].ToString
        ()); // Which means we ALSO used a char of
        sequence b
        strB.Insert(0, sequenceB.Sequence[b - 1].ToString());
    } else { // Only came from
        left
        //alignment[1].Insert(0,
        "-"); // Used a
        char of sequence a but not b
        strB.Insert(0, "-");
    }
} else {
    //alignment[0].Insert(0, "-"); // Did not come
    from left nor diagonal; must have been from top
    strA.Insert(0, "-");
    //alignment[1].Insert(0, sequenceB.Sequence[b-1].ToString());
    strB.Insert(0, sequenceB.Sequence[b - 1].ToString());
}
a = parent_a;
b = parent_b;
}

// Assume we hit the left column. This means a = 0 and b is getting
smaller
// This means we have used all of sequence a already.
while (b != 0) {
    int parent_a = 0;
    int parent_b = calcTable[a, b].parent_y;

    //alignment[0].Insert(0, "-"); // Already used all of a, so it has
    gaps at the beginning
    strA.Insert(0, "-");
    //alignment[1].Insert(0, sequenceB.Sequence[b-1].ToString());
    strB.Insert(0, sequenceB.Sequence[b - 1].ToString());

    a = parent_a;
    b = parent_b;
}

// Assume we hit the top row. This means b = 0 and a is getting smaller
// This means we have used all of sequence b already.
while (a != 0)
{
    int parent_a = calcTable[a, b].parent_x;
    int parent_b = 0;

    //alignment[0] = alignment[0].Insert(0, sequenceA.Sequence[a -
    1].ToString()); // Already used all of a, so it has gaps at the
    beginning
    strA.Insert(0, sequenceA.Sequence[a - 1].ToString());
    //alignment[1] = alignment[1].Insert(0, "-");

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        strB.Insert(0, "-");

        a = parent_a;
        b = parent_b;
    }

    // END final string calculation
    //----->

    //----->

    // From here on out, we are just wrapping up the calculations and
    // returning the results
    // O(1)

    // If we reach here, we should have traced our strings back to the
    // beginning. The alignment strings should be all good and we just need to
    // get the score
    score = calcTable[lengthA, lengthB].score;
    result.Update(score, strA.ToString(), strB.ToString()); //alignment
    [0],alignment[1]); // bundling your results into the
    right object type
    return(result);
}
}

class node {
    public int parent_x;
    public int parent_y;
    public int score;

    public node(int x, int y, int s)
    {
        parent_x = x;
        parent_y = y;
        score = s;
    }
}
}

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