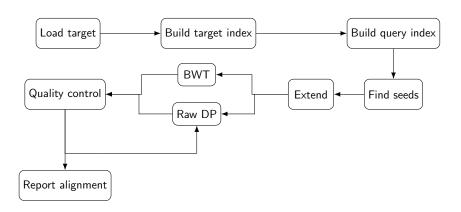
Fast Python sequence aligner

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2024

Seed and Extend approach



Seed extension

Algorithm Standard LIS construction O(n log n)

```
Require: n > 0
 lis len \leftarrow 0
                                                                                             Delight of LIS
 parent \leftarrow \{\infty, \infty, \infty, ..., \infty\}_{n+1}
                                                                          ⊳Mapping to reconstruct LIS
 sub \leftarrow \{\infty, \infty, \infty, ..., \infty\}_{n+1}
                                                   Deliver Array with indices for matches that form LIS
 i \leftarrow 0
 while i < n do
                                                      \trianglerightIterate over all elements i = 0, 1, 2..., n-1
      start \leftarrow 1
     end ← lis_len
      while start < end do
                                                    ⊳Binary search over existing longest sequence
           middle \leftarrow \left| \frac{start + end}{2} \right|
          if matches_{a}[sub[middle]] < matches_{a}[i] then
                start \leftarrow middle + 1
           else
               start \leftarrow middle - 1
      parent[i] \leftarrow sub[start - 1]
                                                        >We pin current value to the found parent
      sub[start] \leftarrow i
      if start > lis len then
           lis_len = start
      i \leftarrow i + 1
```

Seed extension

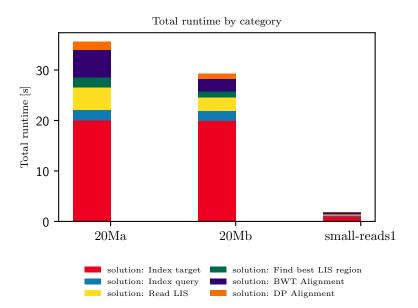
Algorithm Reconstruct LIS by following parent array O(n)

Seed extension

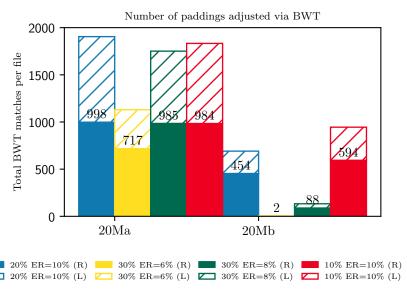
Algorithm Segmented-LIS heuristic O(n log n)

```
Require: n \ge 0
 lis len ← 0
                                                                                                                            DLength of LIS
parent \leftarrow \{\infty, \infty, \infty, ..., \infty\}_{n+1}
                                                                                                           Description Number > Mapping to reconstruct LIS
 sub \leftarrow \{\infty, \infty, \infty, \dots, \infty\}_{n+1}
                                                                                     DArray with indices for matches that form LIS
i \leftarrow 0
while i < n do
                                                                                     \trianglerightIterate over all elements i = 0, 1, 2..., n - 1
     start \leftarrow 1
     end ← lis len
     while start < end do
                                                                                                                        ⊳Binarv search-like
          middle \leftarrow \frac{start + end}{2}
          if matches_{T}[sub[middle]] > matches_{T}[i] - max\_diff then
                                                                                                                  ⊳Encountered old entry
              end \leftarrow start -1
                                                                                                                               ⊳Breaks loop
          else if matches_{\mathcal{O}}[sub[middle]] < matches_{\mathcal{O}}[i] then
               start \leftarrow middle + 1
          else
               start \leftarrow middle - 1
     parent[i] \leftarrow sub[start - 1]
                                                                                         >We pin current value to the found parent
     sub[start] \leftarrow i
     if start > lis_len then
          lis_len = start
     i \leftarrow i + 1
```

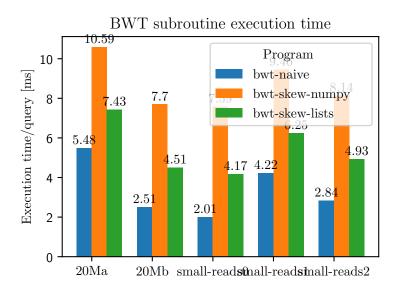
Execution times



Aligner routine effectiveness



BWT routine implementation



Raw DP routine implementation

