```
Hongqi Guo
  LCS (1,1)
a < length ix]
  b < length [y]
  for x to to a do
     for y to to b do.
        if ( x=0 or y=0)
           longth LCS [i,j] to
         else if v [i-1] = b [j-1]
           Length LCS [i-1,j-1]+1.
       else
       length LGS [isj] = mox(length LCS[i-1,j], longeth LCS[i,j-1])
      pointer & length LCS[a,b]
       ito
       i tb.
       while iso and iso.
      do if a [i-1] = b [j-1]
         seq[pointer-1] = x[i-1]
```

```
しゃしー
    j t j-1.
    pointer & pointer -1
  else if length LSS [i-1.j] > langth LCS [i,j-1]
     えせらり
 the jtj-1
  そ と''
for uto to logol [su] -3 ch
    if seq [a] = x and seq [a+1] = y and con [a+2] = y
        w-)ut2
       ,
Z = Z+ (4 [u]
   retan Z.
  Space complexity = O(r)
  Time complexity = 0 (n.n)
```

2. We go note all the subsequences of a sequence, then we compare each subsequences with other sequences to check if they are common and then find the subsequences of minimum length.

Then we have a stray of length n. subsequences of length

1,2,3--n, so ue can use promutation and combination.

5ub-sougenes: nC, + nCz+nCr.

So a stry of length n has 2^n-1 differed possible subsequences since we don't consider the subsequence with layth 0.

to check if a subsequence is common to beh the early or met.

then O(n) thre.

to to tal the needed to check all the sub-sequences = $n\cdot 2^n + 1$.

The complexity be brake force method is $O(n\cdot 2^n - 1)$