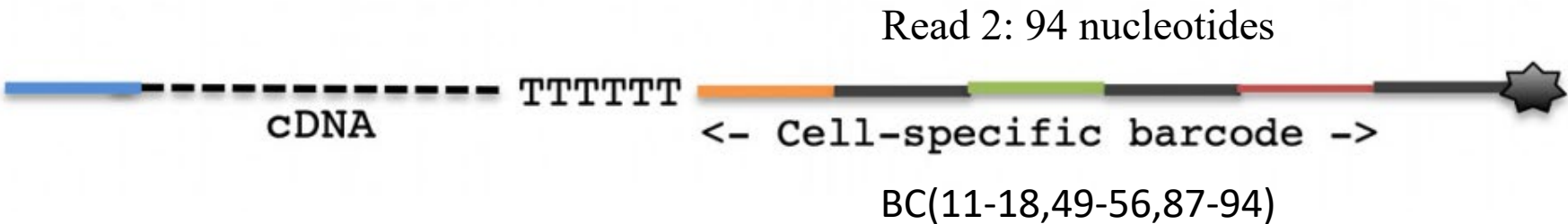


SPLiT-seq barcodes

Three rounds of barcoding (96 barcodes used for each round):



442,368 x AGTGGTCAAGTGGTCAAAACATCG



AGTGGTCAAGTGGTCAGGGGGGGGG
GGGGGGGGGGGGGGGGGGGGGGGGGGGG

- oligo(dT)
- Random hexamers

Reads the lists of barcodes used in each round from .txt files



```
list1 = barcodes1.read().split("\n")  
list2 = barcodes2.read().split("\n")  
list3 = barcodes3.read().split("\n")
```



```
all_list = [list1, list2, list3]
```



Combinatoric iterators: `itertools.product(*all_list)`



Output the final barcode sequences to .txt file

Global Pairwise sequence alignment

```
SeqIO.parse("seq1.fasta", "fasta"))
SeqIO.parse("seq2.fasta", "fasta"))
```

	A	C	G	T
A	0	5	2	5
C	5	0	5	2
G	2	5	0	5
T	5	2	5	0

Gap cost: 5

```
for i in range(1, m+1):
    for j in range(1, n+1):
        Table[i][j] = min(
            Table[i][j - 1] + gap,
            Table[i - 1][j] + gap,
            Table[i - 1][j - 1] +
            cost[tags[str2[i - 1]], tags[str1[j - 1]]]
        )
```



Alignment graph:

		A	A	T	A	A	T	
	[0	5	10	15	20	25	30]
A	[5	0	5	10	15	20	25]
A	[10	5	0	5	10	15	20]
G	[15	10	5	5	7	12	17]
G	[20	15	10	10	7	9	14]
C	[25	20	15	12	12	12	11]
A	[30	25	20	17	12	12	16]

Optimal cost:
16

Optimal alignment:
AATAAT-
AA-GGCA

Global Pairwise sequence alignment

```
SeqIO.parse("seq1.fasta", "fasta")
SeqIO.parse("seq2.fasta", "fasta")
```

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Alignment graph:

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	[0	5	10	15	20	25	30]
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A	[10	5	0	5	10	15	20]
G	[15	10	5	5	7	12	17]
G	[20	15	10	10	7	9	14]
C	[25	20	15	12	12	12	11]
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Optimal alignment:
AATAAT-
AA-GGCA