

# CSE 4065 - Introduction to Computational Genomics

## Project 1 - Report

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WUHAN - Hamming Distance 0

```
WUHAN:          Hamming Distance:0
-----
5-mer
-----
k-mer:   aacaa
rev comp: ttggt
freq:    201

6-mer
-----
k-mer:   taacaa
rev comp: ttgtta
freq:    70

7-mer
-----
k-mer:   ttgttaa
rev comp: ttaacaa
freq:    28

8-mer
-----
k-mer:   aaattggt
rev comp: aacaattt
freq:    12

9-mer
-----
k-mer:   taaacgaac
rev comp: gttcgttta
freq:    7
```

```
10-mer
-----
k-mer:   ctaaacgaac
rev comp: gttcgtttag
freq:    5

11-mer
-----
k-mer:   tctaaacgaac
rev comp: gttcgtttaga
freq:    3

12-mer
-----
k-mer:   gttgatgggtgtt
rev comp: aacaccatcaac
freq:    3

13-mer
-----
k-mer:   ggctactaacaat
rev comp: attgtagtagcc
freq:    2

14-mer
-----
k-mer:   caacctgaagaaga
rev comp: tcttctcaggttg
freq:    2

15-mer
-----
k-mer:   atcagacaactacta
rev comp: tagtagttgtctgat
freq:    2
```

```
16-mer
-----
k-mer:   atcagacaactactat
rev comp: atagtagttgtctgat
freq:    2

17-mer
-----
k-mer:   atcagacaactactatt
rev comp: aatagtagttgtctgat
freq:    2

18-mer
-----
k-mer:   acactggtaattaccagt
rev comp: actggtaattaccagtg
freq:    2

19-mer
-----
k-mer:   acactggtaattaccagtg
rev comp: cactggtaattaccagtg
freq:    2

20-mer
-----
k-mer:   acactggtaattaccagtg
rev comp: acactggtaattaccagtg
freq:    2
```

WUHAN - Hamming Distance 1

```
WUHAN:          Hamming Distance:1
-----
5-mer
-----
k-mer:   aaaaa
rev comp: ttttt
freq:    2315

6-mer
-----
k-mer:   aaaaaa
rev comp: tttttt
freq:    761

7-mer
-----
k-mer:   tgttggt
rev comp: aacaaca
freq:    290

8-mer
-----
k-mer:   tgttggtta
rev comp: taacaaca
freq:    113

9-mer
-----
k-mer:   tgttggttaa
rev comp: ttaacaaca
freq:    47

10-mer
-----
k-mer:   tgatgttggt
rev comp: aacaacatca
freq:    25
```

```
11-mer
-----
k-mer:   aatttaggtga
rev comp: tcacctaaatt
freq:    9

12-mer
-----
k-mer:   tatgaagatttt
rev comp: aaaaatttcata
freq:    5

13-mer
-----
k-mer:   ttatgaagatttt
rev comp: aaaaatttcataa
freq:    4

14-mer
-----
k-mer:   ttgtacagaaagt
rev comp: cactttctgtacaa
freq:    3

15-mer
-----
k-mer:   gtaacaaaccaacca
rev comp: tgggttggtttgttac
freq:    2

16-mer
-----
k-mer:   ctactagtgaagctgt
rev comp: acagcttcactagtag
freq:    2
```

```
17-mer
-----
k-mer:   ctactagtgaagctgtt
rev comp: aacagcttcactagtag
freq:    2

18-mer
-----
k-mer:   ctactagtgaagctgttg
rev comp: caacagcttcactagtag
freq:    2

19-mer
-----
k-mer:   tacactggtaattaccagt
rev comp: actggtaattaccagtgta
freq:    2

20-mer
-----
k-mer:   tacactggtaattaccagtg
rev comp: cactggtaattaccagtgta
freq:    2

21-mer
-----
k-mer:   tacactggtaattaccagtg
rev comp: acactggtaattaccagtgta
freq:    2
```

## WUHAN - Hamming Distance 2

```
WUHAN:      Hamming Distance:2
-----
5-mer
-----
k-mer:      aaaaaa
rev comp:   ttttt
freq:       11057

6-mer
-----
k-mer:      aaaaaa
rev comp:   ttttt
freq:       4903

7-mer
-----
k-mer:      taaaaaa
rev comp:   ttttta
freq:       1973

8-mer
-----
k-mer:      tttaaaat
rev comp:   attttaa
freq:       777

9-mer
-----
k-mer:      aattttaaa
rev comp:   tttaaaatt
freq:       312

10-mer
-----
k-mer:      aacttttaaa
rev comp:   ttttaagtt
freq:       126
```

```
11-mer
-----
k-mer:      taatgggtgta
rev comp:   taacaccatta
freq:       50

12-mer
-----
k-mer:      ttaatgggtgta
rev comp:   taacaccattaa
freq:       27

13-mer
-----
k-mer:      tgттаatgggtgtt
rev comp:   aacaccattaaca
freq:       15

14-mer
-----
k-mer:      ttgттаatgggtgtt
rev comp:   aacaccattaacaa
freq:       9

15-mer
-----
k-mer:      tgggtgtttattctgt
rev comp:   acagaataaacacca
freq:       6

16-mer
-----
k-mer:      aaggcattttgatgaa
rev comp:   ttcacataaaatgcctt
freq:       4
```

```
17-mer
-----
k-mer:      tgttgggtgattattttg
rev comp:   caaaataatcaccaaca
freq:       3

18-mer
-----
k-mer:      aaagtcaacatcaatatt
rev comp:   aatattgatgttgacttt
freq:       2

19-mer
-----
k-mer:      aaagtcaacatcaatattg
rev comp:   caatattgatgttgacttt
freq:       2

20-mer
-----
k-mer:      gctcttcaacctgaagaaga
rev comp:   tcttcttcagggtgaagagc
freq:       2

21-mer
-----
k-mer:      tagtgagtacactggtaatta
rev comp:   taattaccagtgtactcacta
freq:       2

22-mer
-----
k-mer:      tagtgagtacactggtaattac
rev comp:   gtaattaccagtgtactcacta
freq:       2
```

```
23-mer
-----
k-mer:      tagtgagtacactggtaattacc
rev comp:   ggtaattaccagtgtactcacta
freq:       2

24-mer
-----
k-mer:      tagtgagtacactggtaattacca
rev comp:   tggttaattaccagtgtactcacta
freq:       2

25-mer
-----
k-mer:      tagtgagtacactggtaattaccag
rev comp:   ctggtaattaccagtgtactcacta
freq:       2

26-mer
-----
k-mer:      tagtgagtacactggtaattaccagt
rev comp:   actggtaattaccagtgtactcacta
freq:       2

27-mer
-----
k-mer:      tagtgagtacactggtaattaccagtg
rev comp:   cactggtaattaccagtgtactcacta
freq:       2

28-mer
-----
k-mer:      tagtgagtacactggtaattaccagtgt
rev comp:   acactggtaattaccagtgtactcacta
freq:       2
```

## NEPAL - Hamming Distance 0

```
NEPAL:      Hamming Distance:0
-----
5-mer
-----
k-mer:  aacaa
rev comp: ttgtt
freq:    201

6-mer
-----
k-mer:  taacaa
rev comp: ttgtta
freq:    70

7-mer
-----
k-mer:  ttgttaa
rev comp: ttaacaa
freq:    28

8-mer
-----
k-mer:  aaattggt
rev comp: aacaattt
freq:    12

9-mer
-----
k-mer:  taacgaac
rev comp: gttcgttta
freq:    7

10-mer
-----
k-mer:  ctaacgaac
rev comp: gttcgtttag
freq:    5
```

```
11-mer
-----
k-mer:  tctaaacgaac
rev comp: gttcgtttaga
freq:    3

12-mer
-----
k-mer:  gttgatggtgtt
rev comp: aacacatcaac
freq:    3

13-mer
-----
k-mer:  ggctactaacaat
rev comp: attgttagtagcc
freq:    2

14-mer
-----
k-mer:  caacctgaagaaga
rev comp: tctcttcaggittg
freq:    2

15-mer
-----
k-mer:  atcagacaactacta
rev comp: tagtagttgtctgat
freq:    2

16-mer
-----
k-mer:  atcagacaactactat
rev comp: atagtagttgtctgat
freq:    2
```

```
17-mer
-----
k-mer:  atcagacaactactatt
rev comp: aatagtagttgtctgat
freq:    2

18-mer
-----
k-mer:  acactggtaattaccagt
rev comp: actggtaattaccagtgt
freq:    2

19-mer
-----
k-mer:  acactggtaattaccagtg
rev comp: cactggtaattaccagigt
freq:    2

20-mer
-----
k-mer:  acactggtaattaccagtgt
rev comp: acactggtaattaccagtgt
freq:    2
```

## NEPAL - Hamming Distance 1

```
NEPAL:      Hamming Distance:1
-----
5-mer
-----
k-mer:  aaaaa
rev comp: ttttt
freq:    2315

6-mer
-----
k-mer:  aaaaaa
rev comp: tttttt
freq:    761

7-mer
-----
k-mer:  tgttgtt
rev comp: aacaaca
freq:    290

8-mer
-----
k-mer:  tgttgta
rev comp: taacaaca
freq:    113

9-mer
-----
k-mer:  tgttgtaaa
rev comp: ttaacaaca
freq:    47

10-mer
-----
k-mer:  tgatgttgtt
rev comp: aacaacatca
freq:    25
```

```
11-mer
-----
k-mer:  aatttaggtga
rev comp: tcacctaatt
freq:    9

12-mer
-----
k-mer:  tatgaagatttt
rev comp: aaaaatcttcata
freq:    5

13-mer
-----
k-mer:  ttatgaagatttt
rev comp: aaaaatcttcataa
freq:    4

14-mer
-----
k-mer:  ttgtacagaaagt
rev comp: cactttctgtacaa
freq:    3

15-mer
-----
k-mer:  gtaacaaccaaacca
rev comp: tgggtgggtttgttac
freq:    2

16-mer
-----
k-mer:  ctactagtgaagctgt
rev comp: acagcttcactagtag
freq:    2
```

```
17-mer
-----
k-mer:  ctactagtgaagctgtt
rev comp: aacagcttcactagtag
freq:    2

18-mer
-----
k-mer:  ctactagtgaagctgttg
rev comp: caacagcttcactagtag
freq:    2

19-mer
-----
k-mer:  tacactggtaattaccagt
rev comp: actggtaattaccagtgtg
freq:    2

20-mer
-----
k-mer:  tacactggtaattaccagtg
rev comp: cactggtaattaccagtgta
freq:    2

21-mer
-----
k-mer:  tacactggtaattaccagtgt
rev comp: acactggtaattaccagtgtg
freq:    2
```

## NEPAL - Hamming Distance 2

```
NEPAL:      Hamming Distance:2
-----
5-mer
-----
k-mer:      aaaaa
rev comp:   ttttt
freq:       11057

6-mer
-----
k-mer:      aaaaaa
rev comp:   tttttt
freq:       4903

7-mer
-----
k-mer:      taaaaaa
rev comp:   tttttt
freq:       1973

8-mer
-----
k-mer:      tttaaat
rev comp:   attttaa
freq:       777

9-mer
-----
k-mer:      aattttaaa
rev comp:   tttaaaatt
freq:       312

10-mer
-----
k-mer:      aactttaaaa
rev comp:   ttttaaagtt
freq:       126
```

```
11-mer
-----
k-mer:      taatggtgta
rev comp:   taacaccatta
freq:       50

12-mer
-----
k-mer:      ttaatggtgta
rev comp:   taacaccattaa
freq:       27

13-mer
-----
k-mer:      tgtaatggtgtt
rev comp:   aacaccattaaca
freq:       15

14-mer
-----
k-mer:      ttgtaatggtgtt
rev comp:   aacaccattaacaa
freq:       9

15-mer
-----
k-mer:      tgggtgtttattctgt
rev comp:   acagaataaacacca
freq:       6

16-mer
-----
k-mer:      aaggcattttgaatgaa
rev comp:   ttcacaaaatgcctt
freq:       4
```

```
17-mer
-----
k-mer:      tgttggtgattattttg
rev comp:   caaataatcaccaaca
freq:       3

18-mer
-----
k-mer:      aaagtcaacatcaatatt
rev comp:   aatattgatgttgacttt
freq:       2

19-mer
-----
k-mer:      aaagtaacatcaatattg
rev comp:   caatattgatgttgacttt
freq:       2

20-mer
-----
k-mer:      gctcttcaacctgaagaaga
rev comp:   tcttctcagggtgaagagc
freq:       2

21-mer
-----
k-mer:      tagtgagtacactggtaatta
rev comp:   taattaccagtgactcacta
freq:       2

22-mer
-----
k-mer:      tagtgagtacactggtaattac
rev comp:   gtaattaccagtgactcacta
freq:       2
```

```
23-mer
-----
k-mer:      tagtgagtacactggtaattacc
rev comp:   ggtaattaccagtgactcacta
freq:       2

24-mer
-----
k-mer:      tagtgagtacactggtaattacca
rev comp:   tggtaattaccagtgactcacta
freq:       2

25-mer
-----
k-mer:      tagtgagtacactggtaattaccag
rev comp:   ctggtaattaccagtgactcacta
freq:       2

26-mer
-----
k-mer:      tagtgagtacactggtaattaccagt
rev comp:   actggtaattaccagtgactcacta
freq:       2

27-mer
-----
k-mer:      tagtgagtacactggtaattaccagtg
rev comp:   cactggtaattaccagtgactcacta
freq:       2

28-mer
-----
k-mer:      tagtgagtacactggtaattaccagtg
rev comp:   acactggtaattaccagtgactcacta
freq:       2
```

- The algorithm finds the most frequent k-mer from k is 5 to no frequent k-mer. We run the algorithm for both Nepal and Wuhan with hamming distance 0, 1 and 2

For Wuhan

- It finds with hamming distance 0 until 20-mer
- It finds with hamming distance 1 until 21-mer
- It finds with hamming distance 2 until 28-mer

For Nepal

- It finds with hamming distance 0 until 20-mer
- It finds with hamming distance 1 until 21-mer
- It finds with hamming distance 2 until 28-mer

- The most frequent k-mer changes with hamming distance change. Some of them may not change but mostly it changes.
- Also, the number of occurrence of the most frequent k-mer increases with hamming distance increases.
- Wuhan and Nepal, the algorithm find exactly same the most frequent k-mer and number of occurrence for both of them with same hamming distance. So, we can say that there are not mutations in the virus.