

Question 1

(i)
SARS-CoV2 is similar to MERS-COV rather than SARS-CoV1. This can be observed from the plots below.

(ii)
It is easier to identify the similarity using protein sequences rather than DNA sequences. DNA only consists of 4 components, so the chance of accidentally being similar is much higher. On the other hand, there are 20 amino acids. Thus, only similar proteins will have similar protein sequences.

(iii)

Dottup Parameters

- Word Size (k-tuple) = 10

Dotmatcher Parameters

- Window Size = 15
- Threshold Value = 50

Plots

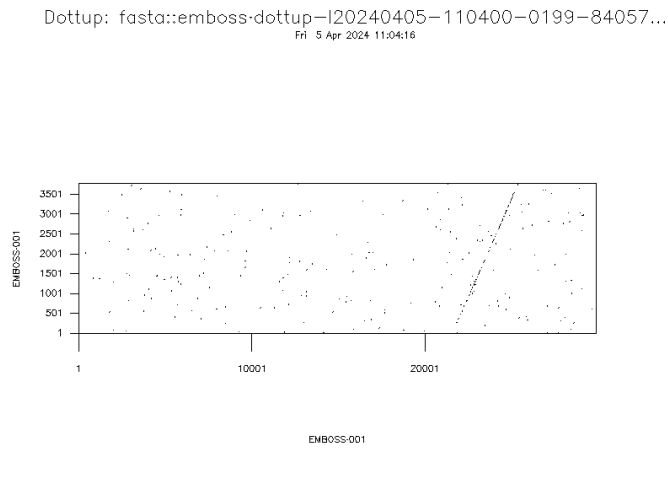


Figure 1: Dottup SARSCoV-SARSCoV2 DNA

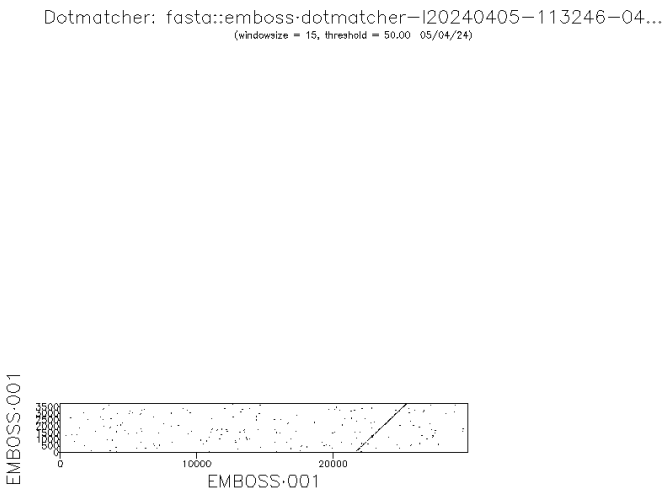


Figure 2: Dotmatcher SARSCoV-SARSCoV2 DNA

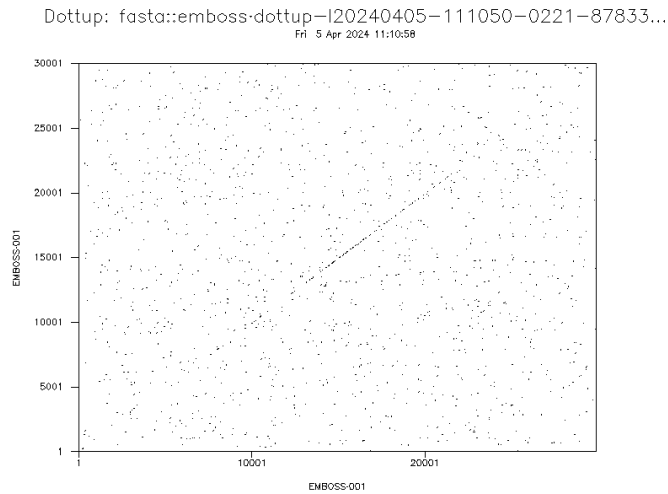


Figure 3: Dottup MERSCoV-SARSCoV2 DNA

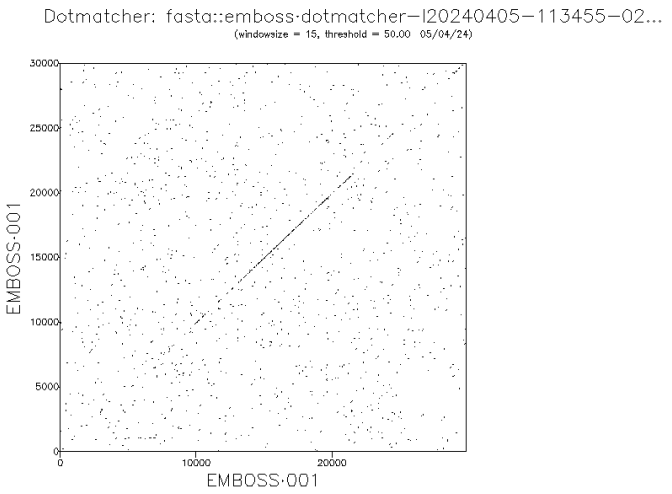


Figure 4: Dotmatcher MERSCoV-SARSCoV2 DNA

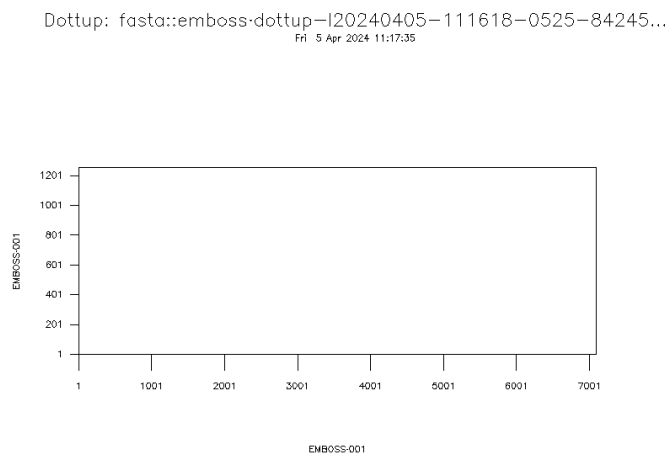


Figure 5: Dottup SARSCoV-SARSCoV2 Protein

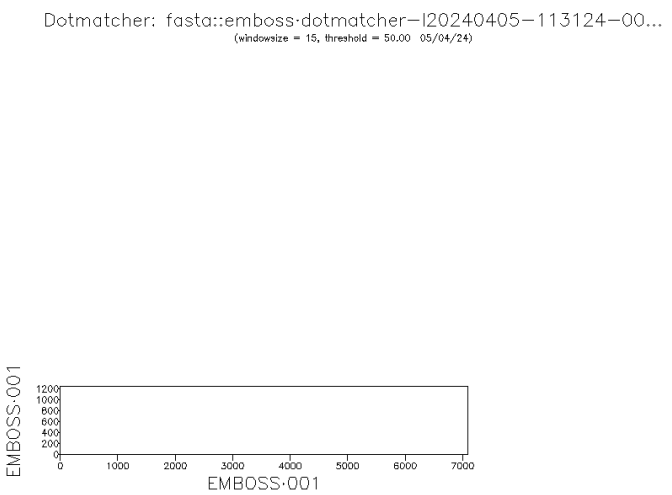


Figure 6: Dotmatcher SARSCoV-SARSCoV2 Protein

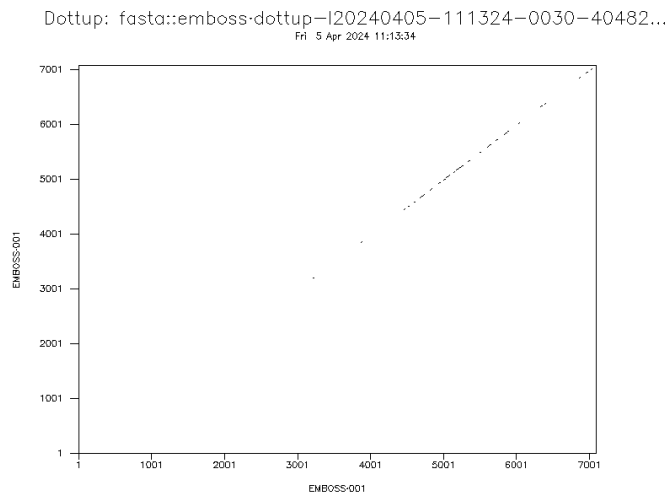


Figure 7: Dottup MERSCoV-SARSCoV2 Protein

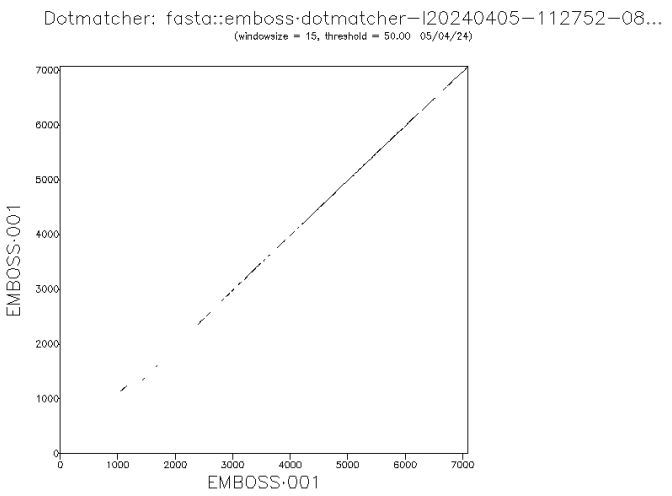


Figure 8: Dotmatcher MERSCoV-SARSCoV2 Protein

Question 2

Question 3

Question 4

Question 5