

The average Silhouette Score is: 0.7804818134155452

Number of clusters: 7

Cluster 0: 2474 sequences

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Cluster 1: 2474 sequences

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CDC-QDX33680550/2022', 'USA/WA-CDC-QDX34073979/2022', 'USA/SC-CDC-LC0529680/2022',
'USA/OH-CDC-2-5508186/2022', 'USA/FL-CDC-STM-VCEDCJVVH/2022', 'USA/LA-EVTL13664/2022',
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'USA/FL-BPHL-5275/2022', 'USA/OR-CDC-LC0553283/2022', 'USA/WI-CDC-ASC210571526/2021',
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'USA/UT-UPHL-220226682751/2022', 'USA/PA-CDC-QDX33353832/2022', 'USA/UT-UPHL-
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NMTFEAE7H/2021', 'USA/UT-UPHL-220204062894/2021', 'USA/WA-S18117/2022', 'USA/NY-CDC-QDX33478568/2022', 'USA/TX-CDC-QDX35136075/2022', 'USA/CA-LACPHL-AF06

Cluster 2: 16 sequences

['USA/TN-CDC-ASC210818198/2022', 'USA/SD-CDC-ASC210575907/2021', 'USA/IL-CDC-ASC210642130/2021', 'USA/AL-CDC-ASC210682435/2022', 'USA/TN-CDC-ASC210584349/2022', 'USA/CA-CDC-ASC210556751/2022', 'USA/NJ-CDC-ASC210569082/2021', 'USA/FL-CDC-ASC210626458/2021', 'USA/NH-CDC-ASC210591083/2022', 'USA/HI-CDC-ASC210559647/2021', 'USA/ME-CDC-ASC210684139/2022', 'USA/IL-CDC-ASC210822446/2022', 'USA/IN-CDC-ASC210683762/2022', 'USA/OH-CDC-ASC210569490/2021', 'USA/CO-CDC-ASC210560238/2021', 'USA/MD-CDC-ASC210641291/2021']

Cluster 3: 2 sequences

['BHR/22120150009_S58_L001/2022', 'BHR/22970088641/2022']

Cluster 4: 8 sequences

['USA/UT-UPHL-220326895757/2022', 'USA/UT-UPHL-220129813536/2022', 'USA/UT-UPHL-220202565360/2022', 'USA/UT-UPHL-220331615904/2022', 'USA/UT-UPHL-220403574879/2022', 'USA/UT-UPHL-220202662018/2022', 'USA/GA-CDC-ASC210629401/2021', 'USA/UT-UPHL-220414614183/2022']

Cluster 5: 6 sequences

['USA/UT-UPHL-220109479721/2022', 'USA/UT-UPHL-220109902039/2022', 'USA/UT-UPHL-220103175466/2021', 'USA/UT-UPHL-220103607989/2021', 'USA/OH-CDC-ASC210566338/2022', 'USA/UT-UPHL-220103275553/2021']

Cluster 6: 5 sequences

['USA/UT-UPHL-220708686006/2022', 'USA/UT-UPHL-220708090362/2022', 'USA/UT-UPHL-220326654464/2022', 'USA/UT-UPHL-220308456806/2022', 'USA/UT-UPHL-220716506106/2022']

Outliers :15 sequences

Total Number of data points in all the clusters and outliers is: 5000.