Mview
Multiple Sequence Alignment (MSA)

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Results for Job ID   mview-I20250609-145946-0041-85960689-p1m

Submission Details

Result Files

Tool Output

Tool output

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Mview

```
Reference sequence (1): 6WX6_A
Identities normalised by aligned length.
Colored by: identity
```



```
                        cov     pid
  1 [

80
  1 6WX6_A         100.0% 100.0%
  2 NP_000137.2     77.1%  99.4%
  3 AAP36762.1      77.5%  89.2%
  4 2FG4_A          76.7%  99.4%
  5 CAE11873.1      91.2%  74.7%
  6 CAG32996.1      77.1%  98.9%
  7 AAA35831.1      77.1%  98.9%
  8 AKI70338.1      77.1%  98.9%
  9 AKI70336.1      77.1%  98.9%
 10 AIC54405.1      77.1%  98.9%
    consensus/100%
    consensus/90%
    consensus/80%
    consensus/70%

                        cov     pid
81
160
  1 6WX6_A         100.0% 100.0%
  2 NP_000137.2     77.1%  99.4%
  3 AAP36762.1      77.5%  89.2%
  4 2FG4_A          76.7%  99.4%
  5 CAE11873.1      91.2%  74.7%
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  8 AKI70338.1      77.1%  98.9%
  9 AKI70336.1      77.1%  98.9%
 10 AIC54405.1      77.1%  98.9%
    consensus/100%
    consensus/90%
    consensus/80%
    consensus/70%

                        cov     pid
161
240
  1 6WX6_A         100.0% 100.0%
  2 NP_000137.2     77.1%  99.4%
  3 AAP36762.1      77.5%  89.2%
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  8 AKI70338.1      77.1%  98.9%
  9 AKI70336.1      77.1%  98.9%
 10 AIC54405.1      77.1%  98.9%
    consensus/100%
    consensus/90%
    consensus/80%
    consensus/70%

                        cov     pid
241
  1 6WX6_A         100.0% 100.0%
```

```
 2 NP_000137.2    77.1%  99.4%  D-------------------------
 3 AAP36762.1     77.5%  89.2%  D---------------------L
 4 2FG4_A         76.7%  99.4%  D-------------------------
 5 CAE11873.1     91.2%  74.7%  D-------------------------
 6 CAG32996.1     77.1%  98.9%  D-------------------------
 7 AAA35831.1     77.1%  98.9%  D-------------------------
 8 AKI70338.1     77.1%  98.9%  D-------------------------
 9 AKI70336.1     77.1%  98.9%  D-------------------------
10 AIC54405.1     77.1%  98.9%  D-------------------------
   consensus/100%                ..........................
   consensus/90%                 ..........................
   consensus/80%                 ..........................
   consensus/70%                 ..........................
```

If you use this service, please consider citing the following publication: The EMBL-EBI Job Dispatcher sequence analysis tools framework in 2024. More information about this bioinformatics application can be found in its bio.tools record.

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