Jpred 4
Incorporating Jnet

Batch submission

Submitting batch job

Jpred 4
Incorporating Jnet

A Protein Secondary Structure Prediction Server

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1. Click

Input seque... VSVSVAAAGETYYIAQCAPVGGQDACNPATATSFTTDASGA...

Advanced options (click to show/hide)

2. Upload file (see tool-tip help for details and example)

3. Click to show options

...or upload a file (?) Choose File batch2.txt

4. Pick batch mode

Select type of input (?) Single Sequence (click to select format): ○ Raw/Fasta ● Batch Mode
Multiple Alignment (click to select format):

Skip searching PDB before prediction (?) ☑ Check to skip

5. Select if needed

Email address (optional) (?) a.drozdetskiy@dundee.ac.uk

6. Provide email (obligatory in batch mode)

Query name (optional) (?) myMDev3

7. Provide name of the batch (obligatory in batch mode)

Make Prediction   Reset Form

8. Push to submit...

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Summary email

From www-jpred@compbio.dundee.ac.uk ⭐
Subject **Jpred batch job testBatch completed**
To Alexey Drozdetskiy⭐

Reply | Forward

```
Your batch job testBatch has completed (with 2 successes out of 2 submitted jobs).

An archive with all the results can be downloaded from the following link:
http://www.compbio.dundee.ac.uk/jpred4/results/jp_batch_1419243898_testBatch_79gWGIi/jp_batch_1419243898_testBatch_ALL_JOBS_ARCHIVE.tar.gz

Results for individual queries are available from the links below:
SUCCESS my_seq3   Link to results: http://www.compbio.dundee.ac.uk/jpred4/results/jp_batch_1419243898_testBatch_hwLYPoh
SUCCESS my_seq2   Link to results: http://www.compbio.dundee.ac.uk/jpred4/results/jp_batch_1419243898_testBatch_79gWGIi
```

Links to batch job results will be sent via email with a subject line including the name of the job (if submitted), number of successful jobs out of all submitted, a link to a *.TAR.GZ archive with all results with directory structure preserved – one directory for each query sequence from the batch, and finally, with links to individual results directories (see the single sequence tutorial for details).