

1. Data Source

Mascot dat Files Folder

☒ Memory
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F:\target\datfiles_324

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2. Agent Summary Table

Name	Active	Veto	Parameters
Delta threshold	<input checked="" type="checkbox"/>	<input type="checkbox"/>	delta : 10
Homology	<input checked="" type="checkbox"/>	<input type="checkbox"/>	NA
Immonium	<input checked="" type="checkbox"/>	<input type="checkbox"/>	error : 0.5 aminoacids : WYFH
Length	<input checked="" type="checkbox"/>	<input type="checkbox"/>	length : 9
Modification	<input type="checkbox"/>	<input type="checkbox"/>	modification : mox
More Confident Hits	<input checked="" type="checkbox"/>	<input type="checkbox"/>	delta : 0
N Term Acetylation	<input checked="" type="checkbox"/>	<input type="checkbox"/>	NA
Proline Peak	<input checked="" type="checkbox"/>	<input type="checkbox"/>	intensity : 0.4
Protein Accession	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	accession : SHUFFLED
Sequence RegExp	<input type="checkbox"/>	<input type="checkbox"/>	regular expression : .*[li]3.*
Single Proteinhit	<input checked="" type="checkbox"/>	<input type="checkbox"/>	NA
Start Site	<input checked="" type="checkbox"/>	<input type="checkbox"/>	low : 2 high : 200
SubSequence	<input type="checkbox"/>	<input type="checkbox"/>	Subsequence : TESTPEPTIDE
Suspect Residue	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	sites : R;H
b-ion coverage	<input checked="" type="checkbox"/>	<input type="checkbox"/>	percentage : 0.30
y-ion coverage	<input checked="" type="checkbox"/>	<input type="checkbox"/>	percentage : 0.30

Load Agents

Save Agents

Clear

3. AgentAggregator Selection Table

Best Hit Agent Aggregator

Info

Properties Table

Name	Value
Theshold Score	2

Confidence (input alpha value 0.05, results in 95% confidence)

0.05

Load Task

Save Task

Start Task

Cancel