

## 1. Select the task

Task 1 : 0 out of 12 validated (0%).

## 2. Save to

Save to CSV

## Content

Property	Output
Suspect Residue	<input type="checkbox"/>
b-ion coverage	<input checked="" type="checkbox"/>
Delta threshold	<input checked="" type="checkbox"/>
y-ion coverage	<input checked="" type="checkbox"/>
Length	<input checked="" type="checkbox"/>
Start Site	<input type="checkbox"/>
Protein Accession	<input checked="" type="checkbox"/>
More Confident Hits	<input type="checkbox"/>
Single Proteinhit	<input type="checkbox"/>
Proline Peak	<input checked="" type="checkbox"/>
Homology	<input checked="" type="checkbox"/>
N Term Acetylation	<input type="checkbox"/>
BTag	<input checked="" type="checkbox"/>
Sequence Database	<input checked="" type="checkbox"/>
Delta rank n - (n-1)	<input checked="" type="checkbox"/>
E-Value	<input checked="" type="checkbox"/>
Homology	<input checked="" type="checkbox"/>

## Options

- ☒ Include validation comments
- ☐ Include confident identifications that did not match the profile
- ☐ Include non confident identifications
- ☐ Include non primary ranked hits

## Target

C:\validation.csv

Browse

Save

Cancel