

Scan	Charge	relPos	modScore	precursor mass	matched ions	TIC	%TIC	BPI
2875	4	1	80.0556	1496.13	94	2.05357e+06	0.851656	108720

Display all charge states analyzed for this scan number

How did this charge state compare with the other ones analyzed for this scan

Display all scans with a precursor mass within 1Da of the current scan

Total ion current for this scan

Base peak intensity for this scan

Display all scans with this charge state

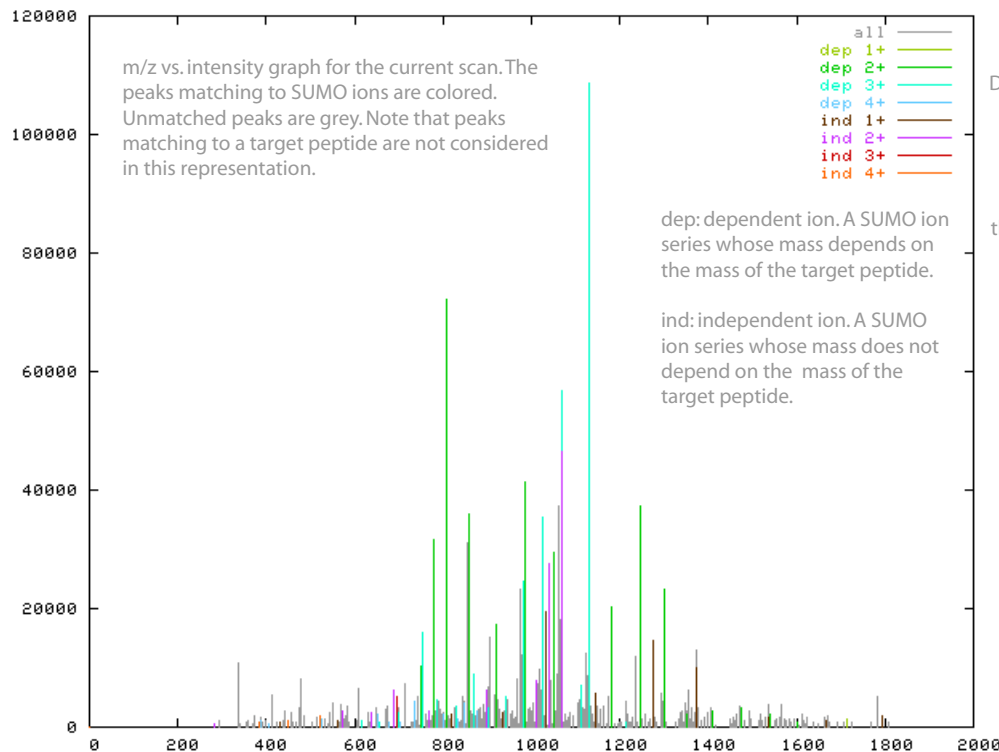
Display all scans with a modScore higher-equal than the one for this scan

Display all scans with number matched ions higher-equal than the one for this scan

Display all scans with %TIC higher-equal than the one for this scan

Expand/Collapse target peptide view

- ◊ 2876 — Unique ID for this target peptide
- ◊ >IPI00129470 Ran GTPase-activating protein 1 — Description of the protein this peptide was derived from
- ◊ LLIHMGLLK(O)SEDK — Peptide sequence. (O) indicate SUMO attachment site
- ◊ 1496.83 — Theoretical mass of this peptide
- ◊ matched ions: 16 — Number of matched ions used to calculate this peptide score
- ◊ %TIC: 0.0678297 — Percentage TIC accounted for by the ions used to calculate the peptide score
- ◊ target score: **1.08527** — Score associated with this target peptide



Display only this scan number

Display only this charge state

Display only scans with a modification score higher-equal than

Display only scans with a target score higher-equal than

Display only scans with a scan score (i.e. the sum of the modification score and the target score) higher-equal than

Display only scans with this precursor mass + or - this tolerance

Display only scans with at least this number of matched ions

Display only scans with at least this % matched TIC

Remove scans with this precursor m/z + or - this tolerance

Sort the selected scans by any of these parameters

Expand the target peptide description for all the selected scans

Scan number	
Charge state	
Modification score	
Target score	
Scan score	
Precursor mass	
Precursor tolerance	
Minimum ions matched	
Minimum % TIC	
Remove m/z	
Remove m/z tolerance	
Sort by	modificationScore
Expand all candidates	<input type="checkbox"/>

GO