## Full data analysis

### Digestion Parameters

#### Cleavage Parameters

* Enzyme=2644
* Whole-Protein=0
* Un-Specific=2418

#### Enzymes

* Lys-N=6
* LysargiNase=1
* Arg-N=10
* Glu-C=0
* Lys-C=1572
* Asp-N=22
* Asp-N (ambic)=44
* Chymotrypsin (no P rule)=1
* Arg-C (no P rule)=1371
* Trypsin=2586
* Chymotrypsin=2
* Lys-C (no P rule)=1534
* Thermolysin=6
* Pepsin A=0
* Arg-C=1393
* CNBr=1
* Trypsin (no P rule)=2560

#### Specificity

* specific=2644
* specificCTermOnly=2543
* specificNTermOnly=2543
* semiSpecific=2543

#### Maximum missed cleavages

* 2 (2644)