`topdownr` - systematic optimisation of protein sequencing for top-down proteomics

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December 2017

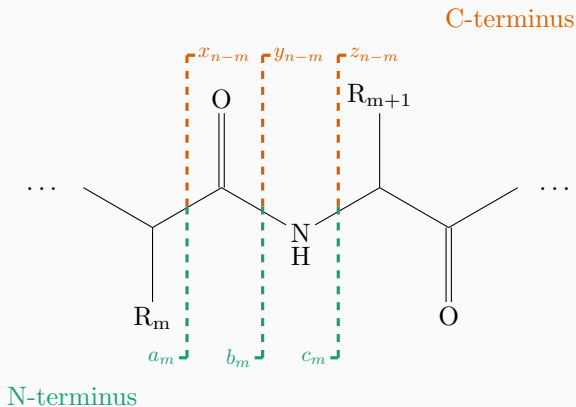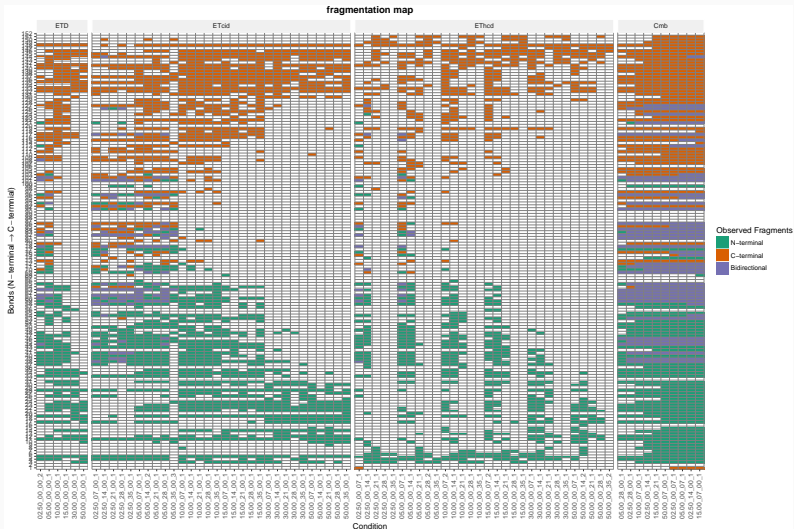**Figure 1:** amino bond, abc and xyz fragments

fragmentation map

Collaborative Work

- Pavel V. Shliaha, University of Southern Denmark.
- Ole Nørregaard Jensen, University of Southern Denmark.
- Malena Schack Jespersen, University of Southern Denmark.
- Veit Schwämmle, University of Southern Denmark.

Thank you for your attention!

```
biocLite("topdownr")
```

(this talk: https://github.com/sgibb/EuroBioc2017-topdownr/)