

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

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Introduction

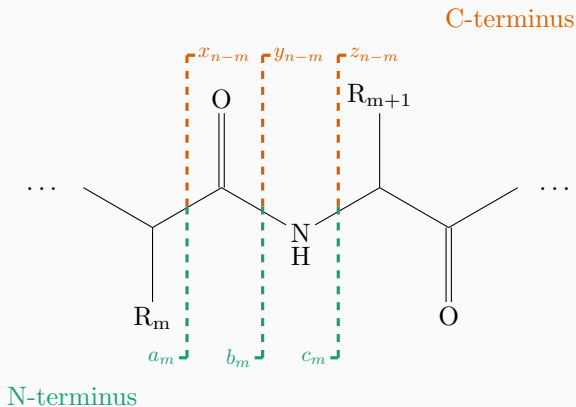


Figure 1: amino bond, abc and xyz fragments

- 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/>)