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Jaroslav Ko?a, Radka Svobodová Va?eková, Lukáš Pravda, Karel Berka, Stanislav Geidl, David Sehnal, Michal Otyepka

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The book describes the individual steps necessary for biomacromolecular fragments analysis, as well as a list of essential software tools. For each step, it also shows corresponding web-based tools in detail and provides practical examples of their use.

All tools and databases mentioned in the examples are available free of charge, platform-independent, web-based, user-friendly and do not require a prior IT background to be fully used.

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