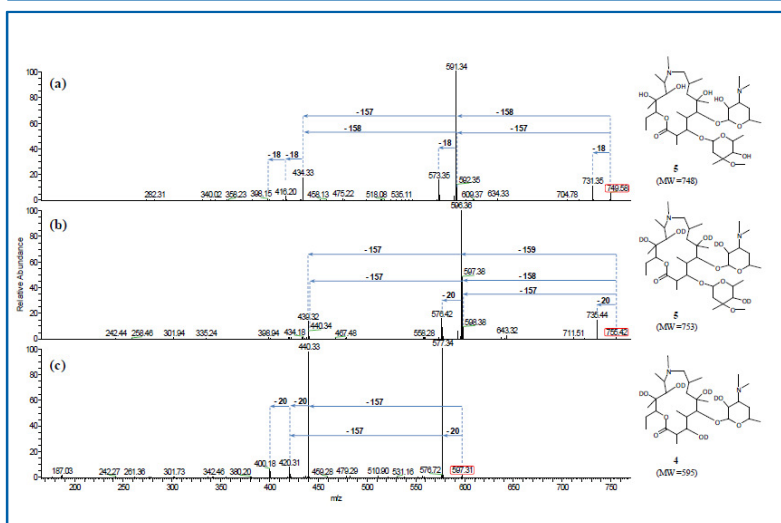


Fragmentation Study Using H/D Exchange and ESI-MSⁿ to Facilitate Structural Elucidation of Novel Macrocycles

Objective

- Detailed fragmentation study of azithromycin aglycone and its derivatives by means of ESI-MSⁿ and H/D exchange – detailed understanding of fragmentation pathways of different 15-membered azalides
- H/D exchange experiments - insight into fragmentation routes of analysed compounds
- Comparison of fragmentation patterns of macrocyclic [M+H]⁺ ions and sodium adduct ions [M+Na]⁺ -influence of an alkali metal interacting with the aglycone ring on the product-ion spectra



- The synthesis of new macrolide antibiotics involving structural modifications of azithromycin leads to novel classes of compounds. New chemical series were based on an aglycone ring as a core structure.

- Understanding the fragmentation of the different structural modifications of 15-membered azalides enables easier structure elucidations of newly synthesised compounds, proces impurities, related substances and degradation products.

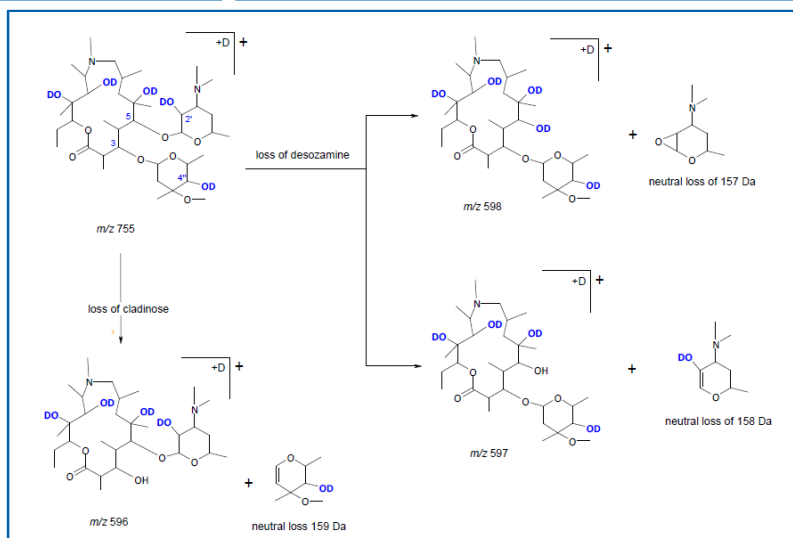
- Comprehensive fragmentation study on azithromycin and it's analogues using ESI-MSⁿ and H/D exchange on both [M+H]⁺ and [M+Na]⁺ gave insight characteristic fragments.

- Characteristic elimination of sugars from azithromycin in deuterated solvent gave ions at m/z 596, 597 and 598.

- The most intense signal at m/z 596 corresponded to the elimination of cladinose sugar moiety.

- The elimination of desosamine sugar from compound azithromycin gave two ions with signals at m/z 597 and m/z 598.

- Possible structures representing the elimination of sugars from azithromycin in deuterated solvent are given in scheme.



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