

Fragmentation Outcome Modelling: Prototype software for prediction of CID fragment ions for small molecule structures

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1. Introduction

Fragmentation Outcome Modelling is a new approach to the prediction of CID fragment ions and has been successfully applied to a series of pharmaceutical structures generating complete fragmentation including rearrangement mechanisms. This expert system uses a series of prediction models that encapsulate the outcome of fragmentation mechanisms that could result in the

breaking of one, two or three bonds. In this paper, midazolam and its 1-hydroxy and 4-hydroxy metabolites were used as test compounds to highlight the advantages of fragmentation outcome modelling compared to other commercially available fragmentation prediction software products.

2. Method and Materials

MS/MS spectra of Midazolam and its metabolites were generated on high mass accuracy systems including Shimadzu LCMS-IT-TOF, Waters Synapt G2 and Thermo LTQ-Orbitrap. Fragmentation Outcome Modelling software was developed by Shimadzu Corporation, Japan. The input file for each structure was supplied as an MDL V2000 MOL file. Comparative analyses were performed using three

commercially available software products which are described as 'Product A', 'Product B' and 'Product C'. Each software application was optimised to give the greatest number of predicted fragment ions. Fragment Outcome Modelling software functions equally for positive and negative ion data and can also be applied to either accurate or nominal mass data.

3. Results

MS/MS of Midazolam, 1-OH Midazolam and 4-OH Midazolam gave 8, 11 and 10 fragment ions respectively. The fragmentation of each molecule was unique with each fragment ion being a unique mass. For comparative

purposes the total number of fragment ions successfully predicted for each software package is expressed as a percentage of the total number of experimental fragment ions.

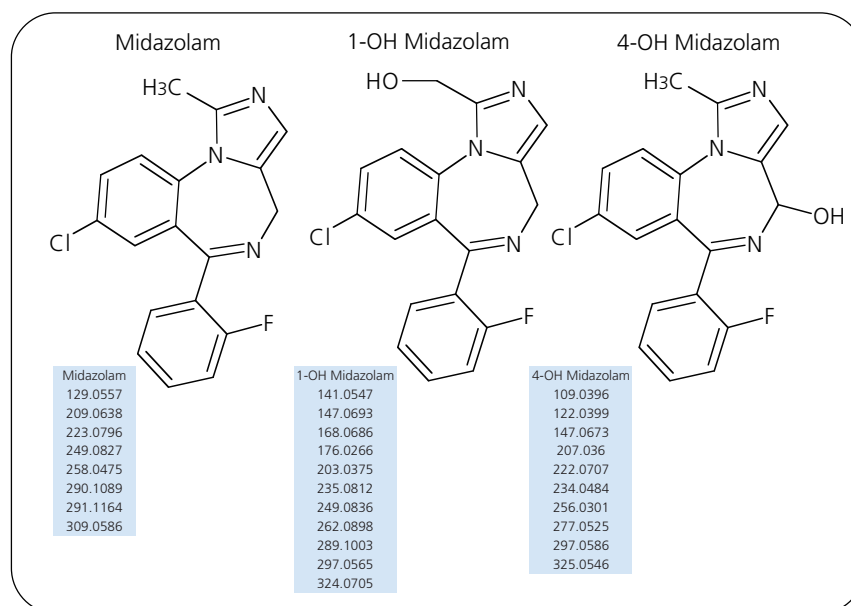


Fig. 1 Structures of Midazolam, 1-OH Midazolam and 4-OH Midazolam with corresponding experimental mass fragmentation data

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3-1. Midazolam

Fragmentation Outcome Modelling prototype software examples are shown as yellow shadowed representations of the parent molecule image. Atoms covered by the shadow image are present in the proposed fragment species. The left hand table gives data relating to the fragments with theoretical m/z , formula, electron type (odd or even), number of steps taken to generate the fragment and the

number of unique images that can be created to represent the selected fragment species. The right hand table gives data on the precursor species with the total number of precursors and a line of example data for the current precursor with precursor formula, neutral loss formula and precursor theoretical m/z . Clicking on cells in the table allows the user to iterate through data for each precursor.

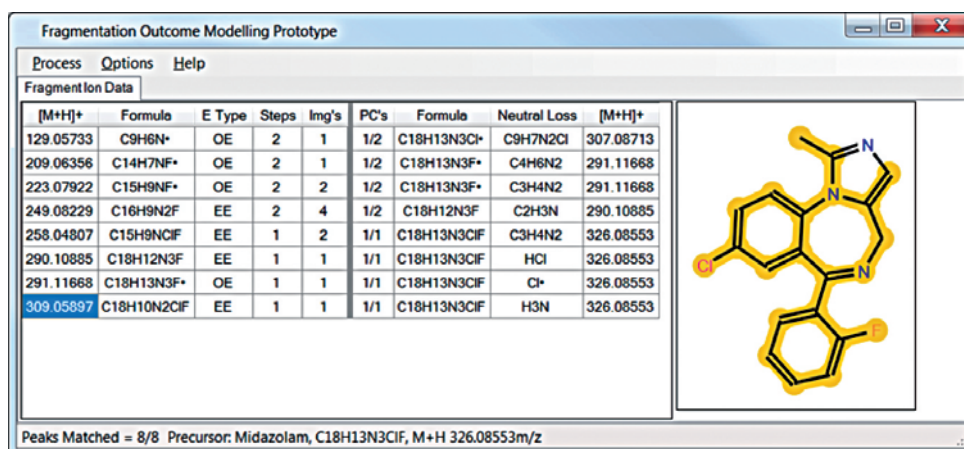


Fig. 2 Protonated fragment formed by loss of NH₃ from the imidazole ring

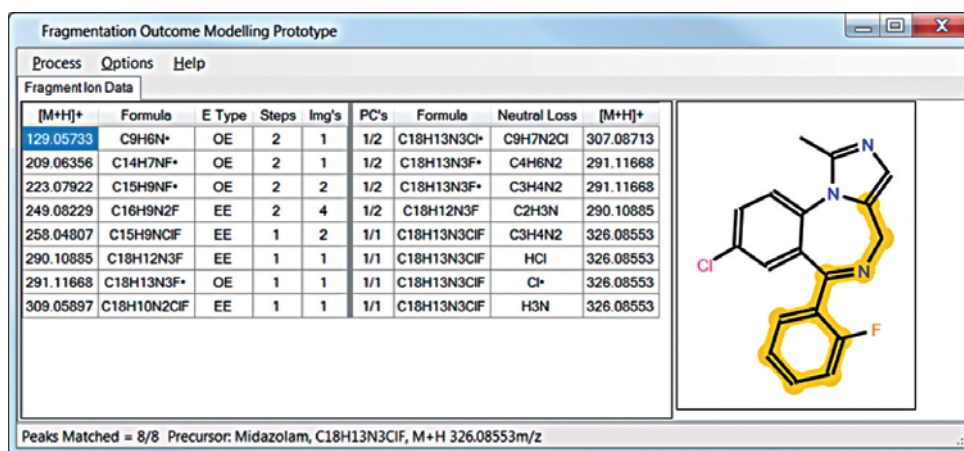


Fig. 3 Protonated radical fragment formed by ring opening and cleavage across imidazole and diazepine rings combined with the loss of a fluorine radical

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3-2. 1-OH Midazolam

Fragmentation Outcome Modelling Prototype

Process Options Help

Fragment Ion Data

[M+H] ⁺	Formula	E Type	Steps	Img's	PC's	Formula	Neutral Loss	[M+H] ⁺
141.05733	C10H6N ⁺	OE	2	1	1/2	C18H13N3OCi ⁺	C8H7N2OCi	323.08204
147.0679	C9H8NO ⁺	OE	3	2	1/3	C17H12N2OF ⁺	C8H4NF	280.10069
168.06823	C11H7N2 ⁺	OE	3	2	1/7	C17H11N3Ci ⁺	C6H4NCi	293.07147
176.02618	C10H6NCi	EE	2	1	1/4	C16H11NCiF	C6H5F	272.06373
203.03708	C11H7N2Ci	EE	2	1	1/2	C18H11N3CiF	C7H4NF	324.06987
235.07922	C16H9NF ⁺	OE	2	1	1/2	C18H13N3OF ⁺	C2H4N2O	307.11159
249.08229	C16H9N2F	EE	2	2	1/2	C18H12N3OF	C2H3NO	306.10376
262.09012	C17H10N2F ⁺	OE	3	2	1/5	C17H11N2CiF ⁺	HCl	298.0668
289.10102	C18H11N3F ⁺	OE	2	1	1/4	C18H12N3CiF ⁺	HCl	325.0777
297.05897	C17H10N2CiF	EE	2	2	1/2	C18H11N3CiF	CHN	324.06987
324.06987	C18H11N3CiF	EE	1	1	1/1	C18H13N3OCiF	H2O	342.08044

Peaks Matched = 11/11 Precursor: 1-OH-Medazolam, C18H13N3OCiF, M+H 342.08044m/z

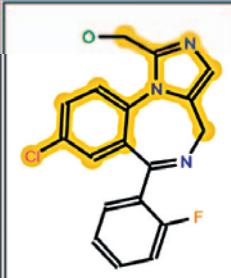


Fig. 4 Protonated fragment formed by ring opening and cleavage across the diazepine ring combined with the loss of water from the hydroxyl group

Fragmentation Outcome Modelling Prototype

Process Options Help

Fragment Ion Data

[M+H] ⁺	Formula	E Type	Steps	Img's	PC's	Formula	Neutral Loss	[M+H] ⁺
141.05733	C10H6N ⁺	OE	2	1	1/2	C18H13N3OCi ⁺	C8H7N2OCi	323.08204
147.0679	C9H8NO ⁺	OE	3	2	1/3	C17H12N2OF ⁺	C8H4NF	280.10069
168.06823	C11H7N2 ⁺	OE	3	2	1/7	C17H11N3Ci ⁺	C6H4NCi	293.07147
176.02618	C10H6NCi	EE	2	1	1/4	C16H11NCiF	C6H5F	272.06373
203.03708	C11H7N2Ci	EE	2	1	1/2	C18H11N3CiF	C7H4NF	324.06987
235.07922	C16H9NF ⁺	OE	2	1	1/2	C18H13N3OF ⁺	C2H4N2O	307.11159
249.08229	C16H9N2F	EE	2	2	1/2	C18H12N3OF	C2H3NO	306.10376
262.09012	C17H10N2F ⁺	OE	3	2	1/5	C17H11N2CiF ⁺	HCl	298.0668
289.10102	C18H11N3F ⁺	OE	2	1	1/4	C18H12N3CiF ⁺	HCl	325.0777
297.05897	C17H10N2CiF	EE	2	2	1/2	C18H11N3CiF	CHN	324.06987
324.06987	C18H11N3CiF	EE	1	1	1/1	C18H13N3OCiF	H2O	342.08044

Peaks Matched = 11/11 Structure 1/2 Precursor: 1-OH-Medazolam, C18H13N3OCiF, M+H 342.08044m/z

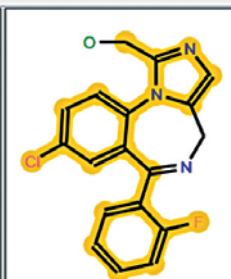


Fig. 5 Protonated fragment formed by ring opening and loss of cyanide from the diazepine ring combined with the loss of water from the hydroxyl group

3-3. 4-OH Midazolam

Fragmentation Outcome Modelling Prototype

Process Options Help

Fragment Ion Data

[M+H] ⁺	Formula	E Type	Steps	Img's	PC's	Formula	Neutral Loss	[M+H] ⁺
109.03965	C5H4N2O	EE	1	2	1/1	C18H13N3OCiF	C13H9NCiF	342.08044
122.04007	C7H4NF	EE	1	1	1/1	C18H13N3OCiF	C11H9N2OCi	342.08044
147.0679	C9H8NO ⁺	OE	2	1	1/2	C18H13N3OCi ⁺	C9H5N2Ci	323.08204
207.03199	C10H7N2OCi	EE	2	5	1/2	C17H9N3OCiF	C7H2NF	326.04912
222.07139	C15H8NF	EE	2	1	1/2	C18H12N3OF	C3H4N2O	306.10376
234.04807	C13H9NCiF	EE	1	2	1/1	C18H13N3OCiF	C5H4N2O	342.08044
256.03241	C15H7NCiF	EE	1	1	1/1	C18H13N3OCiF	C3H6N2O	342.08044
277.05274	C17H9N2Ci	EE	2	1	1/2	C18H12N3OCi	CH3NO	322.07421
277.05388	C14H10N2OCiF	EE	1	1	1/1	C18H13N3OCiF	C4H3N	342.08044
297.05897	C17H10N2CiF	EE	1	1	1/1	C18H13N3OCiF	CH3NO	342.08044
325.05388	C18H10N2OCiF	EE	1	1	1/1	C18H13N3OCiF	H3N	342.08044

Peaks Matched = 10/10 Precursor: 4-OH-Medazolam, C18H13N3OCiF, M+H 342.08044m/z

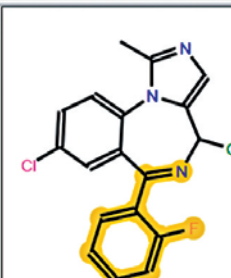


Fig. 6 Protonated fragment formed by ring opening and cleavage across the diazepine ring

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3-4. Summary Table

Results are presented in the order of Midazolam, 1-OH Midazolam & 4-OH Midazolam: Product A 75%, 46%, 70%; Product B 25%, 9%, 50%; Product C 63%, 55%, 80%.

80%; Fragmentation Outcome Modelling 100%, 100%, 100%.

	Fragmentation Outcome Modelling	Product A	Product B	Product C
Midazolam				
129.0557	√			√
209.0638	√	√		
223.0796	√			
249.0827	√	√		√
258.0475	√	√	√	√
290.1089	√	√	√	√
291.1164	√	√		√
309.0586	√	√		√
	n	6	2	5
	%	75	25	63
1-OH Midazolam				
141.0547	√			
147.0693	√			
168.0686	√			√
176.0266	√			
203.0375	√	√		√
235.0812	√			
249.0836	√	√		
262.0898	√			√
289.1003	√	√		√
297.0565	√	√		√
324.0705	√	√	√	√
	n	5	1	6
	%	46	9	55
4-OH Midazolam				
109.0396	√	√	√	√
122.0399	√	√	√	√
147.0673	√			
207.036	√	√	√	√
222.0707	√			
234.0484	√	√	√	√
256.0301	√			√
277.0525	√	√	√	√
297.0586	√	√		√
325.0546	√	√		√
	n	7	5	8
	%	70	50	80

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4. Conclusions

Fragmentation Outcome Modelling was the only application to successfully predict every fragment ion for each of the three examples given here. Typical processing time for each structure was approximately ten seconds.

There were marked differences between the capabilities of each respective software product tested and in general most packages were less successful with fragment ion predictions for the 1-OH metabolite.