Overview

The aerosols from a common commercial mentholated electronic cigarette (Ecigarette) were found to contain mainly glycerin, propylene glycol (PG), nicotine, and menthol. In addition to these compounds, five well defined chromatographic peaks were not possible to assign based on their mass spectra. The molecular formulas for the compounds generating these peaks were obtained with the help of a computer package that provides high mass accuracy from the data acquired on a conventional mass spectrometer of unit mass resolution (MassWorks 3.0.0.0, Cerno Bioscience LLC). The resulting molecular formulas were searched against a database of GRAS flavors (Flavor-Base 2004, Leffingwell & Associates). This allowed the identification in the aerosols of N-ethyl-*p*-menthane-3-carboxamide, menthone glycerol ketal (two isomers), L-menthyl lactate, and 2-secbutylcyclohexanone. The spectra of these compounds were further confirmed using MassFrontier 7.0 software (Thermo Scientific).

Introduction

Electronic cigarettes (E-cigarettes) are battery-powered devices which simulate tobacco smoking. They use a heating element that vaporizes a liquid solution. The aerosols from a common commercial mentholated E-cigarette were analyzed, and some compounds had mass spectra not available in common mass spectral libraries. These compounds needed to be identified.

Method

1. The aerosols from the E-cigarette were collected on 44 mm Cambridge pads with a linear smoking machine (Cerulean SM 450). The pads were extracted with methanol.

2. The samples were analyzed on a 6890/5975 GC/MS (Agilent) equipped with a DB-WAXetr column 30m x 0.32mm with 0.5mm film.

3.Perfluorotributylamine (PFTBA) was automatically infused at the end of the chromatograms during data acquisition and used to generate correction factors for accurate mass. Mass spectra were collected as "raw scan" and with the threshold set at "zero" counts.

4. The spectra of unknowns were corrected to 0.01 Da mass precision using MassWorks package and an elemental composition was generated for unknowns.

5. The molecular formulas were searched against a GRAS flavor database (Flavor Base 2004) to identify the potential compounds

6. The spectrum of each compound was further evaluated using MassFrontier 7.0 software for validation.

Identification of cooling agents in aerosols of an E-cigarette from unit mass resolution spectra enhanced to high mass accuracy

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Results

1. The total ion chromatogram for the aerosol sample is shown in Figure 1.



Figure 1. Total ion chromatogram of the sample, plus PFTBA.

The chromatogram shows the main peaks, and the unknown peaks.

2.Example of average area selected for generating the spectrum of one unknown peak is shown in Figure 2.



Figure 2. Window of the chromatogram showing the area selected for generating an average spectrum (from MassWorks 3.0.0.0.).

3. Average spectrum of the selected peak is shown in Figure 3.



Figure 3. Average spectrum for the peak at 60.9 minutes (from MassWorks 3.0.0.0.).

4. T ir	he res 1 Table
Table m	1. Mo nin.
Row	Fc
1	C13
2	C10 C9H
4	C6H
5	C5H C3H
7	C2H
5. F cu F se	lavor- orresp EMA ensatio
6. T st p	The spo tructur redicte
Abundance	
100	
90	HC
80	
0 IIII	
50	
40	
30	
20	
10	29
0	
m/z>	20
Figur a	e 4. N nd frag

sults of molecular formulas resulting from MassWorks are shown

olecular formulas from MassWorks for the peak eluting at 60.9

ormula	Mono Isotope	Mass Error (mDa)	Mass Error (ppm)	Spectral Accuracy
BH25ON	211.1936	0.7144	3.3828	98.3374
)H27O4	211.1909	-1.9656	-9.3073	96.8036
12703N2	211.2022	9.2677	43.8828	96.61
12906N	211.1995	6.5877	31.1927	93.7021
12706N2	211.1869	-5.9884	-28.3551	93.1122
13109	211.1968	3.9076	18.5025	90.683
12909N	211.1842	-8.6685	-41.0452	90.0863

-Base 2004 indicated that formula C13H25NO (first hit) bonds to N-ethyl-*p*-menthane-3-carboxamide (CAS# 39711-79-0, 3455), which is a non-menthol compound producing a cooling on at 200 ppb.



bectrum of peak eluting at 60.9 min is in agreement with the re of N-ethyl-*p*-menthane-3-carboxamide. Various fragmentations ed by MassFrontier 7.0. are shown in Figure 4.



Mass spectrum assigned to N-ethyl-*p*-menthane-3-carboxamide, gmentations predicted with MassFrontier 7.0.



Figure 5. Mass spectrum assigned to menthone glycerol ketal, and fragmentations predicted with MassFrontier 7.0.

Conclusions

1. High mass accuracy can be generated from the data acquired on a conventional mass spectrometer of unit mass resolution (with MassWorks program from Cerno Bioscience LLC).

2. With high mass accuracy information, the correct molecular formulas can be generated for an unknown compound and corroborated with the information on the sample, the molecular structure can be determined.