Investigation of Aging in Beer Using a New Gas Chromatography Time-of-Flight Mass Spectrometry Benchtop System

Introduction

Gas chromatography paired with time-of-flight mass spectrometry (GC-TOFMS) was used to investigate the aging process of beer to determine time course trends for individual analytes within the aroma profile. This type of non-targeted chemical analysis can provide more insight than targeted approaches, and is a good complement to other traditional analyses, such as sensory panels. Rather than specifying target analytes in advance, analytes of interest can be determined from the data. A time course series of samples was prepared by artificially accelerating aging with storage at elevated temperature for various lengths of time. One day of storage at 40 °C was used to represent one month of storage under appropriate conditions. This protocol was used to produce a time course series of beer samples effectively aged an additional 0, 1, 2, 3, 4, 5, 6, 8, 12, and 20 months. The samples were analyzed in triplicate using headspace solid-phase micro-extraction (HS-SPME) to collect and concentrate volatile and semi-volatile analytes in the headspace of each sample. Chemical analysis was then accomplished with a new commerciallyavailable GC-TOFMS benchtop system. The data were analyzed for specific targets and mined for previously unknown trends and differences. Identification information and relative quantitative information were determined for over 300 analytes. From these data, specific time course trends were observed for some analytes, providing a better understanding of the samples and the aging process.

Samples and Aging Experiment A commercially-available IPA was purchased and artificially aged with storage at elevated temperatures. A previously reported protocol calibrated that 1 day of storage at 40 °C was comparable to 1 month of storage under appropriate conditions. (Ref: Marques. ASBC A30. Chicago, IL. June 2014) That information was used to generate a time course set of samples by storing samples at 40 °C for 0, 1, 2, 3, 4, 5, 6, 8, 12, or 20 days. This sample preparation is diagramed in Figure 1.

Methods

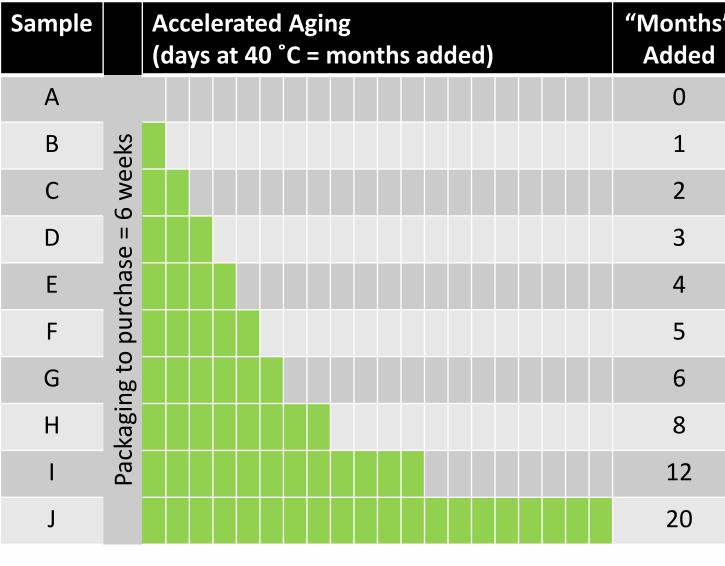


Figure 1. Diagram of forced aging experiment.

Analytical analysis: For each sample, 5 mL of beer were pipetted into a 10 mL glass vial. The samples incubated for 10 min at 60 °C. Extraction was performed with a DVB/CAR/PDMS fiber (Supelco) for 20 min also at 60 °C prior to analysis by GC-TOFMS. Instrument conditions are listed in Table 1

Gas Chromatograph	Agilent 7890 with LECO L-PAL3 Autosampler		
Injection	3 min desorption in 250 °C inlet		
Carrier Gas	He @ 1.0 ml/min, Constant Flow		
Column	Stabilwax, 30 m x 0.25 mm i.d. x 0.25 μ m coating (Restek)		
Oven Program	Hold 4 min at 35 °C, ramp 5 °C/min to 180 °C, ramp 10 °C/min to		
	220 °C and hold 5 min		
Transfer Line	250 °C		
Mass Spectrometer	LECO Pegasus [®] BT		
Ion Source Temp	250 °C		
Mass Range	35-650 m/z		
Acquisition Rate	10 spectra/s		

Table 1. GC-TOFMS Instrument Conditions

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GC-TOFMS

HS-SPME with GC-TOFMS offered an overview of the aging process. These samples have

significant complexity and hundreds of peaks were observed in the aroma profile.

A variety of analytes including terpenes, esters, aromatic compounds, acids, aldehydes,

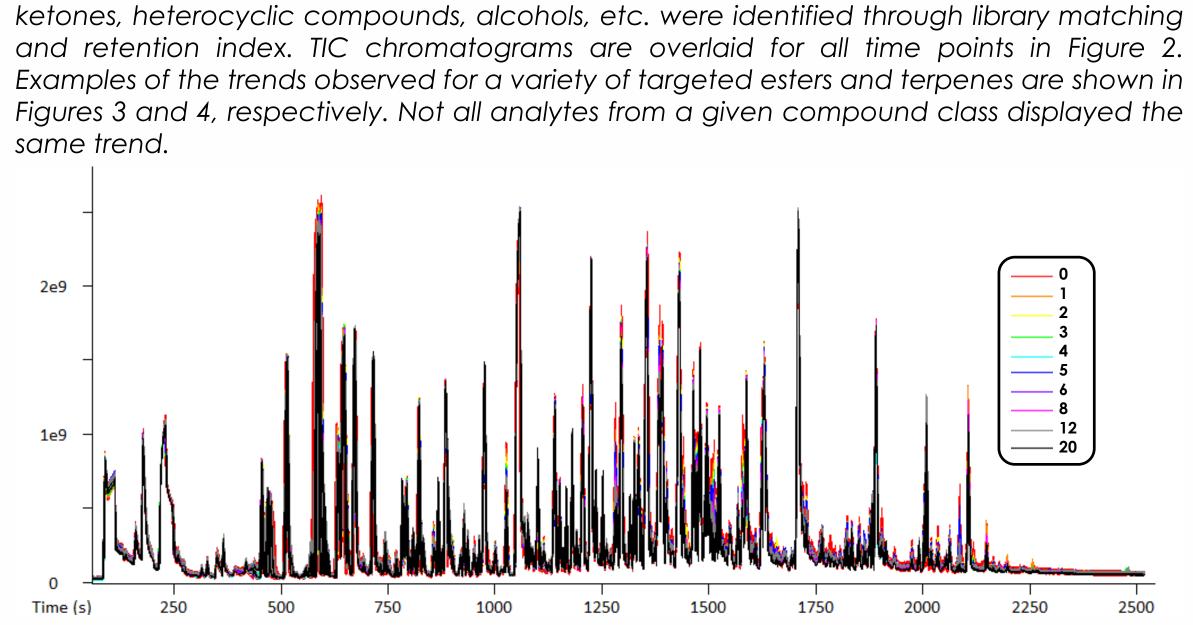


Figure 2. TIC chromatograms are overlaid for all time points in the aging experiment.

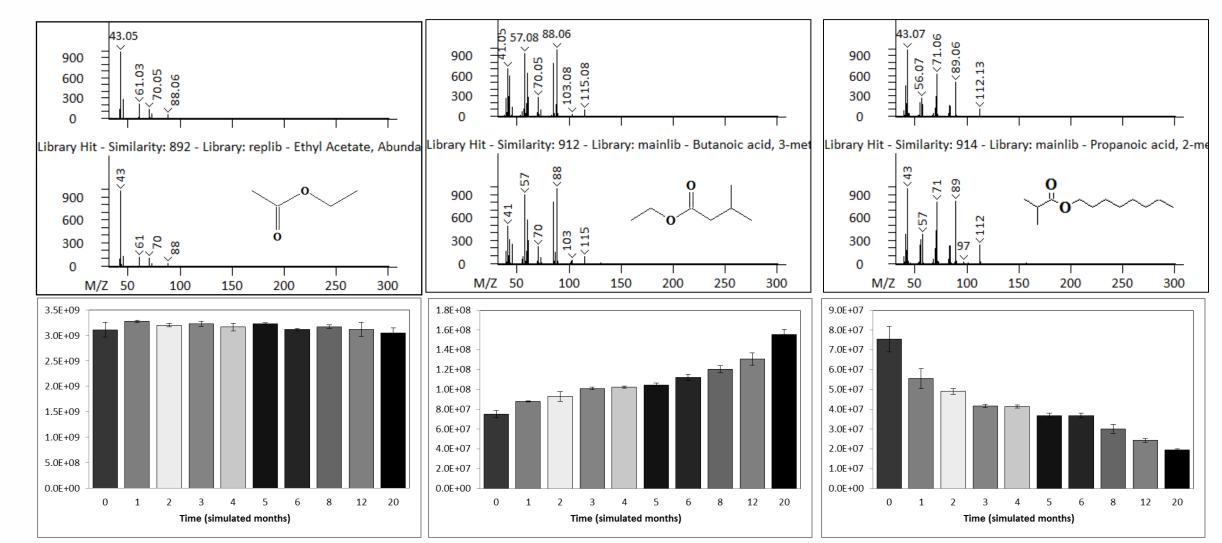


Figure 3. Examples of some of the trends observed for various esters.

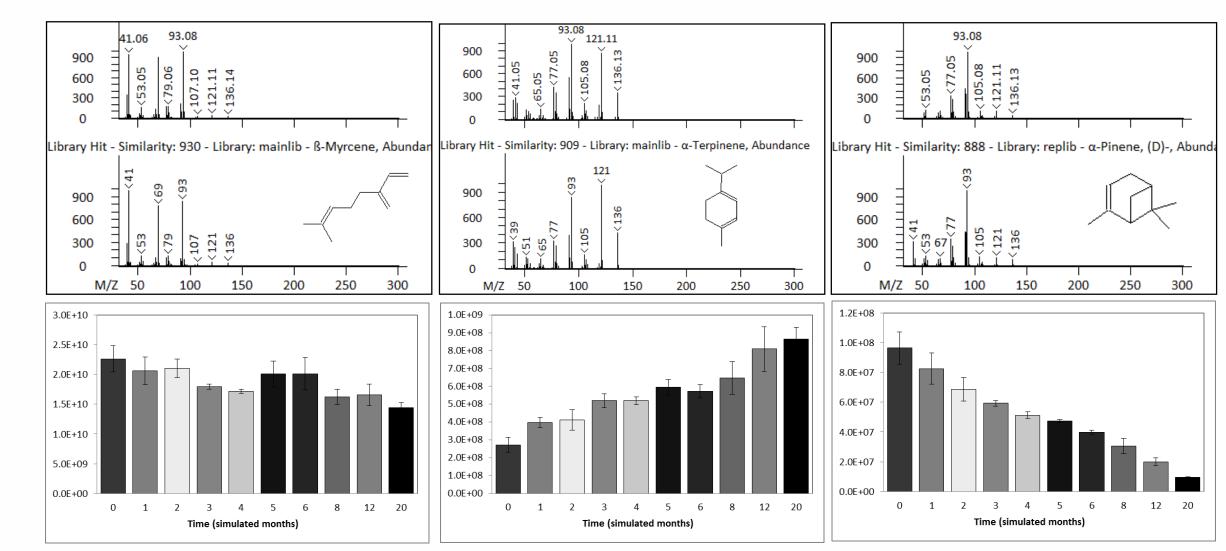


Figure 4. Examples of some of the trends observed for various terpenes



Non-Target Analysis

General peak finding was performed and peak area information for the peaks was aligned between the samples. This resulted in over 300 analytes compiled across the time course. To determine differences and trends within these analytes, a t-test was calculated between peak area data from time 0 and time 20. This highlighted analytes that differed from the beginning to the end of the time course. Regression statistics were then used to distinguish those that were trending with the time course rather than exhibiting random variation between those time points. 108 analytes were found to differ between the beginning and the end of the time course and to correlate with aging. Those analytes are summarized in the heat map, shown in Figure 5. Each row is an analyte, and each column is a time point. The color scale shows relative intensity, so time course trends for each analyte can be observed by the color change across a row. Representative examples are shown in Figure 6 and Table 2.

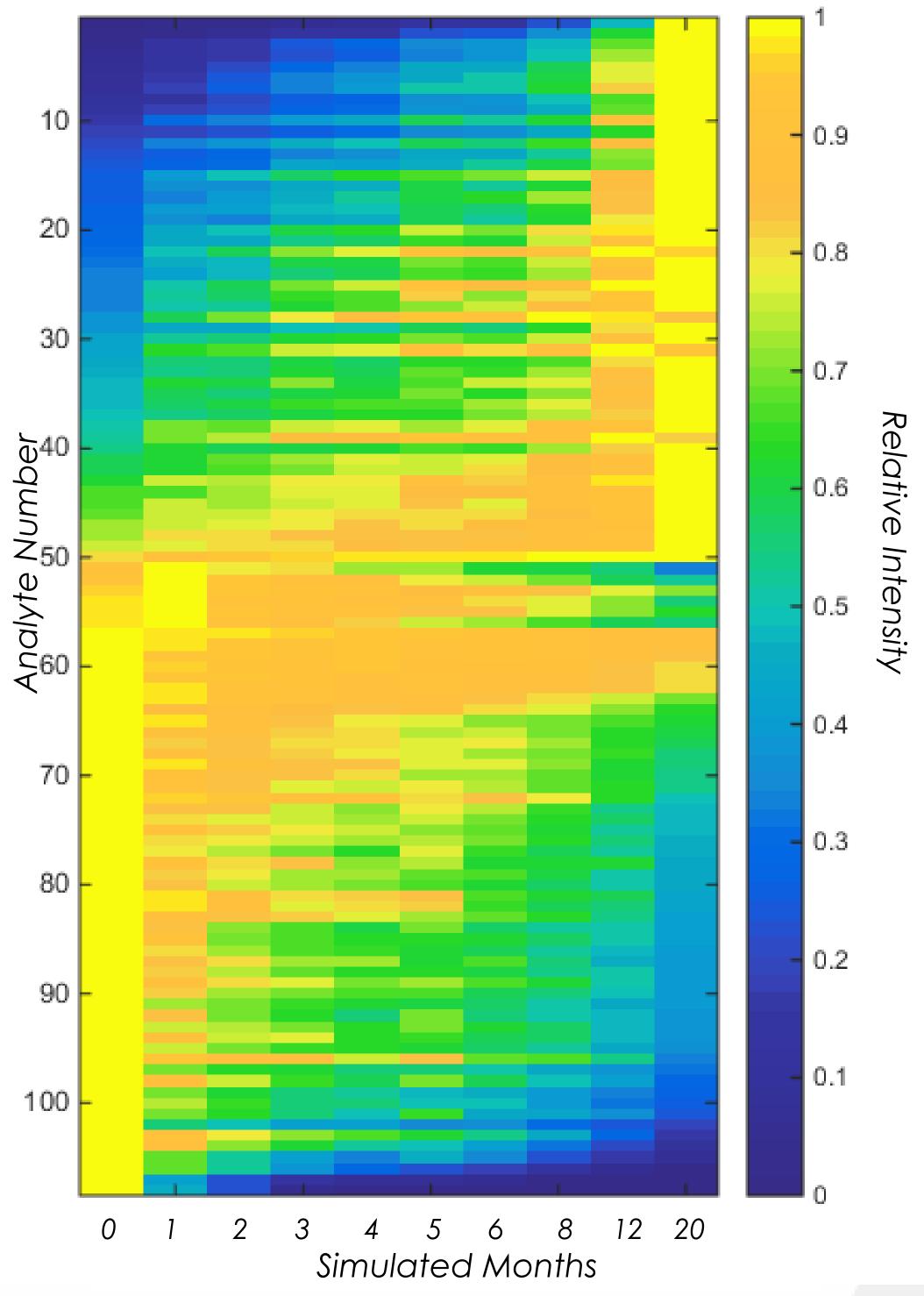
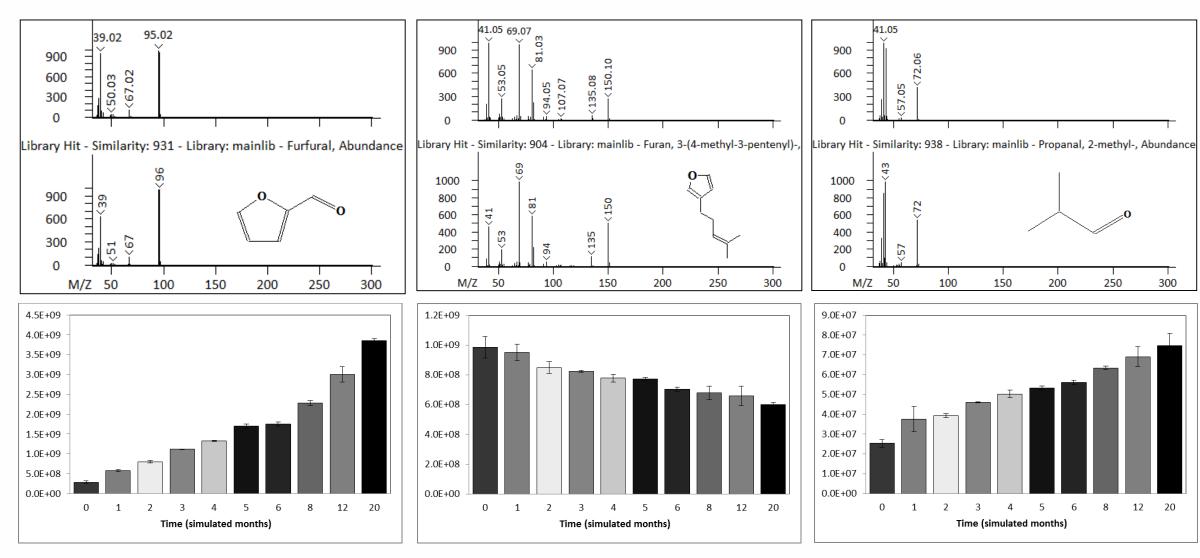


Figure 5. 108 Analytes that trend across the time course/aging process.



Representative Examples

Figure 6. Representative analytes from heat map.

Table 2. Representative analyt	es and their odor properties
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Peak#	Analyte	Similarity	Odor Notes
5	furfural	931	sweet, woody, almond, fragrant baked bread
6	5 2-methyl propanal		pungent, cocoa, musty, green, malty, bready
16	2-pentyl furan	866	fruity, green, earthy, beany, vegetable
19	2-n-butyl furan	837	spicy, fruity, wine, sweet, mild
23	α-terpinene	909	woody, terpene, lemon, herbal, medicinal, citrus
32	furan	906	ethereal
33	2 methyl furan	912	chocolate, ethereal, acetone
36	3-methyl ethyl ester butanoic acid	912	fruity, sweet, apple, pineapple, tutti fruitti
40	dimethyl sulfide	930	sulfur, onion, sweet, corn, vegetable, cabbage, tomato, green, radish
65	3-(4-methyl-3-pentenyl)- furan	904	woody
71	humulene	919	woody
98	caryophyllene	937	sweet, woody, spice, clove, dry
100	2 methyl octyl ester propanoic acid	914	oily, green, waxy, soapy, clean, fruity, creamy
104	α-pinene	888	herbal, fresh, camphor, sweet, pine, earthy, woody
108	caryophyllene oxide	868	sweet, fresh, dry, woody, spicy

Conclusions

This poster has demonstrated GC-TOFMS to monitor and track the aroma profile of aging beer. GC-TOFMS is a non-targeted analytical technique that provides chromatographic separation and full mass range mass spectral detection for the complete chromatographic separation. This comprehensive data can be mined for specific target analytes of interest, and also reviewed for inherent trends and differences in the data. Within a given compound class, various trends (increase, decrease, random variation, or no change) were observed for individual analytes, so non-targeted analysis tools were used to discover specific individual trending analytes. Peak area information for more than 300 analytes was compiled across the time course and 108 analytes that correlated with aging were determined. This analytical approach is useful for these types of discovery analyses.