

INTRODUCTION

Many chemical changes occur during the brewing process as raw materials are taken through various stages to yield a finished product. Understanding these changes and when they happen has the potential to direct process optimization, improve the final product, and improve efficiency of the process. Many of the changes throughout the brewing process can be detected in the aroma profile, which is comprised of the volatile and semi-volatile analytes associated with each sample. Gas chromatography coupled with mass spectrometry (GC-MS) is well-suited for analysis of volatile and semi-volatile analytes and is a powerful way to screen the aroma profile. We use GC-MS in this work to probe five points throughout the brewing process. Samples were collected pre-boil, post-boil, from a full fermenter, at the end of fermentation, and from a full bright beer tank. In this work, GC-MS was paired with headspace solid phase micro-extraction (HS-SPME) as the sampling technique to collect and concentrate the volatile and semi-volatile analytes from the headspace prior to injection. Individual analytes that were collected on the SPME fiber subsequently separate from each other as they travel through GC column. MS detection then provides information for identification and relative quantitation. Hundreds of analytes were detected in these samples and information for representative analytes is presented here. A variety of compound types including esters, terpenes, terpenoids, organic acids, alcohols, aldehydes, ketones, furans, aromatics, nitrogen-containing, and sulfur-containing analytes are presented. Various trends can be observed in these analytes, many that can be connected to the brewing process. For example, some analytes from the malt are observed to decrease during the boil while analytes from the hops are observed to increase during the boil. Ethanol was observed to increase during fermentation. Some esters increase during the boil while others increase during fermentation. Observing changes in the aroma profile overall can provide good insight to the chemical changes occurring throughout the process.

METHOD

Samples were collected at various stages of the brewing process (pre-boil, post-boil, fermenter full, end of fermentation, and bright beer full tank). Each sample (5 mL in a 20 mL vial) was analyzed with HS-SPME coupled to LECO's Pegasus BT GC-TOFMS, with the method conditions listed in Table 1. An alkane standard was also acquired to calculate retention indices.

Table 1. Instrument Conditions

AS	LECO I-PAL3 Autosampler
SPME	10 min incubation, 20 min extraction at 35 °C
SPME fiber	DVB/Car/PDMS (Supelco)
Fiber Conditioning	5 min pre-injection at 250 °C
GC	
Injection	Desorb fiber 3 min at 250 °C, splitless
Columns	Stabilwax 30 m x 0.25 mm i.d. x 0.25 µm coating (Restek)
Carrier Gas	He @ 1.40 mL/min
Oven Program	3 min 40 °C, ramp 10 °C/min to 250 °C, hold 1 min
MS	LECO Pegasus® BT
Ion Source Temp	250 °C
Mass range	33-500 m/z
Acquisition Rate	10 spectra/s

GC-TOFMS Results

Representative chromatograms for each sample from the brewing processing are shown in Figure 1. Many differences are readily apparent, indicating significant chemical changes over the brewing process. Information for four analytes (labeled A-D) that clearly change is compiled in Figure 1. Other important differences may be obscured by coelutions in the total ion chromatogram (TIC) view, but are revealed with automated data processing and deconvolution, as shown in Figure 2.

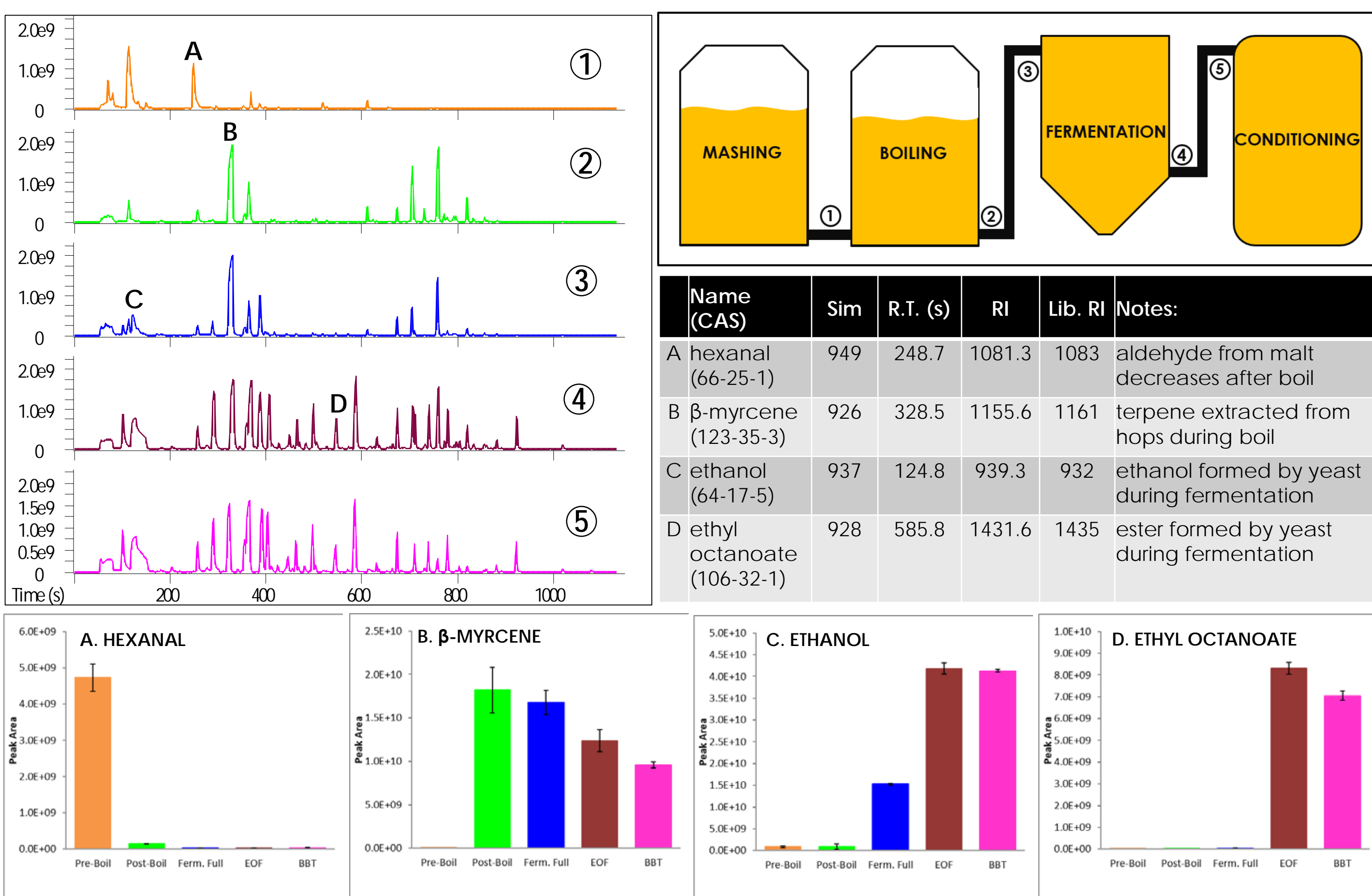


Figure 1. Representative chromatograms and analytes that clearly change during the brewing process.

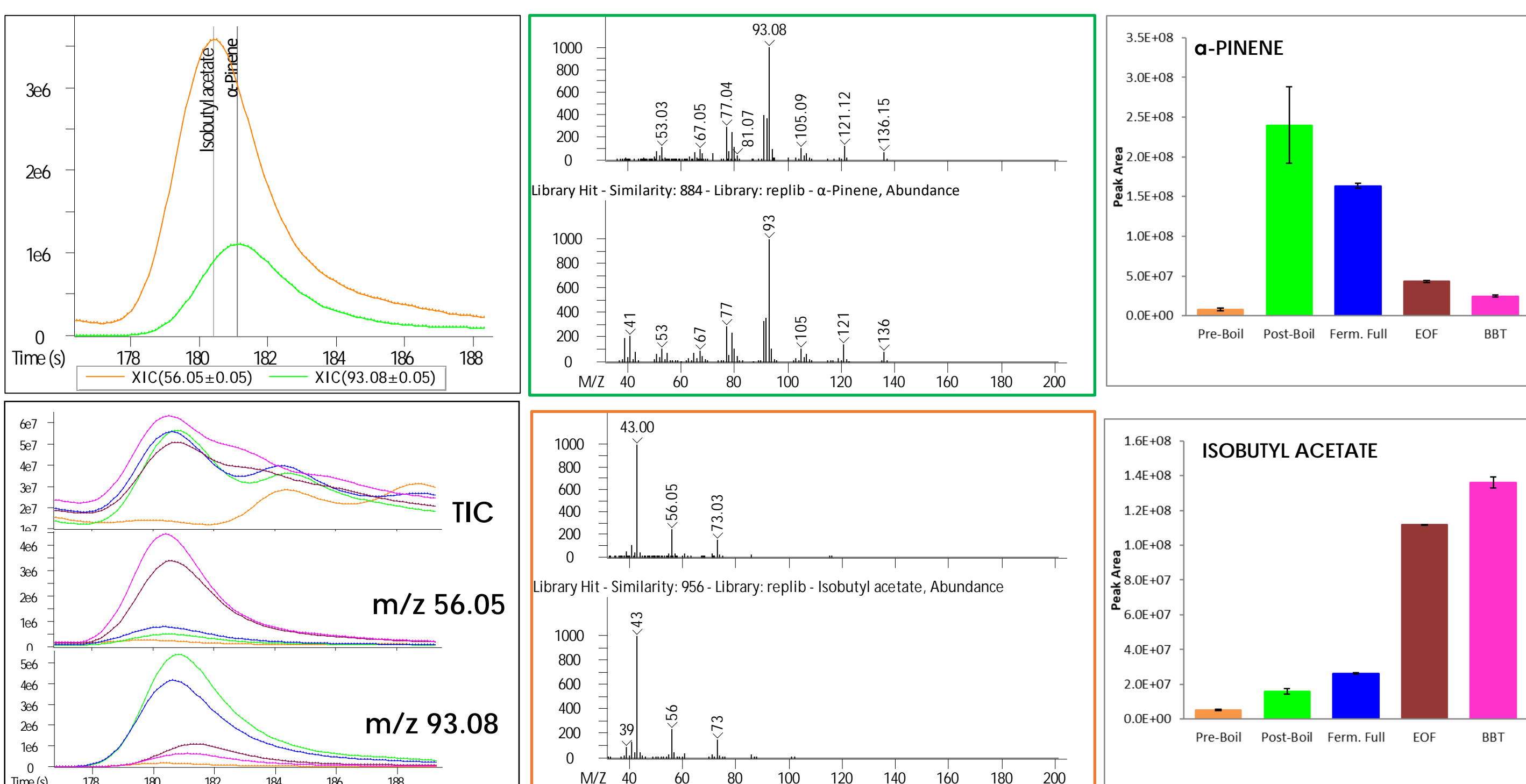


Figure 2. The trends for other analytes may be obscured by coelutions in the TIC view. In these cases, deconvolution is crucial for distinguishing analytes and their changes over the brewing process. Here, an ester increases and a terpene decreases. Their individual differences are not apparent in the TIC view.

ANALYTE TRENDS

A collection of some analytes that change during the brewing process are shown in Figure 3. The analyte identifications were determined by mass spectral and retention index matching (observed compared to NIST library databases). The heat map shows the relative peak area for each analyte at each point in the process, indicating how that analyte changes through the brewing process. Different compound classes have different behaviors. For example, some esters increase during the boil while others increase during fermentation. Terpenes increase during the boil and then gradually decrease throughout the process. Organic acids and alcohols mostly appear after fermentation. This type of information can help track when specific off-flavors have dropped or when various aroma contributors have reached a desired level.

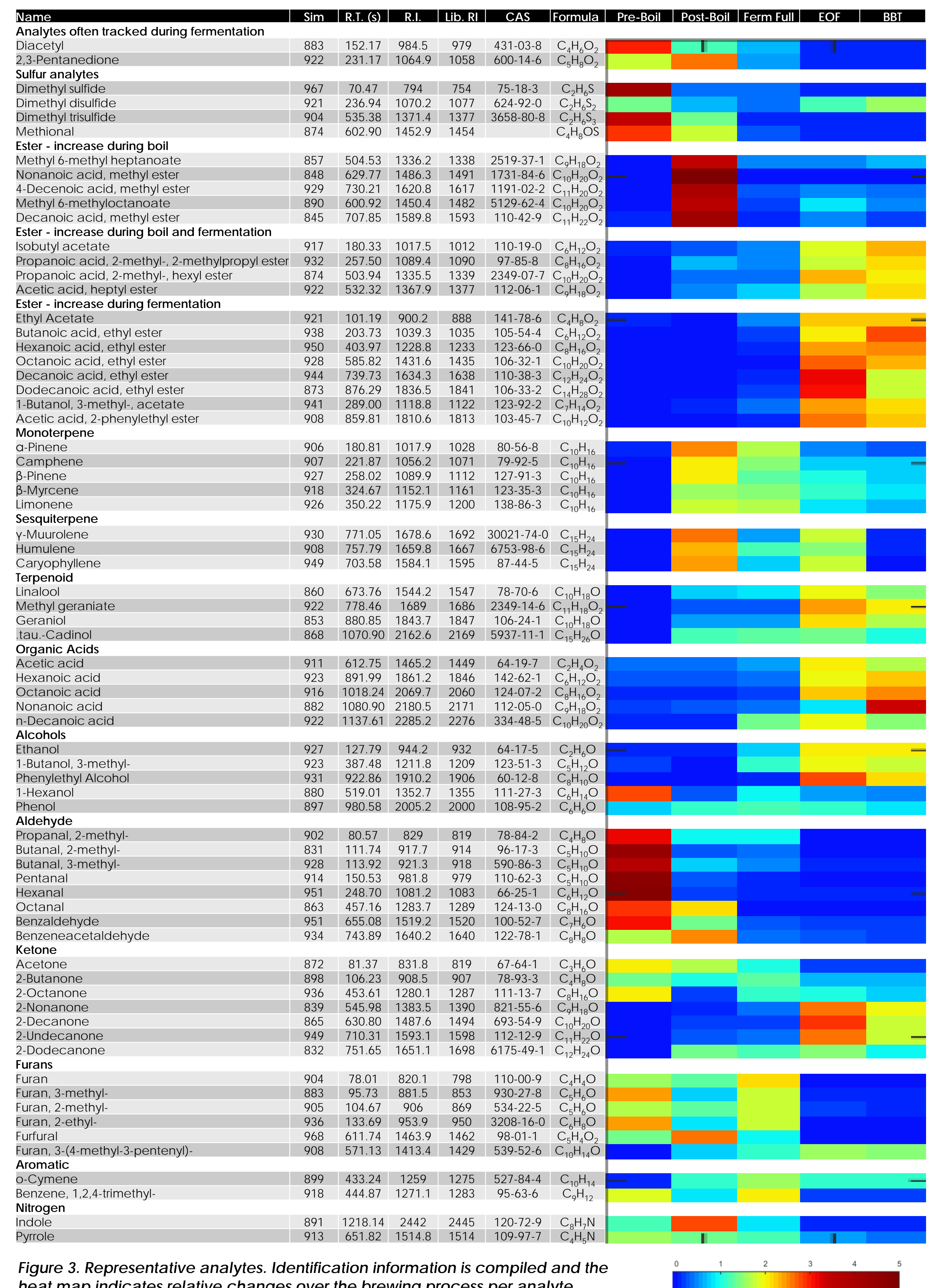


Figure 3. Representative analytes. Identification information is compiled and the heat map indicates relative changes over the brewing process per analyte.

RELATED ANALYTES

This information can also show how analytes may relate to each other. For example, the relationship between furfural, furfuryl alcohol, ethanol, and furfuryl ethyl ether is described in Figure 4. The decreases and increases of individual analyte seem to connect with the expected reactions.

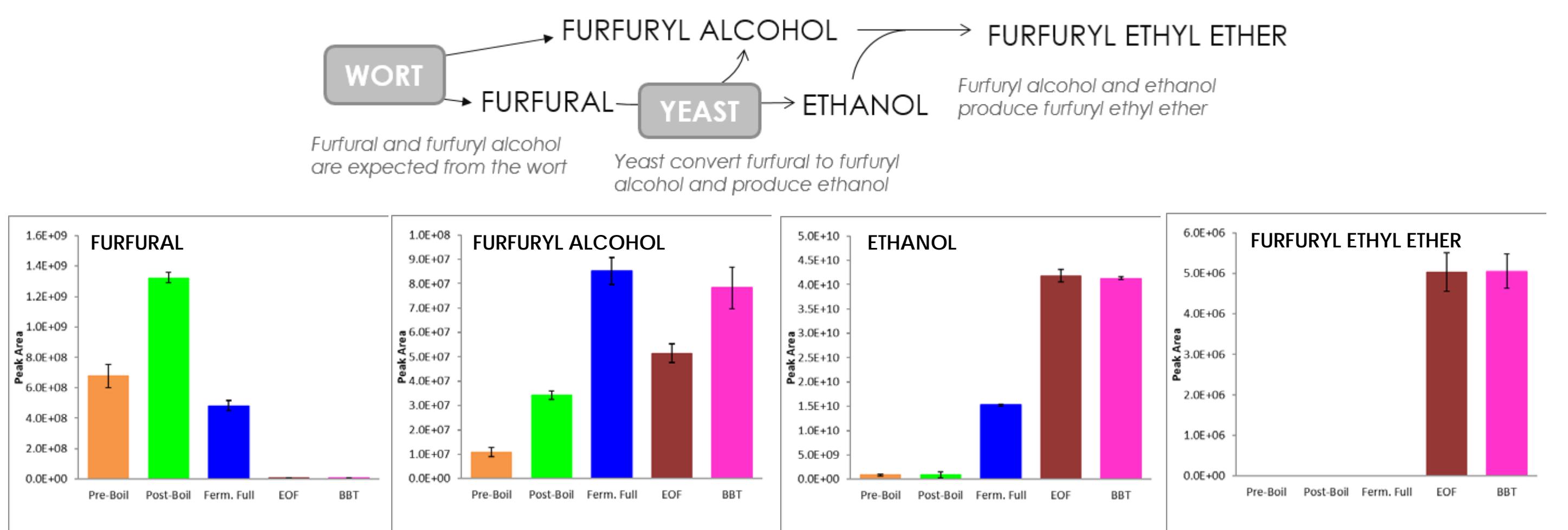


Figure 4. Furfural and furfuryl alcohol, both expected from wort, are observed to increase during the boil. Yeast convert furfural to furfuryl alcohol, and a decrease in furfural along with an increase of furfuryl alcohol is observed after yeast are added at the start of fermentation. Yeast also produce ethanol, which is observed to increase at the start of fermentation. Ethanol and furfuryl alcohol can react with each other to produce furfuryl ethyl ether, which is observed at the end of fermentation along with a corresponding decrease in furfuryl alcohol.

CONCLUSIONS

Many chemical changes occur during the brewing process as raw materials are taken through various stages to yield the finished product. GC-MS is an effective way to determine individual chemicals and to understand the changes they undergo during the brewing process. Hundreds of analytes were detected in these samples and information for representative analytes was presented. Deconvolution was crucial in discerning chromatographic coelutions and provided information on more analytes. Observing changes in the aroma profile provided good insight to the chemical changes occurring throughout the process.