

Application Notes

Gas Chromatography

Analysis of Methyl Laurate Content In Low Concentration Biodiesel Blends (B1, B2) with Multidimensional Gas Chromatography System (Part I)

Introduction

The fossil fuels importing country like Philippines is actively implementing a long-term Alternative Fuels Program to develop indigenous and renewable energy fuels for long term energy security, which will be a pillar for the country's sustainable growth. Biodiesel Program, which is one sub-program of The Alternative Fuels Program, mandates a minimum 1% biodiesel blend (B1) in all diesel fuels sold in the country. It might be increased to 2% biodiesel blend (B2) within two years from the effective date of the Act.^[1]

Biodiesel fuel is produced from renewable sources such as vegetable oil or animal fat through transesterication into fatty acid methyl esters (FAMEs) mixture. Pure biodiesel is called B100. The Draft of Philippines National Standard DPNS/DOE QS 002:2007 specifies that the Methyl Laurate (C12ME) content in the pure (B100) biodiesel should be a minimum of 45% by mass.[2] This is a typical characteristic of biodiesel produced Meanwhile, the standard from coconut oil. DPNS/DOE QS 004:2008 requires that the B2 biodiesel blend (i.e. diesel fuel with 2% v/v biodiesel) that is sold in the country by 2009 should have a minimum of 0.8% by mass of C12ME.[3] This is to ensure that the B100 biodiesel used in the biodiesel blend is produced from the main local indigenous supply, the coconut oil. The FAMEs mixture derived from coconut oil is called as "Coconut Methyl Ester (CME)" or "Coco-Biodiesel" locally.[4] The total FAME content of commercial B1 and B2 biodiesel blend could be easily determined by rapid analysis method with HATR-FTIR technique.[5] However, by only checking the C12ME content in these biodiesel blends, the use of CME in B1 and B2 biodiesel sold in the local market could be confirmed.

B1 & B2 biodiesel blends are complicated mixtures comprising of middle distillates (hydrocarbons range from C9~C23), FAMEs and other minor impurities. At present, to analyze C12ME, the total

FAMEs fraction is first extracted from diesel by atmospheric pressure liquid chromatography on a silica micro column. Then, only the isolated FAMEs fraction is characterized by GC-FID. [6] Besides the target analyte extraction recovery issue, the sample preparation process is time consuming and significantly increases the consumables cost of the current analysis method.

C12ME is commonly co-eluting with other hydrocarbon peaks of diesel in single-dimension (1D) GC separation. Two-dimensional (2D) GC technique offers a superior separation and an alternative for this application. Shimadzu Multi-Dimensional Gas Chromatography system (MDGC) is ideal for this complex mixture. This dual-oven system is equipped with an inert "switching unit" and utilizes the mechanism of Multi-Deans Switching technology. The dual-oven characteristic allows 2D separation by using two columns of different characteristics, each operated under independent temperature programs.

The goal of this experiment is to develop a 2D GC method for direct injection of biodiesel blends employing a heart-cut technique to isolate C12ME from unresolved hydrocarbon components in the B1 and B2 biodiesel blends. The needs of 2D separation and dual oven 2D separation for this application were investigated.



Experimental

Samples and Standards

Four diesel samples named as SS, ES, PS and CS were obtained from different brands in the market and were used as matrices of the spiked samples. C12ME (Methyl Laurate) standard of 98.8% purity was purchased from ChemService, Inc, USA.

Instruments and Analysis Conditions

An MDGC/GCMS-2010 equipped with a liquid auto-injector. AOC-20i and a set GC/GCMS/MDGCsolution Workstation (Shimadzu Corporation, Kyoto, Japan) were used for this experiment. The system configuration is shown in Fig. 1. This system consists of a GC-FID (defined as 1st Instrument), equipped with an inert flow switching element (namely "switching unit") and linked with a heated interface (230°C) to a GC-qMS (defined as 2nd Instrument). A pair of column with combination of non-polar and polar columns were used in these methods development located in 1st and 2nd instrument respectively. The conventional MDGC methods with 0.25mm ID capillary capillary columns analysis conditions are listed in Table 1.

Results

The upper diagram of Fig. 2 shows the chromatogram profiles obtained from the 1st column for 4 different brands of diesel and a particular diesel (CS) spiked with 1% C12ME. Diesel from different brand or sources are slightly different in its hydrocarbons composition. The lower diagram is an enlargement the upper diagram; the highlighted area shows the overlapping zone of C12ME with the matrix. This indicates that the target analyte might be able to be separated from the matrix with an optimized method in a particular diesel sample but it could co-elute with the matrix in other diesel samples. This means, it might not be possible to have a common analysis method by using single dimension separation that is optimized for various types of biodiesel blends.

To enhance the separation of C12ME in B1 and B2 biodiesel blend, the unresolved C12ME elution band from the 1st column (non-polar column) was selectively heart-cut by using the "switching unit" to the 2nd column (polar column) for further, 2nd dimension, separation. The details of the heart-cutting mechanism of the "switching unit" and its operation by MDGC solution software were described in the literature. [7,8,9]

The results of the analysis of CS diesel sample that was spiked with 0.5% C12ME are shown in Fig. 3; the upper diagram shows the chromatograms obtained from the 1st column, and the lower

diagram shows the TIC (total ion chromatograms) obtained from the 2nd column. The data were obtained by using four analysis conditions (listed in Table 1). The unresolved C12ME elution band was transferred to the 2nd column according to the cut time specified in the methods. TIC (a), (b) and (c) in Fig. 3 (lower diagram) were obtained by using analysis conditions (a), (b) and (c), where the oven temperature programming for the 1st and 2nd columns were identical. Under these conditions, C12ME could not be fully separated from the co-eluting hydrocarbons matrix components. This is because C12ME eluted close to the highest density of unresolved hydrocarbons complex mixture region of the diesel sample on a non-polar column. This demonstrates that it is very difficult to optimize the method to achieve the desired 2D separation for a real complex sample, by using a single-oven MDGC system.

On the other hand, TIC (d) in Fig. 3 shows that a perfect separation could be obtained by using a dual-oven system with independent oven temperature programming (see Table 1, conditions (d) for analysis conditions).

Another advantage of using a dual-oven MDGC system for this application is that the high boiling point components from biodiesel blends could be baked out from the non-polar column by using higher temperatures in the 1st oven (> 300°C), while keeping the polar column below its temperature limit (normally 230°C), thus preventing unnecessary column damage in the 2nd oven.

Conclusions

Analysis of C12ME in low concentration biodiesel blends such as B1 or B2 is a great challenge due to the complexity of the sample matrix. Conventional 1D or even single-oven 2D separation could not provide sufficient resolution for the determination of C12ME content using a universal GC detector such as FID. However, it could be well separated by using a dual-oven MDGC system without sample preparation.

Table 1. Analysis Conditions of MDGC

	Conventional MDGC (0.25mm I.D. columns)			
Analytical conditions	a	b	C	d
1st instrument				
1st Column				
type	Rtx-5SilMS; non-polar (Resteks Corp., USA)		Rtx-1; non-polar (Resteks Corp., USA)	
dimension	15 m x 0.25 mm l.D., df=0.1 μm		30 m x 0.25 mm l.D., df=0.25 μm	
Oven temperature				
Initial temperature	50°C (0 min)		70°C (0 min)	
Program 1	50°C @ 20°C/min ~ 230°C (13 min)	50°C @ 10°C/min ~ 230°C (11 min)	70°C @ 10°C/min ~ 250°C (13 min)	70°C @ 10°C/min ~ 220°C (0 min)
Program 2				220°C@35°C/min ~ 300°C (6 min/11 min)
Carrier gas	He			
flush inlet pressure	190 kPa		190 kPa	
Switching gas	He, 1	70 kPa	He, 120 kPa	
heart cut time	7.97 ~ 8.11 min	12.43 ~ 12.58 min	13.74 ~13.94 min	
INJ temperature	250°C			
FID temperature	320°C			
Injection				
mode	Split			
split ratio	200		100	
volume	0.2 µL			
2nd instrument				
2nd Column				
type	Stabilwax; polar (Resteks Corp., USA)			
dimension	30 m x 0.25 mm l.D., df=0.25 μm			
Oven temperature			T	
Initial temperature	50°C (0 min)	50°C (0 min)	70°C (0 min)	50°C (15 min)
	50°C @ 20°C/min ~ 230°C	50°C @ 10°C/min ~ 230°C	70°C @ 10°C/min ~ 250°C	50°C @ 30°C/min ~ 220°C (2
Program 1	(13 min)	(11 min)	(13 min)	min)
GC I/F temperature		2	30°C	
DET, qMS			5000	
I/f temperature	250°C			
IS, temperature	EI, 200°C			
Acquisition	Scan, M/Z 40~350			
Scan speed	1666 amu/sec			

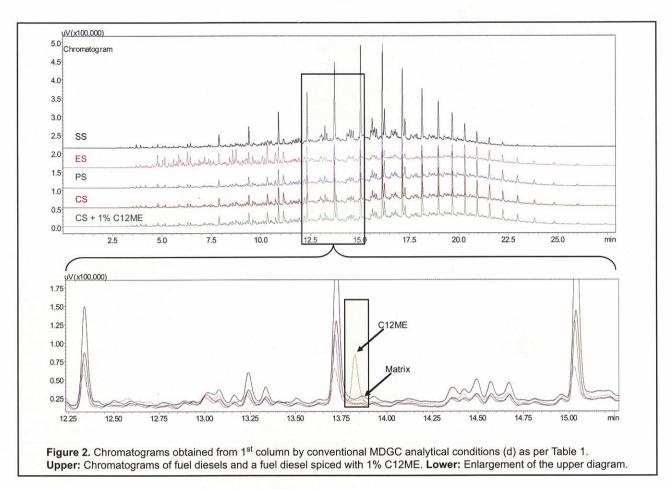
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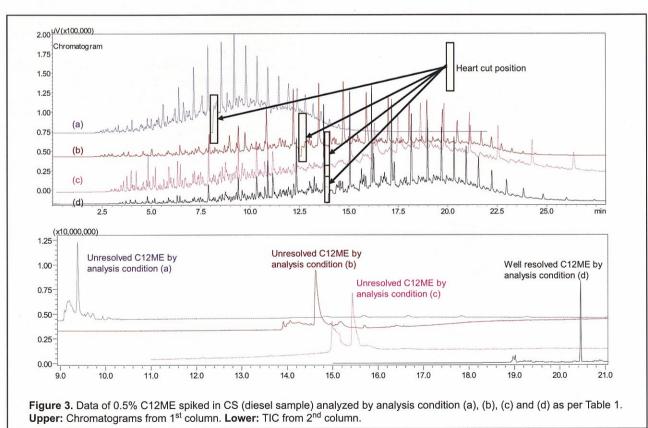
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