

Article

Optimization of Programmed Temperature Vaporization Injection for Determination of Polycyclic Aromatic Hydrocarbons from Diesel Combustion Process

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Abstract: In this study, programmed temperature vaporization in the solvent vent mode (PTV-SV) of gas chromatography-mass spectrometry was optimized and validated for the analysis of particle-phase and gas-phase polycyclic aromatic hydrocarbons from diesel engine combustion. Because of the large number of experimental and response variables to be studied, central composite inscribed design was used to optimize the PTV-SV injection factors, including initial inlet temperature, vaporization flow and time. The optimized PTV-SV method was validated by linearity, accuracy and sensitivity. For the 16 Polycyclic aromatic hydrocarbons (PAHs) studied, the correlation coefficients for the calibration plots of peak areas versus concentrations (0.5–300 ng mL⁻¹) ranged from 0.9812–0.9998. Limits of detection ranged from 0.016–20,130.375 ng mL⁻¹, and limits of quantification ranged from 0.055–1.25 ng mL⁻¹. The optimized method was used for the analysis of real samples collected from a diesel engine, which included particle-phase and gas-phase PAHs. The results showed that the improved PTV-SV method was satisfying for simultaneously identifying and quantifying PAHs produced during diesel combustion.

Keywords: programmed temperature vaporization; solvent vent mode; polycyclic aromatic hydrocarbon; optimization; diesel combustion

1. Introduction

Polycyclic aromatic hydrocarbons (PAHs) are organic compounds which contain two or more aromatic rings. Due to their known or potential mutagenic and carcinogenic properties, PAHs have received more attention. Diesel engines are one of the contributors to the PAHs in atmospheric environment. The diesel-generated PAHs mainly originate from the incomplete combustion of fuel [1,2], and the variety and quantity of PAHs are affected by engine working conditions and fuel properties [3,4]. In addition, they also play important roles in the formation of soot, another undesirable combustion byproduct [3–8]. Therefore, the identification of PAHs is essential for studies on the formation and evolution of diesel pollutants.

Combustion-generated PAHs are generally measured by gas chromatography-mass spectrometry (GC-MS) [2,9,10]. Diesel engine combustion involves complicated physical and chemical processes accompanied by high temperature and pressure. In such processes, the quantity of PAHs is usually a



few micrograms [11]. Wide varieties of intermediate products are also formed during the combustion process and interfere with the analysis of PAHs, complicating PAH identification and quantification. Therefore, using GC-MS to identify most of the PAHs in diesel combustion samples, with low detection and quantification limits, is challenging.

The programmed temperature vaporization (PTV) technique can be widely applied to improve sensitivities for GC-MS identification of trace analytes. PTV not only increases the method sensitivity through the larger volume injection of the final extract, it also simplifies and/or improves offline sample pretreatment procedures, reducing sample loss [12,13]. There are several other advantages to using the PTV technique, such as a decrease in analyte discrimination, improved transfer of thermodegradable compounds and better adaptability to relatively dirty samples [14].

The PTV injector on the GC-MS can be used in several different modes, including splitless injection, pulsed splitless, solvent vent, vapor overflow and solid-phase extraction-thermal desorption [15]. In the classical injection modes, such as split, splitless and pulsed splitless, the maximum injection volume is only approximately 2 μ L [16,17]. By contrast, the programmed temperature vaporization in the solvent vent mode (PTV-SV) significantly increases the sample volume to several hundred μ L and simultaneously improves sensitivity by controlling the time when the split exit is open, as well as the temperature of the inlet port [18]. Because of the numerous experimental variables in the PTV system, an experimental optimization of parameters is needed for different analysis types.

To date, PTV-GC-MS has been successfully applied to the quantitative analysis of PAHs in gasoline samples [19], ambient aerosol samples [20,21], foodstuffs [13] and marine sediments [22]. Literatures on the PAH analysis using PTV-SV-GC-MS is limited in number. The PTV injection system involves many parameters that have complex effects on analyte responses [16]. To improve sensitivity, it is necessary to identify the optimum combination of parameters. The primary aim of this study was to optimize and validate the PTV-SV injection method for the analysis of 16 PAHs produced from diesel engine combustion. Based on literature reviews [21] and our previous experience, three parameters were investigated, including initial inlet temperature (T_A), vaporization flow (F_B) and time (t_C). Optimizations of these three factors for PTV-SV injection were performed using the central composite inscribed (CCI) design. Different performance parameters of the PTV-SV-GC-MS method, such as the linearity, accuracy and sensitivity, were calculated. Analyses of samples from the diesel combustion process were conducted using the optimized method.

2. Materials and Methods

2.1. Preparation of the Real Sample

A real sample was used for validating the optimized programmed temperature vaporization in the solvent vent mode-gas chromatography-mass spectrometry-selected ion monitoring mode (PTV-SV-GC-MS-SIM) method. This real sample was obtained from a total cylinder sampling system of a diesel engine. The diesel engine was operated at an engine speed of 1000 rpm and a fuel-air equivalence ratio of 0.41. A detailed description of the total cylinder sampling system has been reported in the literature [23]. During sampling procedure, an aluminum alloy diaphragm was used to seal the engine cylinder head, making it a sampling valve. At a pre-set crank angle during the sampling cycle, the aluminum alloy diaphragm was instantly cut using an electromagnetic actuated tube-cutter. The cylinder contents rapidly exited from the cylinder into a sampling bag. Simultaneously, the samples were quenched and diluted by mixing with high-pressure nitrogen to achieve a temperature below 52 °C. The cylinder contents collected in the sampling bag were forced to flow through a Teflon®(Polytetrafluoroethylene (PTFE))-coated glass fiber filter (Pall, Ann Arbor MI, USA) inserted into the gas pipe. Particulates deposited on the filter were used for the analysis of particle-phase PAHs. A polyurethane foam (PUF)/XAD/PUF "sandwich" cartridge (SUPELCO, Bellefonte, PA, USA) was placed behind the Teflon®-coated glass fiber filter for sampling gas-phase PAHs. After sampling, the filter and PUF/XAD/PUF cartridge used were immediately Soxhlet extracted

using 60 and 120 mL dichloromethane (DCM), respectively, for 24 h. The DCM used was HPLC-MS grade from Dikma (Beijing, China). By rotary film and vortex evaporation, the resulting extract from the particle-phase sample was concentrated to 10 mL, and the one from the gas-phase was concentrated to 100 mL. The concentration process was performed under nitrogen. The concentrated extracts were stored in sealed bottles at -20 °C in the dark.

2.2. GC-MS Conditions

PAH analyses were performed on an Agilent 7890A gas chromatograph (Agilent Technologies, Santa Clara, California, USA) interfaced to a 5975C mass spectrometer detector (Agilent Technologies, Santa Clara, California, USA) operated in the electron ionization mode. This system was equipped with a PTV injector (Option 130, Agilent Technologies, Santa Clara, California, USA), and the injections were automatically carried out using an ALS 7683B autosampler (Agilent Technologies, Beijing, China). In this study, the maximum volume for the inlet liner is 50 μ L. In addition, too much injection will result in a contamination of the chromatographic column. Therefore, the injection volume was chosen as 25 μ L. The 25 μ L total injection volumes were achieved by five separate injections of 5 μ L at a maximum injection speed of 100 μ L s⁻¹. The GC oven temperature program used for the PTV injection was: The initial temperature of 30 °C was held for 7.5 min, the temperature was increased at a rate of 30 °C min⁻¹ to 280 °C, where it was held for 8 min, and then the temperature was increased at a rate of 60 °C min⁻¹ to 300 °C, where it was held for 6 min. Identification of the target compounds was based on the detection of the corresponding molecular ion and comparisons of retention times with those of relevant PAH standards. Quantification of the PAHs was performed using the SIM mode. The ionization voltage, transfer line temperature and ion source temperature were 70 eV, 280 °C and 280 °C, respectively. All analyses were performed using a capillary column (Agilent HP-5ms, 5% phenyl methyl siloxane, 30 m \times 250 µm \times 0.25 µm film thickness) and helium (99.9995 %) was employed as the carrier gas.

2.3. Calibration

A United States Environmental Protection Agency (USEPA) 16-PAH mixed standard solution (AccuStandard, New Haven, Connecticut, USA), including naphthalene (Nap), acenaphthylene (AcPY), acenaphthene (AcP), fluorene (Flu), phenanthrene (Phe), anthracene (Ant), fluoranthene (FL), pyrene (Pyr), benzo(a)anthracene (BaA), chrysene (Chr), benzo(b)fluoranthene (BbFL), benzo(k)fluoranthene (BkFL), benzo(a)pyrene (BaP), indeno(1,2,3-cd)pyrene (InP), dibenzo(a,h)anthracene (DBA) and benzo (g,h,i)perylene (BghiP), was used for calibration. Standard working solutions for calibration plots were freshly prepared by diluting the stock standard solution in DCM. The 16-PAH mixed standard solution, with an initial concentration of 200 × 103 ng mL⁻¹, was diluted to 300, 200, 50, 10, 5 and 0.5 ng mL⁻¹. All solutions were stored in capped amber vials at -20 °C. The gas chromatography analysis was carried out for these diluted standard solutions using the optimized PTV-SV-GC-MS-SIM method. Calibration curves were prepared using the PAH concentrations and corresponding peak areas. PAH quantification for the real samples could then be performed using the calibration curves.

2.4. Method Validation

The PTV-SV-GC-MS method was validated using linearity, accuracy and sensitivity. The linearity of the method was determined using calibration plots obtained from three replicate analyses of standard solutions with concentrations of 0.5, 5, 10, 50, 200 and 300 ng mL⁻¹. The accuracy of the method was expressed as the ratio of the theoretical value to the average measured value for three concentrations of authentic standards added to the samples. The average recovery of the PAHs was calculated using calibration plots. The limits of detection (LODs) and quantification (LOQs) were calculated based on signal-to-noise ratios of 3:1 and 10:1, respectively, for both the PTV-GC-MS and conventional splitless GC-MS methods. To evaluate the precision of the method, mixed standard solutions (100 ng mL⁻¹)

were added to blank samples and extracted using the sample preparation procedure described in Section 2.1.

3. Results and Discussion

3.1. Optimization of the PTV-SV Injection

To determine the maximum values for the PTV-SV factors, CCI design was performed to estimate the response surfaces. This design used factor settings as the starting points and created factorial or fractional factorial design within the limits. Based upon the previous study [21] and our experience, T_A , F_B and t_C were evaluated for their effects on PTV-SV injection. In this study, each CCI design model selected was composed of a two-cubed full factorial design that included eight cubed points, where six axial and six central points were added, for a total of 20 runs. The 16 PAH peak areas were set to be dependent variables and the three factors were used as input variables. The three factors and their levels are listed in Table 1. A 16-PAH mixed standard solution at a concentration of 100 ng mL⁻¹ was used for these experiments. The experimental conditions designed and responses (chromatographic peak areas) obtained are presented in Table 2. As an example, Figure 1 shows the influence of T_A , F_B and t_C on response surfaces for Phe (25 µL PTV-SV injection). It is obvious that the Phe response areas increase with an increase of T_A , and decrease with decreases of F_B and t_C . Close inspections of Figure 1 and Table 2, however, show that the quantity of experimental data to be analyzed is considerably large when three parameters are optimized simultaneously. Thus, a statistical analysis method is used in the data process to improve the reliability of these results.

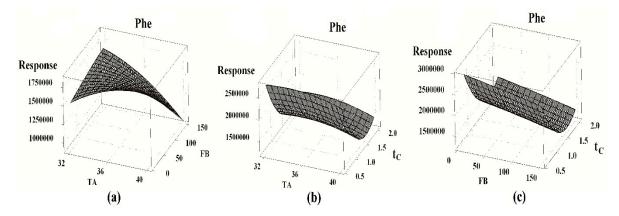


Figure 1. Influence of initial inlet temperature (T_A), vaporization flow (F_B) and vaporization time (t_C) on response surfaces for 25 µL Phe injection: (**a**) Influence of T_A and F_B on response surfaces at $t_C = 1.32$ min; (**b**) Influence of T_A and t_C on response surfaces at $F_B = 80$ mL min⁻¹; (**c**) Influence of F_B and t_C on response surfaces at $T_A = 36$ °C.

Table 1. Factors and corresponding levels for the programmed temperature vaporization (PTV) injection systems.

F (Levels ¹		
Factor	-1	$-1/\alpha$	0	+1/α	+1
Inlet Temperature (°C)	32	34	36	38	40
Vaporization Flow (mL min ⁻¹)	10	38	80	122	150
Vaporization Time (min)	0.6	0.89	1.32	1.75	2.04

¹ The α value is 1.68.

T_A (°C	2)	34	38	34	38	34	38	34	38	32	40	36	36	36	36	36	36	36	36	36	36
<i>F_B</i> (mL m	in ⁻¹)	38	38	122	122	38	38	122	122	80	80	10	150	80	80	80	80	80	80	80	80
t_C (mi	n)	0.89	0.89	0.89	0.89	1.75	1.75	1.75	1.75	1.32	1.32	1.32	1.32	0.6	2.04	1.32	1.32	1.32	1.32	1.32	1.32
Analytes	m/z		Responses (chromatographic peak areas $\times 10^{6}$)																		
Nap	128	0.44	0.40	0.42	0.33	0.41	0.34	0.42	0.36	0.39	0.45	0.57	0.45	1.07	0.36	0.42	0.36	0.51	0.31	0.53	0.36
AcPY	152	0.81	0.54	0.46	0.40	0.57	0.32	0.35	0.35	0.5	0.40	1.06	0.39	1.51	0.37	0.51	0.43	0.49	0.25	0.51	0.41
AcP	153	0.72	0.48	0.38	0.32	0.48	0.26	0.28	0.27	0.41	0.30	0.94	0.30	1.39	0.29	0.42	0.35	0.39	0.21	0.39	0.33
Flu	166	1.08	0.98	0.89	0.63	0.86	0.65	0.61	0.49	0.77	0.65	1.20	0.65	1.82	0.55	0.79	0.65	0.81	0.51	0.83	0.64
Phe	178	1.94	2.02	1.97	1.42	1.78	1.59	1.54	1.23	1.57	1.66	1.82	1.59	3.29	1.33	1.72	1.37	1.85	1.29	1.91	1.42
Ant	178	2.38	2.38	2.42	1.95	2.27	1.99	2.05	1.77	2.05	2.15	2.21	2.04	3.71	1.87	2.24	1.84	2.31	1.67	2.38	1.95
FL	202	2.50	2.69	2.74	2.24	2.46	2.30	2.55	2.14	2.21	2.65	2.16	2.50	4.11	2.14	2.52	1.97	2.71	1.97	2.83	2.18
Pyr	202	2.71	2.92	2.92	2.40	2.66	2.48	2.71	2.25	2.39	2.84	2.35	2.69	4.42	2.29	2.69	2.13	2.92	2.11	3.03	2.35
BaA	228	1.11	1.33	1.45	1.19	1.17	1.15	1.52	1.29	1.05	1.55	0.88	1.39	2.34	1.17	1.32	0.89	1.44	0.97	1.58	1.08
Chr	228	2.33	2.89	3.07	2.63	2.63	2.60	3.21	2.61	2.36	3.25	2.09	3.01	4.28	2.61	2.64	1.92	3.09	2.06	3.22	2.25
BbFL	252	0.94	1.21	1.30	0.98	0.97	1.07	1.39	1.09	0.86	1.39	0.71	1.25	2.07	0.98	1.11	0.74	1.30	0.93	1.42	0.91
BkFL	252	2.08	2.40	2.57	2.17	2.13	2.10	2.72	2.33	1.91	2.71	1.62	2.48	3.95	2.21	2.35	1.66	2.57	1.81	2.77	1.95
BaP	252	0.89	1.15	1.23	1.07	1.00	1.09	1.31	1.03	0.94	1.3	0.75	1.29	2.05	1.04	1.02	0.76	1.21	0.85	1.49	0.96
InP	276	0.54	0.78	0.84	0.56	0.57	0.68	0.91	0.66	0.48	0.88	0.39	0.79	1.40	0.55	0.65	0.40	0.81	0.57	0.89	0.5
DBA	278	1.27	1.48	1.59	1.25	1.31	1.27	1.65	1.53	1.09	1.68	0.86	1.62	2.47	1.17	1.37	0.86	1.53	1.05	1.65	1.09
BghiP	276	1.21	1.54	1.59	1.27	1.25	1.34	1.78	1.31	1.08	1.51	0.90	1.54	2.54	1.25	1.46	0.88	1.57	1.06	1.74	1.14

Table 2. Experimental conditions designed and responses (chromatographic peak areas $\times 10^6$) obtained using the CCI design.

The data from the 25 μ L injection volume are fitted to Equation (1) using a second-degree polynomial equation based on the least squares statistical method.

$$Y = b_0 + b_A T_A + b_B F_B + b_C t_C + b_{AB} T_A F_B + b_{AC} T_A t_C + b_{BC} F_B t_C + b_{AA} T_A^2 + b_{BB} F_B^2 + b_{CC} t_C^2$$
(1)

where T_A , F_B and t_C are the initial inlet temperature, the vaporization flow and the vaporization time, respectively, and b_i , b_{ii} and b_{ii} are the fitting parameters.

The statistical significances of the regression coefficients were determined using the t-test (only significant coefficients with p-value< 0.05 were included). Statistical analysis of the fitted model equation above was checked using analysis of variance (ANOVA) at the 5% significance level. The ANOVA, regression coefficients and their significances in the fitted model are listed in Table 3.

The response optimizer in Minitab 15 (Minitab Inc., USA) was applied to maximize the composite desirability for the 16 PAHs. The optimization of composite desirability was performed using the Derringer's desirability function in the Minitab optimizer [21]. The desirability was set to 0.0 and 1.0 for the lowest and highest responses, respectively, from the CCI designs. Because all the responses had the same importance, a composite desirability was obtained by calculating the geometric average desirability values for the 16 PAHs. The final optimized factors, results from individual desirability for the PAHs and the composite desirability, are provided in Table 4. The samples are introduced into the liner at an initial temperature of 38 $^{\circ}$ C for the 25 μ L injection. The solvent is then vaporized for 0.6 min with the PTV injector temperature raised to 300 °C at a rate of 600 °C min⁻¹ and vented through the split valve at a flow rate of 10 mL min⁻¹. During the cleaning phase, the split valve keeps open for 5 min (50 mL min⁻¹) at a temperature of 300 °C for all the experiments. As an example, Figure 2 shows a PAH standard chromatogram at a concentration of 100 ng mL⁻¹, where the separation of the 16 PAHs is achieved in a total analysis time of 30 min under the optimized PTV-SV injection conditions. The selected conditions for GC-MS in SIM mode are presented in Table 5. Validation of the optimum method from response surface design experiments was determined from six replicate measurements of the 100 ng mL⁻¹ standard solution under the final optimized conditions. The values of relative standard deviation (RSD) for the 16 PAHs range from 1.12%-6.07%, as listed in Table 6.

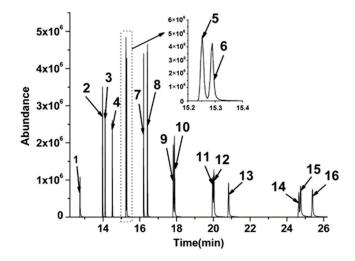


Figure 2. Gas chromatograph mass spectrometry (GC-MS) chromatogram of a 100 ng mL⁻¹ polycyclic aromatic hydrocarbon (PAH) standard sample after optimization. 1. Nap; 2. AcPY; 3. AcP; 4. Flu; 5. Phe; 6. Ant; 7. FL; 8. Pyr; 9. BaA; 10. Chr; 11. BbFL; 12. BkFL; 13. BaP; 14. InP; 15. DBA; 16. BghiP.

	Naj	р	AcP	Ŷ	Ac	P	Flu	1	Pho	e	An	t	FL		Py	r
Terms	Estimate	Р	Estimate	Р	Estimate	Р	Estimate	Р	Estimate	Р	Estimate	Р	Estimate	Р	Estimate	Р
Intercept	429023	< 0.01	500533	< 0.01	358092	< 0.01	699564	< 0.01	1597255	< 0.01	2114094	< 0.01	2428690	< 0.01	2622292	< 0.01
T_A	-11370	0.792	-70979	0.046	-52280	0.022	-65125	0.029	-60225	0.05	-62307	0.441	-10133	0.916	-13937	0.893
F_B	-20222	0.64	-132899	0.013	-129867	< 0.001	-139076	< 0.001	-113585	0.016	-81749	0.318	22198	0.817	6284	0.951
t_C	-91661	0.053	-185495	0.006	-180795	< 0.001	-227695	< 0.001	-330210	0.001	-304792	0.003	-295329	0.01	-324928	0.009
$F_B \times F_B$	-3421	0.935	49839	0.444	49961	0.05	39763	0.034	-5003	0.199	-21732	0.820	-72658	0.444	-79267	0.437
$t_C \times t_C$	68448	0.124	127019	0.142	126211	0.022	132453	0.026	16836	0.614	213158	0.026	228880	0.045	216756	0.051
$T_A \times F_B$	-4572	0.914	43070	0.277	48427	0.164	-7668	0.886	-98224	0.047	-58493	0.577	-118389	0.356	-124337	0.367
ANOVA of	F	Р	F	Р	F	Р	F	Р	F	Р	F	Р	F	Р	F	Р
Model	5.94	0.025	9.78	0.001	4.84	0.025	6.07	0.013	6.00	0.027	11.20	0.004	3.89	0.065	6.25	0.023
Ŧ	Ba	ł	Ch	r	BbB	Ľ	BkF	Ľ	Bal	P	InI	þ	DB	A	Bgh	iP
Terms	Estimate	Р	Estimate	Р	Estimate	Р	Estimate	Р	Estimate	Р	Estimate	Р	Estimate	Р	Estimate	Р
Intercept	1313721	< 0.01	2811816	< 0.01	1123014	< 0.01	2316300	< 0.01	1119073	< 0.01	684561	< 0.01	1385816	< 0.01	1295837	< 0.01
T_A	40445	0.556	71199	0.505	45982	0.487	63193	0.542	37583	0.534	36277	0.477	50330	0.529	25631	0.723
F_B	114236	0.116	191730	0.092	109445	0.116	185456	0.094	104720	0.103	78600	0.141	144740	0.09	123600	0.109
t_C	-140096	0.061	-195245	0.087	-128297	0.072	-210611	0.062	-118243	0.07	-97184	0.076	-147187	0.085	-154754	0.052
$F_B \times F_B$	-87526	0.205	-131722	0.218	-69140	0.291	-129254	0.214	-61606	0.303	-44688	0.372	-66594	0.396	-41246	0.56
$t_C \times t_C$	131432	0.069	184427	0.095	122186	0.077	233996	0.037	123215	0.055	90884	0.086	140501	0.091	197852	0.016
$T_A \times F_B$	-85210	0.349	-196820	0.174	-124487	0.166	-133386	0.332	-99542	0.221	-108560	0.122	-77806	0.458	-150545	0.132
ANOVA of Model	F 5.07	P 0.038	F 2.83	P 0.112	F 4.42	Р 0.051	F 5.13	P 0.038	F 4.13	P 0.058	F 4.09	Р 0.059	F 5.17	Р 0.037	F 9.01	P 0.008

Table 3. Analysis of Variance (ANOVA), regression coefficients and their significances in the fitted model of the 16 PAH peak areas.

FesterJulat Terr		° C)			Opti	mum		
FactorInlet Ten	nperature (()	38					
Vaporization Flo	ow (mL mi	n ⁻¹)			1	0		
Vaporization			0	.6				
Analytes	Nap	AcPY	AcP	Flu	Phe	Ant	FL	Pyr
Desirability (%)	0.60	0.77	0.89	0.97	1	0.91	0.99	0.85
Analytes	BaA	Chr	BbFL	BkFL	BaP	InP	DBA	BghiF
Desirability	0.69	0.88	0.83	0.87	0.77	0.89	0.74	0.97
Ċ	omposite I	Desirability				0.	85	

Table 4. Optimized factor, individual and composite desirability for the 16 PAHs.

Table 5. GC-MS-SIM mode conditions for the studied PAHs.

Analytes	Quantification ion (m/z)	Retention Time (min)	Analytes	Quantification ion (m/z)	Retention Time (min)
Nap	128	12.75	BaA	228	17.87
AcPY	152	13.98	Chr	228	17.92
AcP	153	14.16	BbFL	252	19.96
Flu	166	14.51	BkFL	252	20.05
Phe	178	15.24	BaP	252	20.68
Ant	178	15.29	InP	276	24.66
FL	202	16.23	DBA	278	24.75
Pyr	202	16.41	BghiP	276	25.40

Table 6. Repeatabilities for the PTV method.

NO.	Analytes	RSD (%)	NO.	Analytes	RSD (%)
1	Nap	6.07	9	BaA	1.8
2	AcPY	2.85	10	Chr	1.22
3	AcP	2.99	11	BbFL	1.35
4	Flu	2.08	12	BkFL	1.64
5	Phe	1.34	13	BaP	2.11
6	Ant	1.22	14	InP	1.93
7	FL	1.19	15	DBA	3.77
8	Pyr	1.12	16	BghiP	3.89

3.2. Method Validation

Using the SIM mode, the linearity of the injection method was determined from calibration plots obtained for three replicate analyses with standard solutions of 0.5, 5, 10, 50, 200 and 300 ng mL⁻¹. The coefficient of determination (R^2) for the 16 PAHs were obtained using linear least-squares regression. The correlation coefficients of the calibration plots for the 16 PAHs range from 0.9812–0.9998. The ANOVA of the calibration plots shows that there are no significant differences between the slopes of standard plots (p-value > 0.05). The results in Table 6 indicate that there are good linear relationships between peak areas and concentrations in the 0.5–300 ng mL⁻¹ concentration range.

To evaluate the LOD and LOQ, the 0.5 ng mL⁻¹ standard solution was injected and measured. The results in Table 7 show that for the 25 μ L PTV injection, the LODs and LOQs range from 0.016–0.375 ng mL⁻¹ and 0.055–1.25 ng mL⁻¹, respectively. Under the same GC and MS conditions, 2 μ L standard solution (0.5 ng mL⁻¹) was analyzed using the conventional splitless-GC-MS-SIM method. From Table 7, it can be seen that the LODs and LOQs of the splitless-SIM method range from 0.256–11.811 ng mL⁻¹ and 0.853–39.370 ng mL⁻¹, respectively, much higher than those of the PTV-SIM method.

		PTV-SV-GC-MS-SIM N	lethod			Splitless-GC-M	IS-SIM Method
Analytes	Regression Equation y = ax + b	Coefficient of Determination (R ²)	Linear Range (ng mL ⁻¹)	LOD (ng mL ⁻¹)	LOQ (ng mL ⁻¹)	LOD (ng mL ⁻¹)	LOQ (ng mL ⁻¹)
Nap	y = 27455340x - 288480	0.9949	0.5–300	0.082	0.272	0.592	1.974
AcPY	y = 47113309x - 617805	0.9923	0.5-300	0.038	0.126	0.344	1.148
AcP	y = 39117183x - 355452	0.9957	0.5-300	0.049	0.162	1.322	4.405
Flu	y = 37767418x - 335743	0.9975	0.5-300	0.072	0.242	1.948	6.494
Phe	y = 60532000x - 696491	0.9964	0.5-300	0.023	0.076	0.596	1.988
Ant	y = 61253701x - 357054	0.9987	0.5-300	0.038	0.126	1.020	3.401
FL	y = 64609682x - 547300	0.9976	0.5-300	0.016	0.055	0.256	0.853
Pyr	y = 68056260x - 518969	0.998	0.5-300	0.032	0.106	0.587	1.957
BaA	y = 47673273x - 822250	0.9925	0.5-300	0.067	0.223	1.035	3.448
Chr	y = 57229694x - 250860	0.9998	0.5-300	0.066	0.221	0.959	3.195
BbFL	y = 40717605x - 783554	0.9895	0.5-300	0.375	1.25	1.676	5.587
BkFL	y = 59435860x + 444687	0.9993	0.5-300	0.326	1.087	1.255	4.184
BaP	y = 45868925x - 778709	0.9929	0.5-300	0.288	0.962	5.882	19.608
InP	y = 31717730x - 741731	0.9812	0.5-300	0.263	0.877	7.160	23.866
DBA	y = 63091354x - 1133627	0.9926	0.5-300	0.313	1.042	11.811	39.370
BghiP	y = 50702721x - 742759	0.9964	0.5–300	0.183	0.61	10.830	36.101

Table 7. Calibration plot regression equations, LODs and LOQs for the PTV and splitless methods.

In this study, the recoveries associated with the pre-treatment procedure of the samples were evaluated. A known amount of the analyses at each of three concentrations (100, 50 and 4 ng mL⁻¹) were added to the Teflon®-coated glass fiber filters and the PUF/XAD/PUF cartridges, respectively. The sample preparation procedure was described in Section 2.1, and three samples were prepared at each concentration. Figure 3 shows that the average recoveries range from 86.28%–108.35% for particle-phase PAHs and 71.83%–108.60% for gas-phase PAHs.

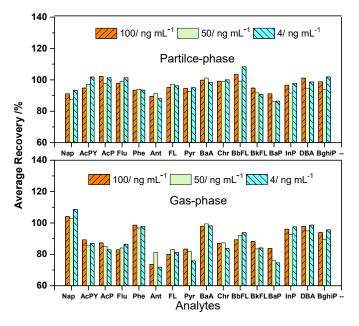


Figure 3. Average recoveries for particle-phase PAHs and gas-phase PAHs.

3.3. Analysis of the Real Sample

The optimized method with a PTV injection volume of 25 μ L was used for the analysis of the real sample obtained by the total cylinder sampling system of the diesel engine. Figures 4 and 5 show the total ion chromatograms in the SIM mode for the real particle-phase and gas-phase extracts, respectively. The 16 PAH peaks in the particle-phase extract are detected simultaneously and are well-separated, as shown in Figure 4. In the gas-phase extract, 10 PAHs are detected and separated. These phenomena indicate that the PTV-SV method can be used for the simultaneous identification and quantification of PAHs present in diesel combustion. Results obtained from the particle-phase and gas-phase extracts are listed in Table 8. The concentrations of the 16 PAHs in the particle-phase extracts range from 2.46–28.53 ng mL⁻¹. The concentrations of BbFL, BkFL, BaP, InP, DBA and BghiP are below their LOQs in the gas-phase sample.

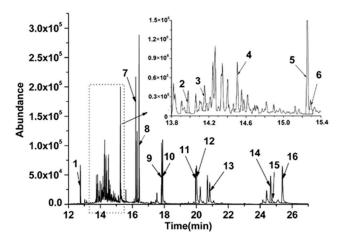


Figure 4. Chromatogram of the particle-phase extracts in SIM mode. 1. Nap; 2. AcPY; 3. AcP; 4. Flu; 5. Phe; 6. Ant; 7. FL; 8. Pyr; 9. BaA; 10. Chr; 11. BbFL; 12. BkFL; 13. BaP; 14. InP; 15. DBA; 16. BghiP.

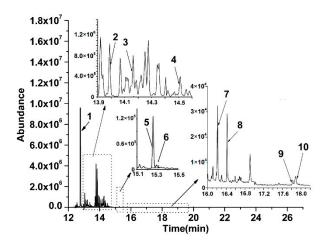


Figure 5. Chromatogram of the gas-phase extracts in SIM mode. 1. Nap; 2. AcPY; 3. AcP; 4. Flu; 5. Phe; 6. Ant; 7. FL; 8. Pyr; 9. BaA; 10. Chr.

A		Sample	(ng mL ⁻¹)	
Analytes	Particle-phase	RSD (%)	Gas-phase	RSD (%)
Nap	19.77	4.33	156.71	3.55
AcPY	5.11	5.77	23.42	4.84
AcP	5.36	3.57	21.55	5.54
Flu	11.18	11.47	15.31	1.48
Phe	18.29	12.29	5.79	2.79
Ant	2.46	13.44	0.58	12.76
FL	16.66	3.27	0.97	9.77
Pyr	21.38	4.13	0.74	8.98
BaA	23.58	12.06	0.23	13.41
Chr	21.42	11.73	0.41	10.97
BbFL	22.14	8.03	-	-
BkFL	18.98	11.53	-	-
BaP	16.57	10.12	-	-
InP	27.13	5.39	-	-
DBA	3.15	12.74	-	-
BghiP	28.53	3.72	-	-

Table 8. Concentrations of PAHs in the real sample.

4. Conclusions

The PTV-SV-GC-MS-SIM injection method has been optimized for the analysis of 16 PAHs using CCI design. The initial inlet temperature, vaporization flow and vaporization time are found to be statistically significant for the 25 μ L PTV injection volume. After the optimization of the PTV injection factors, the initial temperature, vaporization flow and vaporization time are determined to be 38 °C, 10 mL min⁻¹ and 0.6 min, respectively. Validation parameters for the optimized method, such as linearity, accuracy, LODs and LOQs, are satisfactory for the identification of the 16 PAHs. The PTV-SV method in the SIM mode is reliable for the simultaneous identification and quantification of particle-phase and gas-phase samples from diesel combustion.

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