

HD**Measurement of HD Concentration by Gas Chromatography**

, , , , , , , ,

150

(HD)

.

HD

가

HD

가

HD

가

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Abstract

Gas Chromatography has been used for the on-line measurement of deuterium hydride (HD) which is used in a Tritium Removal Pilot Facility for the demonstration of the removal of the tritium mainly generated in CANDU reactors. Two methods with different carrier gases, neon and hydrogen, are tested and compared each other. It was showed that both the mothods could be possible to measure the concentration of H₂ and HD. However, the method with a column packed with alumina showed difficulty in the application due to quite long measurement time and reproducibility. The other method using hydrogen as a carrier gas could measure the concentration accurately within comparably short period.

1.

가 가 가 .

(Tritium Removal Facility) . ,

가 가 ,

(HD) (H₂O)

(TRF) .

HD

1959 Gant Yang [2] Carter Smith [3, 4] (H),

(D) (T)

(retention time)

[5, 6, 7,

8].

Yamanishi [8]

manganese chloride (MnCl₂)

alumina JAERI ()

[9- 11]

manganese chloride

ortho para nuclear spin

[8].

[12]. ,

, MnCl₂

alumina

(-196)

(TCD)

H₂/HD

HD

2.

2.1

가

가

가

()

- 196

molecular

-sieve column trap

(,)

, (: 3mm, : 8m) (:

3mm, : 8m)

activated alumina 10%

manganese chloride(MnCl₂)가

(60-80 Mesh)

molecular sieve 5A (60-80

mesh) 5mm, 1m

150 24

TCD 180 mA 50

40

70ml/min.

D₂ 0.113%, 0.236%, 0.326%

H₂/D₂

가

HD

가

2

,

H₂/HD

가

H₂/HD

가

29

, 1

5

3

H₂

HD

가

HD

가

dsCHROM

, peak width peak threshold

HD

가

H₂

HD

width(=100)

peak threshold(=200)

HD

peak

2.2

, H₂

가

H₂

HD

HD

가 H₂

, molecular sieve

30

2

1

1.

Oven Temperature ()	90
Detector Temperature ()	120
Current (mA)	160
Carrier flow rate (Ml/min)	65
Reference flow rate (Ml/min)	85

3.

3.1

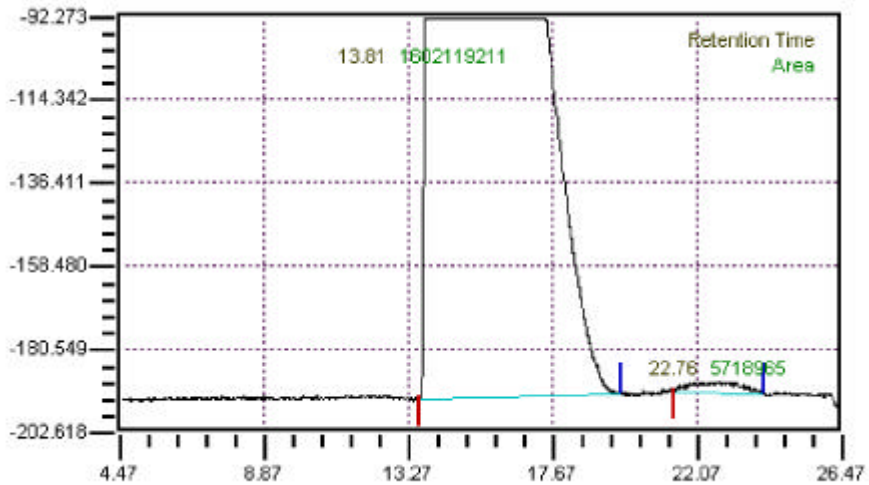
H₂/D₂ 가 HD

. 1

H₂ HD

D₂ . H₂ retention time 13.6 HD

22.7 retention time 가 .



1. H₂ HD

2가 D₂ H₂/HD

. 0.236% 5 , 0.326%

3 2 2 1 .

0.236% D₂/H₂ (5)

H ₂ (A)	HD (B)	B/A (%)	
1602119211	5718965	0.365	1
1605615103	5961152	0.371	
1607855996	5377988	0.334	
1614111488	5368637	0.332	
1607267792	5160538	0.321	
		0.3446	(m)
		0.0220	(s)

* 95% confidence interval, (0.3173, 0.3719)

$0.3446 \pm 2.776 * 0.022 / \sqrt{5} = 0.3446 \pm 0.0273$ (t- distribution, 4 degrees of freedom)

0.326% D₂/H₂ (3)

H ₂ (A)	HD (B)	B/A (%)	
1599509115	6547608	0.409	
1603680484	7036685	0.438	
1611229068	9491177	0.589	
		0.479	(m)
		0.0966	(s)

* 95% confidence interval, (0.1848, 0.7726)

$0.4787 \pm 4.303 * 0.0966 / \sqrt{2} = 0.4787 \pm 0.2939$ (t- distribution, 2 degrees of freedom)

0.326% D₂/H₂ (3)

H ₂ (A)	HD (B)	B/A (%)	
468386599	1876281	0.399	
1230803159	4628907	0.375	
1233936983	6041016	0.489	
		0.421	(m)
		0.060	(s)

* 95% confidence interval, (0.2384, 0.6036)

$0.421 \pm 4.303 * 0.060 / \sqrt{2} = 0.421 \pm 0.1826$ (t- distribution, 2 degrees of freedom)

0.326% D₂/H₂ (2)

H ₂ (A)	HD (B)	B/A (%)	
1020892553	4670801	0.457	
1147562284	5000774	0.436	
		0.447	(m)
		0.0148	(s)

가 HD
 , D₂ 가 D₂가 H₂
 HD
 H₂ HD
 1 30
 1 5 150
 0.326% base line , 3 2 2 1
 3 HD/H₂ (0.479, 0.471, 0.447),

3.2

HD 가 base line
 가 ,
 HD 가
 가 D₂가 HD
 HD 가 HD
 5가

가 5 H₂/D₂
 precision Gas Mass Spectrometer (Finnigan
 MAT 271) 2 ,
 H₂ (HD) 120 ppm, D₂ 가 HD 2300 ppm [1]

(0.23%) :

$$D(\%) = 0.0023 + \frac{0.9977 \times 0.00012 + 0.0023 \times 0.0023}{2} = 0.00236 = 0.236\%$$

2.

	(D %)	(D %)
1	0.23	0.236
2	0.32	0.326
3	0.74	0.747
4	14.85	14.9
5	20.95	21.0

GC

5 가 3 10
 2 . 10
 가 7
 가 4 5 10
 3 7 .
 2
 , multiple sample case
 (7)가 student t-distribution .

$$(m_1 - m_2) \pm t_{\frac{\alpha}{2}} S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}$$

$$S_p^2 = \frac{(n_1 - 1)S_1^2 + (n_2 - 1)S_2^2}{n_1 + n_2 - 2} : \text{pooled sample variance}$$

n_1, n_2

m_1, m_2

S_1, S_2

3.

	1 (x ₁)	2 (x ₂)	(x ₁ - x ₂)	
1	m ₁ = 91.6 S ₁ = 8.34	m ₂ = 87.3 S ₂ = 3.30	(-1.75, 10.35)	89.5
2	m ₁ = 127.1 S ₁ = 3.18	m ₂ = 116.0 S ₂ = 4.16	(7.58, 14.62)	121.6
3	m ₁ = 294.6 S ₁ = 8.34	m ₂ = 299.3 S ₂ = 4.15	(-0.9, 10.3)	297.0
4	m ₁ = 553.9 S ₁ = 1.66			553.9
5	m ₁ = 805.7 S ₁ = 1.89			805.7

가 1, 2, 3 2

1 3 2

가 2

.

2

[-1.44, 4.64],

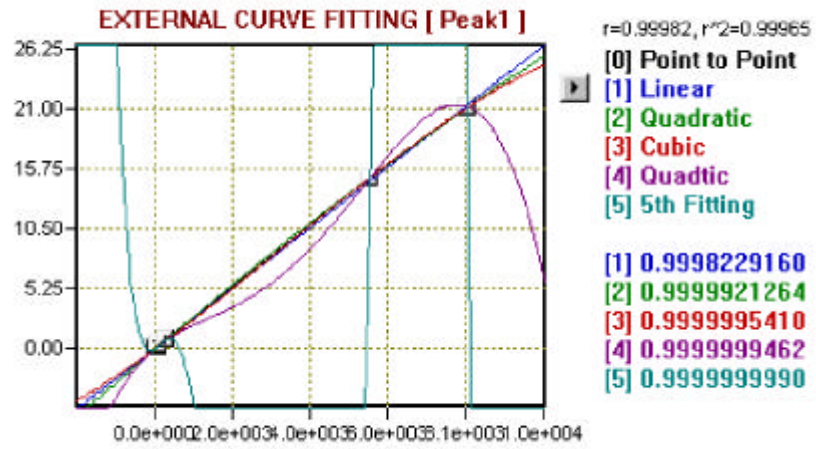
121.6 .

(2).

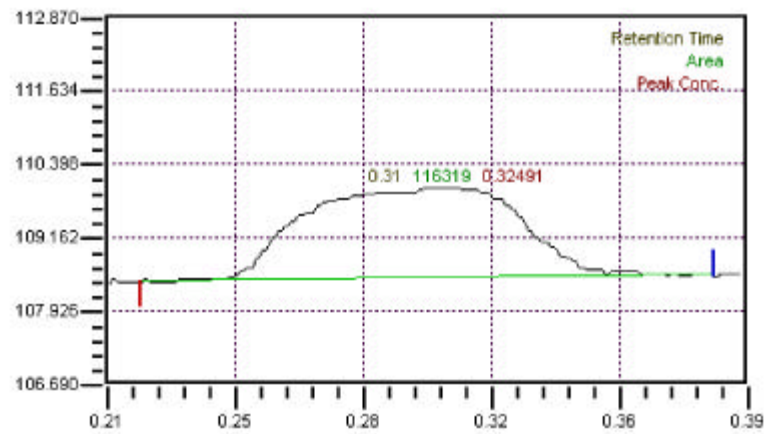
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3

HD



2.



3.

HD

4.

(HD)

(TRF)

HD

HD

가

base line

가

HD

가

HD

가

(2)

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