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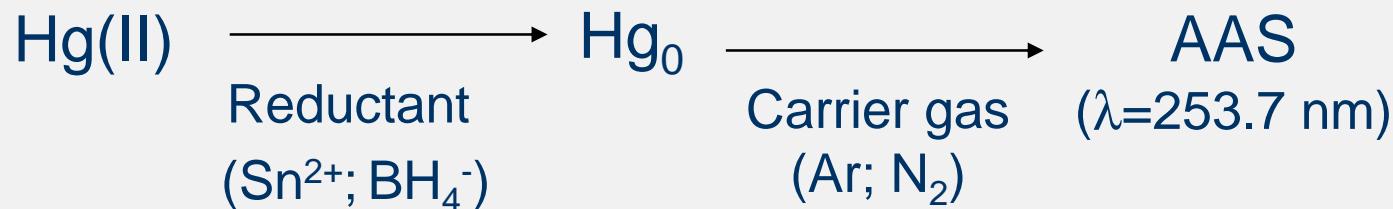
Exploiting the multicommutation approach for calibration using a single standard solution: application to the determination of mercury by CVAAS

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- Hatch and Ott, Anal. Chem. 40 (1968)
 - Standard method for Hg determination (EN 13806, AOAC)

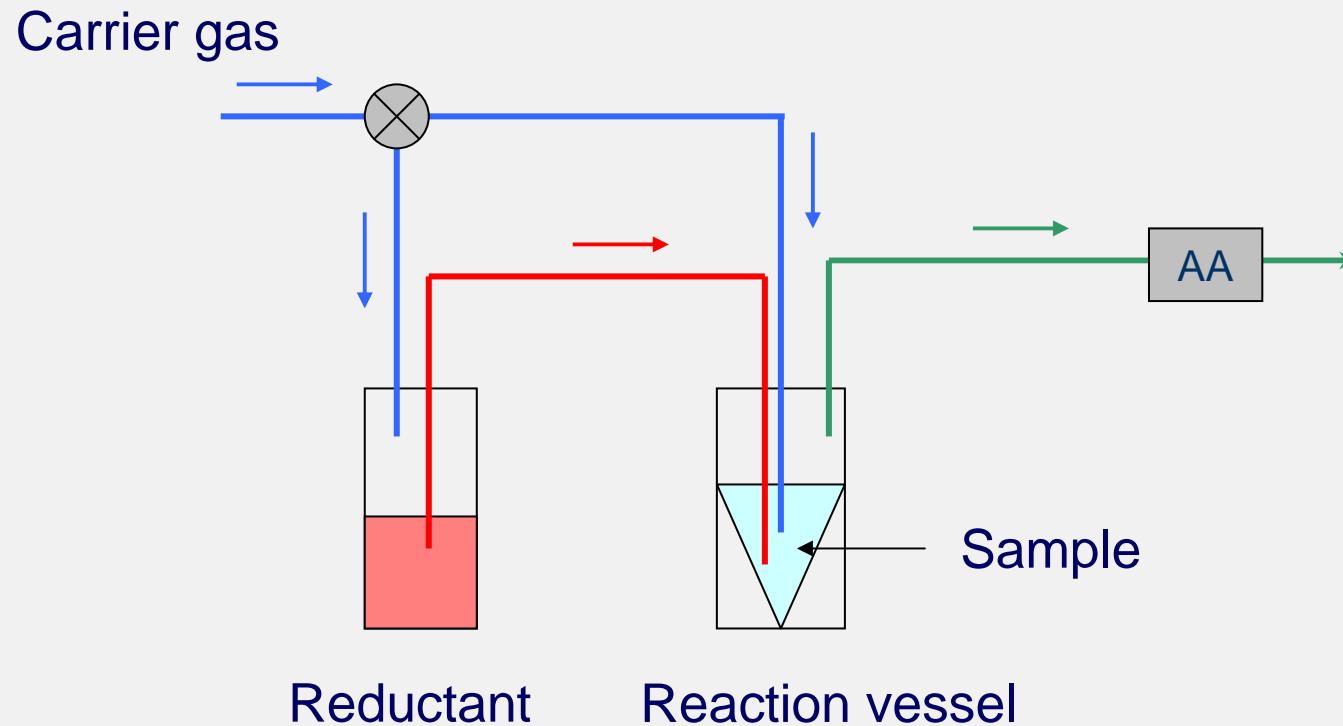


⇒ Batch systems

⇒ Flow systems

CVAAS - Batch system

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Limitations

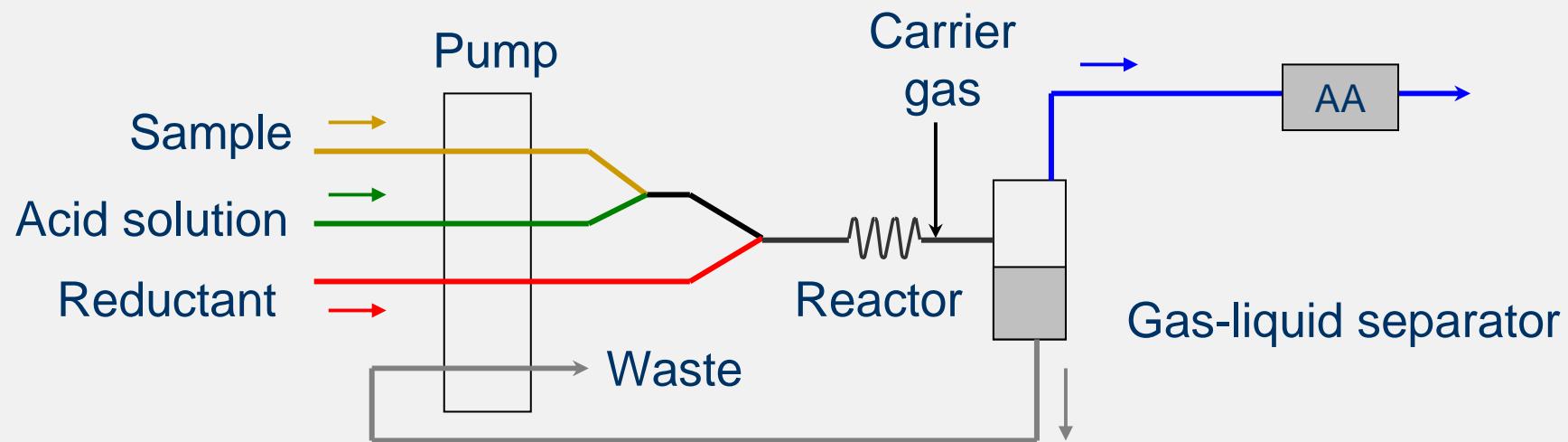
- high reagent/sample consumption
- time consumption

 Batch systems Flow systems

- Segmented flow analysis (SFA)
- Continuous sample flow

CVAAS - Continuous sample flow system

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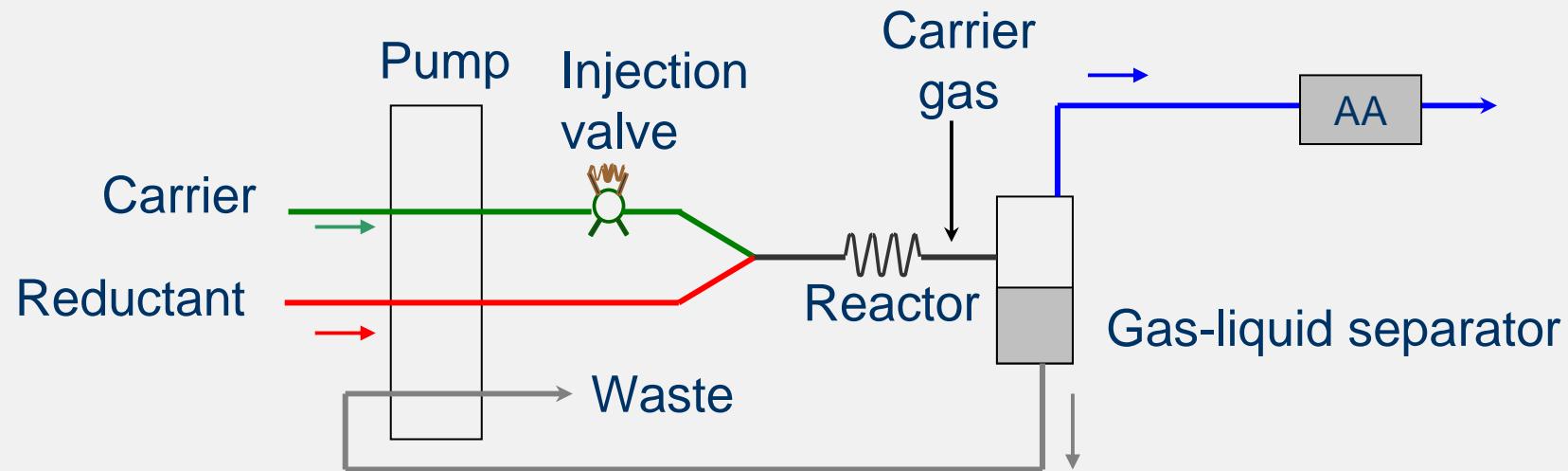
 Batch systems Flow systems

- Segmented flow analysis (SFA)
- Continuous sample flow
- Flow injection analysis (FIA)

Andrade et al, 1983, *Spectrochim. Acta*

CVAAS - Flow injection system (FIA)

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- ➊ sampling rate
- ➋ small and variable sample volumes
- ➌ higher precision

⇒ Batch systems

⇒ Flow systems

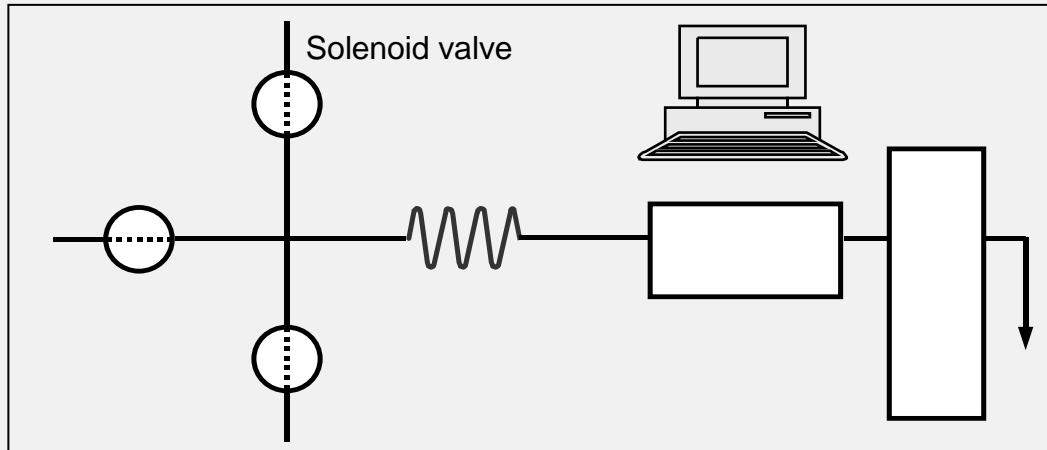
- Segmented flow analysis (SFA)
- Continuous sample flow
- Flow injection analysis (FIA)

Andrade et al, 1983, *Spectrochim. Acta*

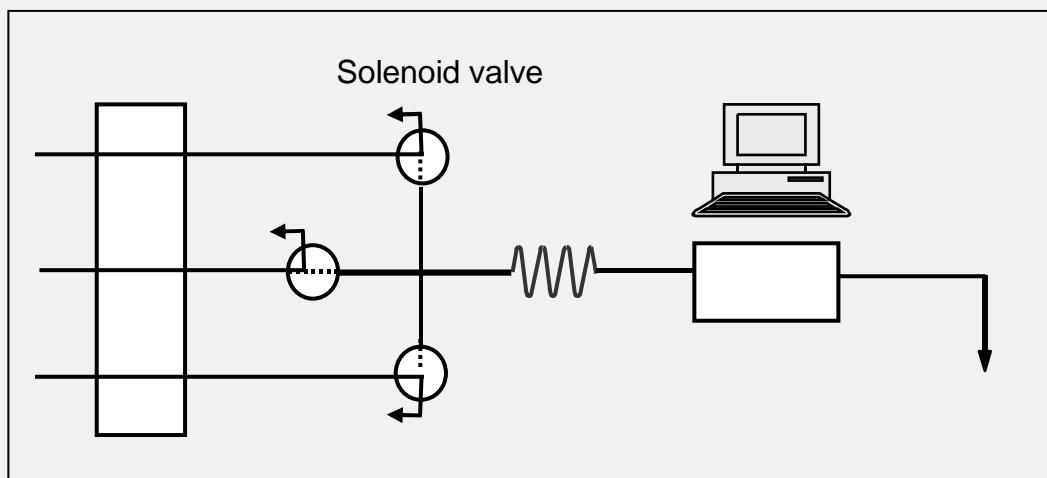
- Sequential injection analysis (SIA)

Reis et al, ACA 293(1994)129.

Aspiration



Propulsion



Advantages

- ⇒ Miniaturisation of flow systems
- ⇒ Improved mixing conditions
- ⇒ Reduced sample and reagent consumption
- ⇒ Flexibility

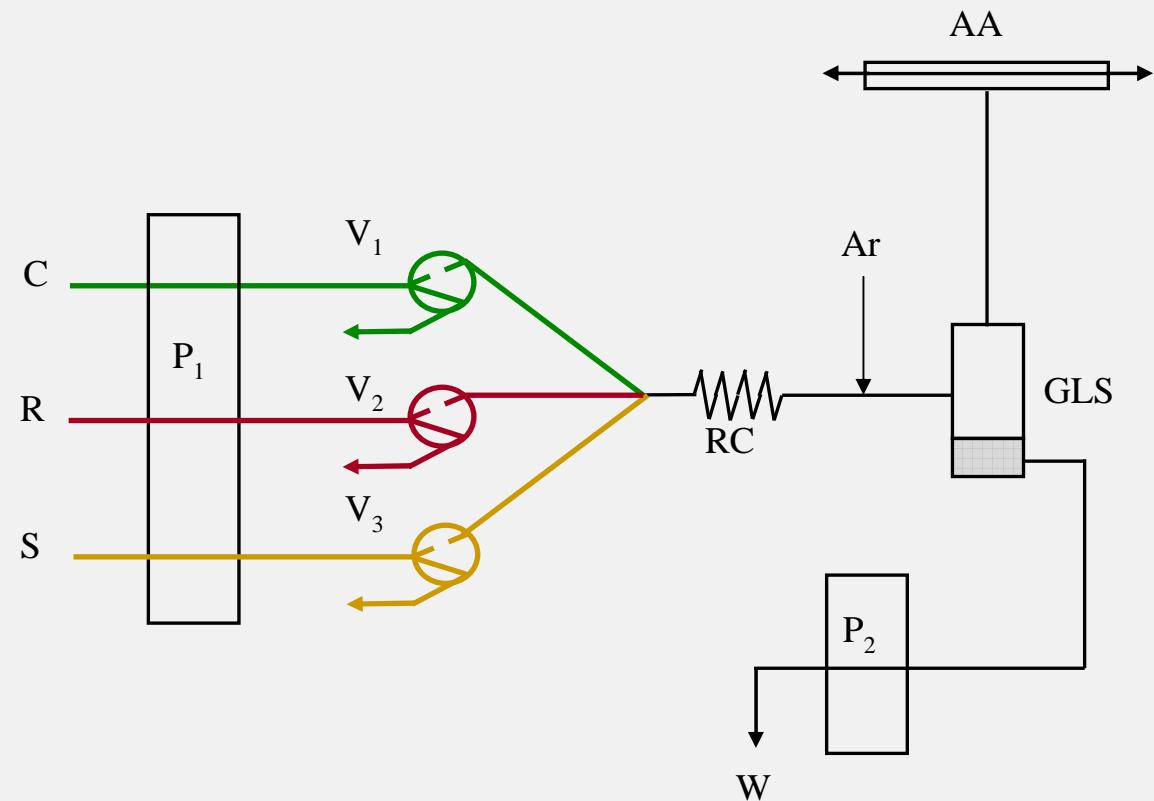
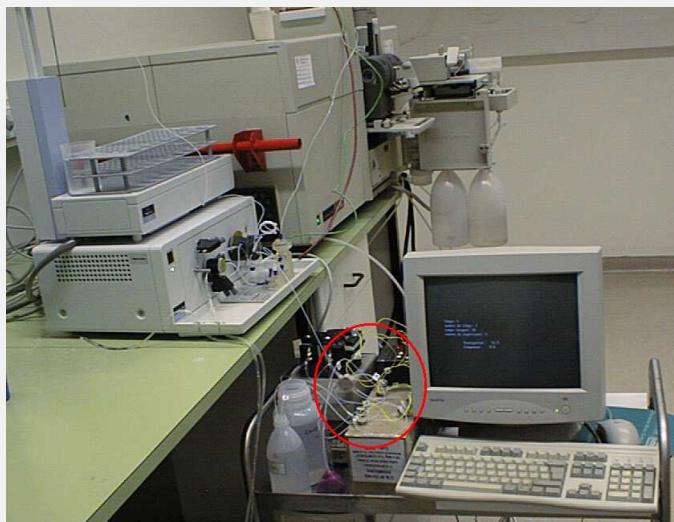
Limitations

- ⇒ The need to aspirate the solutions (or re-circulating the solutions)
- ⇒ Limited commercial availability
- ⇒ Lack of robustness of the valves

- ⇒ Calibration procedure using a single standard solution
- ⇒ Reagent/sample reduction
- ⇒ Waste reduction
- ⇒ Application to the determination of mercury in fish samples

Flow manifold

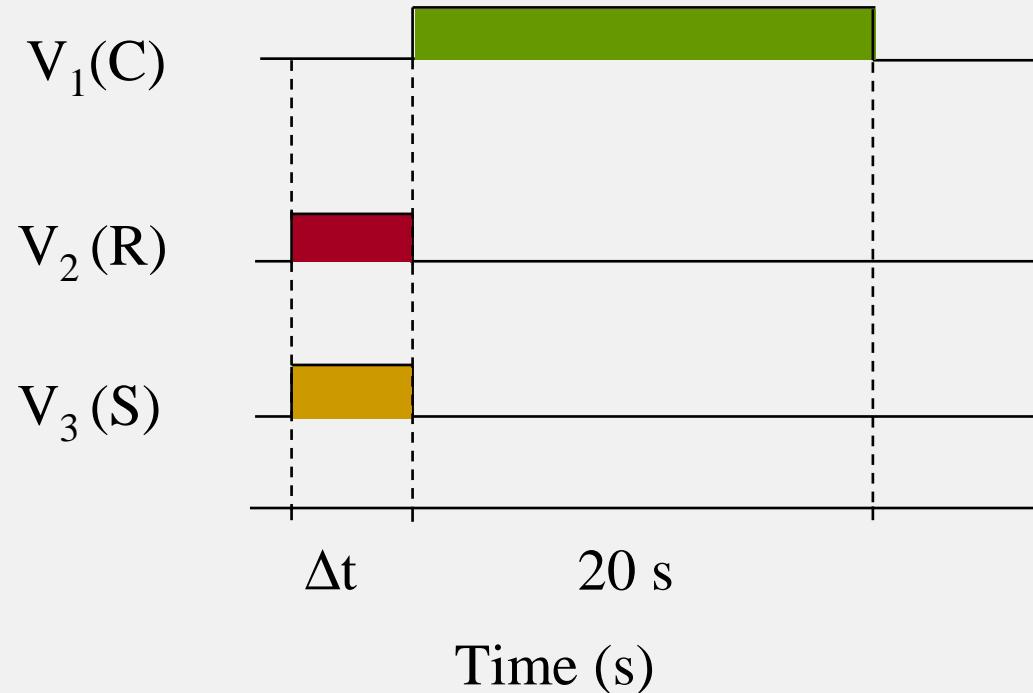
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C: carrier HCl 3% v/v (flow rate: 11 ml min^{-1}); **R:** reducing solution NaBH_4 0.2% m/v in NaOH 0.05% m/v (flow rate: 6 ml min^{-1}); **S:** sample/standard solutions; **RC:** reactor, 30 cm; **Ar:** argon (flow rate: 70 ml min^{-1}); **AA:** atomic absorption spectrophotometer; **V₁-V₃:** solenoid valves; **GLS:** gas-liquid separator; **P₁,P₂**: peristaltic pumps; **W:** waste.

Timing course of the valves

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Single standard calibration → Δt : sampling time (0.5-10 s)

Traditional calibration → Δt : sampling time (5 s)

Figures of merit

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	FIAS 100	MCFIA	MCFIA <i>Single standard calibration (10ppb)</i>
Limit of detection (ng)	0.08	0.13	0.10
Limit of quantification (ng)	0.21	0.34	0.34
RSD %	4.7 (0.83 ng) 1.7 (4.1 ng)	5.8 (0.83 ng) 2.7 (4.2 ng)	6.2 (0.80 ng) 1.9 (4.9 ng)
Linear range (ng)	0 – 10	0 – 10	0 – 10
Regression equation ^b			
m, slope	0.112 (± 0.005)	0.130 (± 0.008)	0.112 (± 0.006)
b, intercept	0.014 (± 0.014)	0.002 (± 0.002)	0.018 (± 0.012)
Correlation coefficient	0.9998	0.9994	0.9988
Sampling rate	90 h ⁻¹	60 h ⁻¹	60 h ⁻¹

^b Mean \pm std deviation of 7 different working days

Figures of merit (cont)

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	FIAS 100	MCFIA	MCFIA Single standard calibration (10 ppb)
Sample consumption ^b	500 µl	500 µl	50 – 1000 µl
HCl consumption ^b	0.25 ml	0.11 ml	0.11 ml
NaBH ₄ consumption ^b	0.9 mg	1.0 mg	0.1 – 2.0 mg
NaOH consumption ^b	0.23 mg	0.25 mg	0.025 – 0.50 mg
Waste ^b	13.3 ml	4.7 ml	3.8 – 5.7 ml

^b Values per assay

Accuracy

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Certified value mg/Kg	MCFIA		Single standard calibration ^b mg/Kg
	MCFIA ^b mg/Kg	Single standard calibration ^b mg/Kg	
Dogfish (DORM-2)	4.64 ± 0.26	4.37 ± 0.13	4.44 ± 0.28

^b average value ± std deviation of two working days with 3 replicates analysed each day (n=6)

Accuracy

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	Without spike (µg/Kg)	Spiked (ng)	MCFIA Recovery (%) ^b	MCFIA Single standard calibration Recovery (%) ^b
Cod	57	200	98.6 ± 0.8	104.6 ± 0.9
		500	99.8 ± 0.3	109.6 ± 0.4
		1000	96.9 ± 2.5	104.5 ± 2.6
Shellfish	≤ LOQ (9 µg/Kg)	200	90.7 ± 5.5	96.6 ± 3.7
		500	94.1 ± 2.7	105.6 ± 3.3
		1000	95.4 ± 1.2	107.1 ± 1.3

^b Mean ± std deviation (n=3)

Different volumes of the same sample/standard solution are intercalated in the system



plotting peak area versus injection time



sample calibration curve

The linear relationship $A = m V + b$
theoretically $b=0$

Beer's law

$A = \epsilon I C$

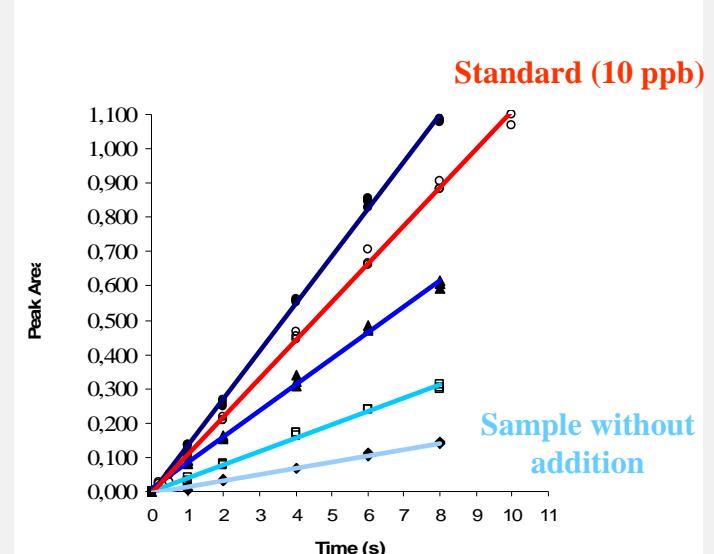
$$C^* = \frac{C_{\text{std}} \times m^*}{m_{\text{std}}}$$

Slope comparasion method

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	Correlation coefficient	Slope ^b	Recovery ^b %
Sample without addition	0.9975	0.018±0.001	-
Sample solution			
Addition 2 ppb	0.9976	0.039±0.001	92.4±7.6
Addition 5 ppb	0.9995	0.076±0.001	103.6±3.7
Addition 10 ppb	0.9994	0.138±0.001	107.2±2.3
Standard 10 ppb	0.9995	0.111±0.002	-

^b mean ± Std deviation (n=3)



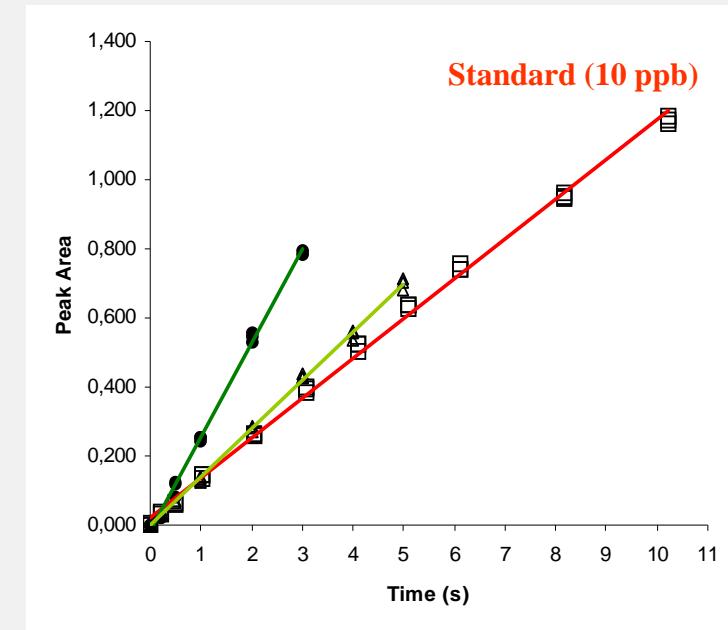
Slope comparasion method

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	Correlation coefficient ^b	Slope ^b	Concentration Hg mg/Kg
Dorm-2 (Dogfish)	0.9982	0.266±0.011	4.54±0.02
Dorm-2 Diluted solution	0.9996	0.140±0.003	4.77±0.09
Standard 10 ppb	0.9987	0.115±0.001	-

^b Mean ± Std deviation (n=3)

Dogfish (Dorm-2): Certified value = 4.64± 0.26 mg/Kg



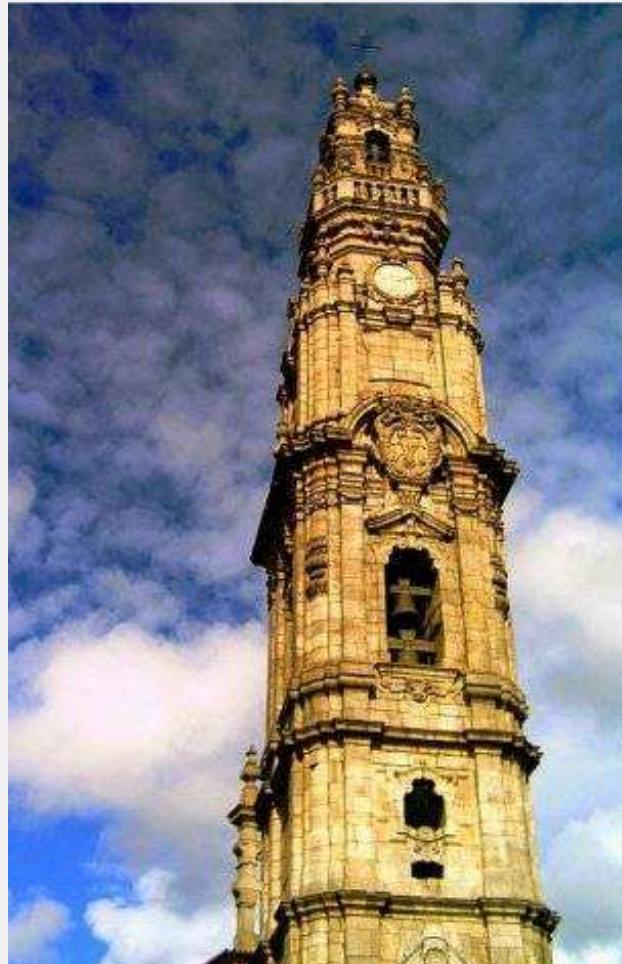
Making use of the linear relation between the signal (peak area) and injection time the concentration of the sample can also be accurately determined.

- ✓ The developed multicommutated flow system is accurate and precise for the determination of mercury in fish samples
- ✓ Advantage over the traditional calibration
 - ⌚ Less time consuming
 - ⌚ Reagent/standard consumption
 - ⌚ Waste production
- ✓ Sample dilution is easily adjusted
- ✓ This methodology can be extended to other elements forming volatile hydrides

Acknowledgements

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