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Validation of an Automated System for Methyl Mercury **Analysis using Cold Vapor Atomic Fluorescence Spectroscopy** with Gas Chromatography Separation

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A system was devised to allow automated analysis of prepared samples for the determination of methyl mercury. This system is based on an established EPA method (1630).



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Prepared samples are buffered in vials and then ethylating reagent in added. The vials are topped off and capped with a septa lid. Preventing any air bubbles is a requirement.

Under computer control, the autosampler will extract the liquid for analysis during the batch run. The liquid is transferred to the purge vessel and then purged $(\sim 5 \text{ minutes}).$

Nitrogen purge gas strips the volatile ethylated species so that they can be adsorbed onto a trap. Drying of a second trap and desorption of a third are all carried out simultaneously. The desorption trap is heated rapidly to release the ethylated compounds. (Cycle time ~6 min.)

Ethylated compounds from the desorption trap are separated from each other in the GC column. All mercury compounds passing through the pyrolytic column are decomposed liberating elemental Hg for detection.

Hg(0) is continuously measured by the CVAFS, giving a relative measure at any given moment. Peaks resulting from GC separated compounds are automatically measured in height or area by software to provide results.

Using the measurement from the peak of interest, the quantity of mercury for a given run is calculated based on standard and blank run results. Results for samples, standards, blanks, and QC runs are processed for the batch under software control.

Addition of ethylating reagent into the vial, ahead of sealing and subsequent purging, was found to be stable and effective as long as no air was trapped within the vial. In the presence of air this highly reactive reagent was found to be adversely affected and lower recoveries resulted. Once careful attention was paid to issues such as this, the system was found to work reliably and provided results as good or better than from manual analysis. Considerably less human labor is required as most repetitious tasks are no longer the responsibility of the technician.

We investigated a number of variables in order to try and discover influential parameters unique to the capped ethylation and postponed purging technique that we employed. A subset of results from preliminary work to investigate these effects and to determine general performance of our automated system are shown below:



| | | Instrument Deter | ction Limit Test | |
|----------------------|-------------|---|-----------------------------|---------------|
| 0.58 | | | | |
| 0.50 | | • 0.56- | 610501 | 0.559695 |
| 0.54 | • 0.5054205 | 05 • 0.527748037 | 0.632663036 | • 0.632663036 |
| 0.5154 0.5 154 | 80539 | | • 0.50 | 80880-4 |
| 2 0.48 | | | | |
| 0.40 | | 8 replicates (0.5 pc RSD = 3.64 Average = 0.535 | MMHg.in.42mL) : % pg. | |
| 0.44 | | Std Dev = 0.0199 IDL = 0.058 IDL conc. = 0.001 | 5 pg. 1 pg. 139 ng/L | |





Summary: We found that by optimizing the system components, providing computer controlled automation, and modifying the standard methodology slightly, it is possible to get excellent results. Our findings up to this point have yet to show any significant limitation that might prevent analysis of a particular type sample matrix that is currently performed successfully with manual procedures.

Advantages over the existing manual method are easier reproducibility of results with greater throughput and reduced requirement of analytical expertise. Maintenance of the system has been found to be minimal and sources of potential errors are well understood. Refinements of every part within the analytical process and elimination of steps that introduce human errors has lead to results that are found to be as good or better than those resulting from analysis using manual procedures.

48 run batch comparison (person hours)

Manual: 8+ hrs. continuous attended operation Automated: <2 hrs. pre-analysis, unattended afterward