

APPLICATIONS INFORMATION USING ADVANCED SAMPLE HANDLING TECHNOLOGY

Quantitation of C_2 to C_4 Carbonyls by Purge & Trap

Purge and trap has been used as an analytical technique for many years, in the analysis of volatile organic compounds in drinking water and waste water. Organic trace volatiles such as C_2 - C_4 aldehydes and ketones have also been quantitated and qualitated using purge and trap. The sample is normally purged from water using an inert carrier gas (N_2 or He), trapped onto an adsorbent bed, and then thermally desorbed to a GC/MS for quantitation.

A 10 μ Mole solution containing acetaldehyde, propanal, butyraldehyde, acetone, and 2butanone was prepared in distilled water. Serial dilutions were made from the stock and consisted of stock (10 μ M), dilutions 5 μ M, 1 μ M, 0.5 μ M, and 0.25 μ M. A 5 ml sample of each dilution was then placed into the 5 ml sparging vessel of a CDS 7000 Sample Concentrator, which was interfaced to a GC/MS (ion trap). Each dilution was sparged for eleven minutes at ambient temperature and collected on a Vocarb mutibed trap. The trap was desorbed for four minutes at 250°C to the GC/MS.

Figure 1 shows the 10 µM chromatogram of the component mixture. The elution order is propanal (1), acetone (2),butyraldehyde (3), 2-butanone (4), and acetaldehyde (tetramer) (5). Figure 2-6 show linearity plots of each compound and each dilution. Each data point is an average area based on three runs. Plot correlation coefficients were 0.9752 (F2), 0.9965 (F3), 0.9987 (F4), 0.9996 (F5), 0.9998 (F6).



Propanal P&T CDS 7000 $f^{R^2} = 0.9752$ $f^{R^2} = 0.9752$ f









Figure 4



Figure 5





CDS 7000 Parameters

Valve Oven:	150°C		
Transfer Line:	150°C		
Purge Ready:	40°C	10ml/min	
Dry Purge:	35°C	2.0/min	200ml/min
Desorb Prehea	t: 245°C	0.0/min	
Trap Desorb:	250°C	4.0/min	250ml/min
Trap Bake:	260°C	10.0/min	400ml/min
GC Time:	15.5/mir	ו	10ml/min

GC/MS Parameters

Column: Varian CP 624 30m , 0.25mm , 1.4µm Flow Rate: 1.3m/min Split Ratio: 20:1 Program: 40°C/2min Ramp 10°C/min Final: 175 MS: Varian Ion Trap

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