

Method 525Date Extracted 9/12/01Page 77

Lab #	Description	Sample vol/wt	Drip rate ml/min	Surr add	Spk add	IS add	Final vol.	T.V. pos.	pH			Count
	LIB	994 ml		1 ml		1000 ml			5.2			
	MDL				1 ml	1000 ml						
	Rel				1 ml	1000 ml						
22166	Sample											4
22167												3
22168												6
22169												2
22170												8
22171												9
QC Reviewed by, <u>[Signature]</u> 9/12/01 Remarks: <u>0.62 mg/L on all 6.</u> drip rate must be between 5-15 ml/min for liq-liq extractors Extraction type - <u>SPE</u> Solv. lot# <u>MC</u> NaCl lot# <u>T51F95</u> Na2SO4 lot# <u>V18617</u> STANDARD INT STD LOG# <u>1855/1853/1854</u> CALIBRATION <u>1855/1853/1865</u> REF/SPIKE <u>1894</u> MDL/MDRF <u>1895</u> Q. CHECK INT. STD. SURROGATE LPC ENDRIN/DDT												

Analyst J. Cai / T. Doherty
revision 10/15/99

Verified by _____

Supv. approved DN
9/18/01

Method 525.2
Instrument 204File Directory E

Storage Directory _____

Lab #	Filename	Description	Vial #		Inj.	Standards used - Lot #			Remarks	Initials
			front	rear		Calibration	Reference	Surrogate		
	DTA0915	DTFP		1	1	1801b/1802c	1972			TB
	DTA0915	DDT		2		1803c				
	C2A0915	C2		3		1894c/1895a/1896a	1970/1971			
	3	3		4						
	4	4		5						
	5	5		6						
	6	6		7						
	7	7		8						
	8	8		9						
	RUN0915	Ref std		10		1894c/1895a/1853a/1855a	1854b	Not Extd		
	MCA0915	DMC		11						
	LB0915	LB		12		1895	1894d			
	MDL0915	MDL		13		1855/1853/1865d				
22166	22166A	Sample		14						
	7B			15						
	8C			16						
	9D			17						
22170	22170E			18						
22171	22171F			19						
	REF0915	Ref std		20		1855/1853/1854c		Extd		
	CSB0915	Cst		21		1894c/1895a/1896a	1970/1971	2nd inj		
	C6B0915	C6		22				" "		

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F **File** **Control** **Util**

Exit

PATH E:

DATA FILE DFTA0915

SAMP

OPER BOHLK

COMM BOHLK; METHOD 525; INST 204; 09/15/01; DFTPP/ENDRIN

Analysis List:

5041

22 entries

**Edit
Current
Method**

**Select
Current
Method**

**Clear
Current
Method**

	<u>DATA FILE</u>	<u>AS METH</u>	<u>VIAL</u>	<u>GC METH</u>	<u>ACQ METH</u>	<u>DS PROC</u>
Add						
Expand >	1) DFTA0915	DFT0621	1	DFTPP8	DFTPP8	
	2) DDTA0915	DFT0621	2	DFTPP8	DFTPP8	
Del	3) C2A0915	DFT0621	3	DFTPP8	DFTPP8	
	4) C3A0915	DFT0621	4	DFTPP8	DFTPP8	
UnDel	5) C4A0915	DFT0621	5	DFTPP8	DFTPP8	
	6) C5A0915	DFT0621	6	DFTPP8	DFTPP8	
	7) C6A0915	DFT0621	7	DFTPP8	DFTPP8	
Home	8) C7A0915	DFT0621	8	DFTPP8	DFTPP8	
	9) C8A0915	DFT0621	9	DFTPP8	DFTPP8	
Scr Up	10) RUN0915	DFT0621	10	DFTPP8	DFTPP8	
	11) MCAA0915	DFT0621	11	DFTPP8	DFTPP8	
Scr Dn	12) LB0915	DFT0621	12	DFTPP8	DFTPP8	
	13) MDL0915	DFT0621	13	DFTPP8	DFTPP8	
End						

F **File** **Control** **Util**

Exit

PATH E:

DATA FILE C6B0915

SAMP

OPER BOHLK

COMM BOHLK; METHOD 525; INST 204; 09/15/01; C6

Analysis List:

5041

22 entries

**Edit
Current
Method**

**Select
Current
Method**

**Clear
Current
Method**

Add	<u>DATA FILE</u>	<u>AS METH</u>	<u>VIAL</u>	<u>GC METH</u>	<u>ACQ METH</u>	<u>DS PROC</u>
Expand	10) RUN0915	DFT0621	10	DFTPP8	DFTPP8	
	11) MCAA0915	DFT0621	11	DFTPP8	DFTPP8	
Del	12) LB0915	DFT0621	12	DFTPP8	DFTPP8	
	13) MDL0915	DFT0621	13	DFTPP8	DFTPP8	
UnDel	14) 22166A	DFT0621	14	DFTPP8	DFTPP8	
	15) 22167B	DFT0621	15	DFTPP8	DFTPP8	
	16) 22168C	DFT0621	16	DFTPP8	DFTPP8	
Home	17) 22169D	DFT0621	17	DFTPP8	DFTPP8	
	18) 22170E	DFT0621	18	DFTPP8	DFTPP8	
Scr Up	19) 22171F	DFT0621	19	DFTPP8	DFTPP8	
	20) REF0915	DFT0621	20	DFTPP8	DFTPP8	
Scr Dn	21) C5B0915	DFT0621	21	DFTPP8	DFTPP8	
	> 22) C6B0915	DFT0621	22	DFTPP8	DFTPP8	
End						

LAB #	DATE EXT.	COMMENT:	NAME											
22166 through 22170	9/12/01	525	Boh											
		① When the C2 cal std. was quantitated, the following compounds deviated from their T.V.'s by more than 30 % <table> <tr> <td>Simazine</td> <td>0.054 mg/L</td> <td rowspan="5">} T.V. = 0.1</td> </tr> <tr> <td>atrazine</td> <td>0.063</td> </tr> <tr> <td>adipate</td> <td>0.057</td> </tr> <tr> <td>BAP</td> <td>0.064</td> </tr> <tr> <td>metribuzin</td> <td>0.067</td> </tr> </table>	Simazine	0.054 mg/L	} T.V. = 0.1	atrazine	0.063	adipate	0.057	BAP	0.064	metribuzin	0.067	
Simazine	0.054 mg/L	} T.V. = 0.1												
atrazine	0.063													
adipate	0.057													
BAP	0.064													
metribuzin	0.067													
		② for the C5 cal std. run at the end of the Sequence, <table> <tr> <td>adipate</td> <td>quantitated at 2.762</td> <td>or 138</td> </tr> <tr> <td>terbacil</td> <td>"</td> <td>" 2.648 or 132</td> </tr> </table> R-analyses ① & ② not required of T.V. which is 2.0 mg/L	adipate	quantitated at 2.762	or 138	terbacil	"	" 2.648 or 132						
adipate	quantitated at 2.762	or 138												
terbacil	"	" 2.648 or 132												
		In each of the extracts (including B) there is an unknown eluting just prior to phthalate. The LB contains a very small amount - with samples 22168, 22170, 22171 containing the largest amounts - However, even these sample peaks are small in size. Refer to list of possible												

Terbacil is based on the 4 cal stds
C5 - C8 only.

Ted & I are of the opinion this peak is insignificant. All fits are low, it identifies differently in each sample.

The sample with the largest amt is still small compared to adjacent phthalate peak.

An peak was seen (similar to this one) in a previous unrelated data file.

D. Vini 9/18/11