

Method 508 ModDate Extracted 1/2/02Page 1

REF

#GCMS-16370

Lab #	Description	Sample vol/wt	Drip rate ml/min	Surr add	Sol add	IS add	Final vol.	T.V. pos.	pH	Count		
LB ✓		500ul		1ml			1ml	8A	7			
REF ✓					0.5ml			7A				
29228 ✓								8B				
29286 ✓	Acidic							6A				
29292 ✓								7A				
29284 ✓	Acidic							6B				
29465 ✓								7B				
29344 ✓								7B				
29343 ✓								6A				
29519 ✓								8A				
29562 ✓								8B				
30631 ✓	see 12/17/01							6B				
30632 ✓								8B				
Remarks: Samples #29284 & 29286 were very Acidic. 2.5 ml of 6M NaOH was added to the samples to bring pH 7.												
Reviewed by, <u>C/S 5/11/02</u>												
drip rate must be between 5-15 ml/min for liq-liq extractors drip rate -												
Extraction type <u>SB flm</u> Solv. lot# <u>V39466</u> NaCl lot# <u>V11599</u> Na2SO4 lot# <u>V39H00</u>												
STANDARD	INT STD LOG#											
CALIBRATION												
REF SPIKE	91759											
MDL/MDRF												
Q. CHECK												
INT. STD.												
SURROGATE	GCMS 121901											
LPC												
ENDRIN/DDT												

Analyst QA
Revision 4/12/01Reviewed by ① 1/2/02

Sequence Name: C:\HPCHEM\1\SEQUENCE\jan0302.S

Comment: INST 209; 30Mx0.25umX0.50

Operator: 209 GALOS

Data Path: C:\HPCHEM\1\data\jan0302\

Pre-Seq Cmd:

Post-Seq Cmd:

Method Sections To Run	On A Barcode Mismatch
(X) Full Method	(X) Inject Anyway
() Reprocessing Only	() Don't Inject

Line Type	Vial	DataFile	Method	Sample Name
1 Sample	1	DFTPP	GCMS1126	
2 Sample	2	CAL5	GCMS1126	mix
3 Sample	3	CAL10	GCMS1126	mix
4 Sample	4	CAL25	GCMS1126	mix
5 Sample	5	CAL40	GCMS1126	mix
6 Sample	6	CAL80	GCMS1126	mix
7 Sample	7	CAL100	GCMS1126	mix
8 Sample	8	CALCHK	GCMS1126	20
9 Sample	9	SURR	GCMS1126	
10 Sample	10	30117	GCMS1126	gcms scan 12/12
11 Sample	11	30118	GCMS1126	gcms scan 12/12
12 Sample	12	29791	GCMS1126	gcms scan 12/10
13 Sample	13	29671	GCMS1126	gcms scan 12/7
14 Sample	14	29734	GCMS1126	gcms scan 12/7
15 Sample	15	29670	GCMS1126	gcms scan 12/7
16 Sample	16	29520	GCMS1126	gcms scan
17 Sample	17	29517	GCMS1126	gcms scan
18 Sample	18	LB	GCMS1126	gcms scan 1/2
19 Sample	19	REF	GCMS1126	gcms scan 1/2
20 Sample	20	29228	GCMS1126	gcms scan 1/2
21 Sample	21	29562	GCMS1126	gcms scan 1/2
22 Sample	22	29519	GCMS1126	gcms scan 1/2
23 Sample	23	29292	GCMS1126	gcms scan 1/2
24 Sample	24	29284	GCMS1126	gcms scan 1/2
25 Sample	25	29343	GCMS1126	gcms scan 1/2
26 Sample	26	29344	GCMS1126	gcms scan 1/2
27 Sample	27	30632	GCMS1126	gcms scan 1/2
28 Sample	28	29465	GCMS1126	gcms scan 1/2
29 Sample	29	29286	GCMS1126	gcms scan 1/2
30 Sample	30	30631	GCMS1126	gcms scan 1/2
31 Sample	31	LBB	GCMS1126	gcms scan 1/2a
32 Sample	32	REFB	GCMS1126	gcms scan 1/2a
33 Sample	33	31642	GCMS1126	gcms scan 1/2a
34 Sample	34	30111	GCMS1126	gcms scan 1/2a
35 Sample	35	30253	GCMS1126	gcms scan 1/2a
36 Sample	36	31640	GCMS1126	gcms scan 1/2a
37 Sample	37	30023	GCMS1126	gcms scan 1/2a
38 Sample	38	31586	GCMS1126	gcms scan 1/2a
39 Sample	39	31641	GCMS1126	gcms scan 1/2a
40 Sample	40	31633	GCMS1126	gcms scan 1/2a
41 Sample	41	30010	GCMS1126	gcms scan 1/2a
42 Sample	42	31632	GCMS1126	gcms scan 1/2a

Last Modified: Tue Jan 08 09:12:14 2002

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DFTPP

Data File : C:\HPCHEM\1\DATA\JAN0302\DFTPP.D

Acq On : 3 Jan 2002 9:35 am

Sample :

Misc :

MS Integration Params: GOOFY.P

Method : C:\HPCHEM\1\METHODS\GCMS1126.M (RTE Integrator)

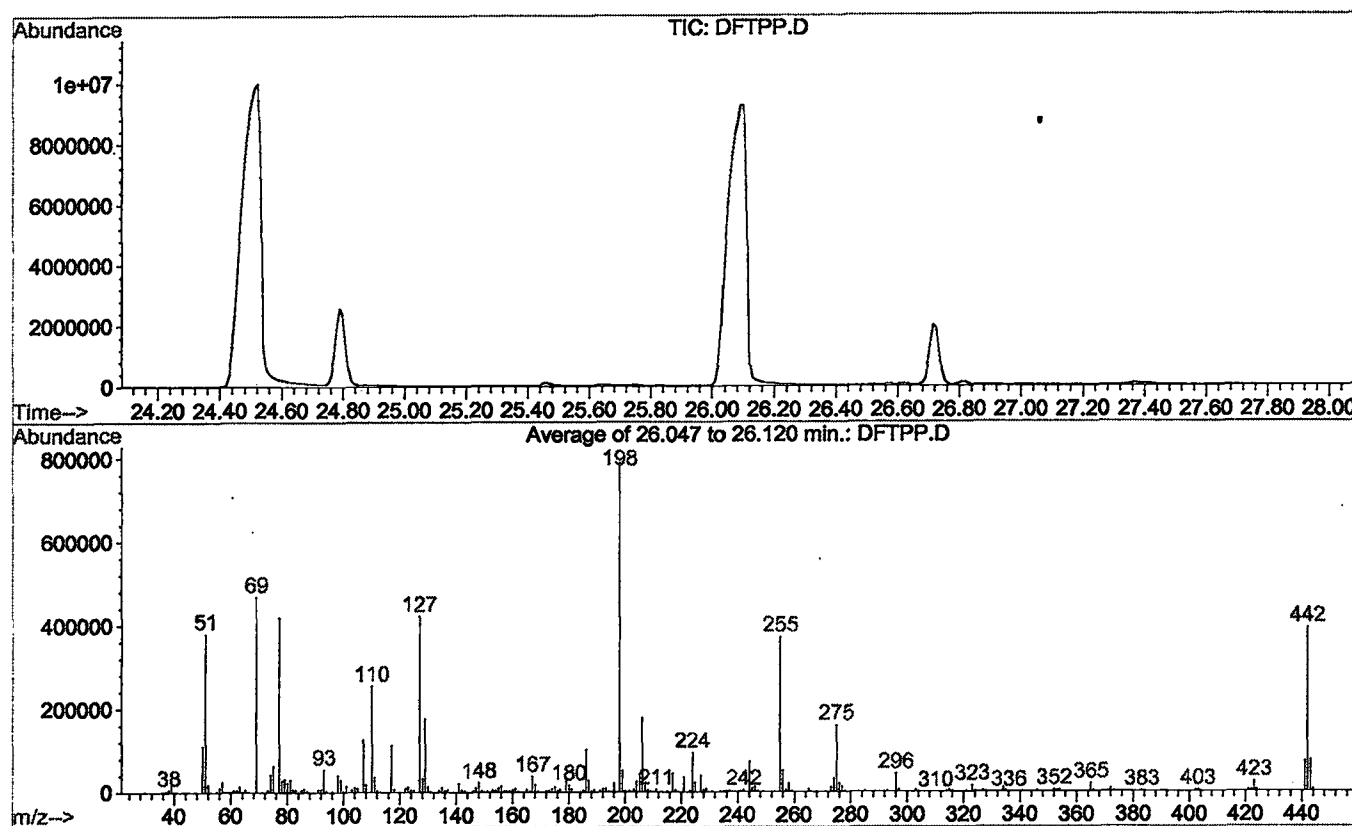
Title : 209 625/TCLP calibration table

Vial: 1

Operator: 209 GALOS

Inst : GC/MS 209

Multiplr: 1.00



Spectrum Information: Average of 26.047 to 26.120 min.

Target Mass	Rel. to Mass	Lower Limit%	Upper Limit%	Rel. Abn%	Raw Abn	Result Pass/Fail
51	198	30	60	48.3	381035	PASS
68	69	0.00	2	0.0	0	PASS
69	198	0.00	100	59.5	469099	PASS
70	69	0.00	2	0.0	211	PASS
127	198	40	60	53.5	422351	PASS
197	198	0.00	1	0.0	0	PASS
198	198	100	100	100.0	788713	PASS
199	198	5	9	6.7	52678	PASS
275	198	10	30	20.3	160060	PASS
365	198	1	100	2.6	20380	PASS
441	443	0.01	100	94.8	72500	PASS
442	198	40	110	49.6	391054	PASS
443	442	17	23	19.6	76478	PASS