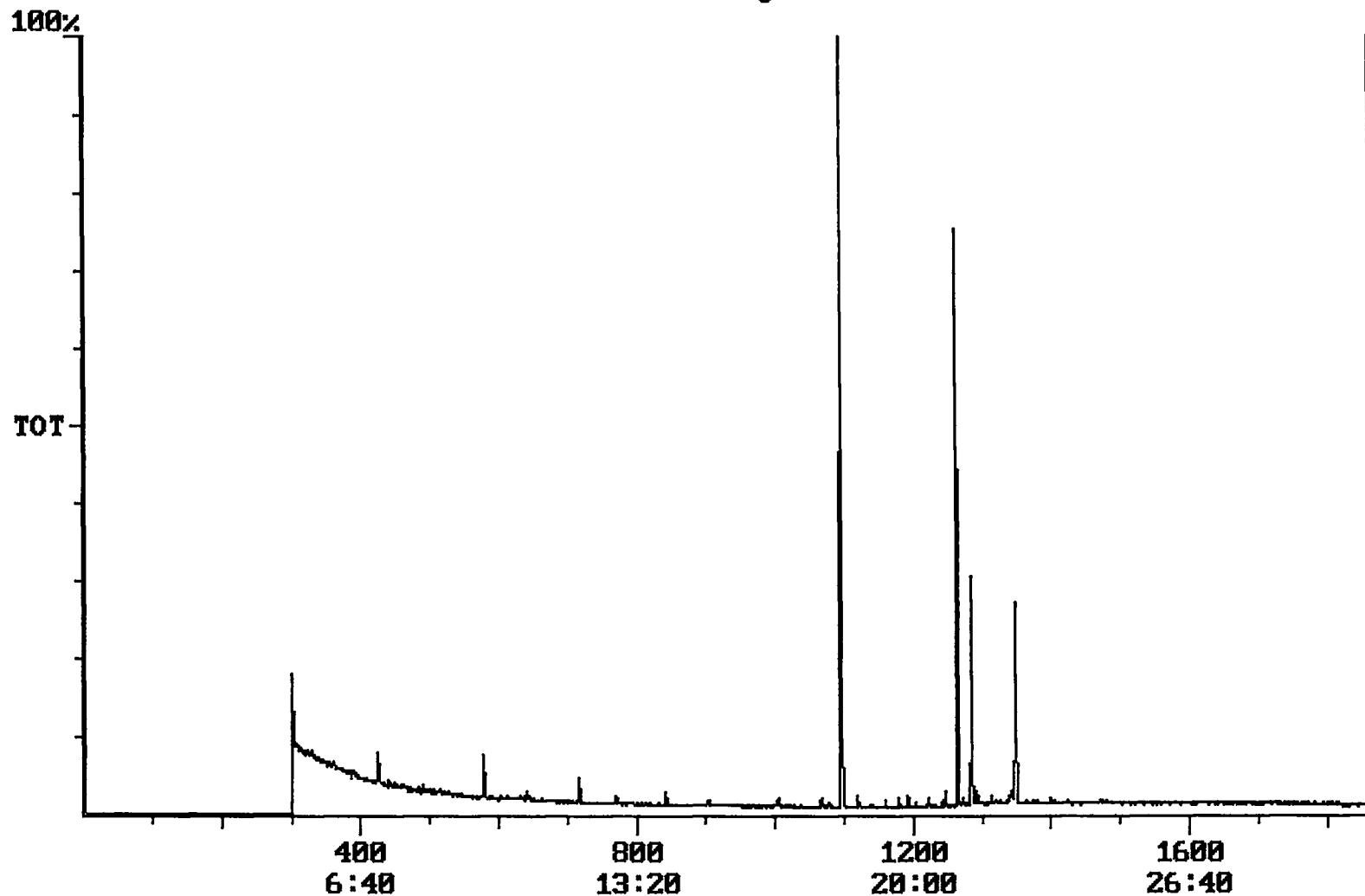


User: Bohlk, Ted
 Westchester Dept. of Labs & Research
 Date: 01/03/2002 Time: 14:50:13

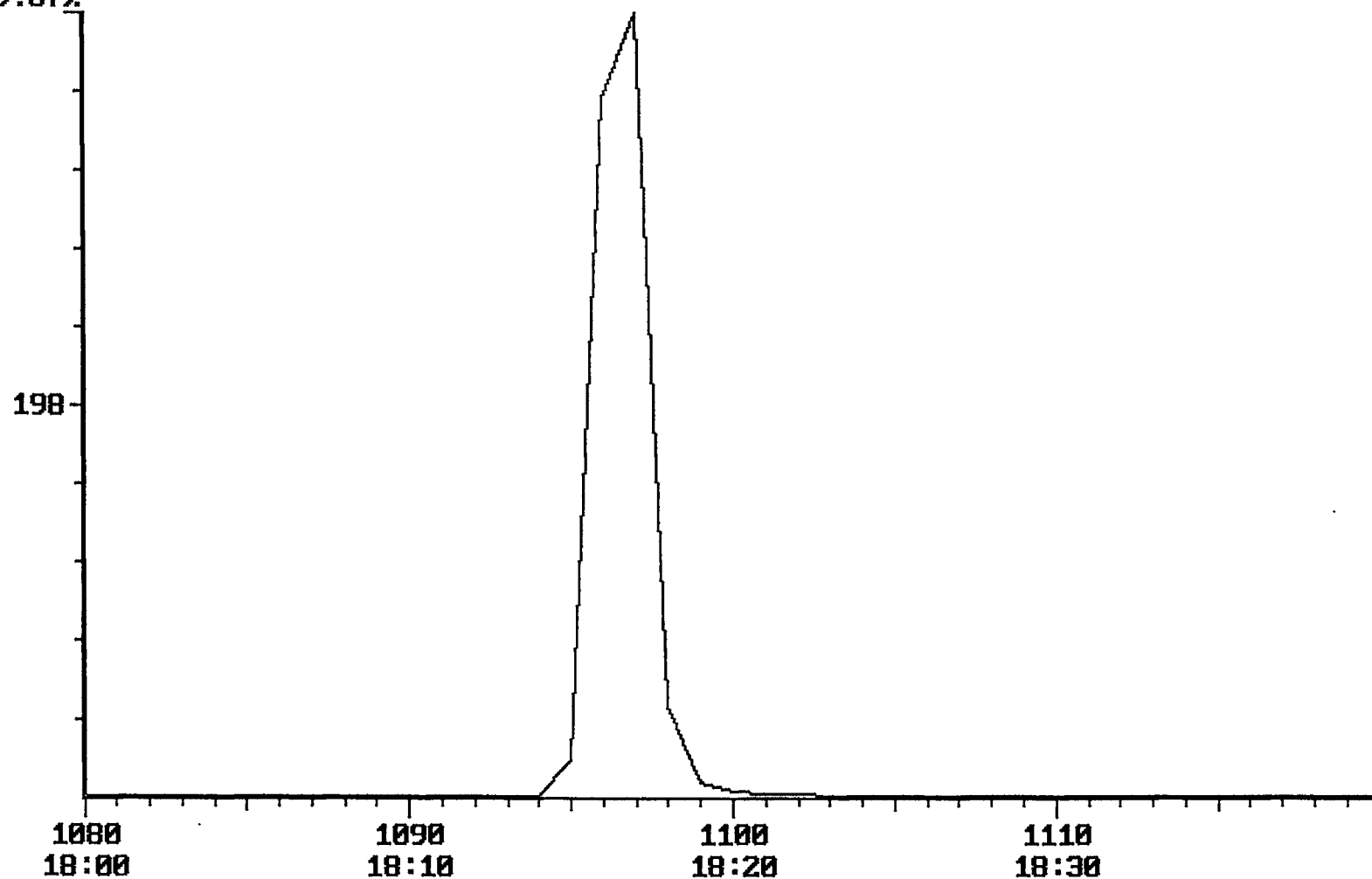
Sample: AD27116
 Analysis: SLP525 Organic Extractables
 Units: ug
 Started: 12/1/01 14:48 Ended: 12/1/01 14:48 Analyst: TB
 Page: 1

	Component Name	Viol	Result
1	DFTPP		PASSED
2	ENDRIN	USPEC	49
3	DDT		12

Chromatogram Plot File: E:\DFTA1201 Date: Dec-01-1991 06:13:55
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP
Scan No: 1 Retention Time: 0:01 RIC: 0 Mass Range: 0 - 0
Plotted: 1 to 1859 Range: 1 to 1859 100% = 3044925



Chromatogram Plot File: E:\DFTA1201 Date: Dec-01-1991 06:13:55
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP
Scan No: 1080 Retention Time: 18:00 RIC: 35997 Mass Range: 47 - 149
Plotted: 1080 to 1120 Range: 1 to 1859 100% = 3044925
9.67%



Mass List

File: E:\DFTA1201 Date: Dec-01-1991 06:13:55

Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP

Scan No: 1097 Retention Time: 18:17 RIC: 3044925 Mass Range: 45 - 444

Peaks: 206 Mass of Base Peak: 442 Int: 370149 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
45	1,368	0.37	80	6,592	1.78	117	66,791	18.04
47	3,233	0.87	81	9,328	2.52	118	2,860	0.77
48	1,243	0.34	82	1,990	0.54	122	3,980	1.08
49	19,029	5.14	83	1,741	0.47	123	3,358	0.91
50	74,626	20.16	84	4,104	1.11	124	2,238	0.60
51	195,771	52.89	85	1,865	0.50	127	107,213	28.96
52	8,582	2.32	86	8,582	2.32	128	8,333	2.25
55	4,850	1.31	87	3,731	1.01	129	48,756	13.17
56	3,482	0.94	91	3,233	0.87	130	2,860	0.77
57	8,333	2.25	92	3,482	0.94	131	1,119	0.30
61	3,606	0.97	93	23,134	6.25	134	2,238	0.60
62	1,741	0.47	97	995	0.27	135	4,726	1.28
63	6,965	1.88	98	21,766	5.88	136	2,487	0.67
64	1,368	0.37	99	6,840	1.85	137	3,233	0.87
65	1,616	0.44	101	3,855	1.04	139	995	0.27
67	3,731	1.01	103	3,980	1.08	140	1,616	0.44
68	2,487	0.67	104	3,109	0.84	141	8,955	2.42
69	140,422	37.94	105	3,358	0.91	142	1,865	0.50
73	3,358	0.91	106	1,616	0.44	143	1,616	0.44
74	21,517	5.81	107	45,398	12.26	144	870	0.24
75	21,890	5.91	108	4,353	1.18	146	2,487	0.67
76	7,089	1.92	109	1,616	0.44	147	5,099	1.38
77	150,870	40.76	110	90,298	24.40	148	8,333	2.25
78	4,975	1.34	111	13,432	3.63	149	2,114	0.57
79	18,283	4.94	116	3,233	0.87	151	2,487	0.67

Mass List

File: E:\DFTA1201 Date: Dec-01-1991 06:13:55

Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP

Scan No: 1097 Retention Time: 18:17 RIC: 3044925 Mass Range: 45 - 444

Peaks: 206 Mass of Base Peak: 442 Int: 370149 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
151	2,487	0.67	180	2,736	0.74	216	1,616	0.44
152	1,119	0.30	181	3,233	0.87	217	15,671	4.23
153	1,741	0.47	182	1,119	0.30	218	2,736	0.74
154	1,616	0.44	184	995	0.27	221	6,716	1.81
155	2,860	0.77	185	3,731	1.01	222	2,238	0.60
156	4,228	1.14	186	42,786	11.56	223	3,731	1.01
157	995	0.27	187	7,462	2.02	224	29,228	7.90
158	1,616	0.44	189	4,228	1.14	225	7,338	1.98
159	2,363	0.64	193	4,477	1.21	226	1,243	0.34
160	2,363	0.64	194	870	0.24	227	16,915	4.57
161	3,109	0.84	196	3,731	1.01	228	3,855	1.04
162	1,616	0.44	197	2,860	0.77	229	2,985	0.81
163	870	0.24	198	294,527	79.57	231	995	0.27
165	3,482	0.94	199	18,656	5.04	233	1,492	0.40
166	1,865	0.50	201	995	0.27	235	2,114	0.57
167	20,398	5.51	202	870	0.24	241	1,243	0.34
168	5,970	1.61	203	3,358	0.91	242	1,616	0.44
169	1,741	0.47	204	8,582	2.32	243	2,487	0.67
172	2,238	0.60	205	21,019	5.68	244	23,383	6.32
173	2,114	0.57	206	58,457	15.79	245	1,243	0.34
174	3,233	0.87	207	9,452	2.55	246	6,965	1.88
175	3,606	0.97	208	3,109	0.84	247	995	0.27
177	4,477	1.21	210	2,363	0.64	249	2,363	0.64
178	1,616	0.44	211	2,736	0.74	254	1,741	0.47
179	7,711	2.08	215	1,243	0.34	255	183,955	49.70

Mass List

File: E:\DFTA1201 Date: Dec-01-1991 06:13:55

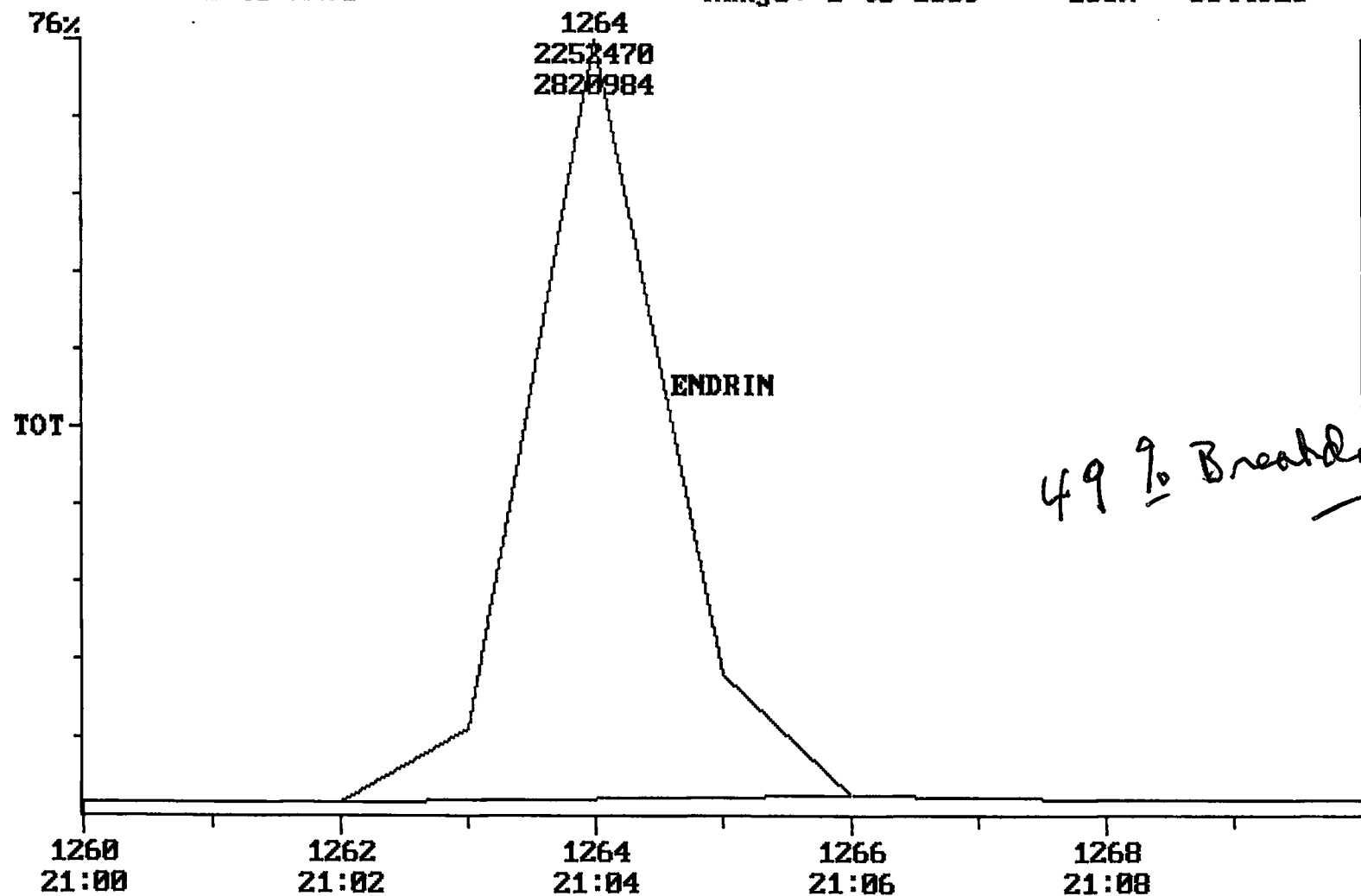
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP

Scan No: 1097 Retention Time: 18:17 RIC: 3044925 Mass Range: 45 - 444

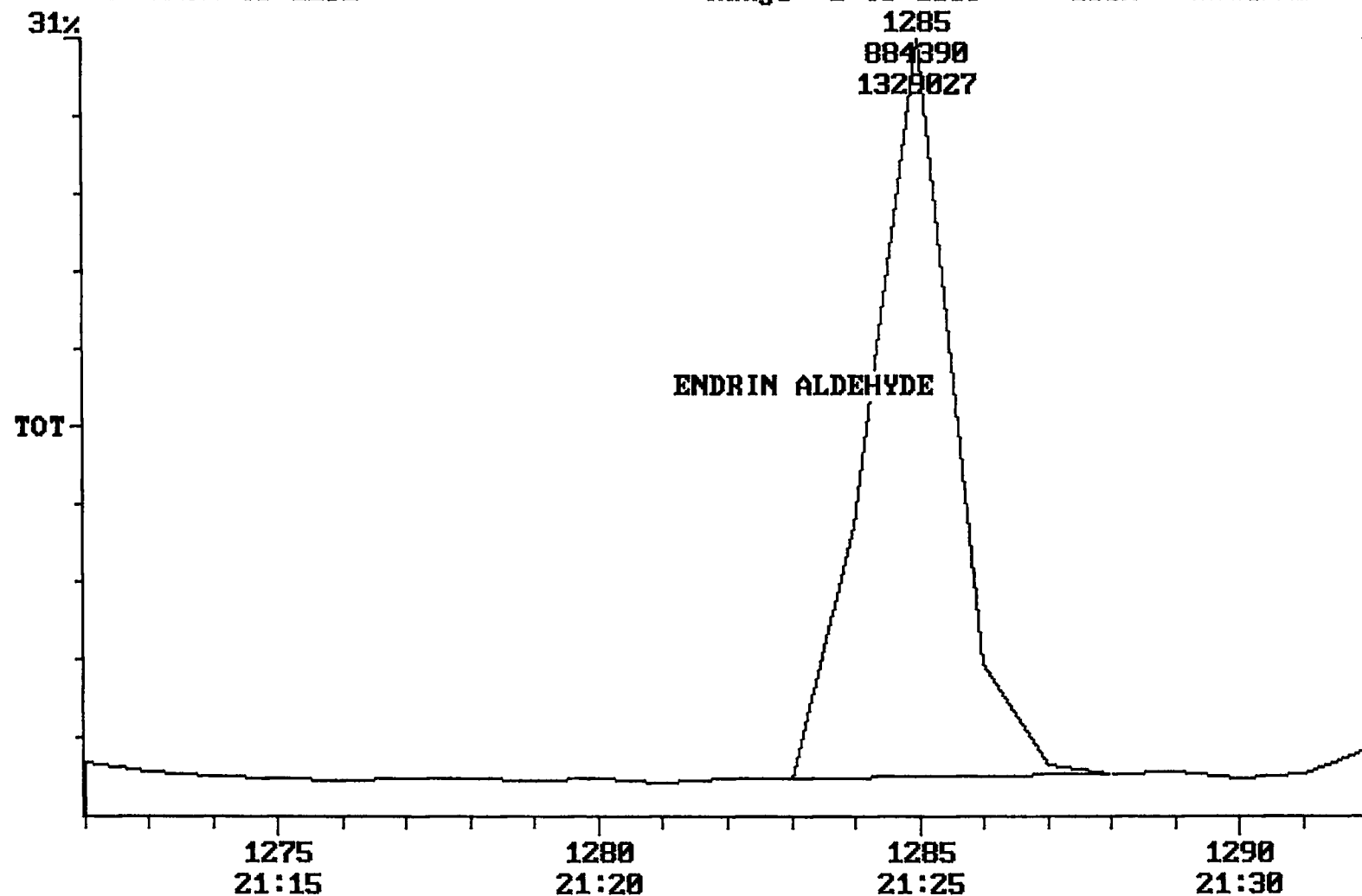
Peaks: 206 Mass of Base Peak: 442 Int: 370149 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
225	7,338	1.98	274	15,174	4.10	340	870	0.24
226	1,243	0.34	275	104,228	28.16	346	2,860	0.77
227	16,915	4.57	276	8,955	2.42	352	1,990	0.54
228	3,855	1.04	277	11,815	3.19	353	2,114	0.57
229	2,985	0.81	278	1,368	0.37	354	4,975	1.34
231	995	0.27	281	1,492	0.40	355	1,243	0.34
233	1,492	0.40	284	995	0.27	365	20,522	5.54
235	2,114	0.57	285	1,865	0.50	366	1,865	0.50
241	1,243	0.34	290	870	0.24	372	6,218	1.68
242	1,616	0.44	291	995	0.27	373	2,611	0.71
243	2,487	0.67	293	4,601	1.24	383	1,741	0.47
244	23,383	6.32	296	19,402	5.24	390	1,368	0.37
245	1,243	0.34	297	1,492	0.40	391	1,243	0.34
246	6,965	1.88	302	995	0.27	392	1,492	0.40
247	995	0.27	303	3,855	1.04	402	2,114	0.57
249	2,363	0.64	304	1,865	0.50	403	4,353	1.18
254	1,741	0.47	313	995	0.27	404	1,492	0.40
255	183,955	49.70	315	2,114	0.57	421	1,616	0.44
256	21,517	5.81	316	1,492	0.40	422	3,980	1.08
257	1,492	0.40	323	7,960	2.15	423	15,174	4.10
258	10,696	2.89	324	2,487	0.67	424	3,980	1.08
259	1,990	0.54	327	2,114	0.57	441	59,328	16.03
260	995	0.27	333	2,363	0.64	442	370,149	100.00
265	4,850	1.31	334	7,089	1.92	443	76,492	20.67
273	5,845	1.58	335	2,238	0.60	444	8,830	2.39

Chromatogram Plot File: E:\DFTA1201 Date: Dec-01-1991 06:13:55
 Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP
 Scan No: 1270 Retention Time: 21:10 RIC: 43116 Mass Range: 47 - 281
 Plotted: 1260 to 1270 Range: 1 to 1859 100% = 3044925



Chromatogram Plot File: E:\DFTA1201 Date: Dec-01-1991 06:13:55
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP
Scan No: 1292 Retention Time: 21:32 RIC: 79145 Mass Range: 45 - 429
Plotted: 1272 to 1292 Range: 1 to 1859 100% = 3044925



Chromatogram Plot

File: E:\DFTA1201 Date: Dec-01-1991 06:13:55

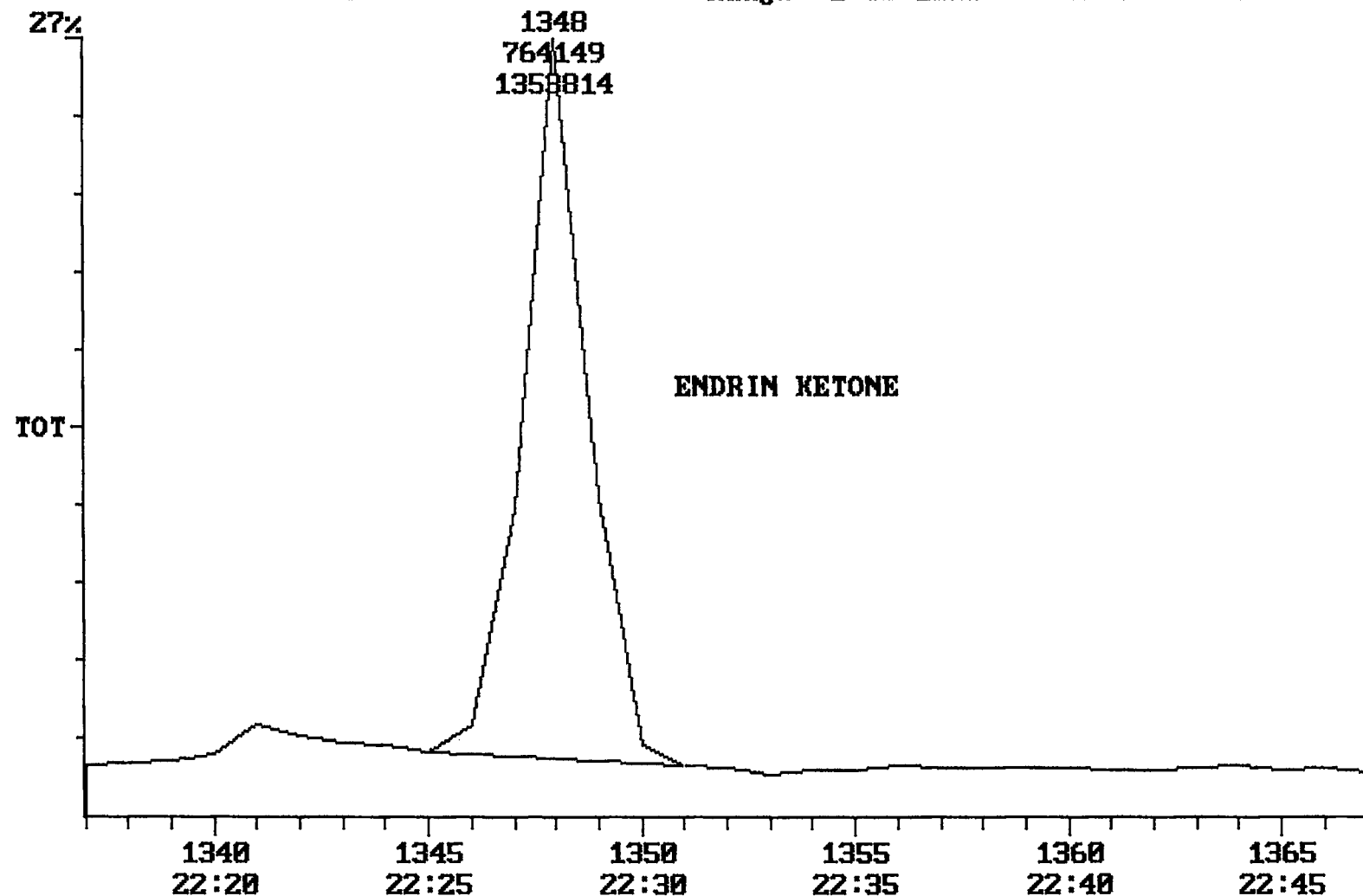
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP

Scan No: 1367 Retention Time: 22:47 RIC: 46405 Mass Range: 45 - 281

Plotted: 1337 to 1367

Range: 1 to 1859

100% = 3044925



Chromatogram Plot

File: E:\DDTA1201

Date: Dec-01-1991 06:48:21

Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DDT

Scan No: 1

Retention Time: 0:01

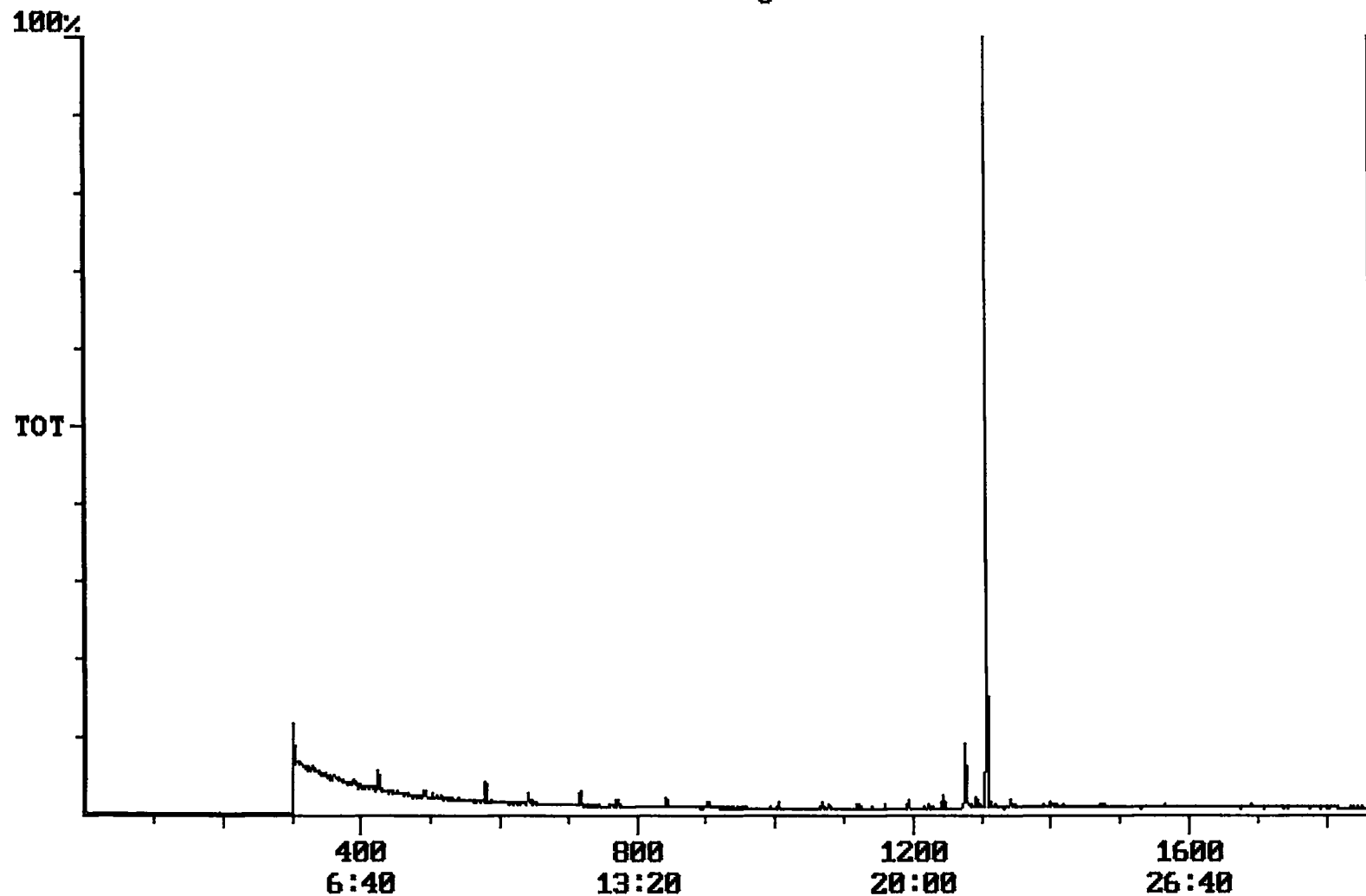
RIC: 0

Mass Range: 0 - 0

Plotted: 1 to 1859

Range: 1 to 1859

100% = 3651289



Chromatogram Plot

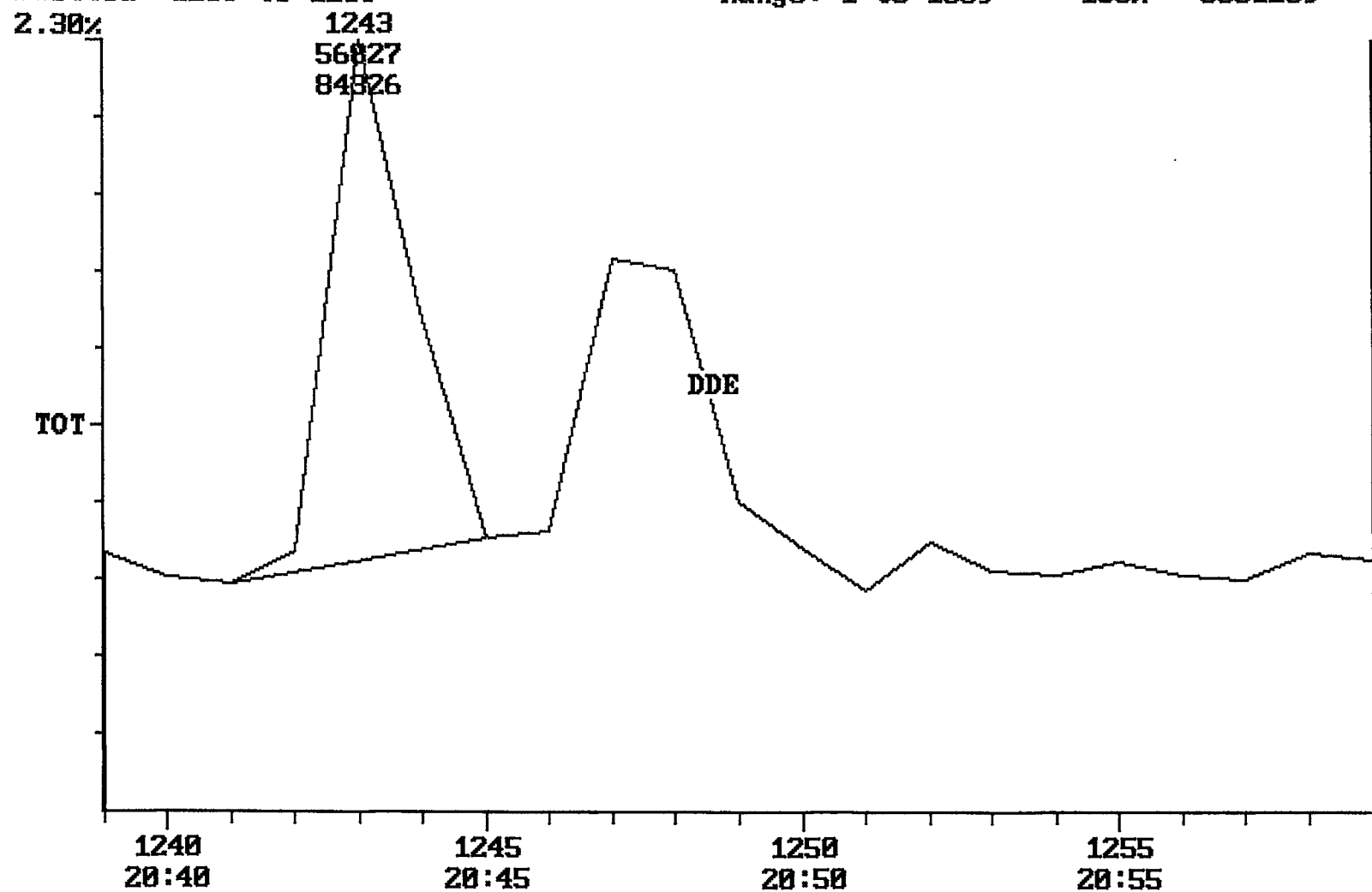
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Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DDT

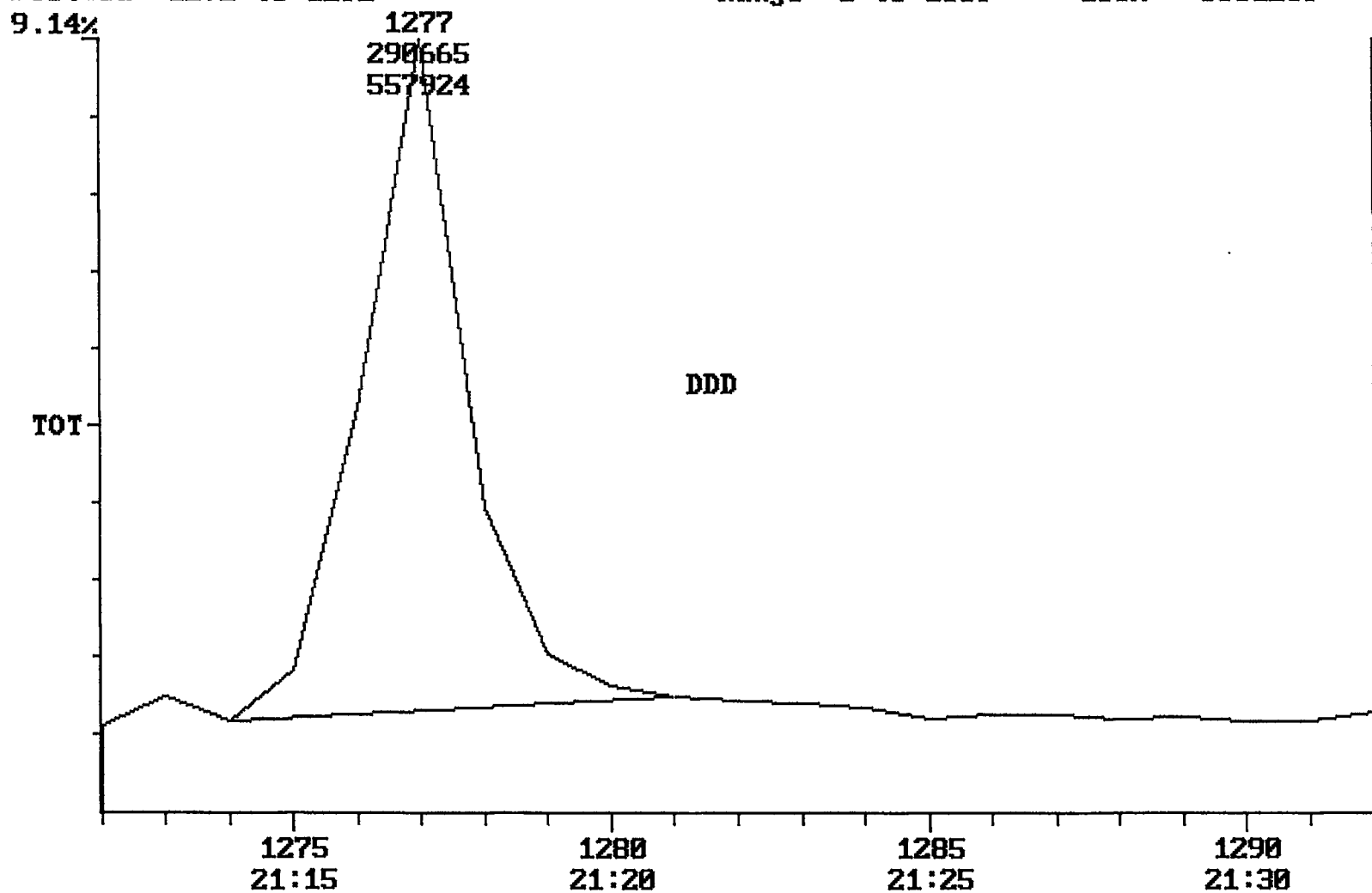
Scan No: 1259 Retention Time: 20:59 RIC: 27317 Mass Range: 47 - 207

Plotted: 1239 to 1259

Range: 1 to 1859 100% = 3651289



Chromatogram Plot File: E:\DDTA1201 Date: Dec-01-1991 06:48:21
 Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DDT
 Scan No: 1292 Retention Time: 21:32 RIC: 42359 Mass Range: 47 - 284
 Plotted: 1272 to 1292 Range: 1 to 1859 100% = 3651289



Chromatogram Plot

File: E:\DDTA1201

Date: Dec-01-1991 06:48:21

Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DDT

Scan No: 1324

Retention Time: 22:04

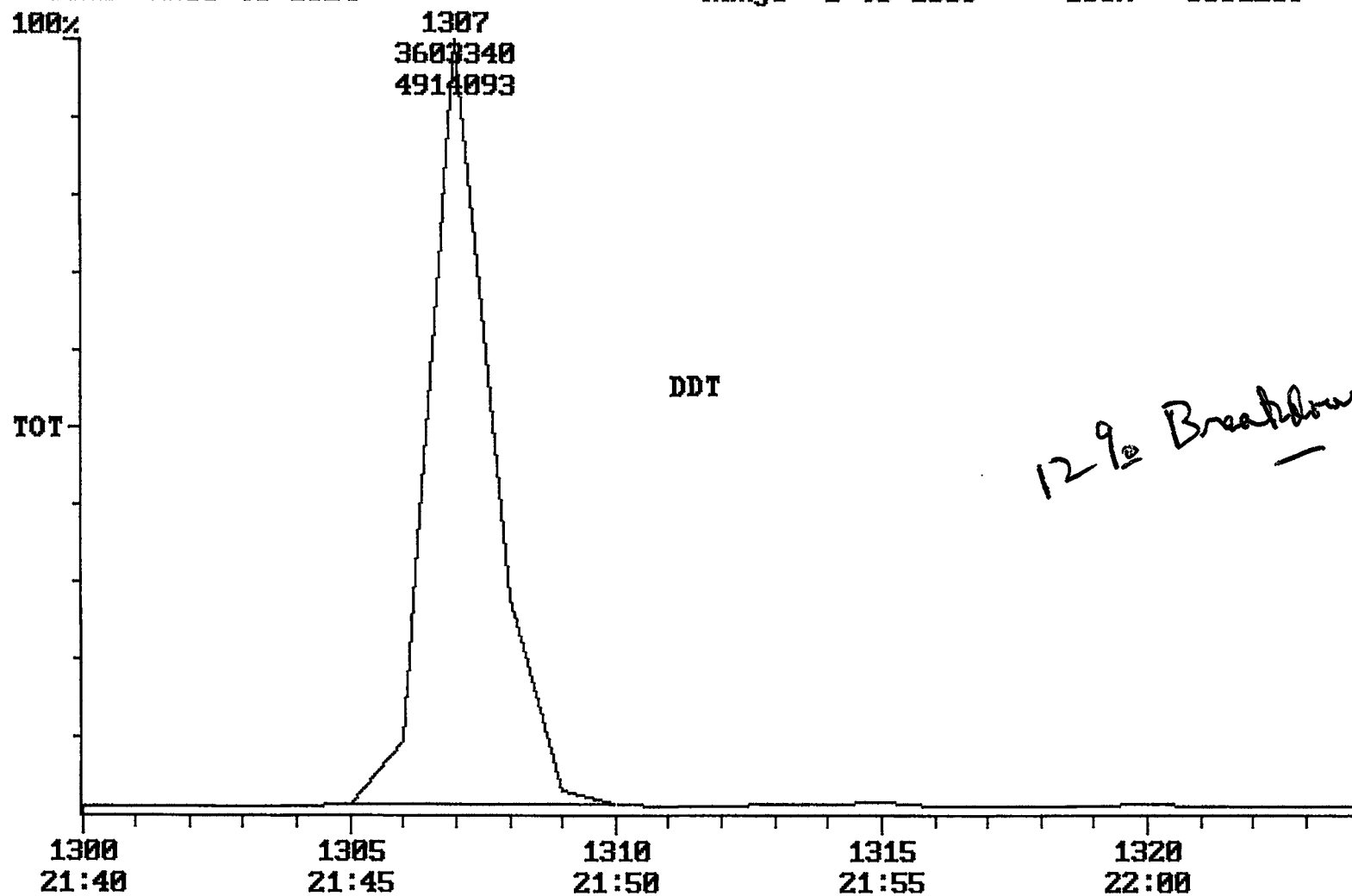
RIC: 35280

Mass Range: 45 - 355

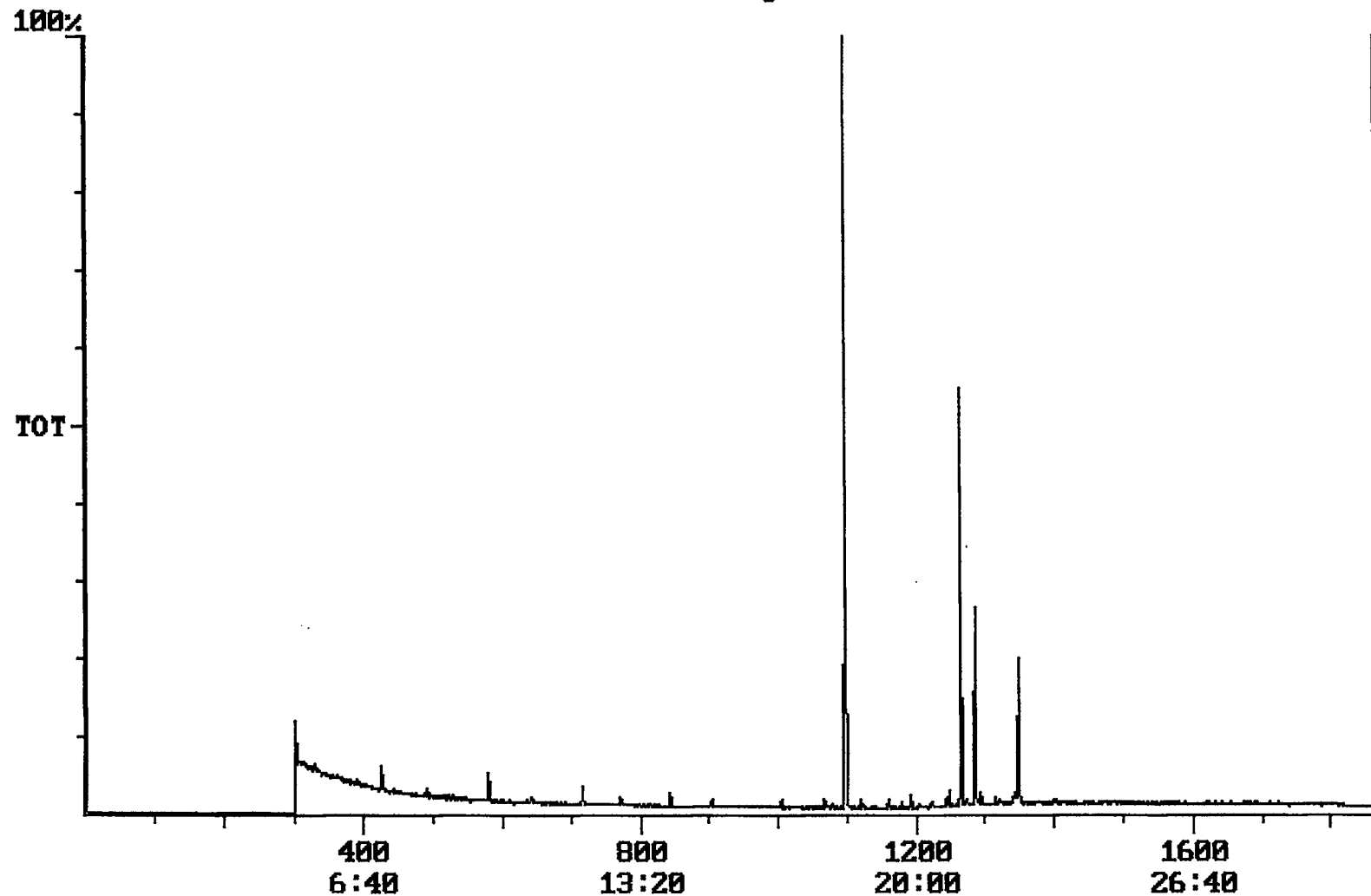
Plotted: 1300 to 1324

Range: 1 to 1859

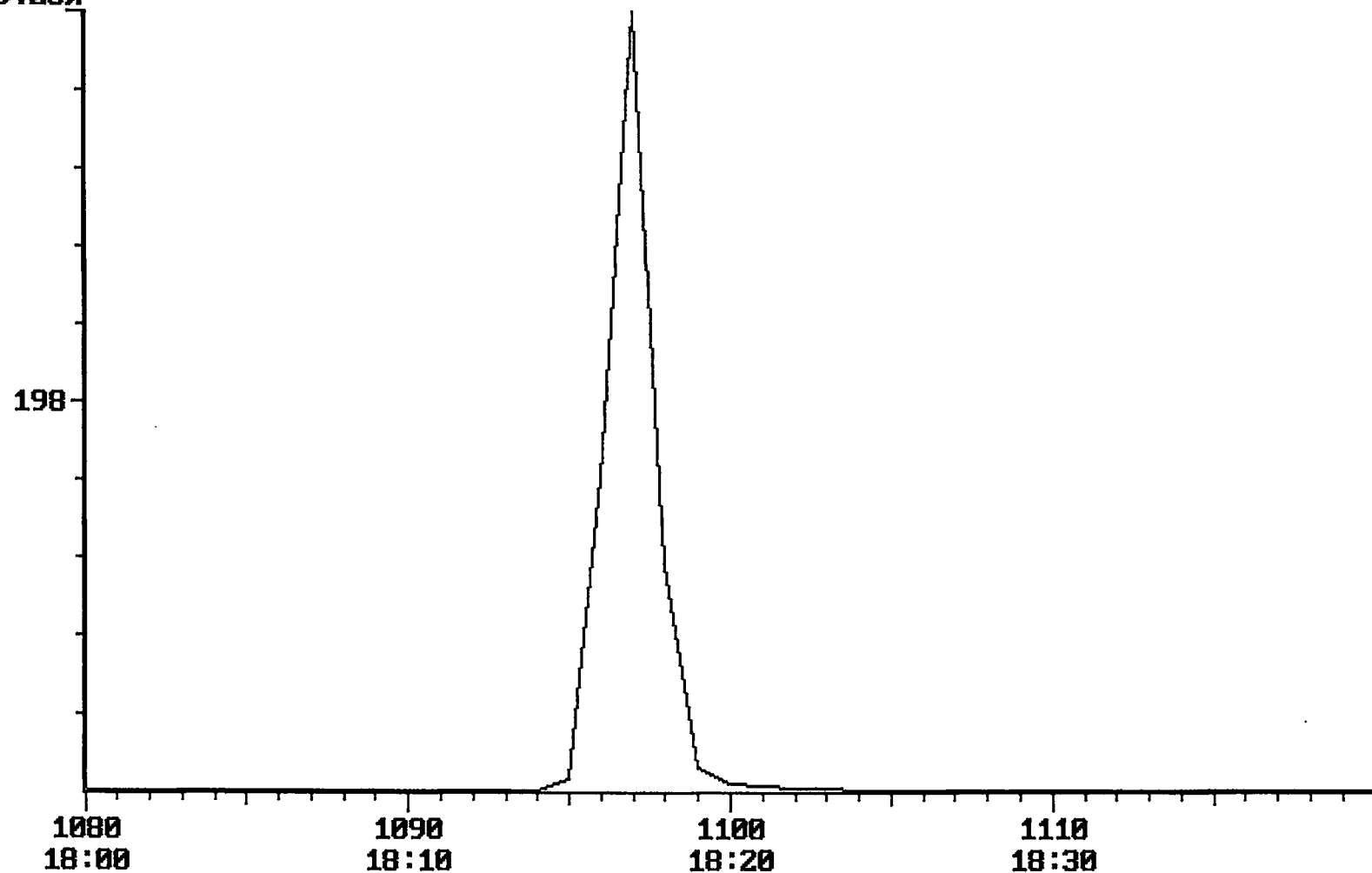
100% = 3651289



Chromatogram Plot File: E:\DFTB1201 Date: Dec-01-1991 16:19:05
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP
Scan No: 1 Retention Time: 0:01 RIC: 0 Mass Range: 0 - 0
Plotted: 1 to 1860 Range: 1 to 1860 100% = 3888273



Chromatogram Plot File: E:\DFTB1201 Date: Dec-01-1991 16:19:05
 Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP
 Scan No: 1080 Retention Time: 18:00 RIC: 37327 Mass Range: 47 - 149
 Plotted: 1080 to 1120 Range: 1 to 1060 100% = 3888273
 9.28%



Mass List

File: E:\DFTB1201 Date: Dec-01-1991 16:19:05

Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP

Scan No: 1097 Retention Time: 18:17 RIC: 3888273 Mass Range: 47 - 444

Peaks: 198 Mass of Base Peak: 442 Int: 522743 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
47	1,388	0.27	82	4,340	0.83	116	5,034	0.96
49	12,152	2.32	83	3,645	0.70	117	72,048	13.78
50	96,527	18.47	84	1,909	0.37	118	3,993	0.76
51	269,444	51.54	86	7,465	1.43	122	3,645	0.70
52	15,277	2.92	87	4,340	0.83	123	7,812	1.49
55	4,861	0.93	88	1,909	0.37	124	2,256	0.43
56	7,465	1.43	91	2,951	0.56	125	1,909	0.37
57	8,680	1.66	92	1,909	0.37	127	137,326	26.27
60	868	0.17	93	31,770	6.08	128	9,201	1.76
61	5,555	1.06	96	2,777	0.53	129	51,388	9.83
62	4,340	0.83	97	2,083	0.40	130	5,034	0.96
63	6,076	1.16	98	23,263	4.45	131	1,736	0.33
65	1,736	0.33	99	6,944	1.33	132	2,777	0.53
67	1,562	0.30	101	5,034	0.96	135	5,208	1.00
68	3,819	0.73	103	2,083	0.40	136	5,381	1.03
69	163,888	31.35	104	9,027	1.73	137	3,993	0.76
73	2,256	0.43	105	4,513	0.86	138	1,909	0.37
74	40,104	7.67	106	2,083	0.40	141	5,208	1.00
75	26,215	5.01	107	65,798	12.59	142	5,729	1.10
76	10,763	2.06	108	6,423	1.23	143	2,604	0.50
77	172,222	32.95	109	1,388	0.27	145	1,736	0.33
78	13,194	2.52	110	104,340	19.96	146	2,256	0.43
79	21,875	4.18	111	19,444	3.72	147	5,034	0.96
80	14,583	2.79	112	1,562	0.30	148	12,500	2.39
81	10,763	2.06	115	1,736	0.33	149	868	0.17

Mass List

File: E:\DFTB1201

Date: Dec-01-1991 16:19:05

Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP

Scan No: 1097

Retention Time: 18:17

RIC: 3888273

Mass Range: 47 - 444

Peaks: 198

Mass of Base Peak: 442

Int: 522743

Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
149	868	0.17	184	1,562	0.30	224	34,722	6.64
152	2,951	0.56	185	7,638	1.46	225	13,194	2.52
153	1,388	0.27	186	59,722	11.42	226	1,562	0.30
155	3,472	0.66	187	13,020	2.49	227	17,534	3.35
156	6,076	1.16	189	2,951	0.56	228	2,604	0.50
158	1,388	0.27	192	4,166	0.80	229	4,687	0.90
159	2,777	0.53	193	3,298	0.63	231	1,388	0.27
160	1,736	0.33	196	6,423	1.23	233	1,909	0.37
161	2,951	0.56	197	2,951	0.56	234	2,777	0.53
165	3,472	0.66	198	360,763	69.01	235	3,645	0.70
166	4,166	0.80	199	20,138	3.85	239	2,083	0.40
167	15,451	2.96	200	4,340	0.83	242	1,736	0.33
168	10,243	1.96	203	2,256	0.43	243	3,472	0.66
170	1,736	0.33	204	8,854	1.69	244	30,555	5.85
171	1,736	0.33	205	15,798	3.02	245	3,298	0.63
172	1,736	0.33	206	77,604	14.85	246	8,333	1.59
173	1,909	0.37	207	12,847	2.46	247	3,472	0.66
174	2,430	0.46	208	2,604	0.50	253	1,215	0.23
175	7,118	1.36	209	1,215	0.23	255	249,305	47.69
176	1,736	0.33	210	1,909	0.37	256	29,687	5.68
177	3,298	0.63	211	2,951	0.56	258	8,333	1.59
178	868	0.17	217	14,930	2.86	259	1,736	0.33
179	10,416	1.99	218	3,993	0.76	265	4,340	0.83
180	6,250	1.20	221	8,333	1.59	273	6,597	1.26
181	5,034	0.96	223	6,597	1.26	274	22,916	4.38

Mass List

File: E:\DFTB1201 Date: Dec-01-1991 16:19:05

Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP

Scan No: 1097 Retention Time: 18:17 RIC: 3088273 Mass Range: 47 - 444

Peaks: 198 Mass of Base Peak: 442 Int: 522743 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
223	6,597	1.26	274	22,916	4.38	346	2,430	0.46
224	34,722	6.64	275	120,486	23.05	351	1,215	0.23
225	13,194	2.52	276	18,923	3.62	352	5,902	1.13
226	1,562	0.30	277	13,541	2.59	353	4,166	0.80
227	17,534	3.35	278	1,562	0.30	354	6,944	1.33
228	2,604	0.50	280	1,041	0.20	355	2,430	0.46
229	4,687	0.90	283	2,083	0.40	359	1,909	0.37
231	1,388	0.27	285	2,777	0.53	364	1,562	0.30
233	1,909	0.37	289	1,388	0.27	365	24,652	4.72
234	2,777	0.53	293	2,256	0.43	366	3,645	0.70
235	3,645	0.70	294	1,909	0.37	370	1,909	0.37
239	2,083	0.40	296	28,645	5.48	372	7,465	1.43
242	1,736	0.33	297	5,208	1.00	383	1,736	0.33
243	3,472	0.66	303	3,125	0.60	384	1,909	0.37
244	30,555	5.85	304	3,645	0.70	396	1,215	0.23
245	3,298	0.63	314	2,604	0.50	402	2,430	0.46
246	8,333	1.59	315	6,423	1.23	403	3,472	0.66
247	3,472	0.66	316	2,083	0.40	421	4,340	0.83
253	1,215	0.23	323	9,201	1.76	422	2,951	0.56
255	249,305	47.69	324	3,645	0.70	423	33,680	6.44
256	29,687	5.68	327	2,430	0.46	424	6,770	1.30
258	8,333	1.59	332	1,562	0.30	441	84,201	16.11
259	1,736	0.33	334	7,118	1.36	442	522,743	100.00
265	4,340	0.83	335	2,256	0.43	443	103,125	19.73
273	6,597	1.26	341	3,125	0.60	444	9,201	1.76

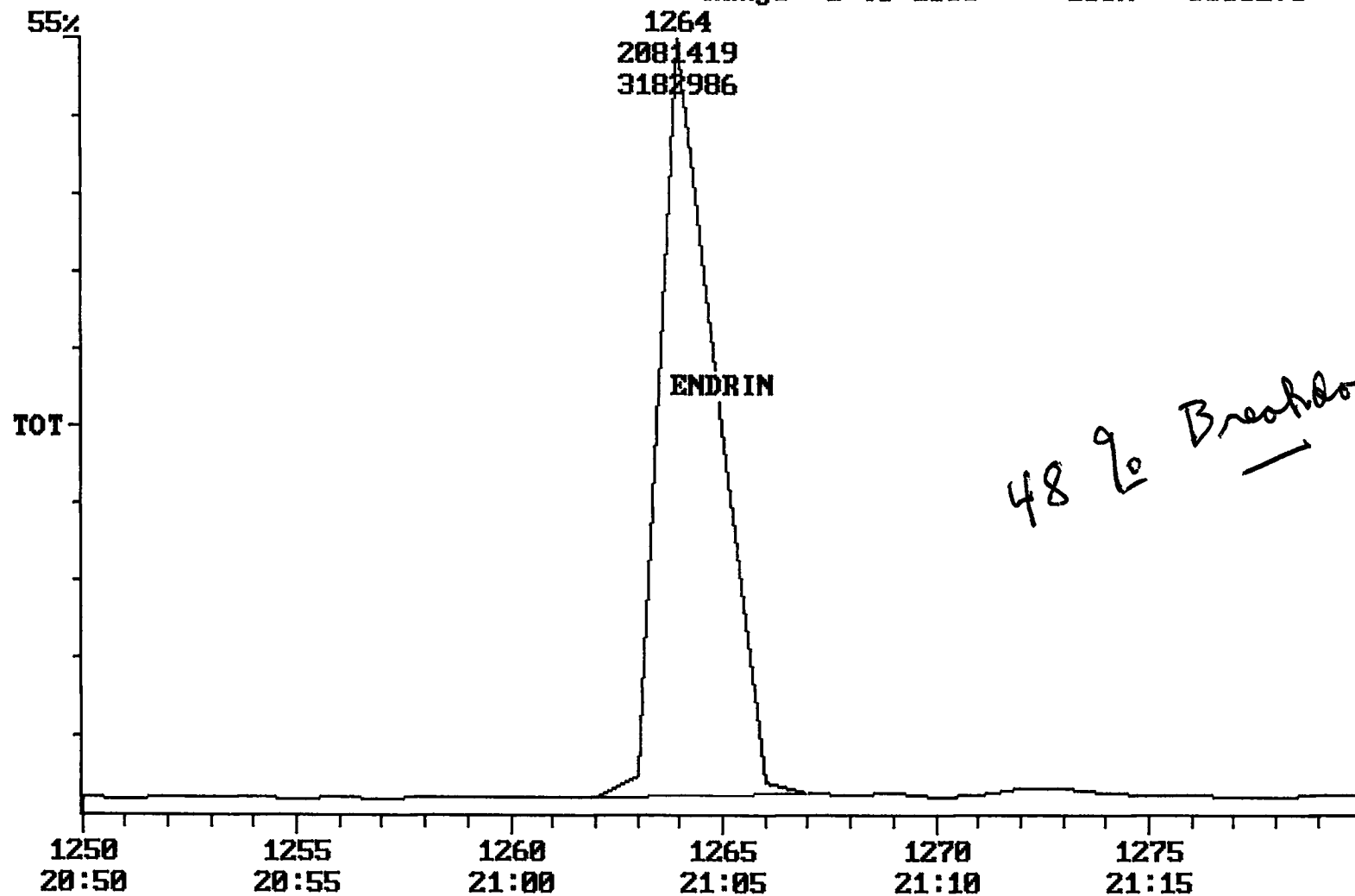
Chromatogram Plot

File: E:\DFTB1201 Date: Dec-01-1991 16:19:05

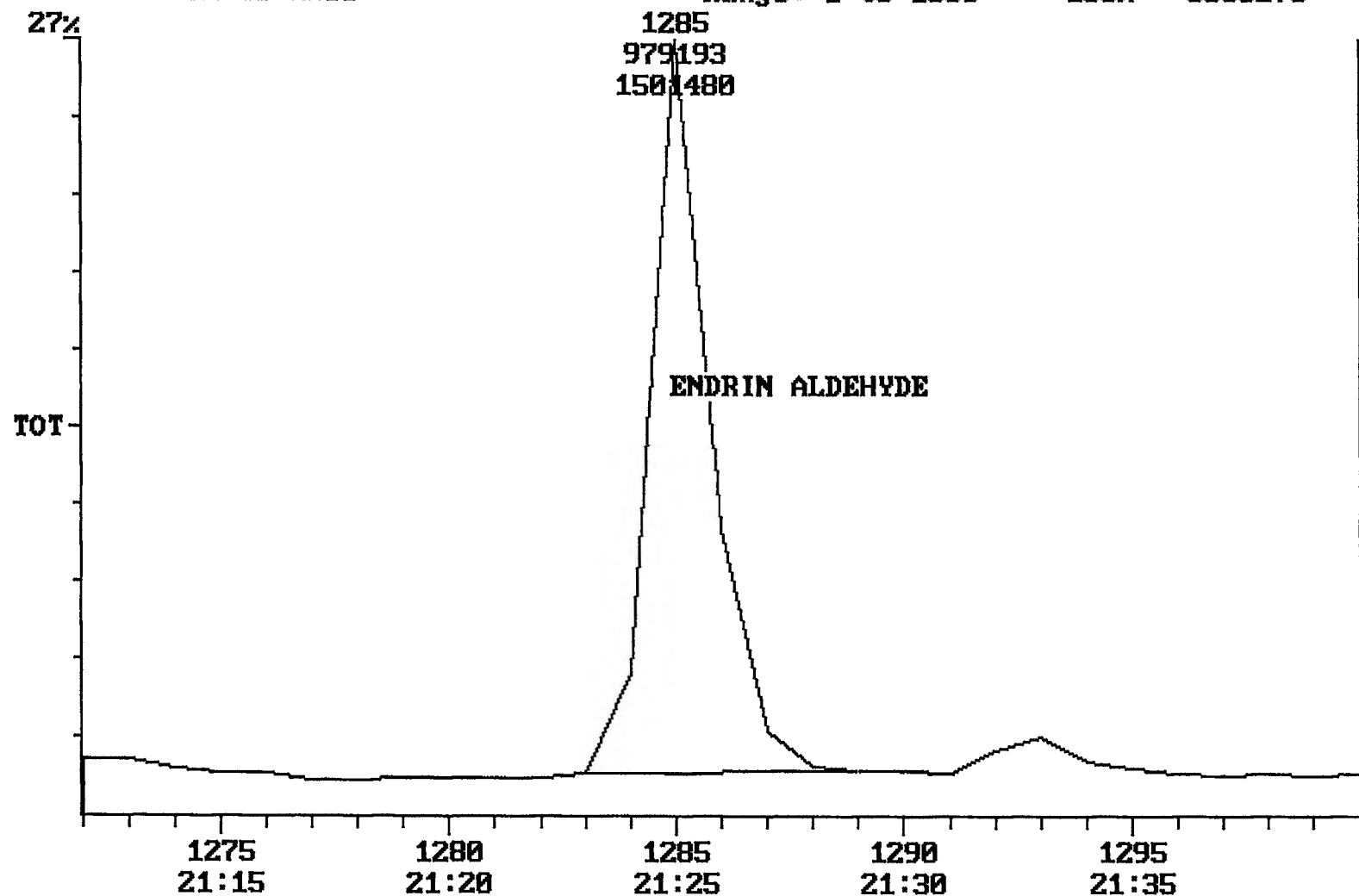
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP

Scan No: 1280 Retention Time: 21:20 RIC: 50217 Mass Range: 45 - 355

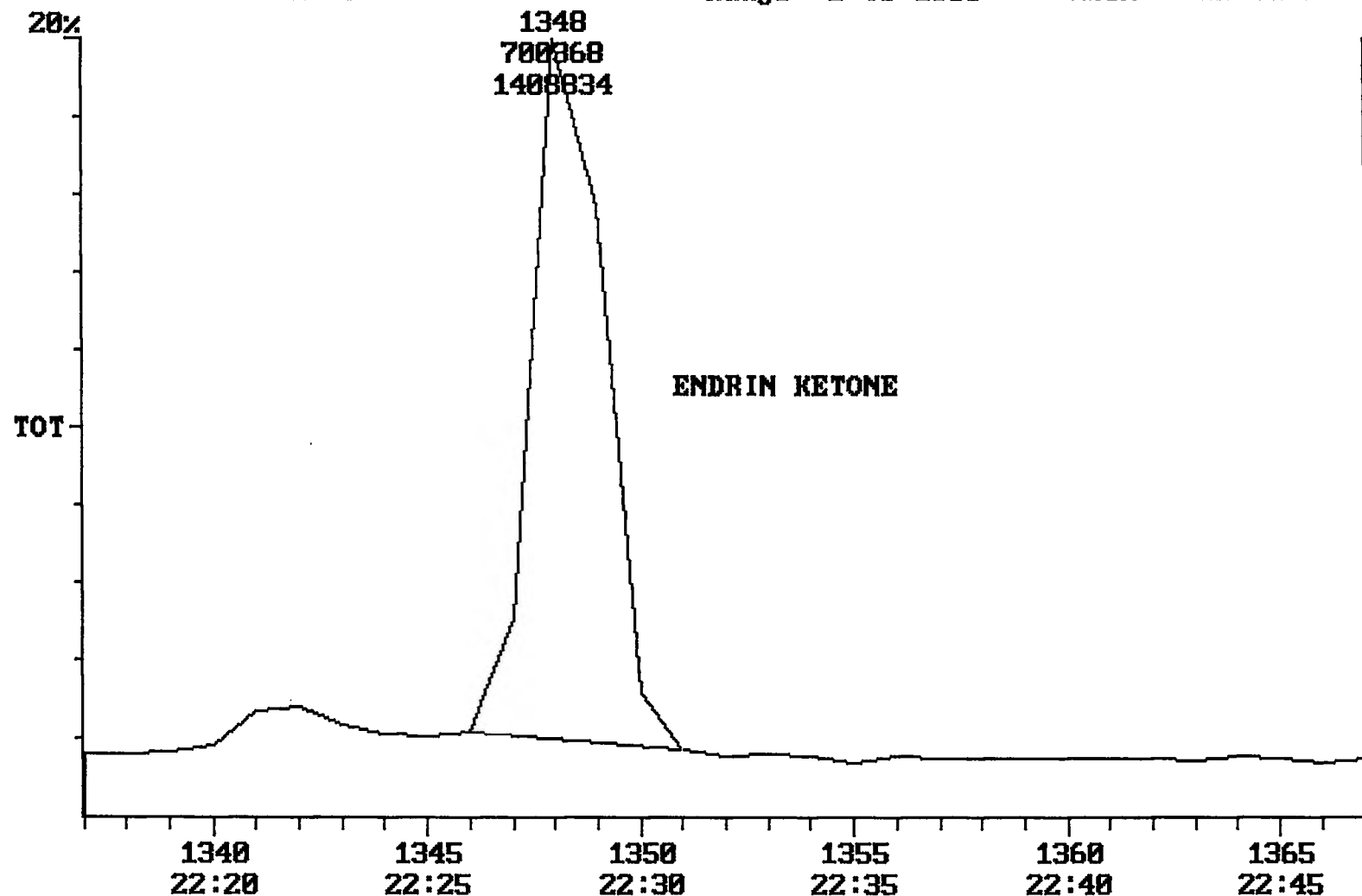
Plotted: 1250 to 1280 Range: 1 to 1860 100% = 3888273



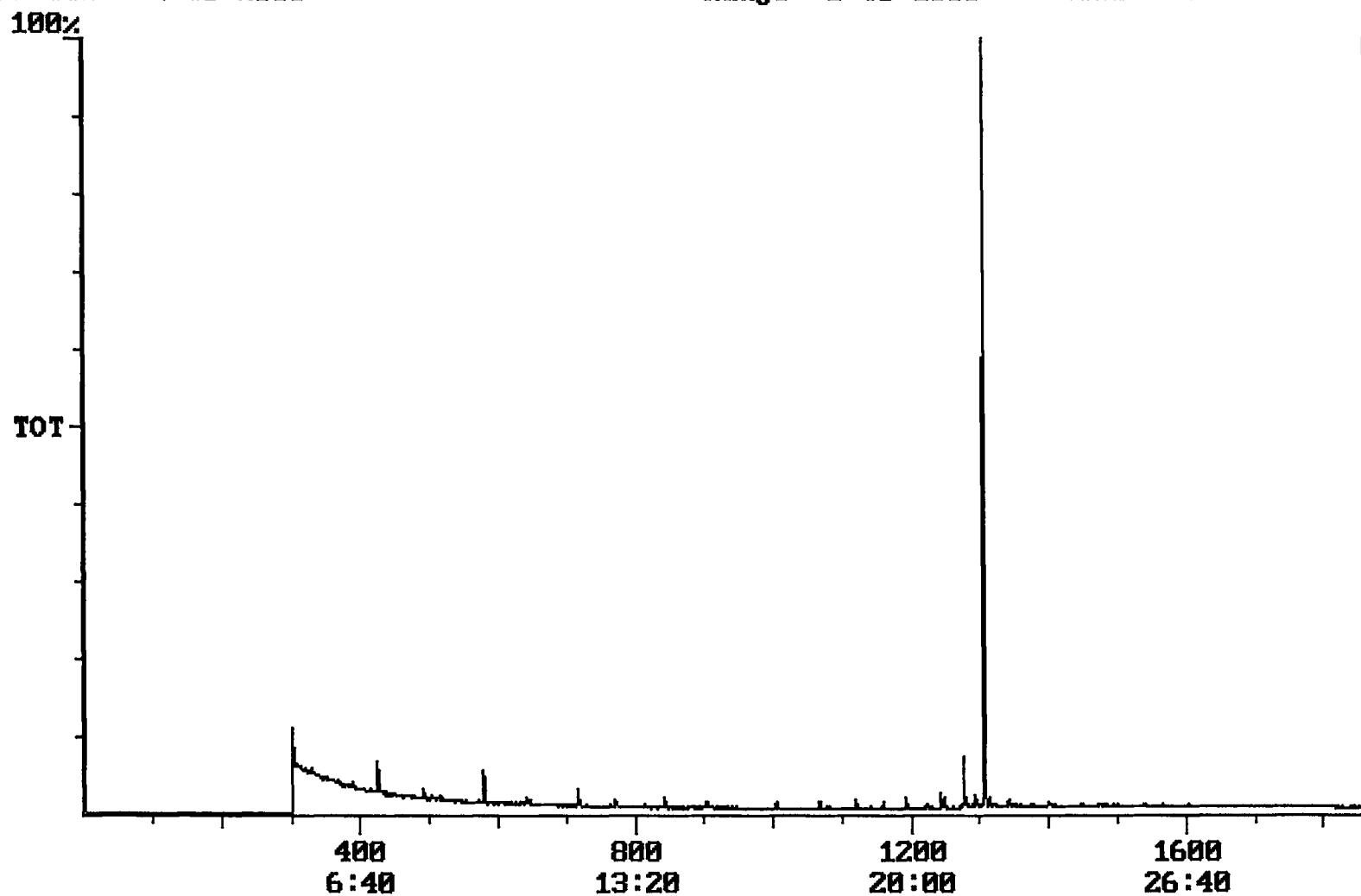
Chromatogram Plot File: E:\DFTB1201 Date: Dec-01-1991 16:19:05
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP
Scan No: 1300 Retention Time: 21:40 RIC: 54232 Mass Range: 45 - 281
Plotted: 1272 to 1300 Range: 1 to 1860 100% = 3888273



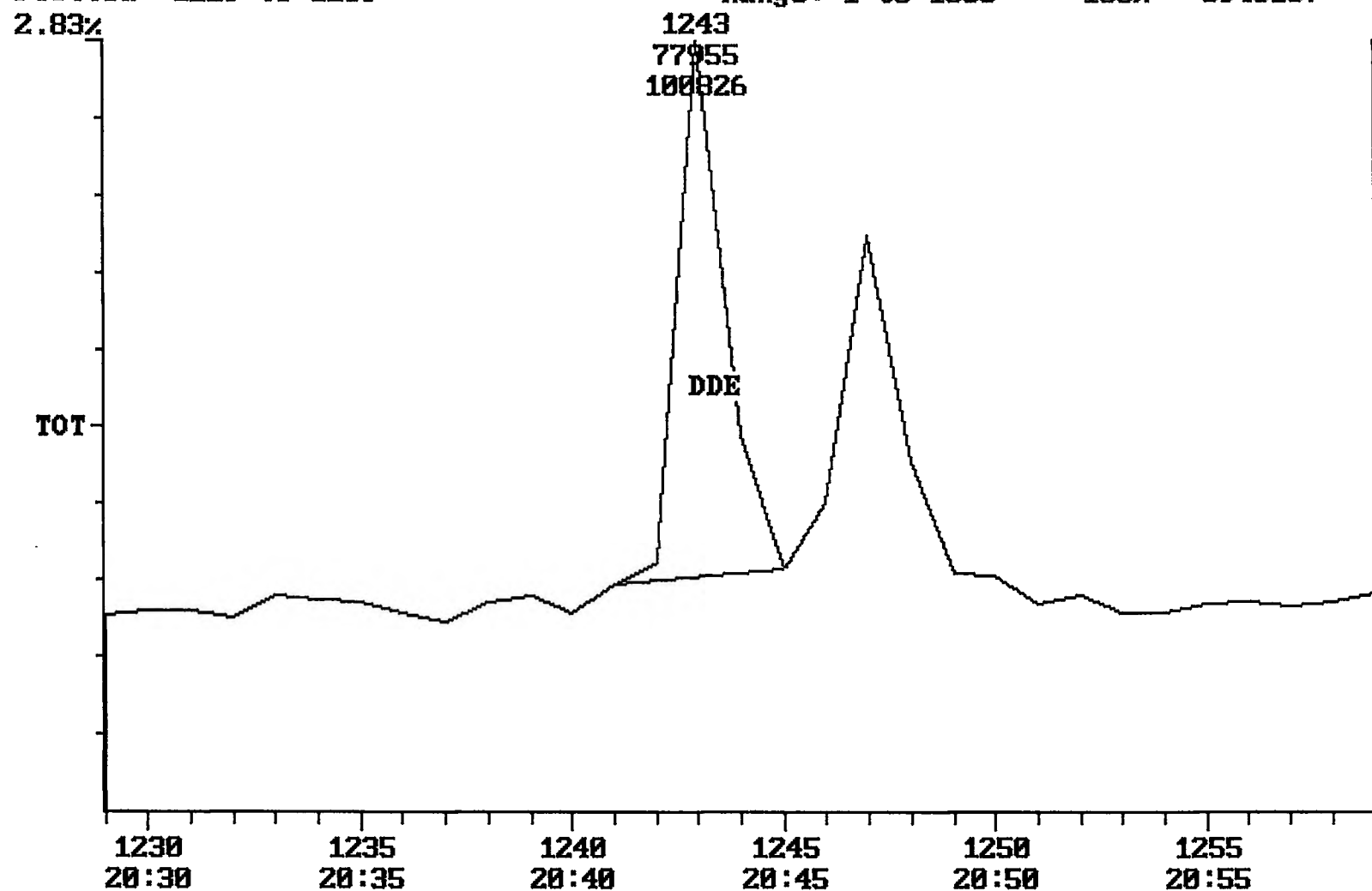
Chromatogram Plot File: E:\DFTB1201 Date: Dec-01-1991 16:19:05
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DFTPP
Scan No: 1367 Retention Time: 22:47 RIC: 56178 Mass Range: 45 - 281
Plotted: 1337 to 1367 Range: 1 to 1860 100% = 3888273



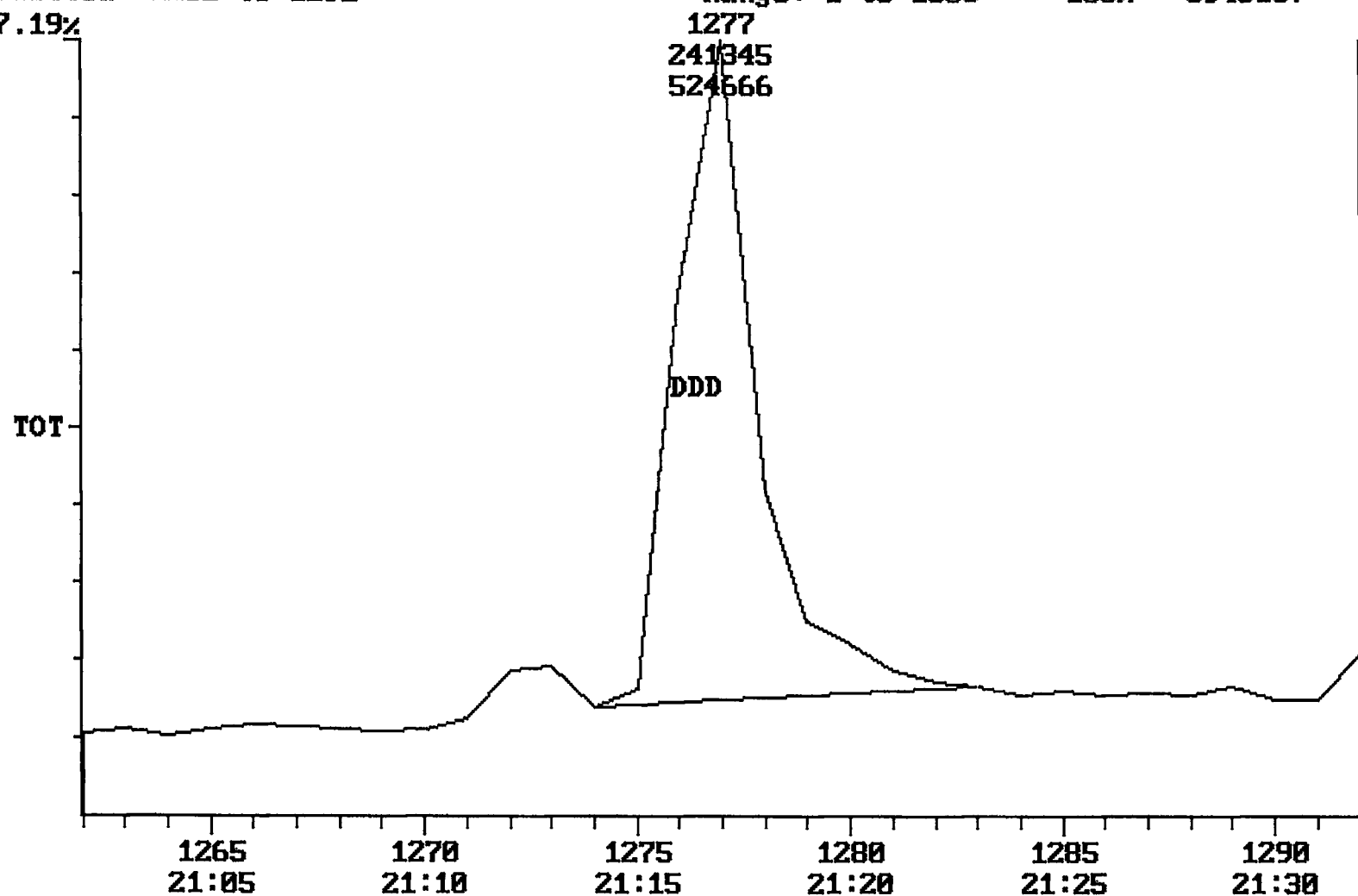
Chromatogram Plot File: E:\DDTB1201 Date: Dec-01-1991 16:53:39
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DDT
Scan No: 1 Retention Time: 0:01 RIC: 0 Mass Range: 0 - 0
Plotted: 1 to 1860 Range: 1 to 1860 100% = 3943187



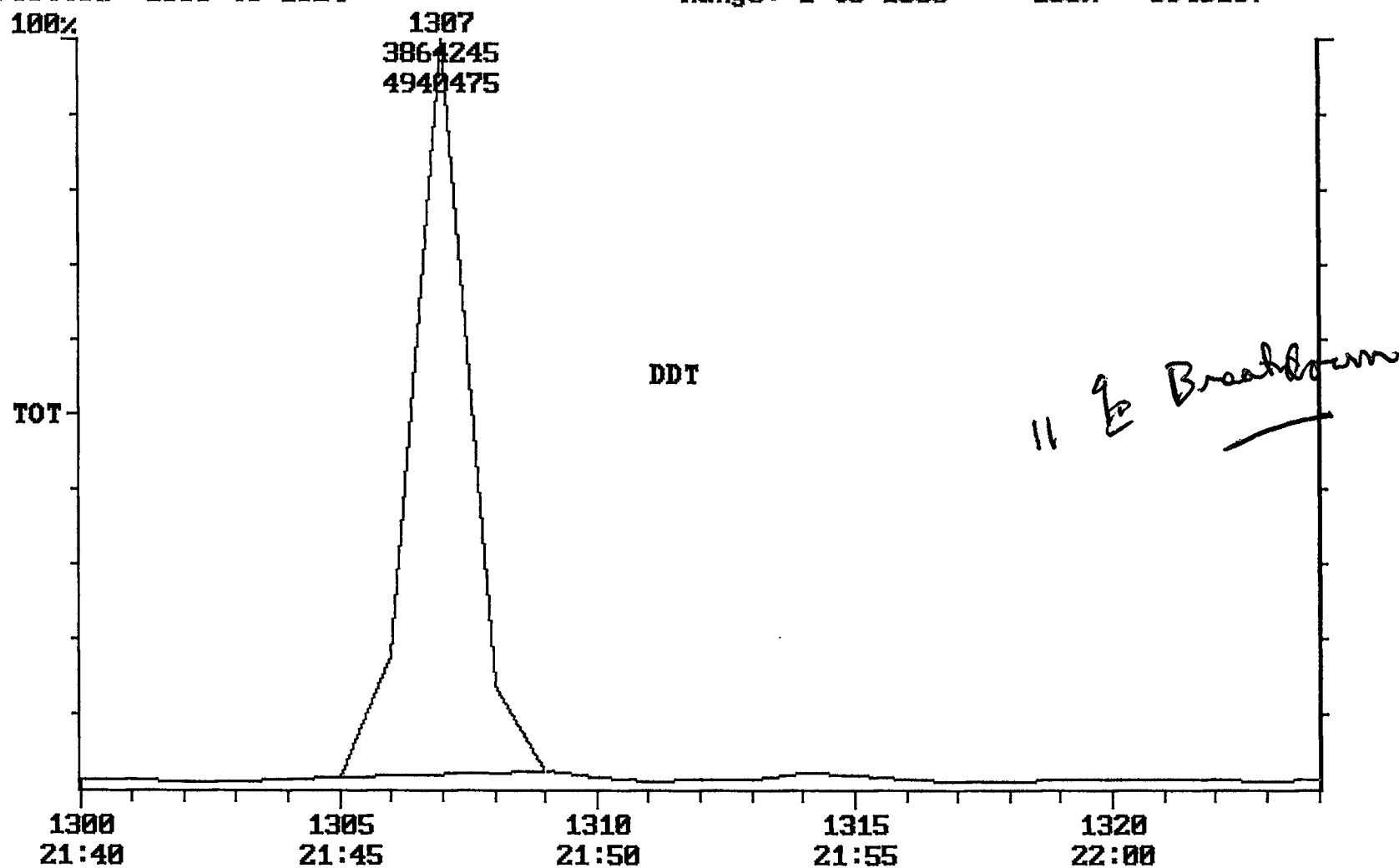
Chromatogram Plot File: E:\DDTB1201 Date: Dec-01-1991 16:53:39
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DDT
Scan No: 1259 Retention Time: 20:59 RIC: 31694 Mass Range: 47 - 207
Plotted: 1229 to 1259 Range: 1 to 1860 100% = 3943187



Chromatogram Plot File: E:\DDTB1201 Date: Dec-01-1991 16:53:39
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DDT
Scan No: 1292 Retention Time: 21:32 RIC: 58695 Mass Range: 47 - 429
Plotted: 1262 to 1292 Range: 1 to 1860 100% = 3943187
7.19%



Chromatogram Plot File: E:\DDTB1201 Date: Dec-01-1991 16:53:39
Comment: BOHLK; METHOD 525; INST 204; 12/01/01; DDT
Scan No: 1324 Retention Time: 22:04 RIC: 46893 Mass Range: 45 - 355
Plotted: 1300 to 1324 Range: 1 to 1860 100% = 3943187



Chromatogram Plot

File: F:\DFTA1203

Date: Dec-03-1991 10:57:57

Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DFTPP/ENDRIN

Scan No: 1

Retention Time: 0:01

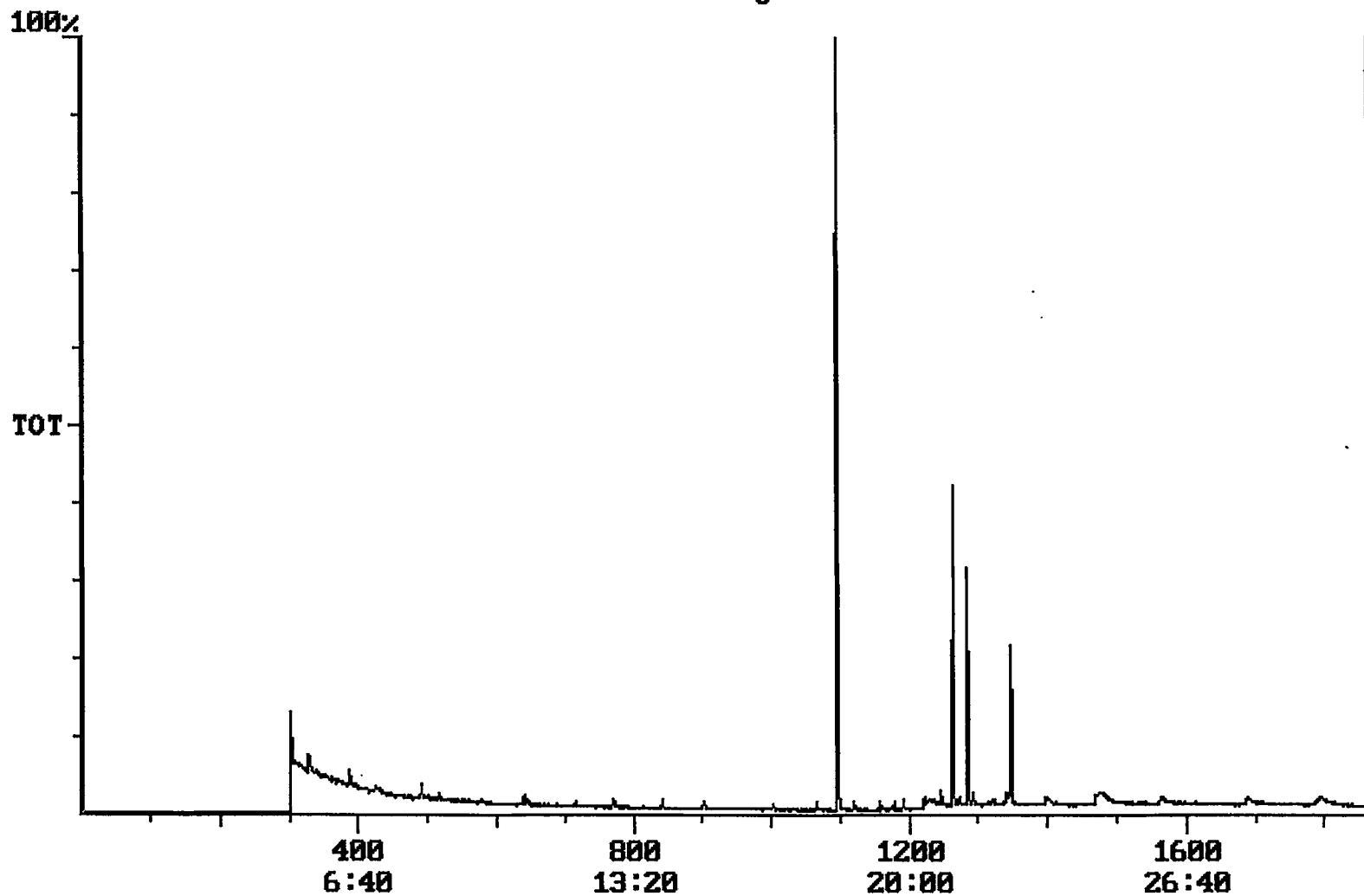
RIC: 0

Mass Range: 0 - 0

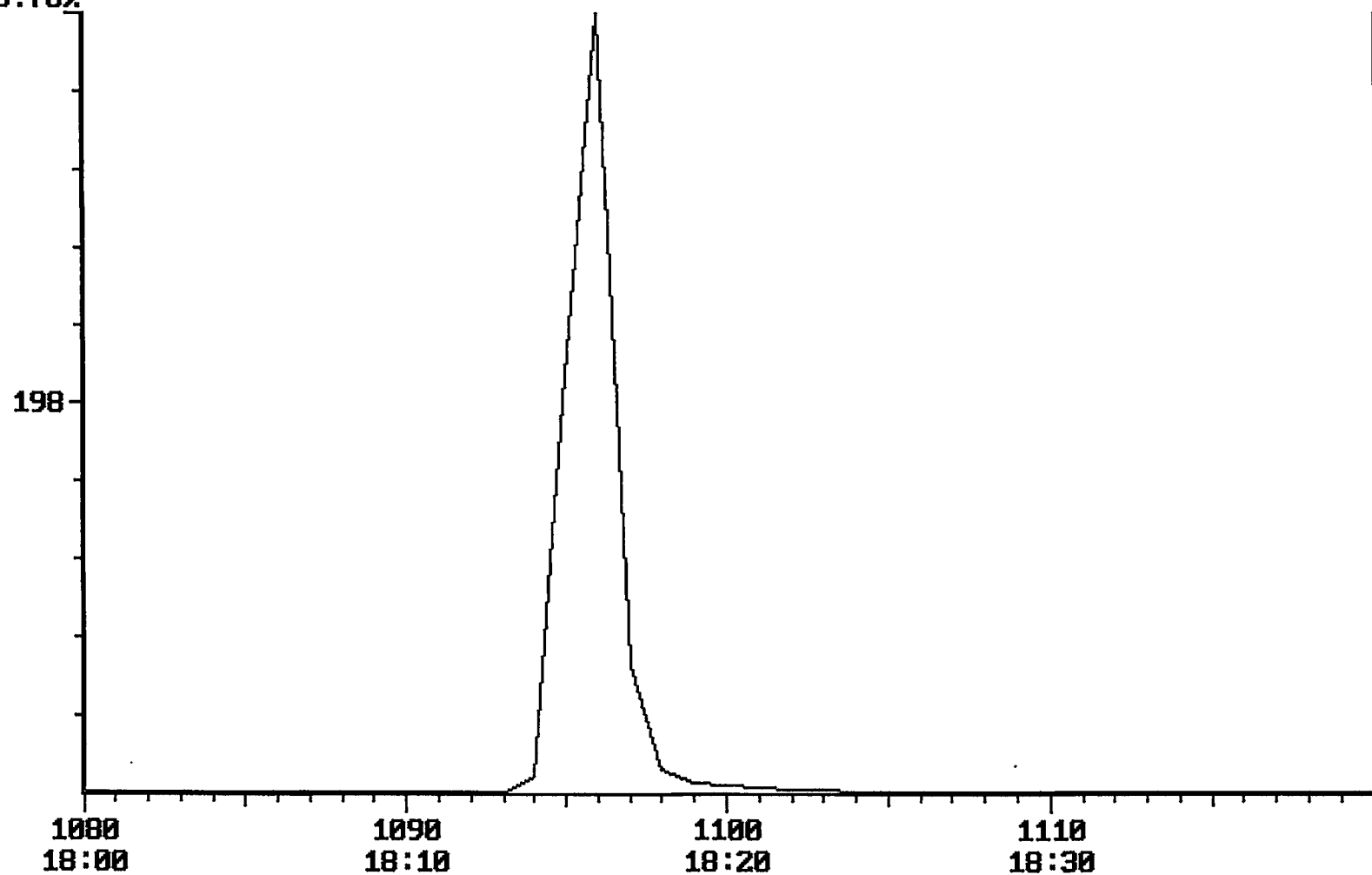
Plotted: 1 to 1860

Range: 1 to 1860

100% = 5006015



Chromatogram Plot File: F:\DFTA1203 Date: Dec-03-1991 10:57:57
Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DFTPP/ENDRIN
Scan No: 1080 Retention Time: 18:00 RIC: 31669 Mass Range: 47 - 281
Plotted: 1080 to 1120 Range: 1 to 1860 100% = 5006015
8.76%



Mass List

File: F:\DFTA1203

Date: Dec-03-1991 10:57:57

Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DFTPP/ENDRIN

Scan No: 1096

Retention Time: 18:16

RIC: 5006015

Mass Range: 48 - 445

Peaks: 202

Mass of Base Peak: 442

Int: 636585

Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
48	1,422	0.22	82	3,861	0.61	116	2,032	0.32
49	11,585	1.82	83	4,878	0.77	117	96,951	15.23
50	109,552	17.21	84	5,487	0.86	118	7,113	1.12
51	385,365	60.54	85	6,097	0.96	119	2,032	0.32
52	10,772	1.69	86	8,536	1.34	120	1,829	0.29
55	3,455	0.54	87	2,439	0.38	122	5,487	0.86
56	7,520	1.18	91	6,300	0.99	123	5,284	0.83
57	12,195	1.92	92	5,081	0.80	124	3,048	0.48
61	7,113	1.12	93	30,894	4.85	127	167,682	26.34
62	3,252	0.51	95	1,829	0.29	128	14,024	2.20
63	5,081	0.80	97	3,048	0.48	129	78,861	12.39
64	1,829	0.29	98	36,178	5.68	130	8,130	1.28
65	6,097	0.96	99	17,886	2.81	131	2,235	0.35
67	3,048	0.48	100	5,894	0.93	135	14,634	2.30
68	3,252	0.51	101	5,284	0.83	136	3,455	0.54
69	253,861	39.88	103	9,146	1.44	137	4,268	0.67
73	7,113	1.12	104	12,804	2.01	141	14,634	2.30
74	35,975	5.65	105	8,943	1.40	142	5,284	0.83
75	32,723	5.14	106	3,048	0.48	146	4,268	0.67
76	13,008	2.04	107	77,845	12.23	147	6,504	1.02
77	247,154	38.82	108	14,837	2.33	148	18,292	2.87
78	13,008	2.04	109	4,674	0.73	151	1,219	0.19
79	30,487	4.79	110	138,617	21.78	153	2,642	0.42
80	9,959	1.56	111	21,951	3.45	154	2,439	0.38
81	12,195	1.92	112	2,032	0.32	155	6,504	1.02

Mass List

File: F:\DFTA1203 Date: Dec-03-1991 10:57:57

Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DFTPP/ENDRIN

Scan No: 1096 Retention Time: 18:16 RIC: 5006015 Mass Range: 48 - 445

Peaks: 202 Mass of Base Peak: 442 Int: 636585 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
155	6,504	1.02	189	4,268	0.67	224	55,487	8.72
156	8,333	1.31	191	4,268	0.67	225	13,821	2.17
157	2,642	0.42	192	2,439	0.38	226	2,439	0.38
158	2,032	0.32	193	5,081	0.80	227	27,439	4.31
159	2,642	0.42	196	9,756	1.53	228	4,268	0.67
160	2,845	0.45	197	3,252	0.51	229	6,504	1.02
161	4,471	0.70	198	438,617	68.90	230	2,032	0.32
162	3,252	0.51	199	25,813	4.05	234	1,219	0.19
164	2,642	0.42	200	2,032	0.32	235	2,642	0.42
165	3,252	0.51	203	4,878	0.77	236	1,829	0.29
166	3,658	0.57	204	12,804	2.01	238	2,235	0.35
167	28,252	4.44	205	21,138	3.32	241	1,219	0.19
168	7,317	1.15	206	88,821	13.95	242	3,048	0.48
173	3,658	0.57	207	12,601	1.98	244	38,821	6.10
174	3,658	0.57	208	4,065	0.64	245	3,861	0.61
175	12,601	1.98	209	2,439	0.38	246	6,097	0.96
176	1,422	0.22	211	2,642	0.42	249	2,439	0.38
177	3,861	0.61	212	2,235	0.35	255	313,617	49.27
178	2,235	0.35	215	3,861	0.61	256	38,821	6.10
179	9,349	1.47	216	2,845	0.45	258	11,178	1.76
180	8,333	1.31	217	24,796	3.90	259	2,845	0.45
181	7,317	1.15	218	3,658	0.57	263	1,829	0.29
185	8,943	1.40	221	7,723	1.21	265	5,691	0.89
186	79,878	12.55	222	3,252	0.51	270	2,642	0.42
187	25,000	3.93	223	3,658	0.57	271	1,219	0.19

Mass List

File: F:\DFTA1203 Date: Dec-03-1991 10:57:57

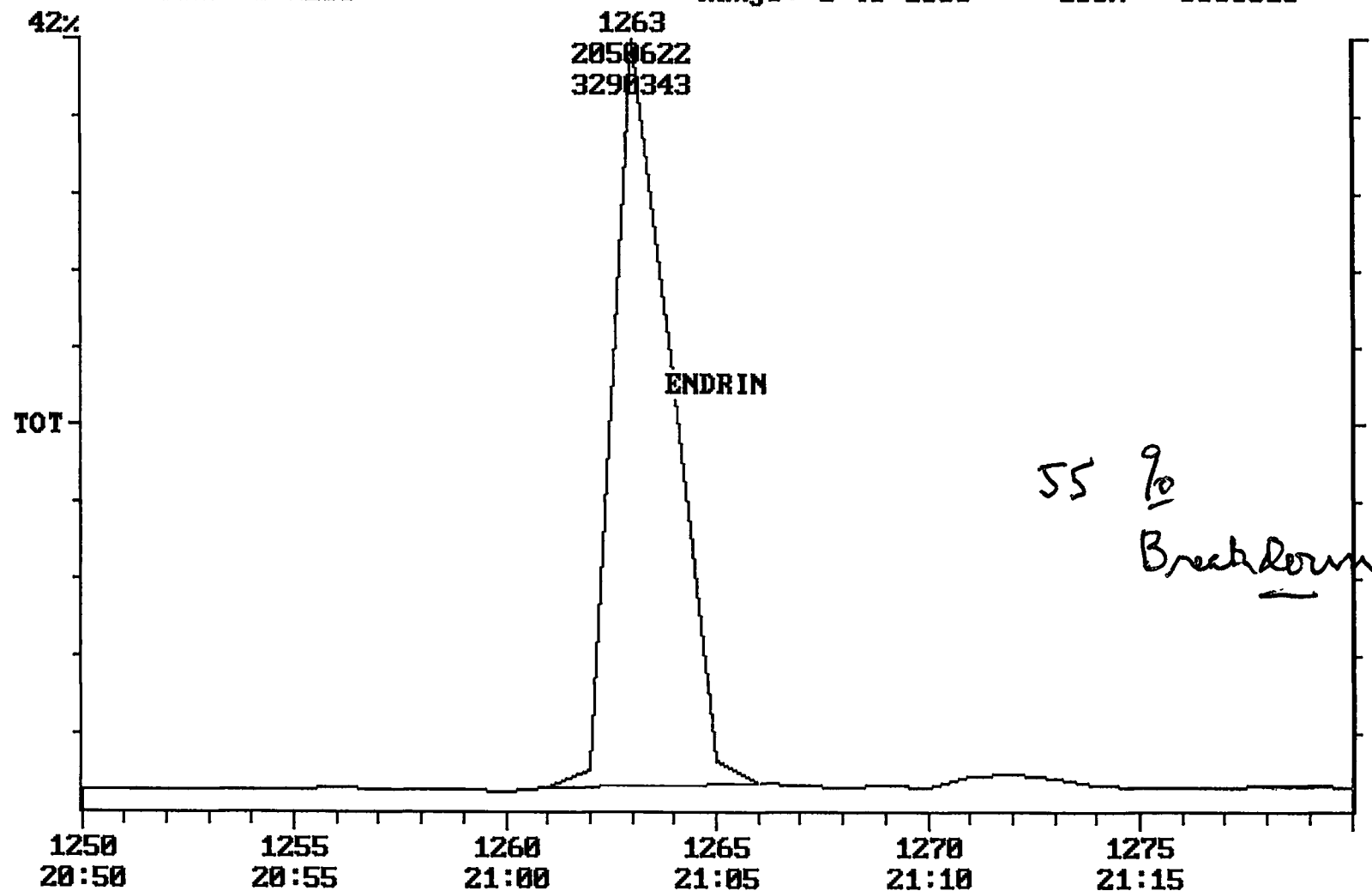
Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DFTPP/ENDRIN

Scan No: 1096 Retention Time: 18:16 RIC: 5006015 Mass Range: 48 - 445

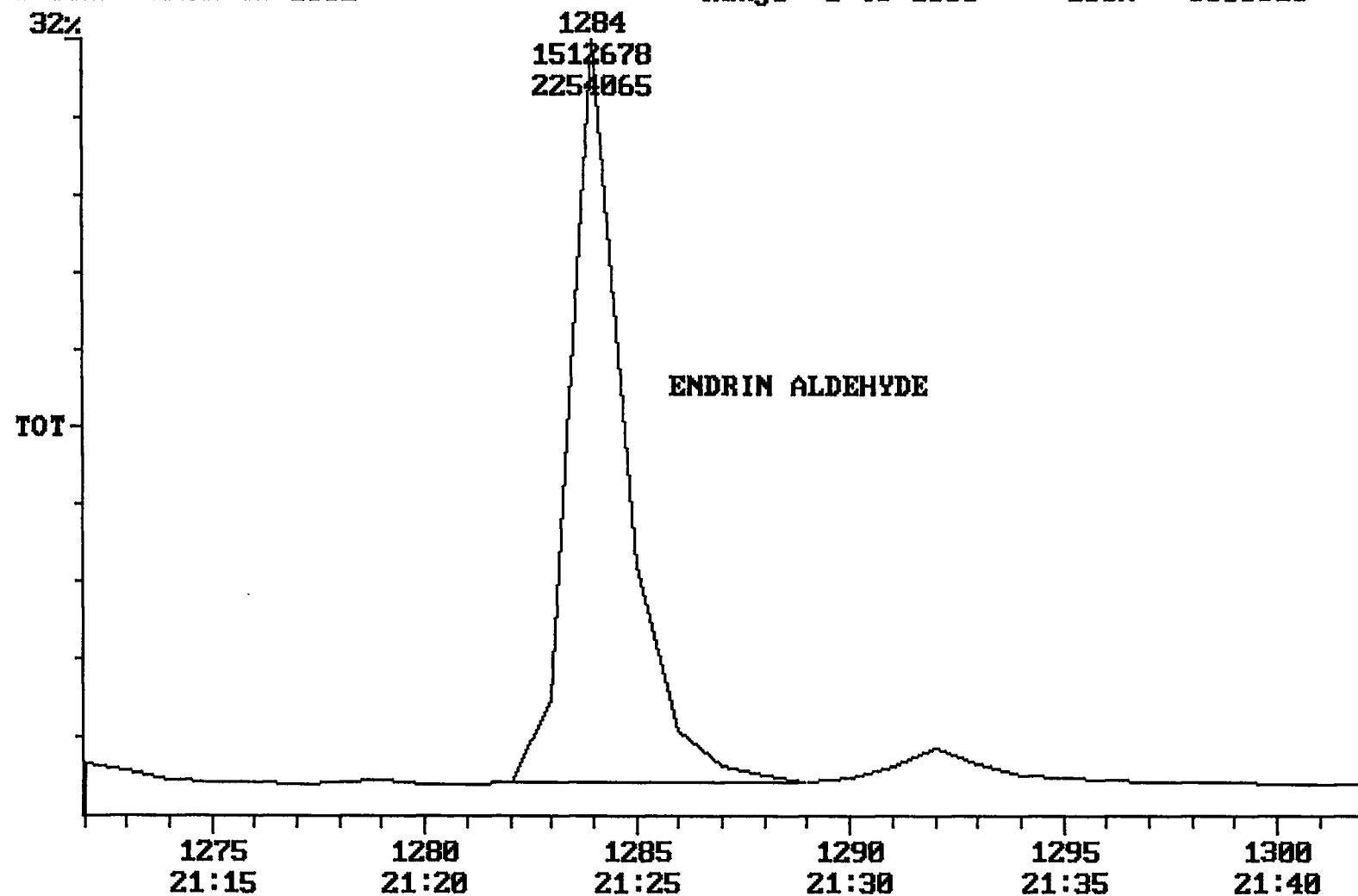
Peaks: 202 Mass of Base Peak: 442 Int: 636585 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
227	27,439	4.31	276	16,869	2.65	353	6,504	1.02
228	4,268	0.67	277	19,105	3.00	354	10,162	1.60
229	6,504	1.02	278	4,674	0.73	355	1,219	0.19
230	2,032	0.32	280	1,219	0.19	365	32,723	5.14
234	1,219	0.19	285	4,268	0.67	366	1,829	0.29
235	2,642	0.42	286	1,626	0.26	371	2,032	0.32
236	1,829	0.29	292	1,829	0.29	372	9,756	1.53
238	2,235	0.35	293	2,439	0.38	373	6,910	1.09
241	1,219	0.19	294	1,219	0.19	383	1,626	0.26
242	3,048	0.48	296	38,821	6.10	390	1,829	0.29
244	38,821	6.10	297	9,959	1.56	391	2,642	0.42
245	3,861	0.61	299	1,626	0.26	402	2,845	0.45
246	6,097	0.96	303	3,252	0.51	403	6,504	1.02
249	2,439	0.38	304	2,235	0.35	404	2,439	0.38
255	313,617	49.27	314	3,048	0.48	405	1,829	0.29
256	38,821	6.10	315	6,504	1.02	410	1,626	0.26
258	11,178	1.76	316	5,894	0.93	421	1,829	0.29
259	2,845	0.45	323	16,463	2.59	422	4,674	0.73
263	1,829	0.29	324	5,081	0.80	423	28,861	4.53
265	5,691	0.89	327	2,642	0.42	424	5,487	0.86
270	2,642	0.42	334	15,243	2.39	441	103,252	16.22
271	1,219	0.19	335	5,894	0.93	442	636,585	100.00
273	7,317	1.15	341	3,455	0.54	443	114,227	17.94
274	33,739	5.30	346	4,878	0.77	444	2,845	0.45
275	158,943	24.97	352	4,674	0.73	445	1,626	0.26

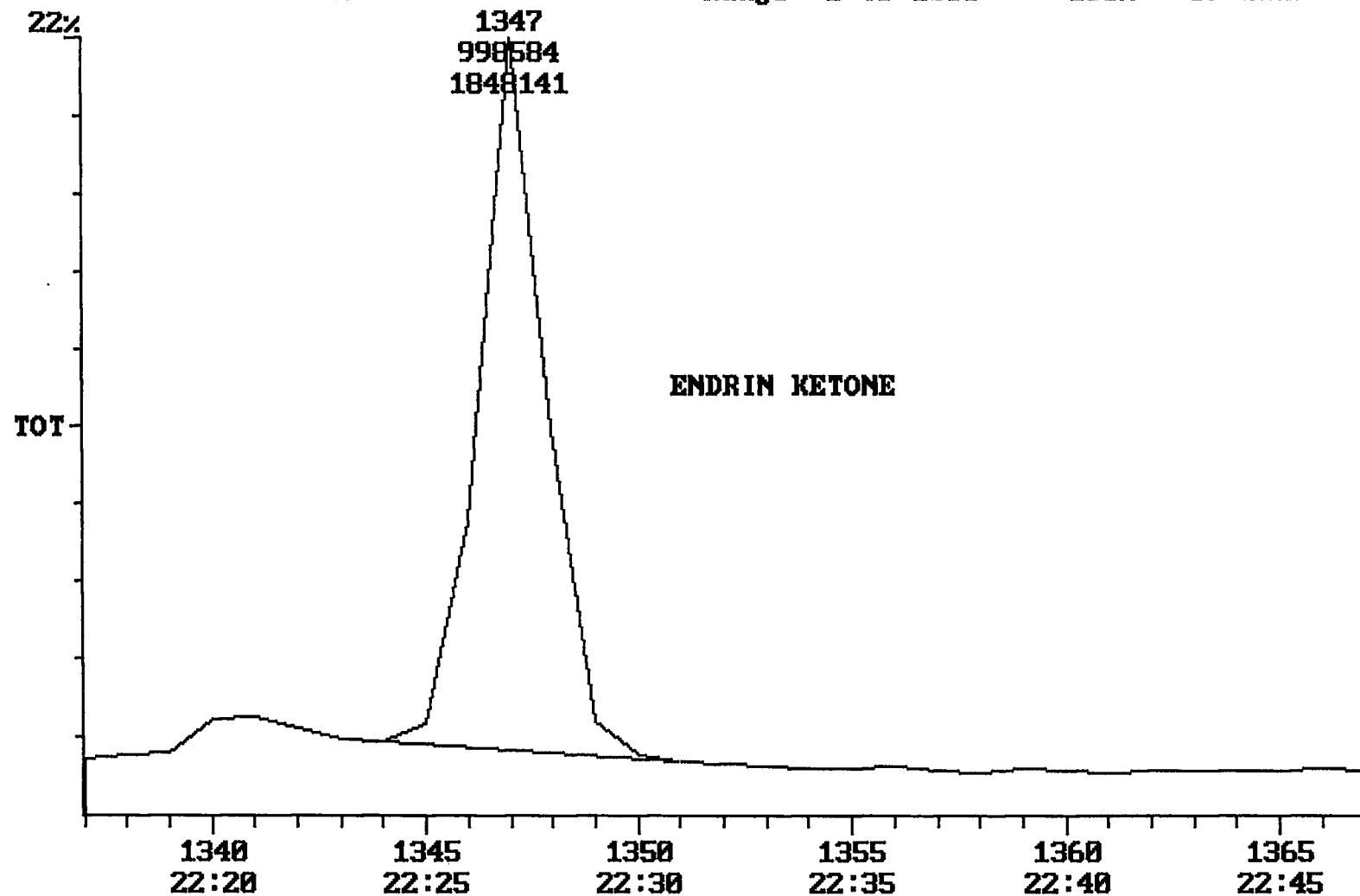
Chromatogram Plot File: F:\DFTA1203 Date: Dec-03-1991 10:57:57
Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DFTPP/ENDRIN
Scan No: 1280 Retention Time: 21:20 RIC: 61206 Mass Range: 45 - 429
Plotted: 1250 to 1280 Range: 1 to 1860 100% = 5006015



Chromatogram Plot File: F:\DFTA1203 Date: Dec-03-1991 10:57:57
Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DFTPP/ENDRIN
Scan No: 1302 Retention Time: 21:42 RIC: 63115 Mass Range: 45 - 429
Plotted: 1272 to 1302 Range: 1 to 1860 100% = 5006015



Chromatogram Plot File: F:\DFTA1203 Date: Dec-03-1991 10:57:57
Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DFTPP/ENDRIN
Scan No: 1367 Retention Time: 22:47 RIC: 61272 Mass Range: 45 - 355
Plotted: 1337 to 1367 Range: 1 to 1860 100% = 5006015



Chromatogram Plot

File: F:\DDTA1203

Date: Dec-03-1991 11:32:26

Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DDT

Scan No: 1

Retention Time: 0:01

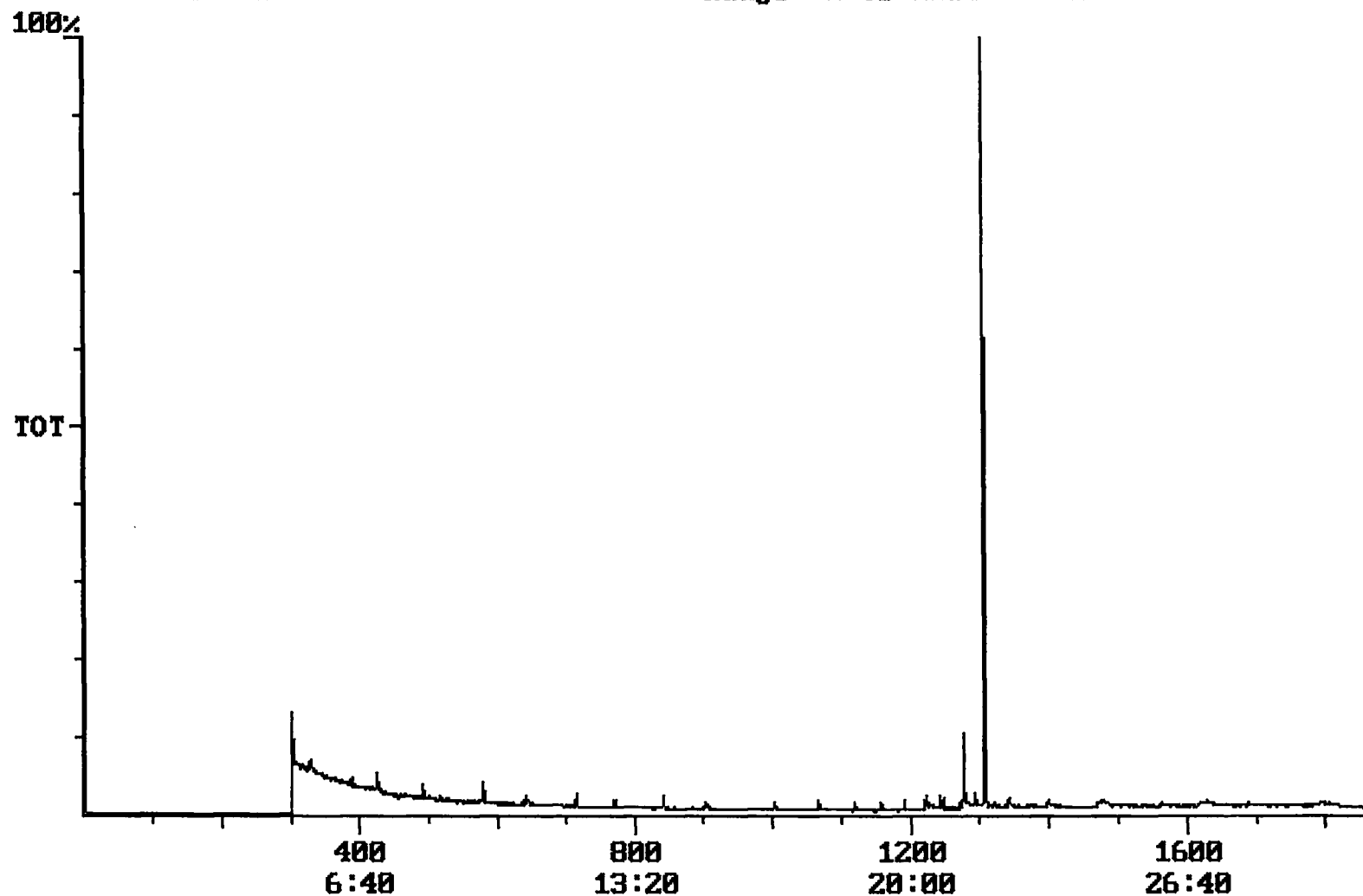
RIC: 0

Mass Range: 0 - 0

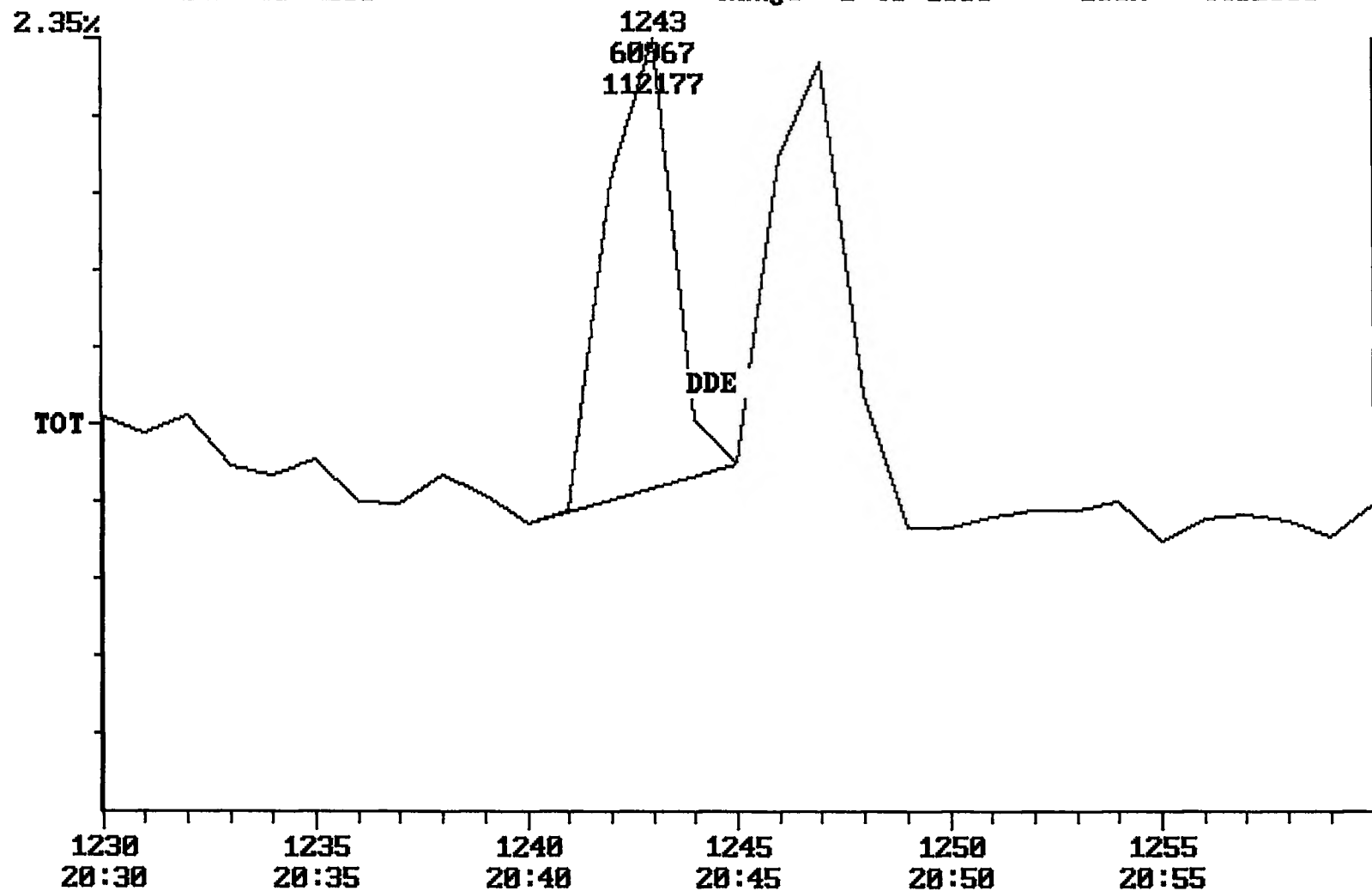
Plotted: 1 to 1860

Range: 1 to 1860

100% = 4462663



Chromatogram Plot File: F:\DDTA1203 Date: Dec-03-1991 11:32:26
Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DDT
Scan No: 1260 Retention Time: 21:00 RIC: 41404 Mass Range: 45 - 429
Plotted: 1230 to 1260 Range: 1 to 1860 100% = 4462663



Chromatogram Plot

File: F:\DDTA1203

Date: Dec-03-1991 11:32:26

Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DDT

Scan No: 1292

Retention Time: 21:32

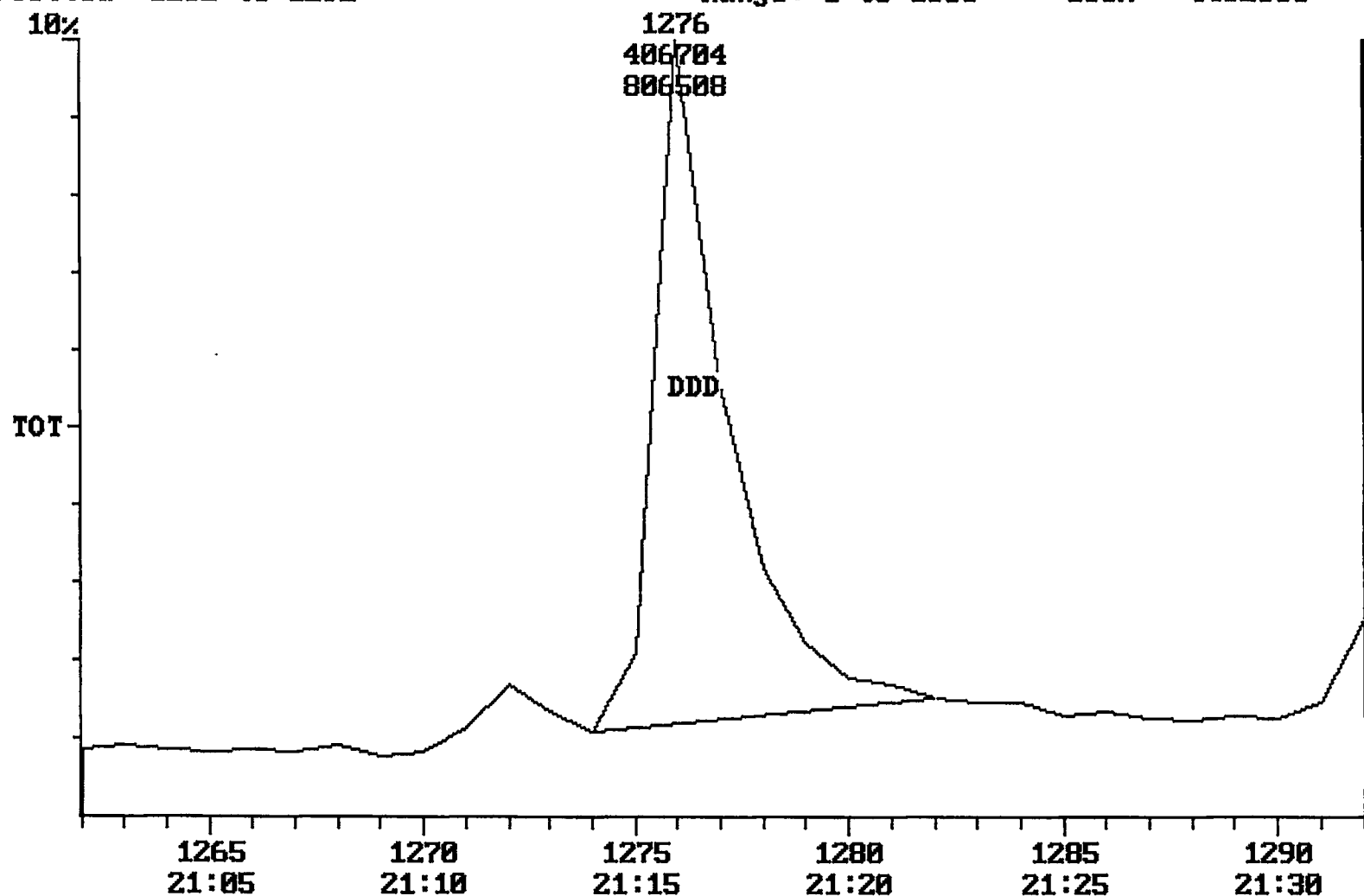
RIC: 115509

Mass Range: 45 - 431

Plotted: 1262 to 1292

Range: 1 to 1860

100% = 4462663



Chromatogram Plot

File: F:\DDTA1203

Date: Dec-03-1991 11:32:26

Comment: BOHLK; METHOD 525; INST 204; 12/03/01; DDT

Scan No: 1324

Retention Time: 22:04

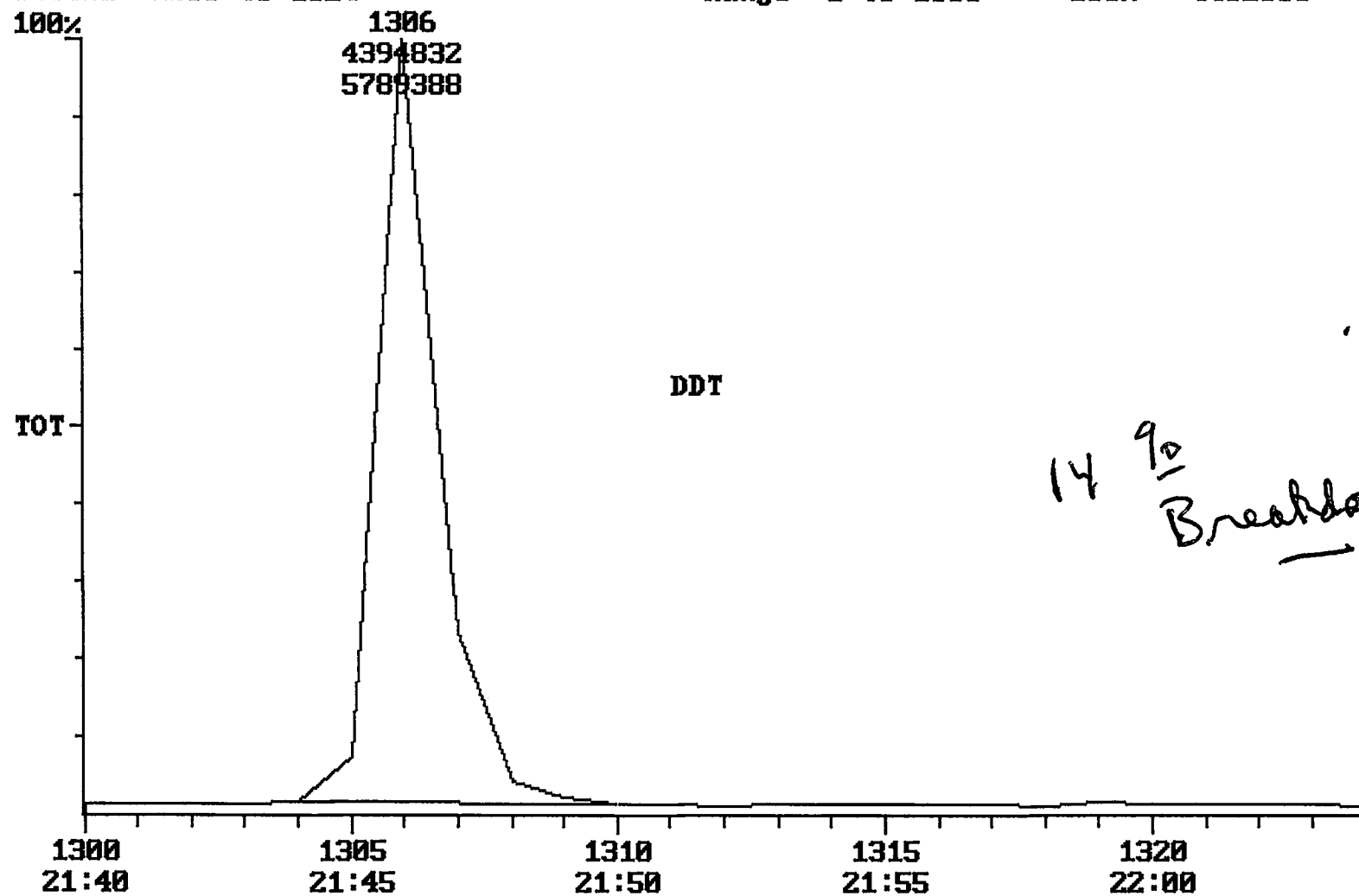
RIC: 49087

Mass Range: 45 - 429

Plotted: 1300 to 1324

Range: 1 to 1860

100% = 4462663



DFTPP

Data File : C:\MSDCHEM\1\DATA\121801\DFTPP3.D
Acq On : 18 Dec 2001 9:38 am
Sample :
Misc :
MS Integration Params: BENZO.P

Vial: 2
Operator: McLean
Inst : Instrumen
Multiplr: 1.00

Method : C:\MSDCHEM\1\5973N\121401.M (RTE Integrator)
Title : Quantitation For Method 525.2

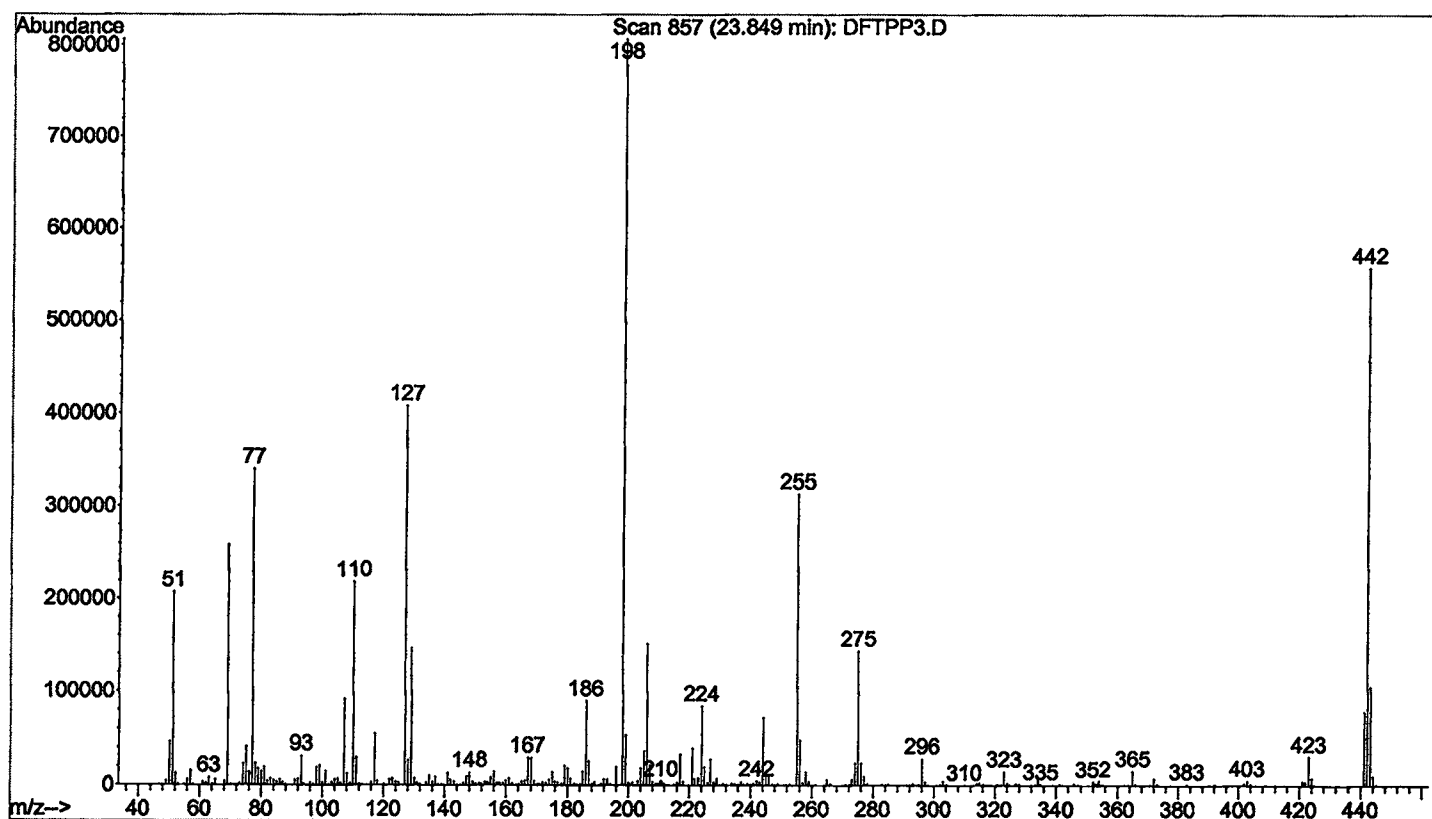
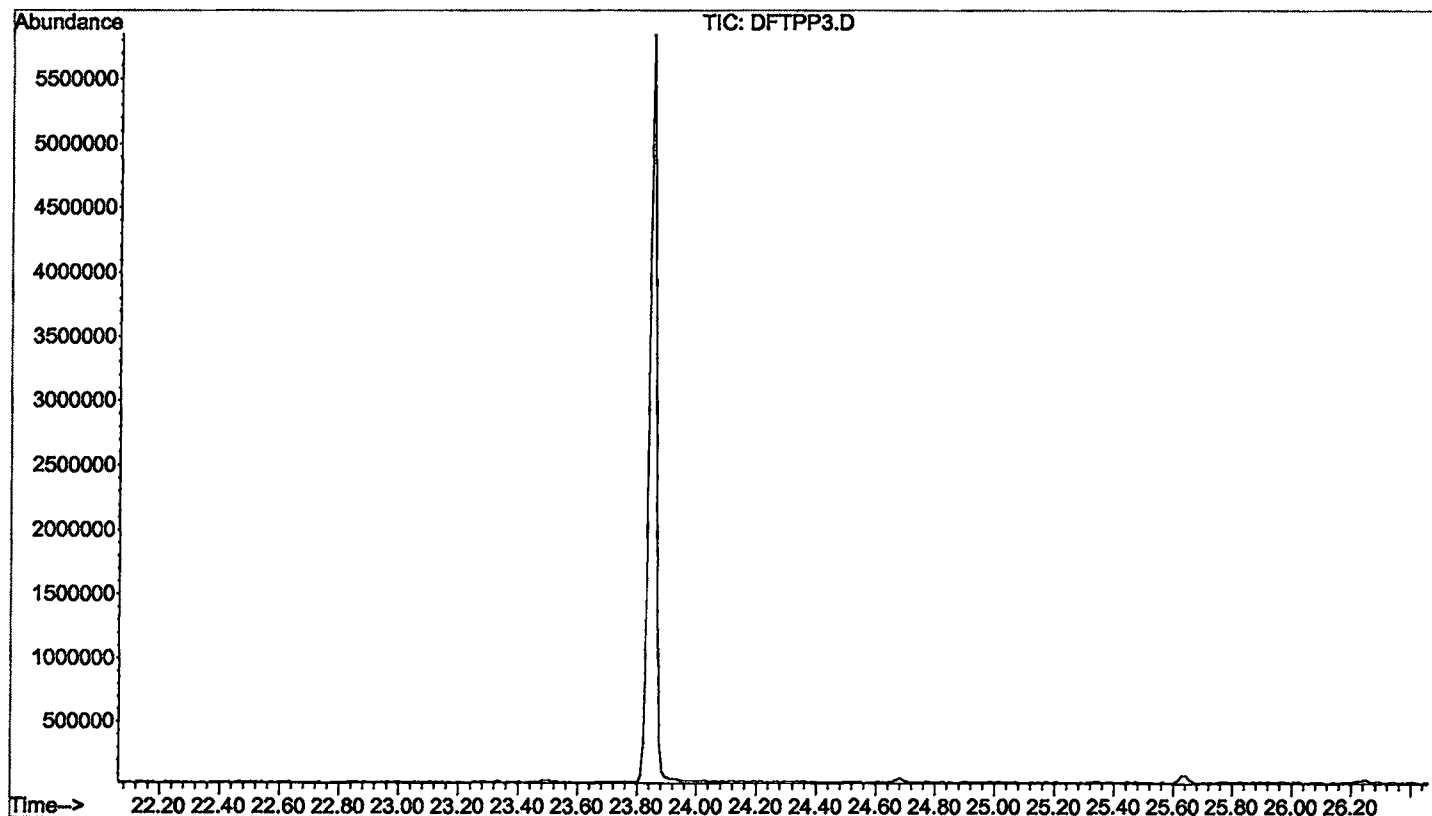
Spectrum Information: Scan 857

Target Mass	Rel. to Mass	Lower Limit%	Upper Limit%	Rel. Abn%	Raw Abn	Result Pass/Fail
51	198	10	80	25.7	207424	PASS
68	69	0.00	2	1.7	4438	PASS
69	198	0.00	100	32.1	258880	PASS
70	69	0.00	2	0.4	1139	PASS
127	198	10	80	50.6	408320	PASS
197	198	0.00	2	0.0	0	PASS
198	198	50	100	100.0	806272	PASS
199	198	5	9	6.6	53416	PASS
275	198	10	60	17.7	143104	PASS
365	198	1	100	1.9	15488	PASS
441	443	0.01	100	74.7	78352	PASS
442	198	50	100	69.1	557184	PASS
443	442	15	24	18.8	104912	PASS

for
Re-exts

DFTPP3.D 121401.M Wed Dec 19 14:47:24 2001

File : C:\MSDCHEM\1\DATA\121801\DFTPP3.D
 Operator : McLean
 Acquired : 18 Dec 2001 9:38 am using AcqMethod BIG3
 Instrument : Instrumen
 Sample Name:
 Misc Info :
 Vial Number: 2



Area Percent Report

Data File : C:\MSDCHEM\1\DATA\121801\DFTPP3.D

Acq On : 18 Dec 2001 9:38 am

Sample :

Misc :

MS Integration Params: BENZO.P

Vial: 2

Operator: McLean

Inst : Instrumen

Multiplr: 1.00

Method : C:\MSDCHEM\1\5973N\121401.M (RTE Integrator)

Title : Quantitation For Method 525.2

Smoothing : ON

Sampling : 1

Start Thrs: 0.15

Stop Thrs : 0

Filtering: 5

Min Area: 2500 Area counts

Max Peaks: 10

Peak Location: TOP

If leading or trailing edge < 100 prefer < Baseline drop else tangent >

Peak separation: 5

Signal : TIC

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	27.876	1296	1301	1307	rM	3367110	6363875	100.00%	93.551%
2	28.475	1365	1367	1369	rM 3	89485	120949	1.90%	1.778%
3	29.527	1479	1483	1486	rM 5	160811	317781	4.99%	4.671%

Sum of corrected areas: 6802605

Area Percent Report

Data File : C:\MSDCHEM\1\DATA\121801\DDT3.D

Acq On : 18 Dec 2001 10:18 am

Sample :

Misc :

MS Integration Params: BENZO.P

Vial: 3

Operator: McLean

Inst : Instrumen

Multiplr: 1.00

Method : C:\MSDCHEM\1\5973N\121401.M (RTE Integrator)

Title : Quantitation For Method 525.2

Smoothing : ON

Sampling : 1

Start Thrs: 0.15

Stop Thrs : 0

Filtering: 5

Min Area: 2500 Area counts

Max Peaks: 10

Peak Location: TOP

If leading or trailing edge < 100 prefer < Baseline drop else tangent >

Peak separation: 5

Signal : TIC

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	28.266	1340	1344	1351	rM 2	603220	1201894	14.80%	12.889%
2	29.146	1433	1441	1447	rM	4580037	8123006	100.00%	87.111%

Sum of corrected areas: 9324900

DFTPP

Data File : C:\MSDCHEM\1\DATA\121901\DFTPP1.D
Acq On : 19 Dec 2001 12:24 pm
Sample : dftpp +end
Misc :
MS Integration Params: BENZO.P

Vial: 1
Operator: McLean
Inst : Instrumen
Multiplr: 1.00

Method : C:\MSDCHEM\1\5973N\121401.M (RTE Integrator)
Title : Quantitation For Method 525.2

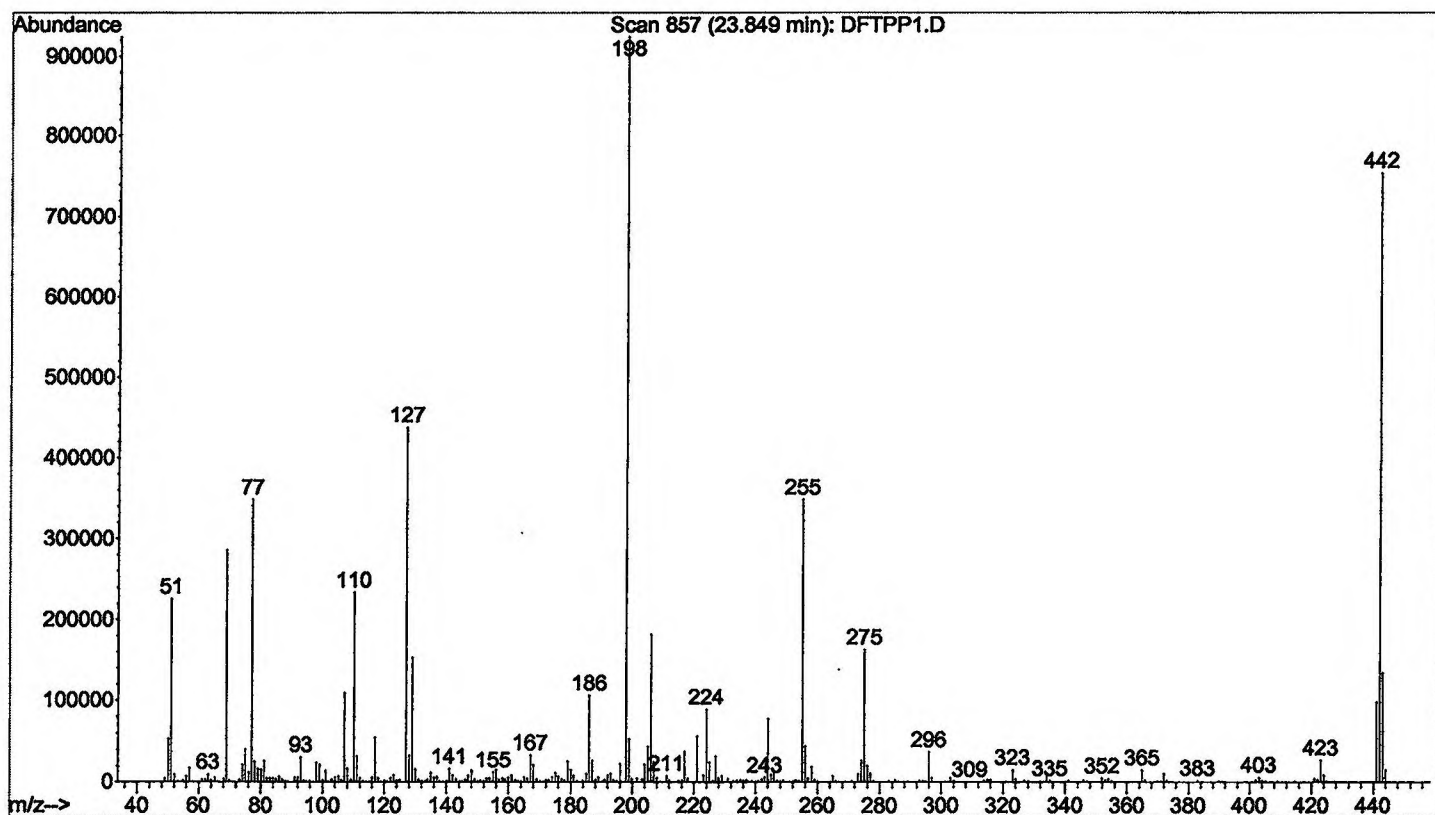
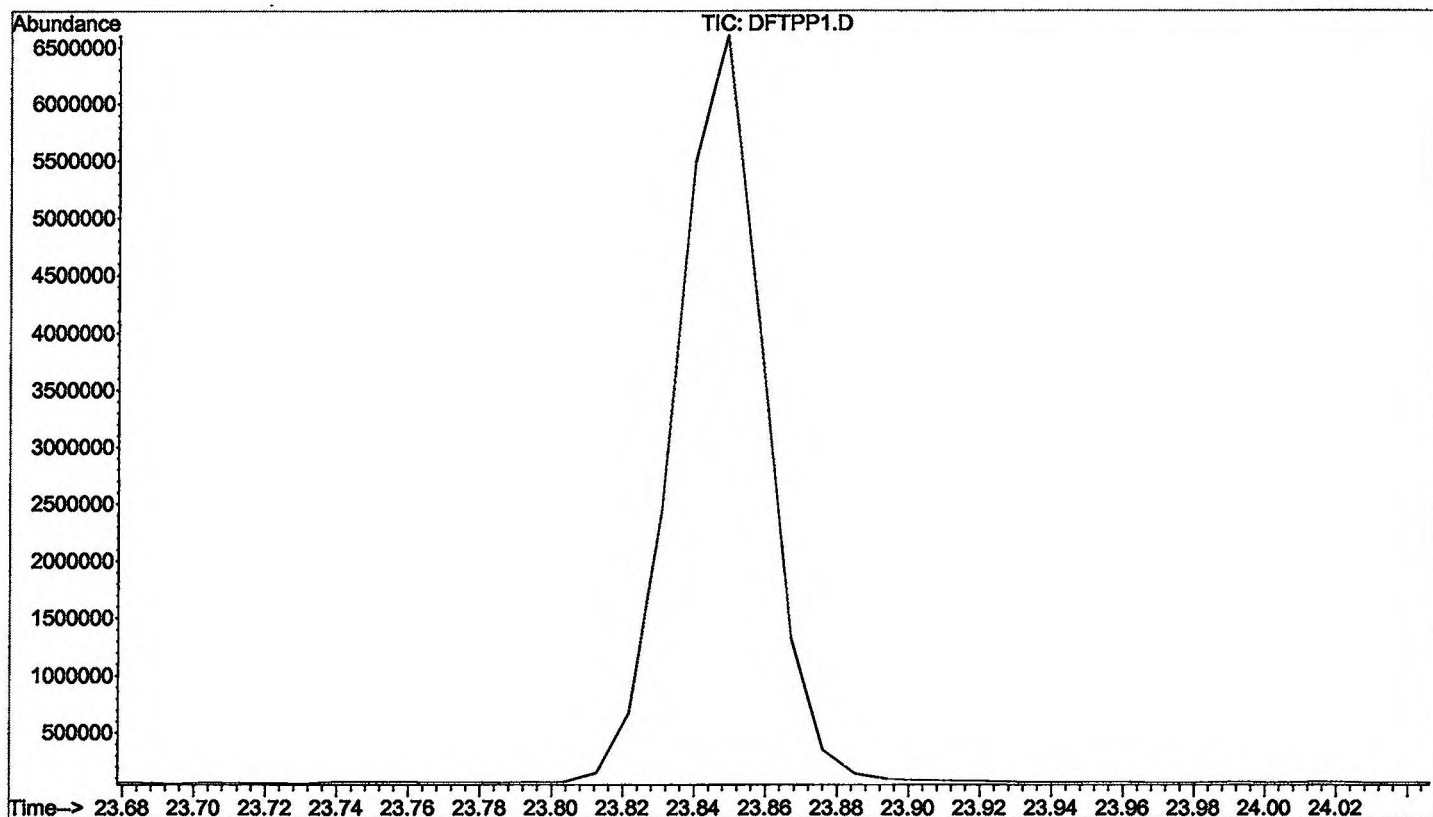
Spectrum Information: Scan 857

Target Mass	Rel. to Mass	Lower Limit%	Upper Limit%	Rel. Abn%	Raw Abn	Result Pass/Fail
51	198	10	80	24.5	225984	PASS
68	69	0.00	2	1.5	4381	PASS
69	198	0.00	100	31.0	286144	PASS
70	69	0.00	2	0.8	2422	PASS
127	198	10	80	47.4	437888	PASS
197	198	0.00	2	0.0	0	PASS
198	198	50	100	100.0	924096	PASS
199	198	5	9	5.8	53464	PASS
275	198	10	60	17.7	163264	PASS
365	198	1	100	1.6	14974	PASS
441	443	0.01	100	73.0	98280	PASS
442	198	50	100	81.5	753600	PASS
443	442	15	24	17.9	134592	PASS

for
Re-ents

DFTPP1.D 121401.M Wed Dec 19 13:38:21 2001

File : C:\MSDCHEM\1\DATA\121901\DFTPP1.D
 Operator : McLean
 Acquired : 19 Dec 2001 12:24 pm using AcqMethod BIG3
 Instrument : Instrumen
 Sample Name: dftpp +end
 Misc Info :
 Vial Number: 1



Area Percent Report

Data File : C:\MSDCHEM\1\DATA\121901\DFTPP1.D

Acq On : 19 Dec 2001 12:24 pm

Sample : dftpp +end

Misc :

MS Integration Params: BENZO.P

Vial: 1

Operator: McLean

Inst : Instrumen

Multiplr: 1.00

Method : C:\MSDCHEM\1\5973N\121401.M (RTE Integrator)

Title : Quantitation For Method 525.2

Smoothing : ON

Filtering: 5

Sampling : 1

Min Area: 2500 Area counts

Start Thrs: 0.15

Max Peaks: 10

Stop Thrs : 0

Peak Location: TOP

If leading or trailing edge < 100 prefer < Baseline drop else tangent >

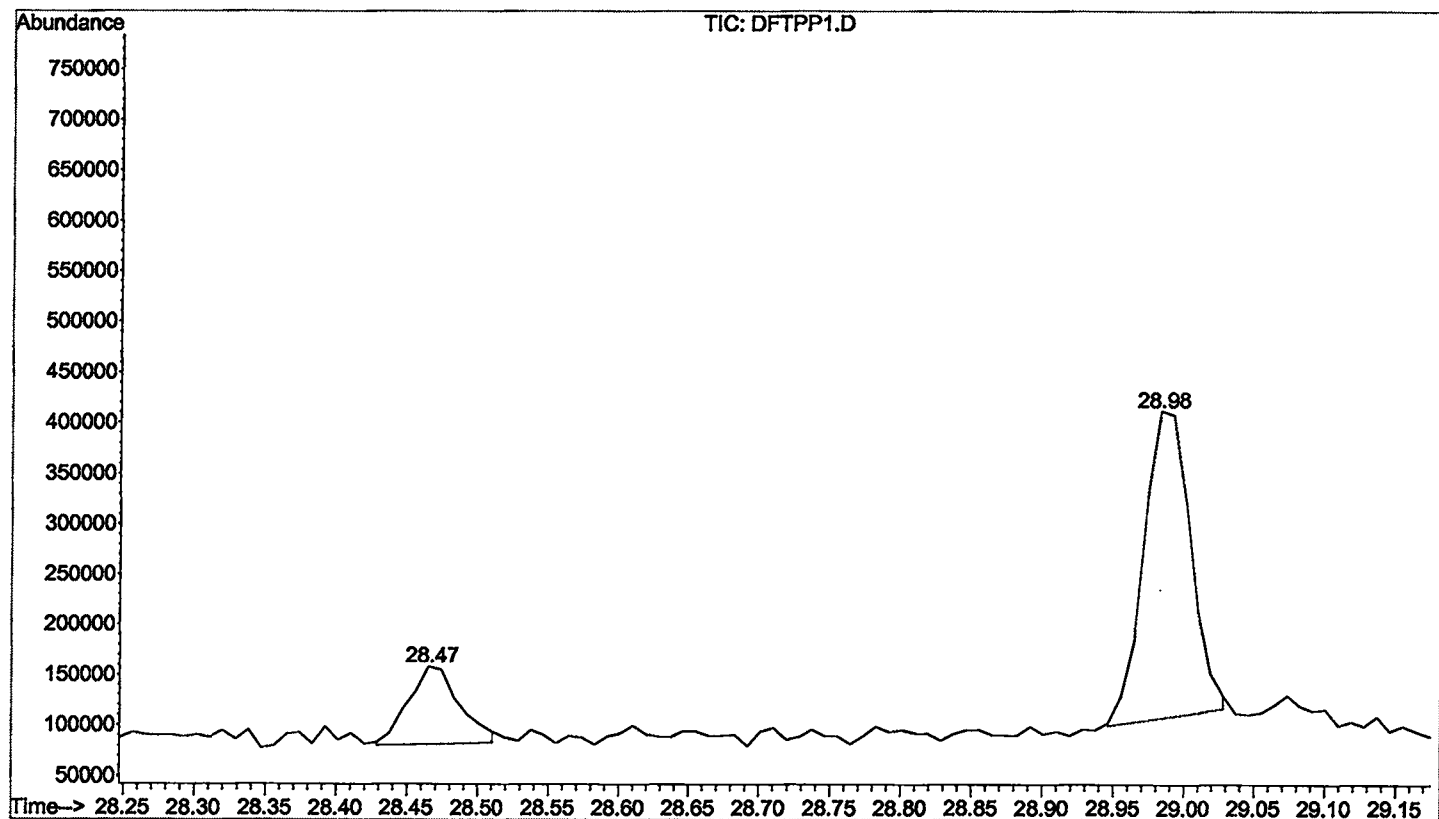
Peak separation: 5

Signal : TIC

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	27.876	1296	1301	1306	rM 2	4096887	7534737	100.00%	89.308%
2	28.466	1362	1366	1371	rM 8	76719	193608	2.57%	2.295%
3	28.983	1419	1423	1428	rM 2	303891	708422	9.40%	8.397%

Sum of corrected areas: 8436767

File : C:\MSDCHEM\1\DATA\121901\DFTPP1.D
 Operator : McLean
 Acquired : 19 Dec 2001 12:24 pm using AcqMethod BIG3
 Instrument : Instrumen
 Sample Name: dftpp +end
 Misc Info :
 Vial Number: 1



Area Percent Report

Data File : C:\MSDCHEM\1\DATA\121901\DDT1.D

Acq On : 19 Dec 2001 1:04 pm

Sample : ddt

Misc :

MS Integration Params: BENZO.P

Vial: 2

Operator: McLean

Inst : Instrumen

Multiplr: 1.00

Method : C:\MSDCHEM\1\5973N\121401.M (RTE Integrator)

Title : Quantitation For Method 525.2

Smoothing : ON

Filtering: 5

Sampling : 1

Min Area: 2500 Area counts

Start Thrs: 0.15

Max Peaks: 10

Stop Thrs : 0

Peak Location: TOP

If leading or trailing edge < 100 prefer < Baseline drop else tangent >

Peak separation: 5

Signal : TIC

peak #	R.T. min	first scan	max scan	last scan	PK TY	peak height	corr. area	corr. % max.	% of total
1	28.992	1420	1424	1428	rM	546353	1191855	24.33%	19.570%
2	29.137	1435	1440	1446	rM	2538453	4898386	100.00%	80.430%

Sum of corrected areas: 6090241

File : C:\MSDCHEM\1\DATA\121901\DDT1.D
 Operator : McLean
 Acquired : 19 Dec 2001 1:04 pm using AcqMethod BIG3
 Instrument : Instrumen
 Sample Name: ddt
 Misc Info :
 Vial Number: 2

