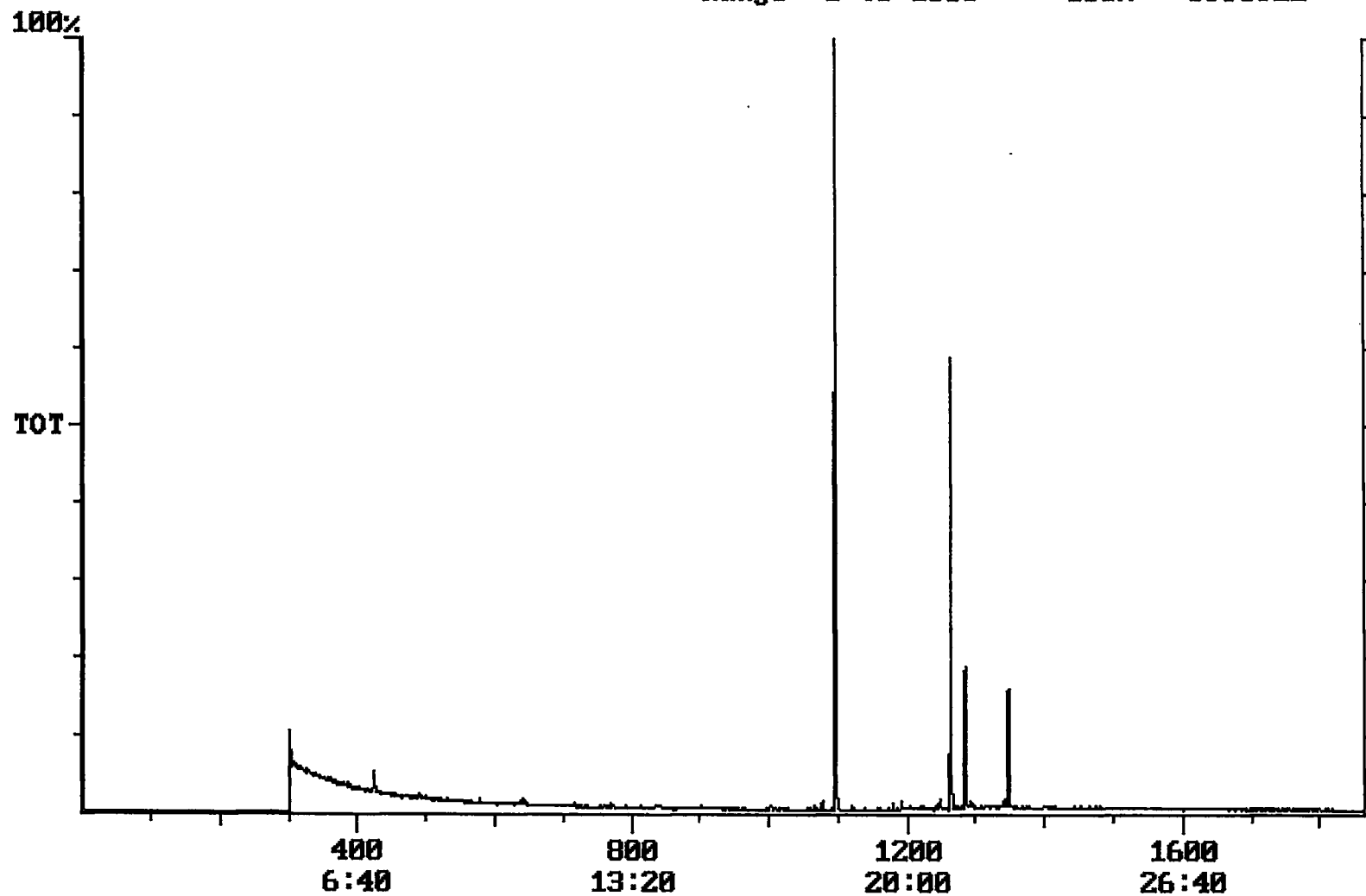


User: Bohlk, Ted
Westchester Dept. of Labs & Research
Date: 01/03/2002 Time: 09:56:10

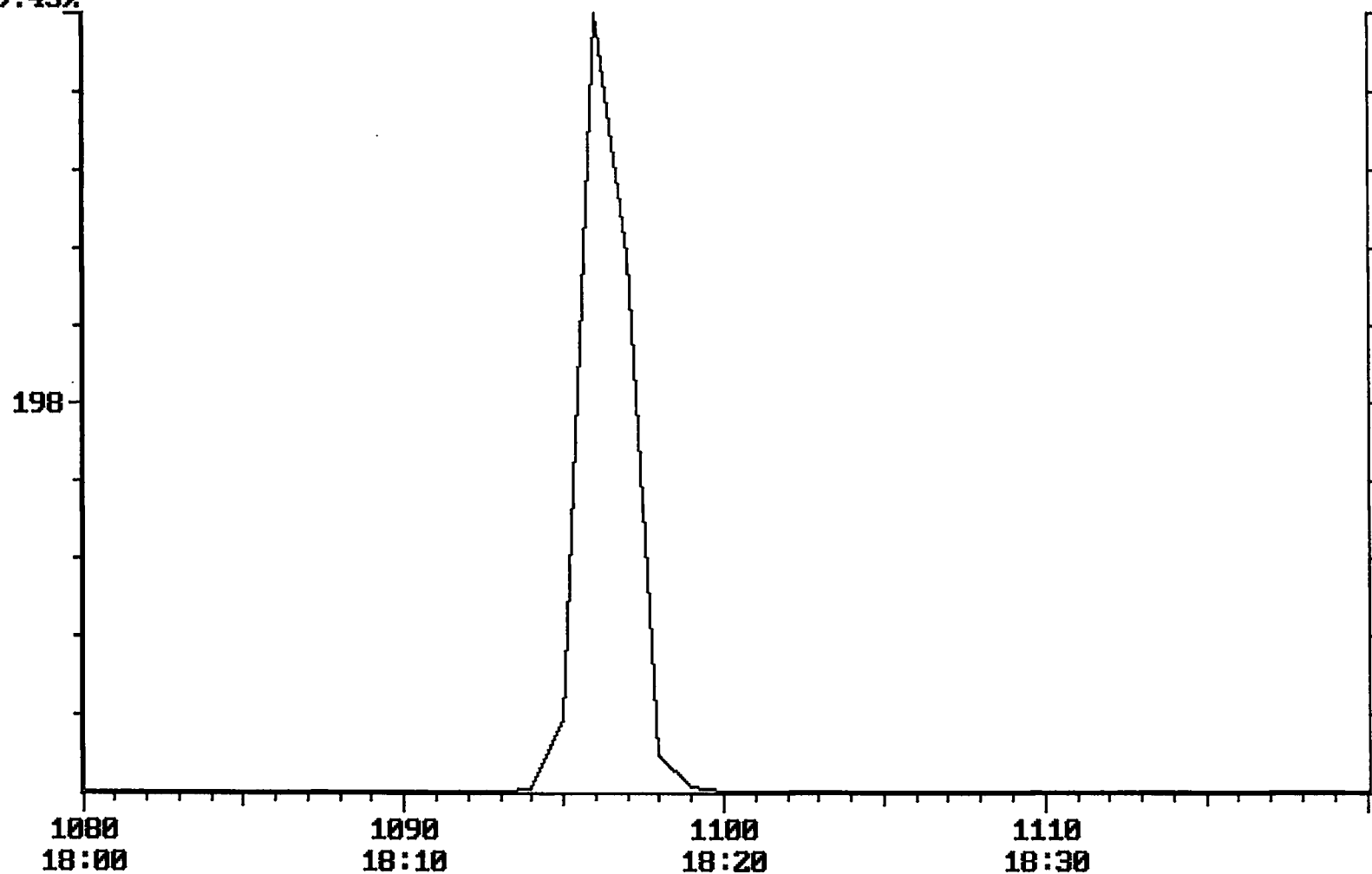
Sample: AD27009
Analysis: \$LP525 Organic Extractables
Units: ug
Started: 11/30/01 09:54 Ended: 11/30/01 09:54 Analyst: TB
Page: 1

	Component Name	Viol	Result
1	DFTPP		PASSED
2	ENDRIN	USPEC	48
3	DDT		3.9

Chromatogram Plot File: E:\DFT11130 Date: Nov-30-1991 09:36:05
Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP
Scan No: 1 Retention Time: 0:01 RIC: 0 Mass Range: 0 - 0
Plotted: 1 to 1860 Range: 1 to 1860 100% = 3996922



Chromatogram Plot File: E:\DFT11130 Date: Nov-30-1991 09:36:05
Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP
Scan No: 1080 Retention Time: 18:00 RIC: 21908 Mass Range: 47 - 149
Plotted: 1080 to 1120 Range: 1 to 1860 100% = 3996922
9.43%



Mass List

File: E:\DFT11130 Date: Nov-30-1991 09:36:05

Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP

Scan No: 1096 Retention Time: 18:16 RIC: 3996922 Mass Range: 48 - 444

Peaks: 190 Mass of Base Peak: 442 Int: 559105 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
48	1,324	0.24	79	19,867	3.55	118	3,642	0.65
49	9,271	1.66	80	16,059	2.87	122	3,642	0.65
50	89,403	15.99	81	17,384	3.11	123	6,456	1.15
51	272,019	48.65	82	3,476	0.62	124	1,655	0.30
52	12,086	2.16	83	3,145	0.56	127	124,834	22.33
53	1,324	0.24	85	2,317	0.41	128	9,437	1.69
55	5,629	1.01	86	7,450	1.33	129	60,927	10.90
56	4,470	0.80	91	1,490	0.27	130	4,801	0.86
57	10,099	1.81	92	5,629	1.01	131	3,311	0.59
58	1,490	0.27	93	31,953	5.72	134	2,980	0.53
60	1,324	0.24	94	2,317	0.41	135	6,125	1.10
61	2,980	0.53	95	2,317	0.41	136	4,304	0.77
62	3,311	0.59	98	33,940	6.07	137	6,125	1.10
63	4,635	0.83	99	13,576	2.43	141	9,105	1.63
64	3,145	0.56	101	6,622	1.18	142	3,311	0.59
65	1,986	0.36	103	3,476	0.62	143	2,980	0.53
67	1,986	0.36	104	3,642	0.65	147	7,119	1.27
68	4,635	0.83	105	4,801	0.86	148	13,245	2.37
69	186,589	33.37	107	67,715	12.11	149	1,655	0.30
73	3,642	0.65	108	6,125	1.10	150	2,317	0.41
74	29,966	5.36	110	116,390	20.82	153	2,483	0.44
75	26,655	4.77	111	14,403	2.58	155	7,284	1.30
76	10,430	1.87	115	1,655	0.30	156	5,629	1.01
77	185,264	33.14	116	4,470	0.80	160	2,317	0.41
78	8,609	1.54	117	68,211	12.20	161	2,980	0.53

Mass List

File: E:\DFT11130 Date: Nov-30-1991 09:36:05

Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP

Scan No: 1096 Retention Time: 18:16 RIC: 3996922 Mass Range: 48 - 444

Peaks: 190 Mass of Base Peak: 442 Int: 559105 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
161	2,980	0.53	200	1,324	0.24	237	1,821	0.33
164	1,324	0.24	202	1,821	0.33	239	1,821	0.33
165	1,986	0.36	203	2,980	0.53	241	1,490	0.27
166	2,152	0.38	204	12,251	2.19	242	2,483	0.44
167	26,158	4.68	205	15,728	2.81	243	1,986	0.36
168	5,298	0.95	206	61,258	10.96	244	28,642	5.12
171	1,324	0.24	207	8,278	1.48	245	3,973	0.71
172	1,986	0.36	208	3,807	0.68	246	11,589	2.07
174	2,649	0.47	209	1,655	0.30	247	1,324	0.24
175	4,801	0.86	210	1,821	0.33	253	1,324	0.24
177	2,814	0.50	211	2,814	0.50	254	2,649	0.47
178	1,986	0.36	212	993	0.18	255	256,291	45.84
179	6,291	1.13	217	20,198	3.61	256	28,476	5.09
180	7,947	1.42	218	2,317	0.41	257	3,311	0.59
181	2,152	0.38	220	993	0.18	258	13,410	2.40
185	8,940	1.60	221	5,463	0.98	259	2,814	0.50
186	51,986	9.30	222	2,814	0.50	265	3,476	0.62
187	19,370	3.46	223	3,311	0.59	267	1,655	0.30
188	2,152	0.38	224	28,476	5.09	273	5,629	1.01
189	4,139	0.74	225	10,927	1.95	274	29,304	5.24
192	3,807	0.68	227	12,417	2.22	275	144,701	25.88
193	3,973	0.71	228	1,655	0.30	276	11,589	2.07
196	4,635	0.83	234	1,158	0.21	277	10,264	1.84
198 ✓	376,986	67.43	235	1,655	0.30	278	3,145	0.56
199	21,854	3.91	236	3,311	0.59	282	1,490	0.27

Mass List

File: E:\DFT11130 Date: Nov-30-1991 09:36:05

Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP

Scan No: 1096 Retention Time: 18:16 RIC: 3996922 Mass Range: 48 - 444

Peaks: 190 Mass of Base Peak: 442 Int: 559105 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
161	2,980	0.53	200	1,324	0.24	237	1,821	0.33
164	1,324	0.24	202	1,821	0.33	239	1,821	0.33
165	1,986	0.36	203	2,980	0.53	241	1,490	0.27
166	2,152	0.38	204	12,251	2.19	242	2,483	0.44
167	26,158	4.68	205	15,728	2.81	243	1,986	0.36
168	5,298	0.95	206	61,258	10.96	244	28,642	5.12
171	1,324	0.24	207	8,278	1.48	245	3,973	0.71
172	1,986	0.36	208	3,807	0.68	246	11,589	2.07
174	2,649	0.47	209	1,655	0.30	247	1,324	0.24
175	4,801	0.86	210	1,821	0.33	253	1,324	0.24
177	2,814	0.50	211	2,814	0.50	254	2,649	0.47
178	1,986	0.36	212	993	0.18	255	256,291	45.84
179	6,291	1.13	217	20,198	3.61	256	28,476	5.09
180	7,947	1.42	218	2,317	0.41	257	3,311	0.59
181	2,152	0.38	220	993	0.18	258	13,410	2.40
185	8,940	1.60	221	5,463	0.98	259	2,814	0.50
186	51,986	9.30	222	2,814	0.50	265	3,476	0.62
187	19,370	3.46	223	3,311	0.59	267	1,655	0.30
188	2,152	0.38	224	28,476	5.09	273	5,629	1.01
189	4,139	0.74	225	10,927	1.95	274	29,304	5.24
192	3,807	0.68	227	12,417	2.22	275	144,701	25.88
193	3,973	0.71	228	1,655	0.30	276	11,589	2.07
196	4,635	0.83	234	1,158	0.21	277	10,264	1.84
198	376,986	67.43	235	1,655	0.30	278	3,145	0.56
199	21,854	3.91	236	3,311	0.59	282	1,490	0.27

Mass List

File: E:\DFT11130 Date: Nov-30-1991 09:36:05

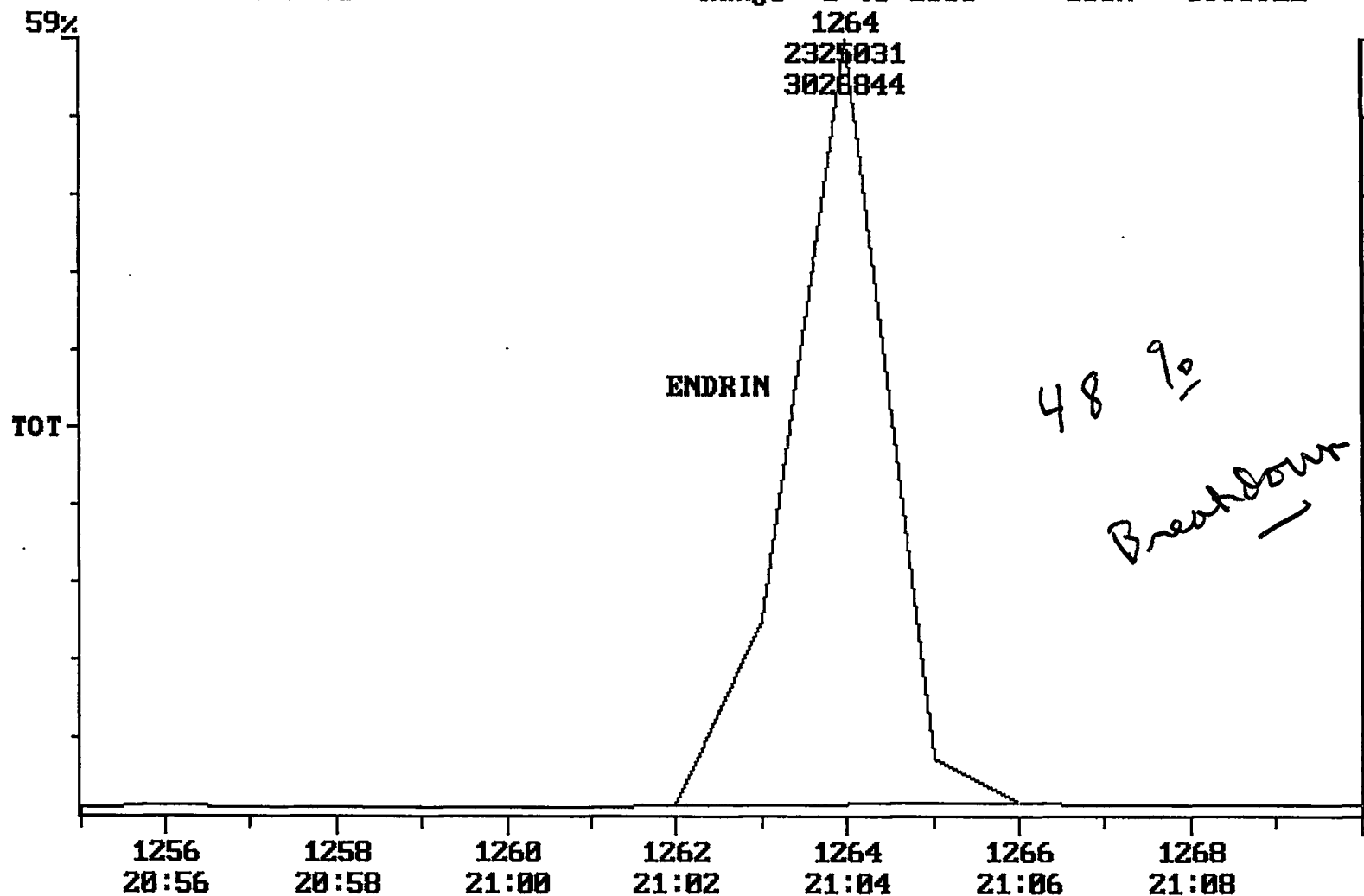
Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP

Scan No: 1096 Retention Time: 18:16 RIC: 3996922 Mass Range: 48 - 444

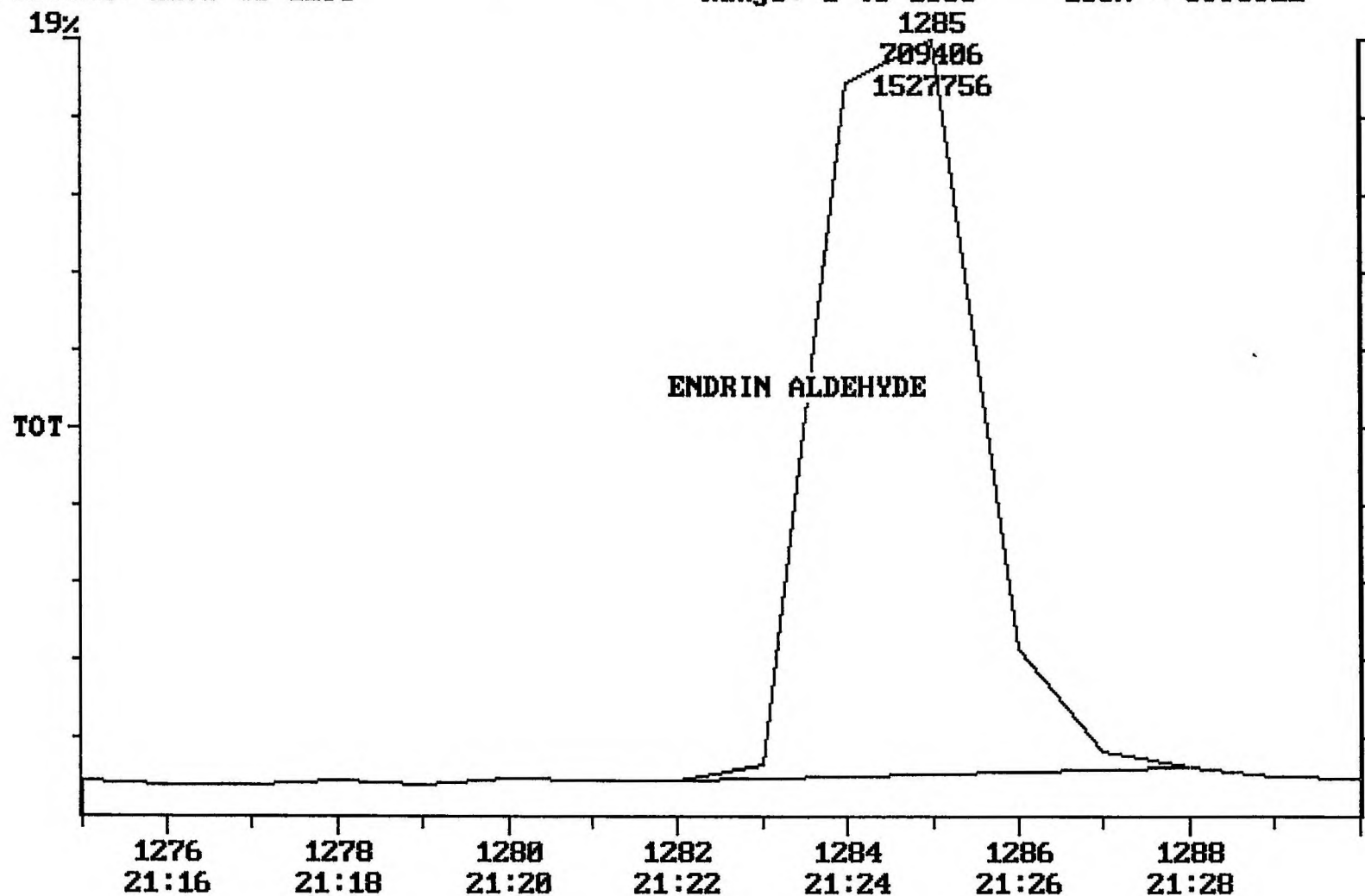
Peaks: 190 Mass of Base Peak: 442 Int: 559105 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
222	2,814	0.50	265	3,476	0.62	336	1,158	0.21
223	3,311	0.59	267	1,655	0.30	341	3,145	0.56
224	28,476	5.09	273	5,629	1.01	342	1,324	0.24
225	10,927	1.95	274	29,304	5.24	346	3,311	0.59
227	12,417	2.22	275	144,701	25.88	352	3,973	0.71
228	1,655	0.30	276	11,589	2.07	353	4,470	0.80
234	1,158	0.21	277	10,264	1.84	354	6,456	1.15
235	1,655	0.30	278	3,145	0.56	355	1,821	0.33
236	3,311	0.59	282	1,490	0.27	365	24,668	4.41 ✓
237	1,821	0.33	285	1,655	0.30	366	3,807	0.68
239	1,821	0.33	293	3,145	0.56	371	1,324	0.24
241	1,490	0.27	296	33,443	5.98	372	5,463	0.98
242	2,483	0.44	297	4,635	0.83	373	3,807	0.68
243	1,986	0.36	303	3,642	0.65	383	2,483	0.44
244	28,642	5.12	304	1,324	0.24	402	2,483	0.44
245	3,973	0.71	314	1,655	0.30	403	4,635	0.83
246	11,589	2.07	315	5,794	1.04	404	2,980	0.53
247	1,324	0.24	316	2,814	0.50	421	2,483	0.44
253	1,324	0.24	323	13,907	2.49	422	4,635	0.83
254	2,649	0.47	324	1,986	0.36	423	29,801	5.33
255	256,291	45.84	327	2,317	0.41	424	6,125	1.10
256	28,476	5.09	328	1,986	0.36	441	106,291	19.01 ✓
257	3,311	0.59	331	993	0.18	442	559,105	100.00 ✓
258	13,410	2.40	334	9,768	1.75	443	116,059	20.76 ✓
259	2,814	0.50	335	3,311	0.59	444	12,582	2.25

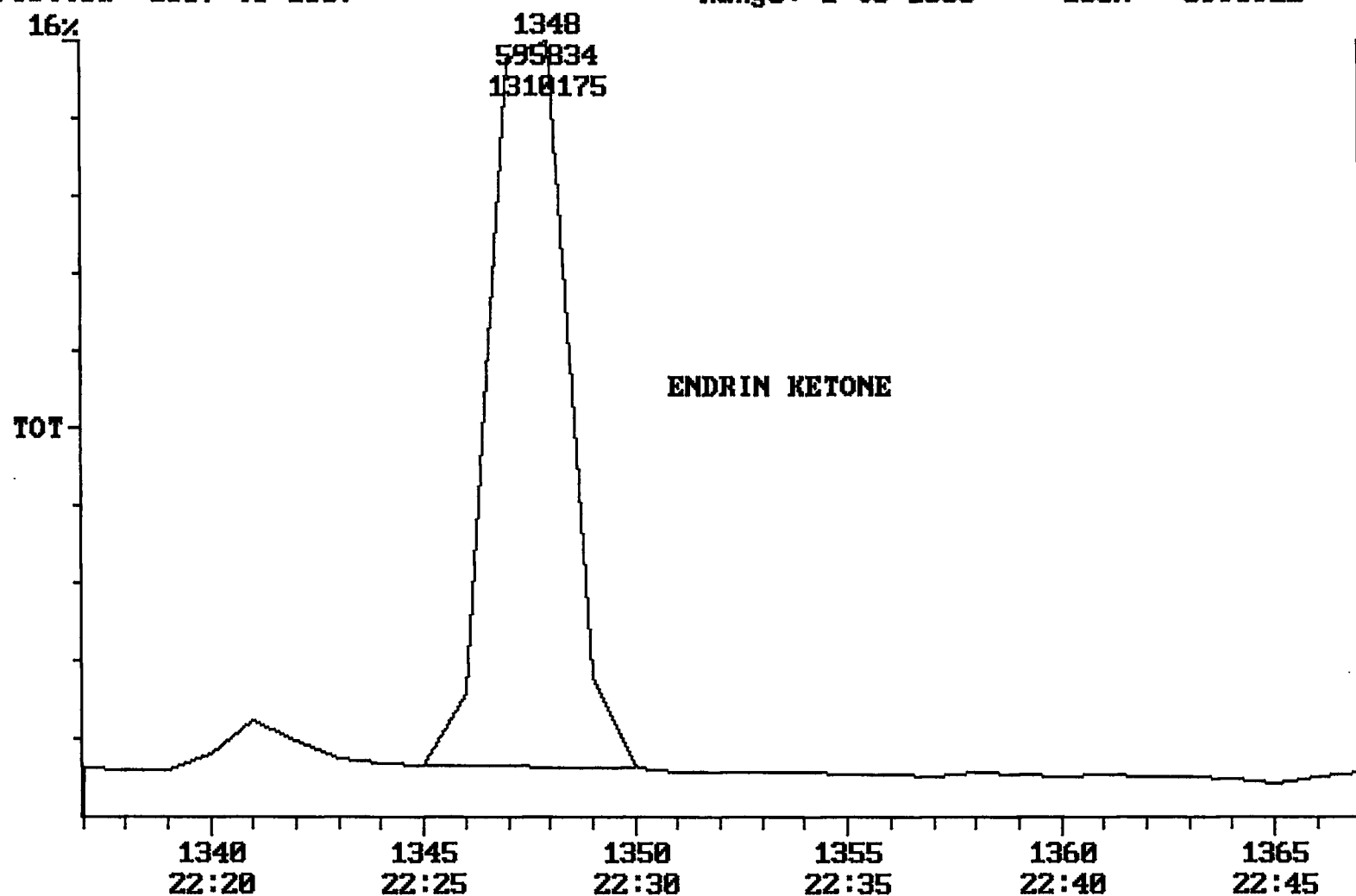
Chromatogram Plot File: E:\DFT11130 Date: Nov-30-1991 09:36:05
 Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP
 Scan No: 1270 Retention Time: 21:10 RIC: 30413 Mass Range: 47 - 281
 Plotted: 1255 to 1270 Range: 1 to 1860 100% = 3996922



Chromatogram Plot File: E:\DFT11130 Date: Nov-30-1991 09:36:05
Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP
Scan No: 1290 Retention Time: 21:30 RIC: 34604 Mass Range: 47 - 281
Plotted: 1275 to 1290 Range: 1 to 1860 100% = 3996922



Chromatogram Plot File: E:\DFT11130 Date: Nov-30-1991 09:36:05
Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP
Scan No: 1367 Retention Time: 22:47 RIC: 35485 Mass Range: 45 - 291
Plotted: 1337 to 1367 Range: 1 to 1860 100% = 3996922



Chromatogram Plot

File: E:\DDTA1130

Date: Nov-30-1991 10:13:47

Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DDT

Scan No: 1

Retention Time: 0:01

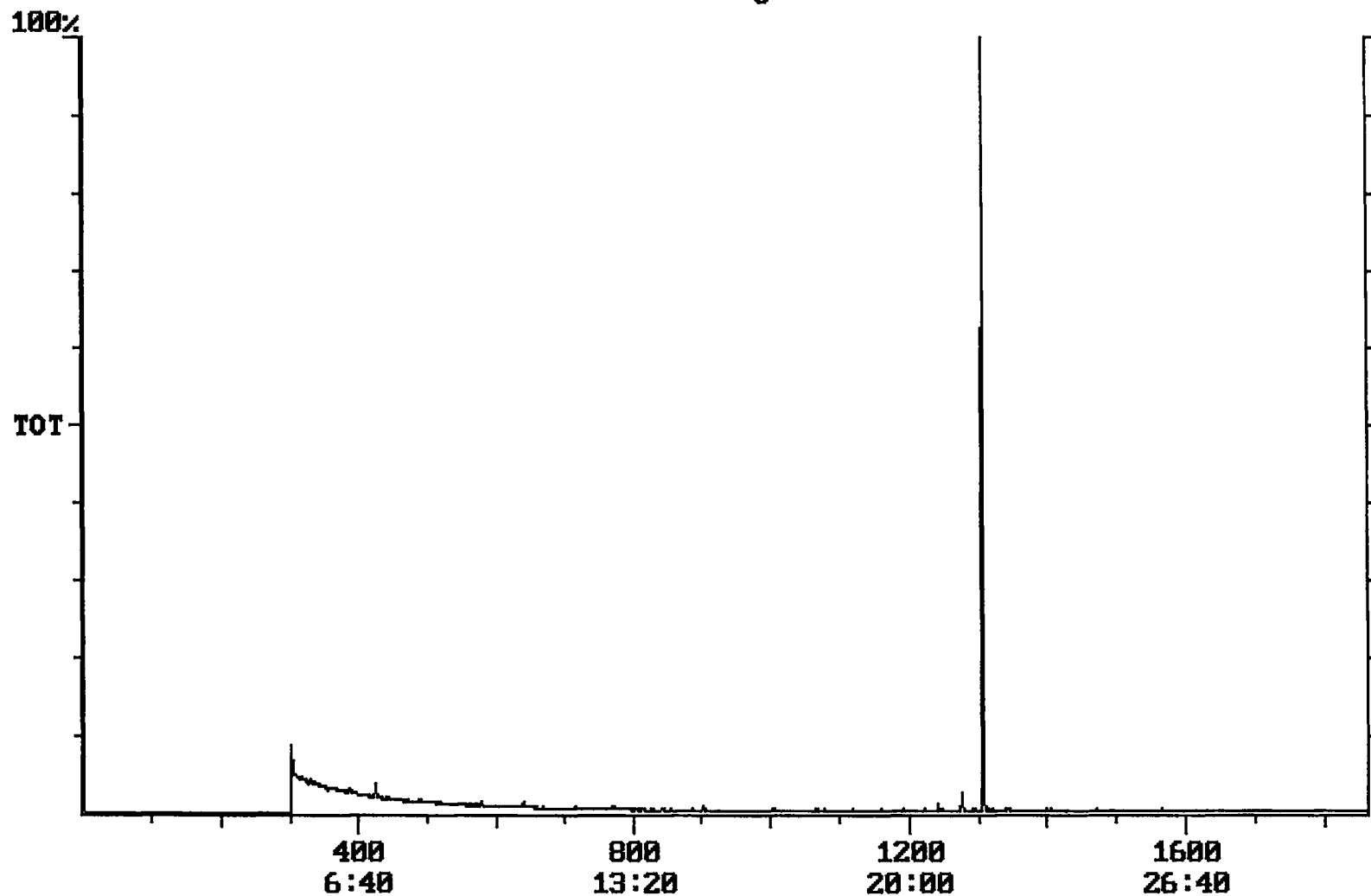
RIC: 0

Mass Range: 0 - 0

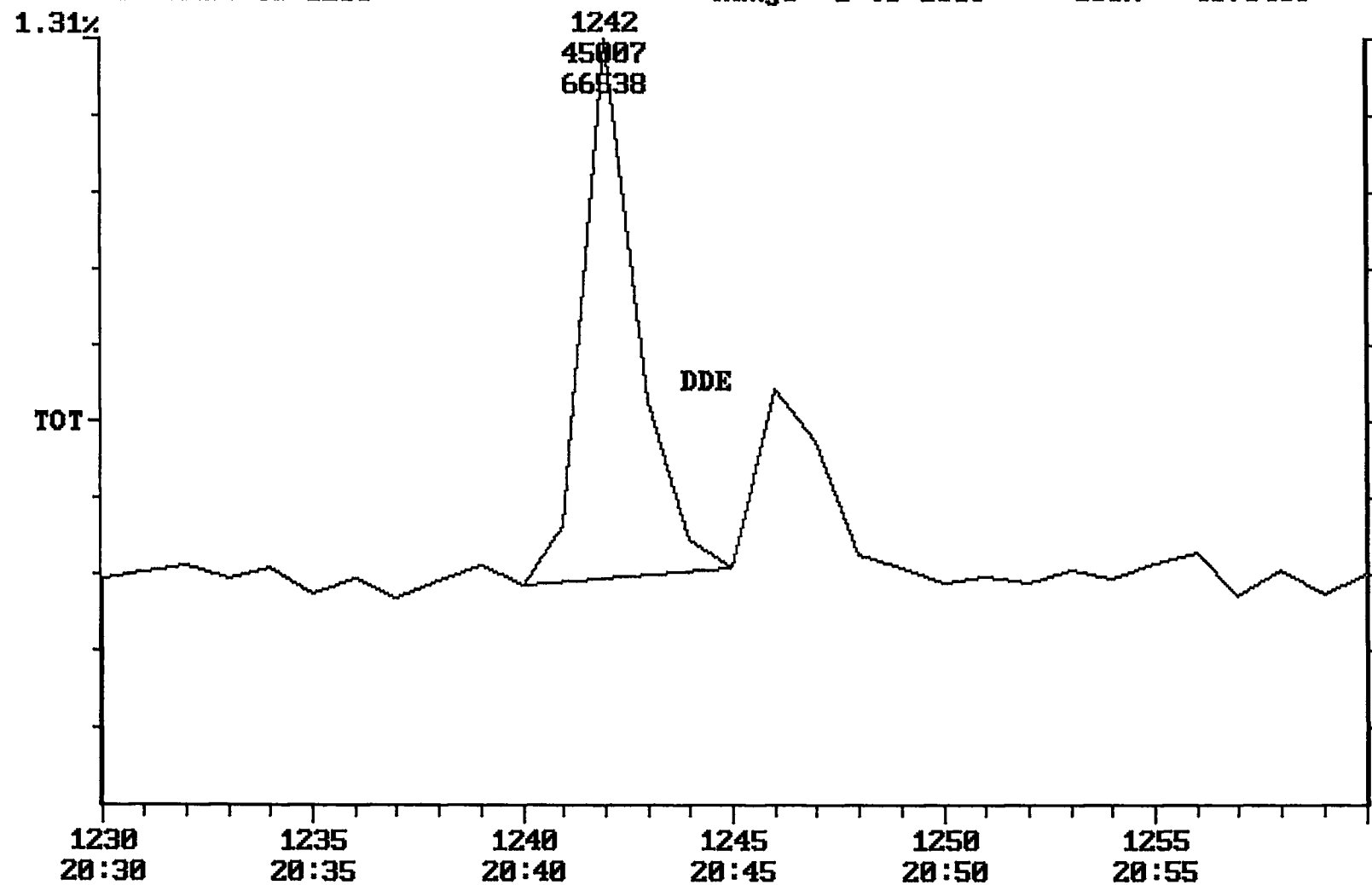
Plotted: 1 to 1859

Range: 1 to 1859

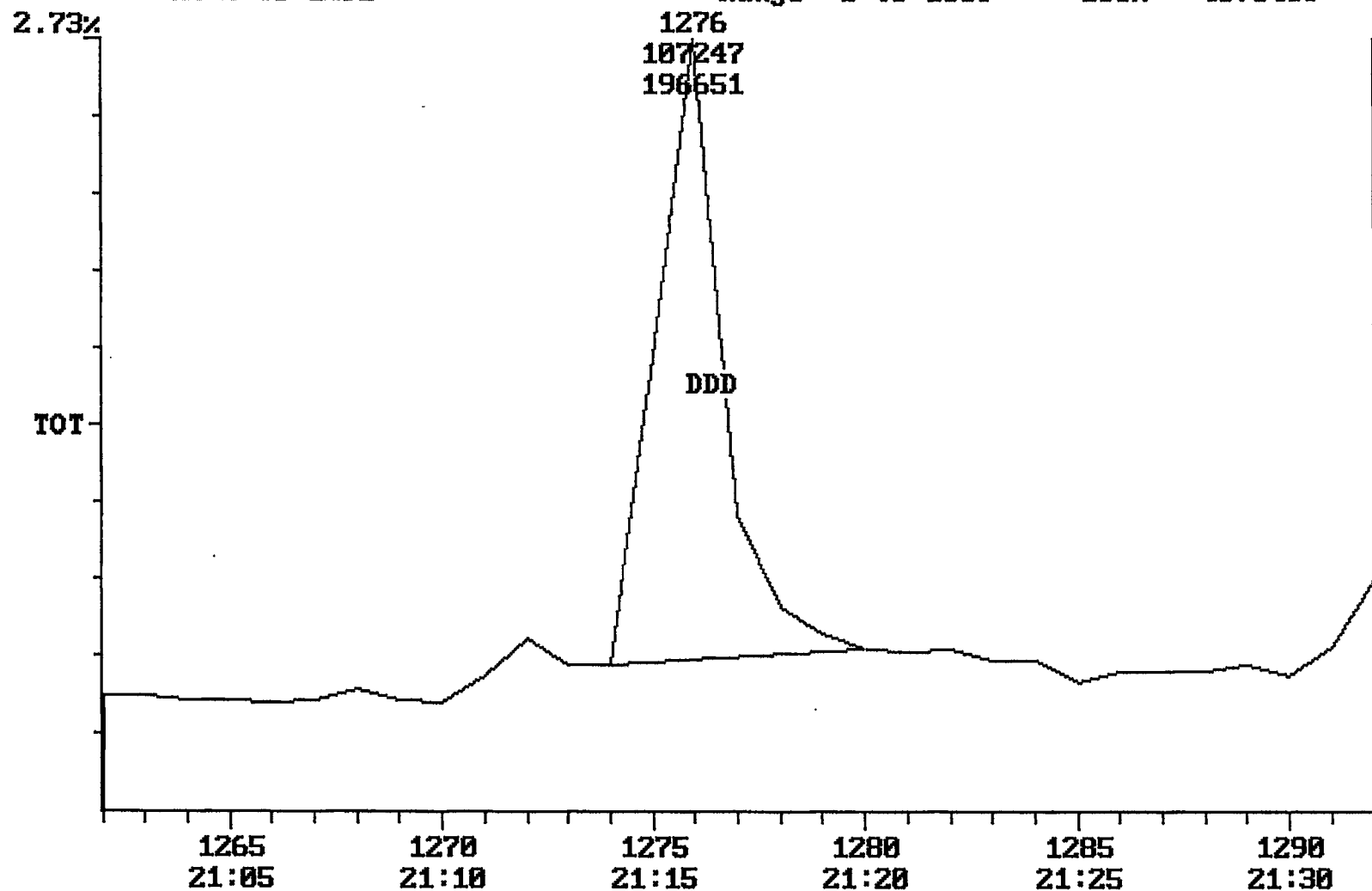
100% = 4870439



Chromatogram Plot File: E:\DDTA1130 Date: Nov-30-1991 10:13:47
Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DDT
Scan No: 1260 Retention Time: 21:00 RIC: 19156 Mass Range: 47 - 207
Plotted: 1230 to 1260 Range: 1 to 1859 100% = 4870439



Chromatogram Plot File: E:\DDTA1130 Date: Nov-30-1991 10:13:47
Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DDT
Scan No: 1292 Retention Time: 21:32 RIC: 39654 Mass Range: 45 - 429
Plotted: 1262 to 1292 Range: 1 to 1859 100% = 4870439



Chromatogram Plot

File: E:\DDTA1130

Date: Nov-30-1991 10:13:47

Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DDT

Scan No: 1324

Retention Time: 22:04

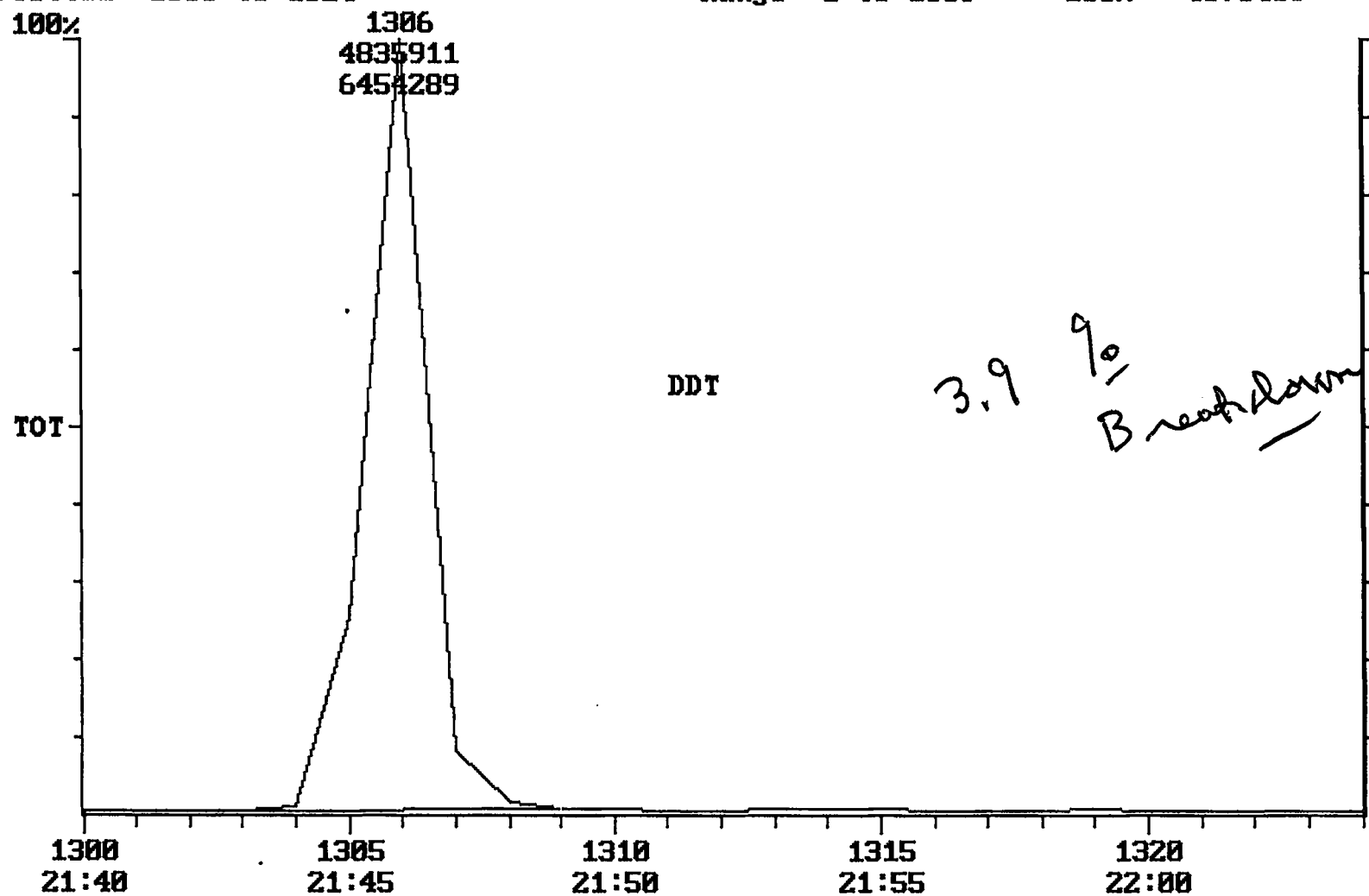
RIC: 26018

Mass Range: 47 - 281

Plotted: 1300 to 1324

Range: 1 to 1859

100% = 4870439



Chromatogram Plot

File: E:\DFTB1130

Date: Nov-30-1991 18:03:41

Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP/ENDRIN

Scan No: 1

Retention Time: 0:01

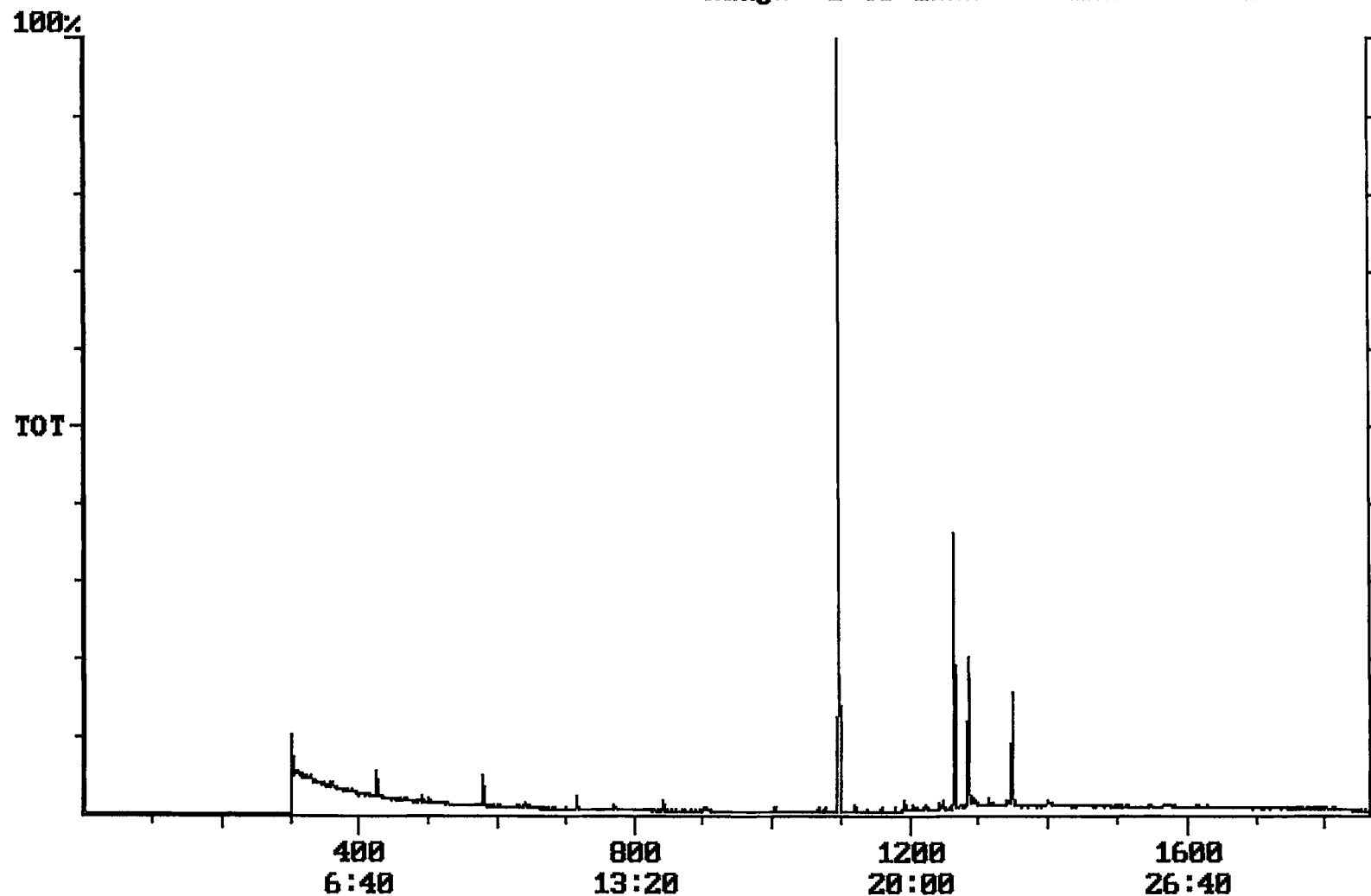
RIC: 0

Mass Range: 0 - 0

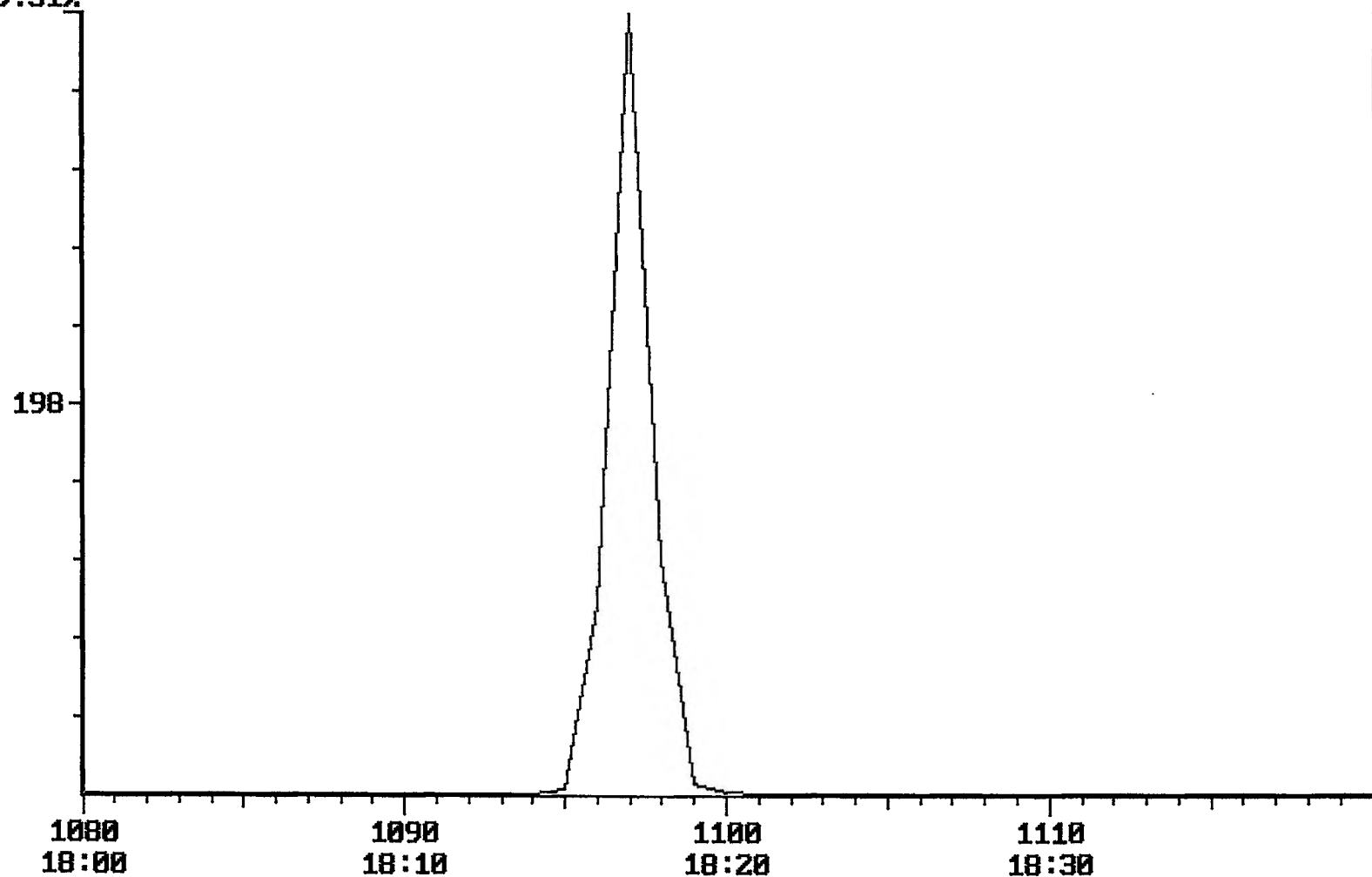
Plotted: 1 to 1860

Range: 1 to 1860

100% = 5069934



Chromatogram Plot File: E:\DFTB1130 Date: Nov-30-1991 18:03:41
Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP/ENDRIN
Scan No: 1080 Retention Time: 18:00 RIC: 24375 Mass Range: 47 - 207
Plotted: 1080 to 1120 Range: 1 to 1860 100% = 5069934
9.31%



Mass List

File: E:\DFTB1130 Date: Nov-30-1991 18:03:41

Comment: BOHLK; METHOD 525; INST 204; 11/30/01: DFTPP/ENDRIN

Scan No: 1097 Retention Time: 18:17 RIC: 5069934 Mass Range: 49 - 444

Peaks: 207 Mass of Base Peak: 442 Int: 697083 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
49	7,500	1.08	81	23,333	3.35	113	1,666	0.24
50	110,000	15.78	82	4,166	0.60	116	1,875	0.27
51	326,250	46.80	83	2,708	0.39	117	83,958	12.04
52	13,750	1.97	85	3,750	0.54	118	6,458	0.93
53	2,083	0.30	86	8,333	1.20	122	7,083	1.02
55	7,708	1.11	91	2,916	0.42	123	7,916	1.14
56	7,916	1.14	92	6,458	0.93	127	177,916	25.52
57	10,208	1.46	93	35,416	5.08	128	15,000	2.15
59	1,458	0.21	94	5,625	0.81	129	88,958	12.76
61	8,750	1.26	96	2,291	0.33	130	5,625	0.81
62	7,500	1.08	97	3,333	0.48	131	1,458	0.21
63	9,583	1.37	98	41,875	6.01	132	3,958	0.57
65	6,041	0.87	99	18,333	2.63	133	1,875	0.27
66	1,875	0.27	100	2,291	0.33	134	2,083	0.30
68	3,750	0.54	101	7,291	1.05	135	10,000	1.43
69	262,916	37.72	103	2,916	0.42	136	2,708	0.39
70	2,916	0.42	104	8,125	1.17	137	1,875	0.27
73	2,500	0.36	105	9,791	1.40	140	2,291	0.33
74	45,208	6.49	106	3,541	0.51	141	11,458	1.64
75	26,875	3.86	107	95,208	13.66	142	2,291	0.33
76	11,458	1.64	108	9,583	1.37	143	4,791	0.69
77	236,458	33.92	109	7,500	1.08	145	2,083	0.30
78	18,750	2.69	110	119,583	17.15	146	2,500	0.36
79	28,333	4.06	111	17,500	2.51	147	11,250	1.61
80	11,875	1.70	112	3,125	0.45	148	13,958	2.00

Mass List

File: E:\DFTB1130 Date: Nov-30-1991 18:03:41

Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP/ENDRIN

Scan No: 1097 Retention Time: 18:17 RIC: 5069934 Mass Range: 49 - 444

Peaks: 207 Mass of Base Peak: 442 Int: 697083 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
149	3,333	0.48	184	4,583	0.66	222	6,875	0.99
150	1,458	0.21	185	10,833	1.55	223	6,875	0.99
151	2,500	0.36	186	88,125	12.64	224	52,708	7.56
154	3,750	0.54	187	19,375	2.78	225	12,500	1.79
155	3,541	0.51	189	5,625	0.81	227	21,041	3.02
156	7,291	1.05	191	1,666	0.24	228	3,541	0.51
157	2,291	0.33	192	1,666	0.24	230	2,291	0.33
158	4,583	0.66	193	5,000	0.72	231	2,708	0.39
159	3,333	0.48	194	1,875	0.27	235	2,083	0.30
161	6,666	0.96	196	8,333	1.20	236	2,708	0.39
163	2,083	0.30	198	472,083	67.72	237	3,958	0.57
165	3,125	0.45	199	29,166	4.18	238	2,708	0.39
166	6,041	0.87	200	2,083	0.30	239	2,708	0.39
167	28,333	4.06	202	1,458	0.21	240	2,916	0.42
168	8,333	1.20	203	1,458	0.21	241	4,375	0.63
169	2,500	0.36	204	14,166	2.03	243	1,875	0.27
172	6,041	0.87	205	31,666	4.54	244	27,500	3.95
173	2,500	0.36	206	99,583	14.29	245	2,500	0.36
174	4,375	0.63	207	12,500	1.79	246	8,541	1.23
175	9,583	1.37	211	4,791	0.69	247	2,291	0.33
176	1,666	0.24	215	2,916	0.42	249	2,708	0.39
179	9,375	1.34	216	2,708	0.39	250	2,083	0.30
180	7,291	1.05	217	33,958	4.87	255	298,750	42.86
181	4,791	0.69	218	2,916	0.42	256	36,875	5.29
182	1,666	0.24	221	7,083	1.02	257	4,791	0.69

Mass List

File: E:\DFTB1130 Date: Nov-30-1991 18:03:41

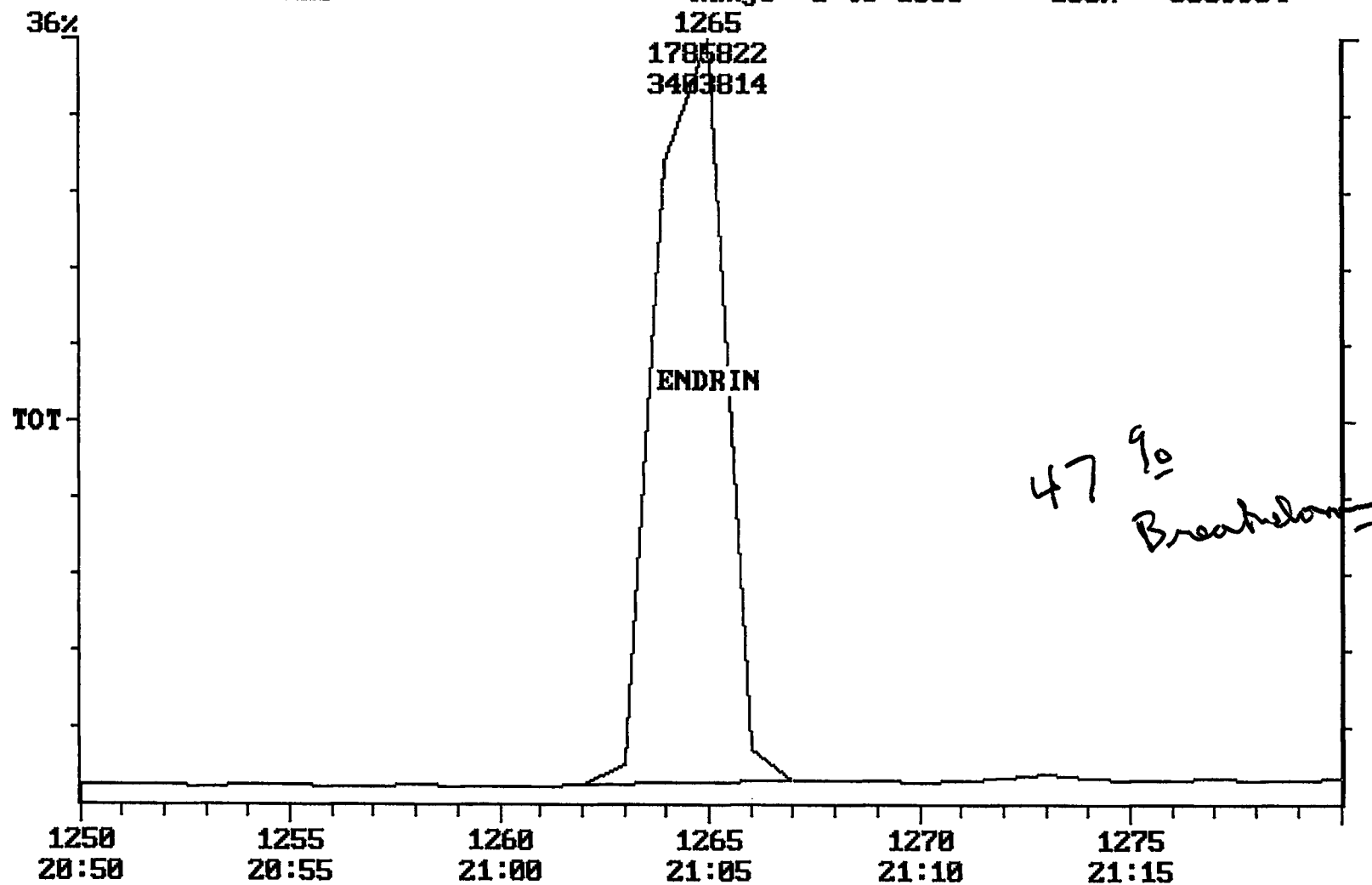
Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP/ENDRIN

Scan No: 1097 Retention Time: 18:17 RIC: 5069934 Mass Range: 49 - 444

Peaks: 207 Mass of Base Peak: 442 Int: 697083 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
231	2,708	0.39	273	9,583	1.37	341	5,000	0.72
235	2,083	0.30	274	25,833	3.71	347	2,500	0.36
236	2,708	0.39	275	158,958	22.80	352	4,583	0.66
237	3,958	0.57	276	19,166	2.75	353	6,250	0.90
238	2,708	0.39	277	20,833	2.99	354	9,166	1.31
239	2,708	0.39	278	3,125	0.45	361	1,250	0.18
240	2,916	0.42	285	3,125	0.45	365	36,666	5.26
241	4,375	0.63	293	4,166	0.60	366	4,375	0.63
243	1,875	0.27	294	1,666	0.24	371	2,500	0.36
244	27,500	3.95	296	31,666	4.54	372	12,291	1.76
245	2,500	0.36	297	5,833	0.84	373	3,750	0.54
246	8,541	1.23	303	8,125	1.17	383	5,625	0.81
247	2,291	0.33	304	2,083	0.30	384	1,458	0.21
249	2,708	0.39	314	3,958	0.57	391	2,291	0.33
250	2,083	0.30	315	2,291	0.33	392	3,125	0.45
255	298,750	42.86	316	3,958	0.57	403	5,208	0.75
256	36,875	5.29	317	2,083	0.30	404	1,666	0.24
257	4,791	0.69	321	3,541	0.51	421	1,875	0.27
258	12,291	1.76	323	20,000	2.87	422	4,166	0.60
259	3,125	0.45	324	2,500	0.36	423	37,708	5.41
261	2,916	0.42	327	4,375	0.63	424	6,875	0.99
263	1,458	0.21	332	1,666	0.24	441	93,750	13.45
265	8,125	1.17	333	2,916	0.42	442	697,083	100.00
271	1,875	0.27	334	8,541	1.23	443	118,333	16.98
272	1,458	0.21	335	5,833	0.84	444	7,708	1.11

Chromatogram Plot File: E:\DFTB1130 Date: Nov-30-1991 18:03:41
Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP/ENDRIN
Scan No: 1280 Retention Time: 21:20 RIC: 61750 Mass Range: 45 - 281
Plotted: 1250 to 1280 Range: 1 to 1860 100% = 5069934



Chromatogram Plot

File: E:\DFTB1130

Date: Nov-30-1991 18:03:41

Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP/ENDRIN

Scan No: 1302

Retention Time: 21:42

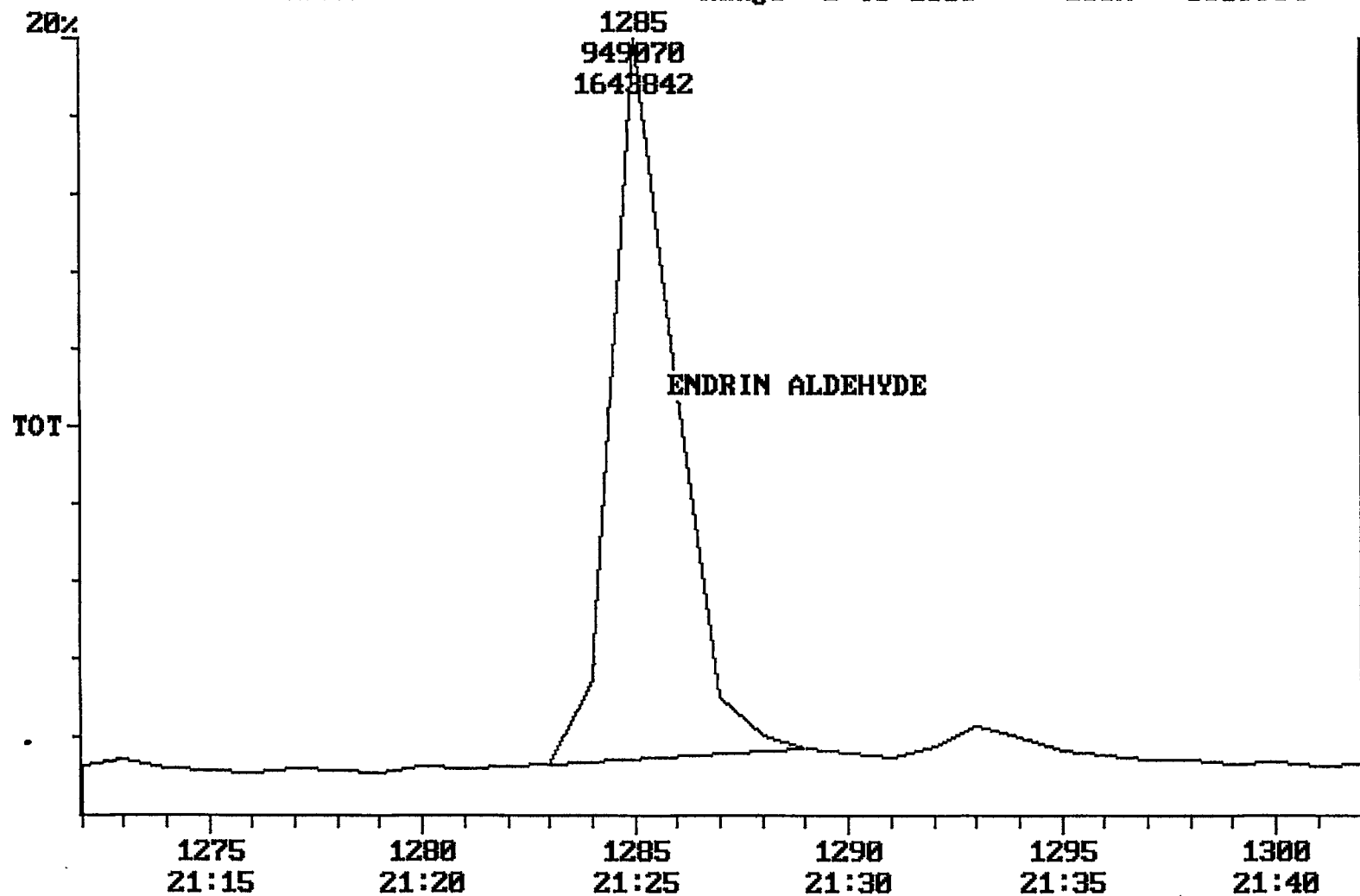
RIC: 67144

Mass Range: 45 - 281

Plotted: 1272 to 1302

Range: 1 to 1860

100% = 5069934



Chromatogram Plot

File: E:\DFTB1130 Date: Nov-30-1991 18:03:41

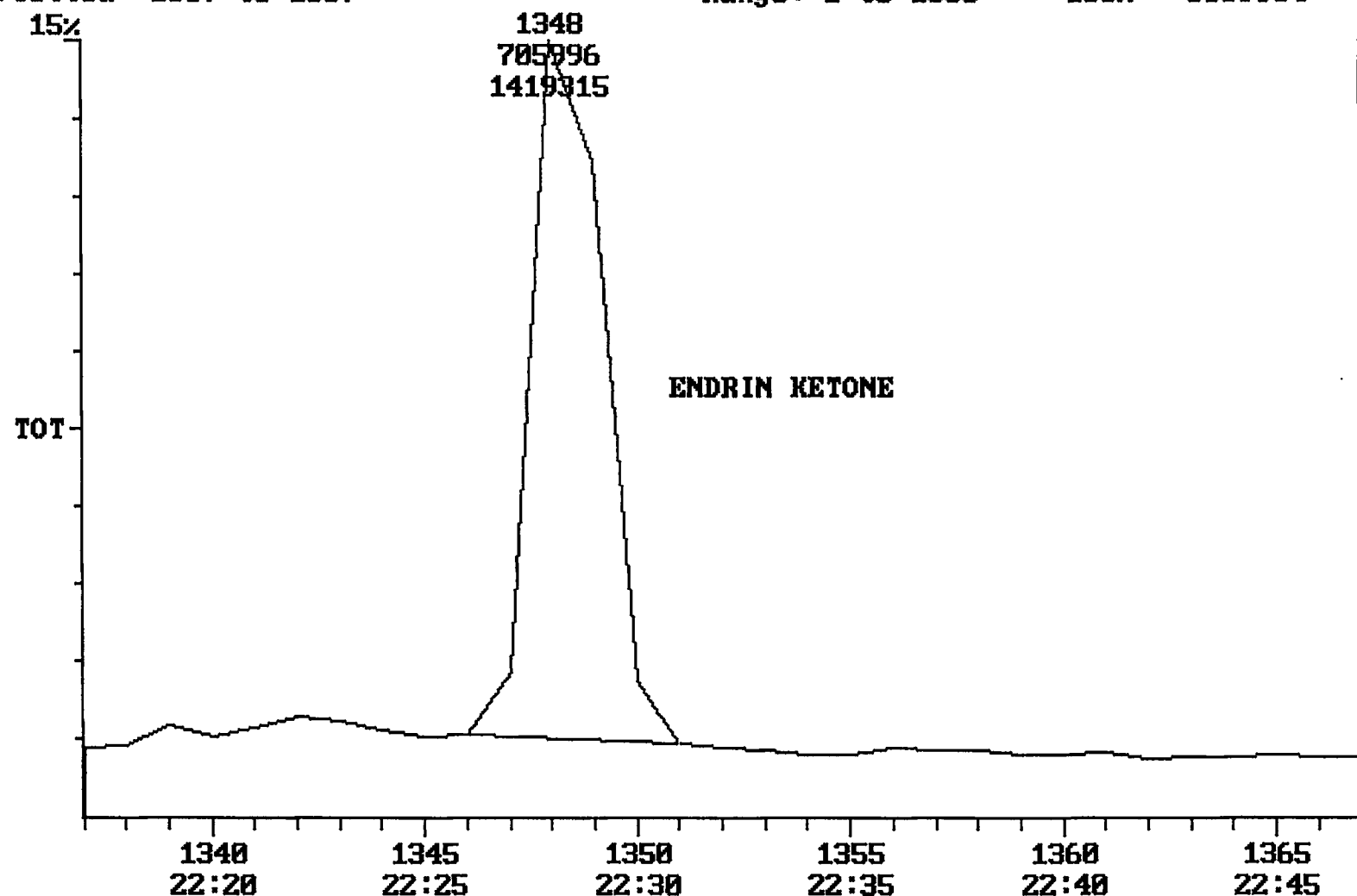
Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DFTPP/ENDRIN

Scan No: 1367 Retention Time: 22:47 RIC: 59639 Mass Range: 45 - 281

Plotted: 1337 to 1367

Range: 1 to 1860

100% = 5069934



Chromatogram Plot

File: E:\DDTB1130

Date: Nov-30-1991 18:38:31

Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DDT

Scan No: 1

Retention Time: 0:01

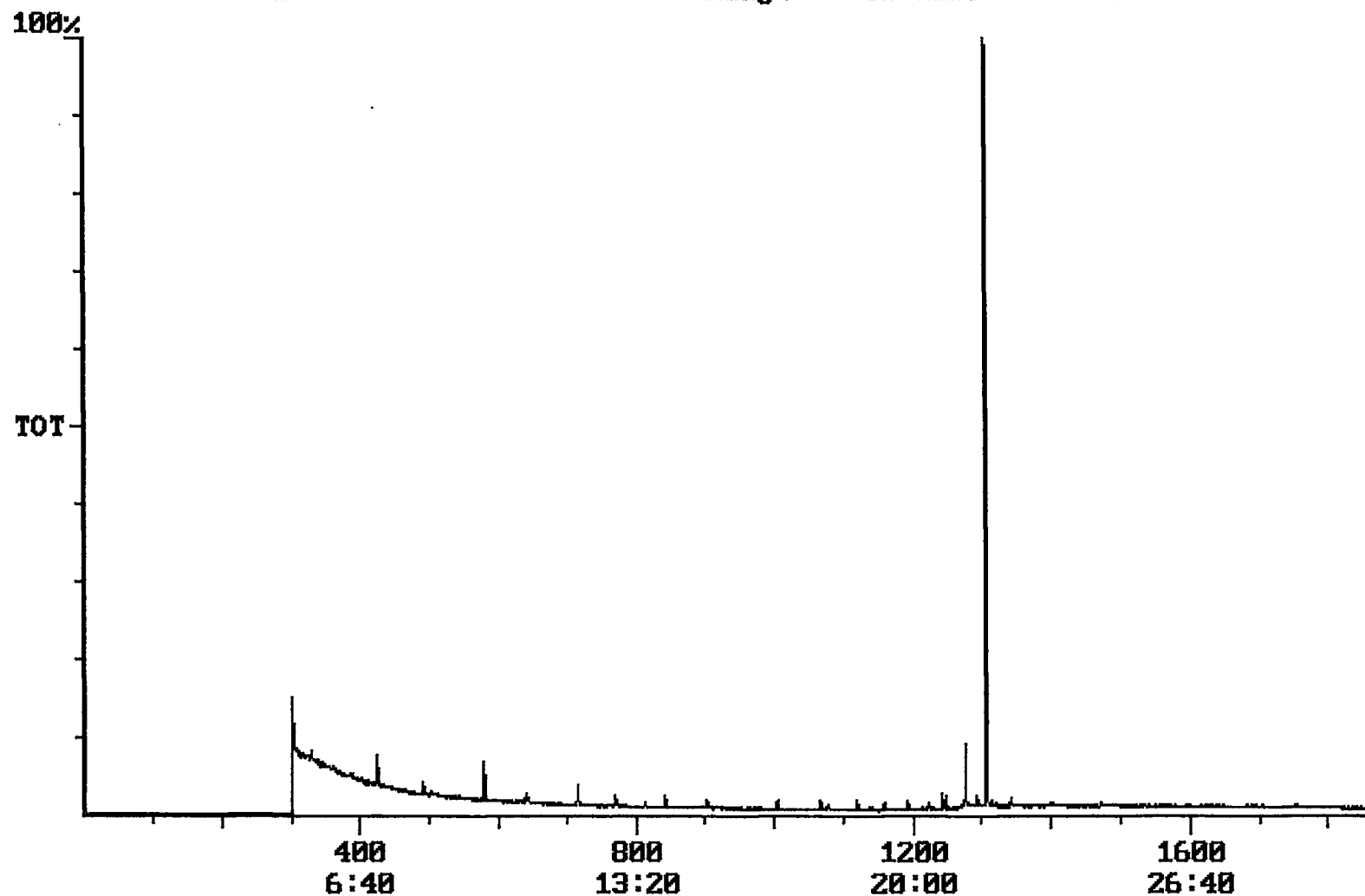
RIC: 0

Mass Range: 0 - 0

Plotted: 1 to 1860

Range: 1 to 1860

100% = 3040537



Chromatogram Plot

File: E:\DDTB1130

Date: Nov-30-1991 18:38:31

Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DDT

Scan No: 1269

Retention Time: 21:09

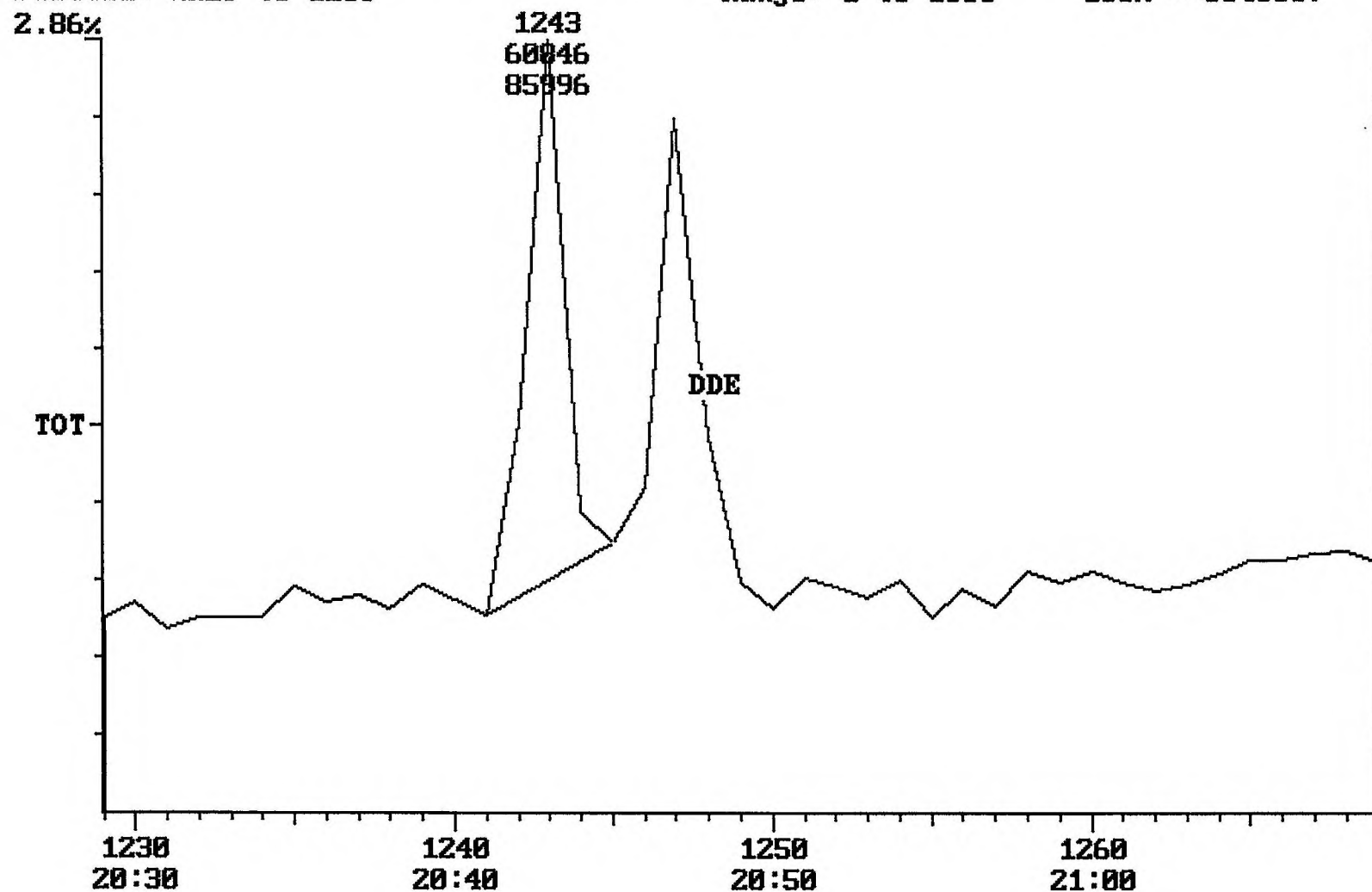
RIC: 28044

Mass Range: 47 - 207

Plotted: 1229 to 1269

Range: 1 to 1860

100% = 3040537



Chromatogram Plot

File: E:\DDTB1130

Date: Nov-30-1991 18:38:31

Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DDT

Scan No: 1292

Retention Time: 21:32

RIC: 65275

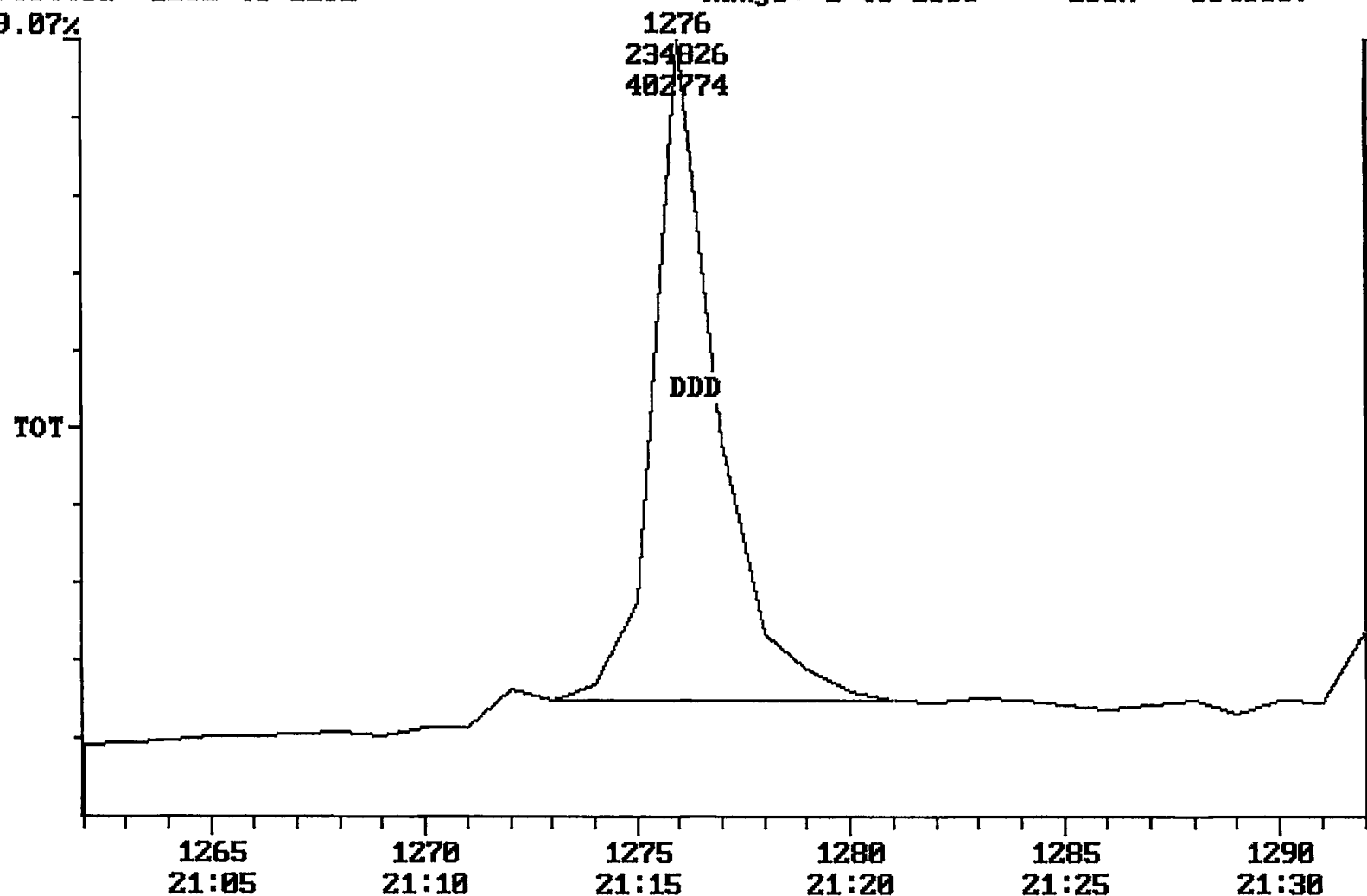
Mass Range: 45 - 429

Plotted: 1262 to 1292

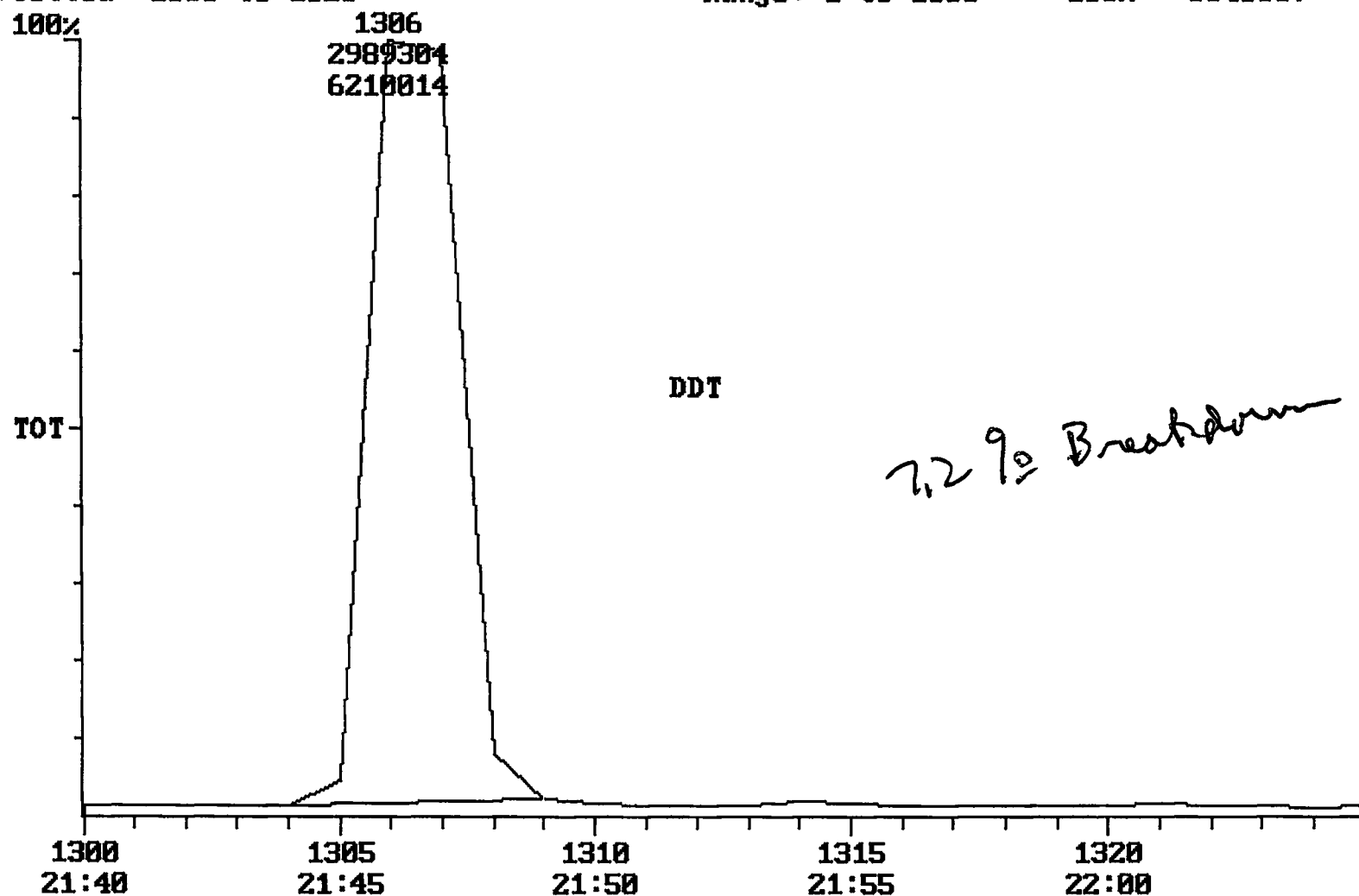
Range: 1 to 1860

100% = 3040537

9.07%



Chromatogram Plot File: E:\DDTB1130 Date: Nov-30-1991 18:38:31
 Comment: BOHLK; METHOD 525; INST 204; 11/30/01; DDT
 Scan No: 1325 Retention Time: 22:05 RIC: 36411 Mass Range: 45 - 281
 Plotted: 1300 to 1325 Range: 1 to 1860 100% = 3040537



DFTPP

Data File : C:\MSDCHEM\1\DATA\121901\DFTP4.D
Acq On : 20 Dec 2001 9:00 am
Sample :
Misc :
MS Integration Params: BENZO.P

Vial: 14
Operator: McLean
Inst : Instrumen
Multiplr: 1.00

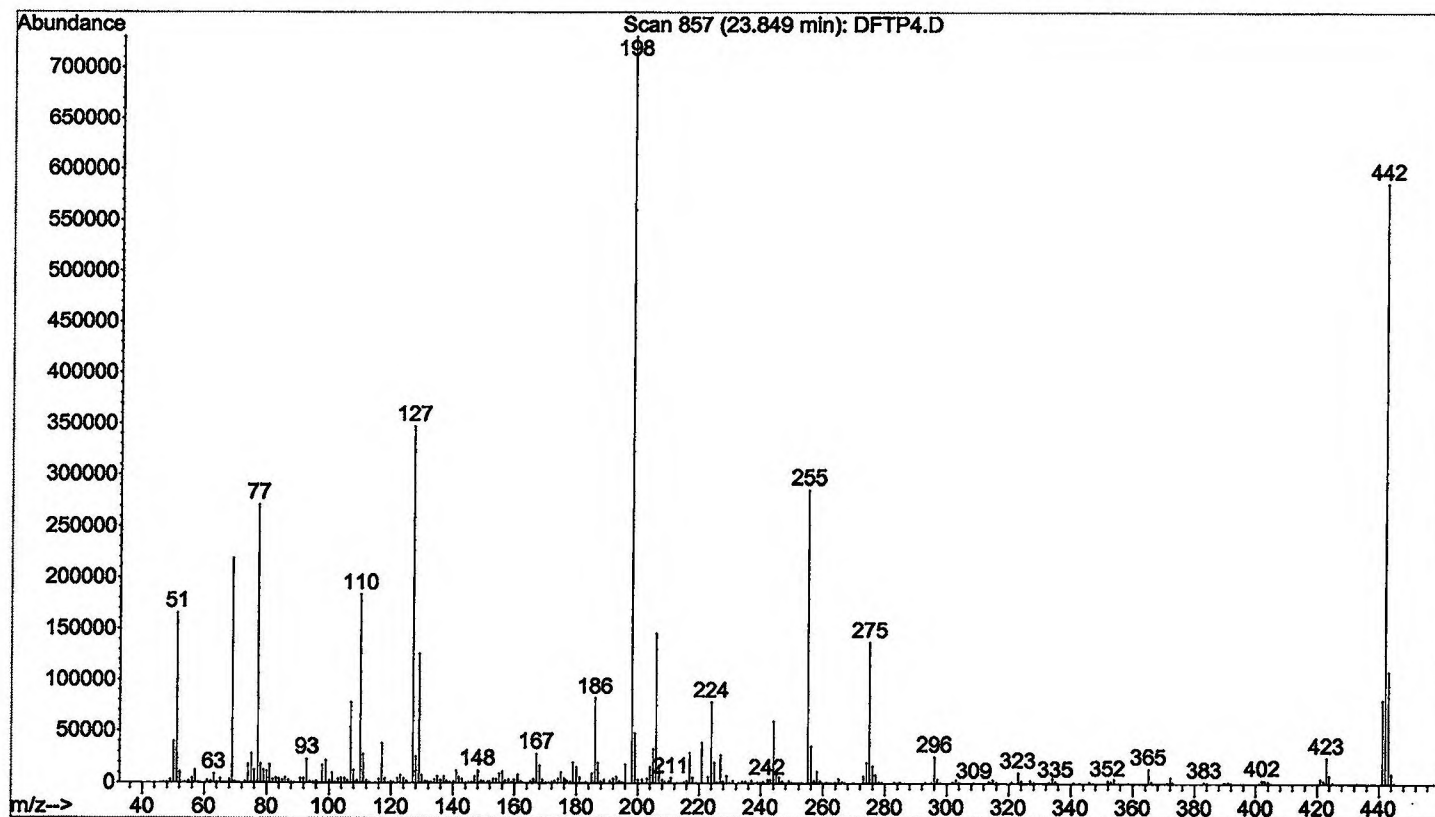
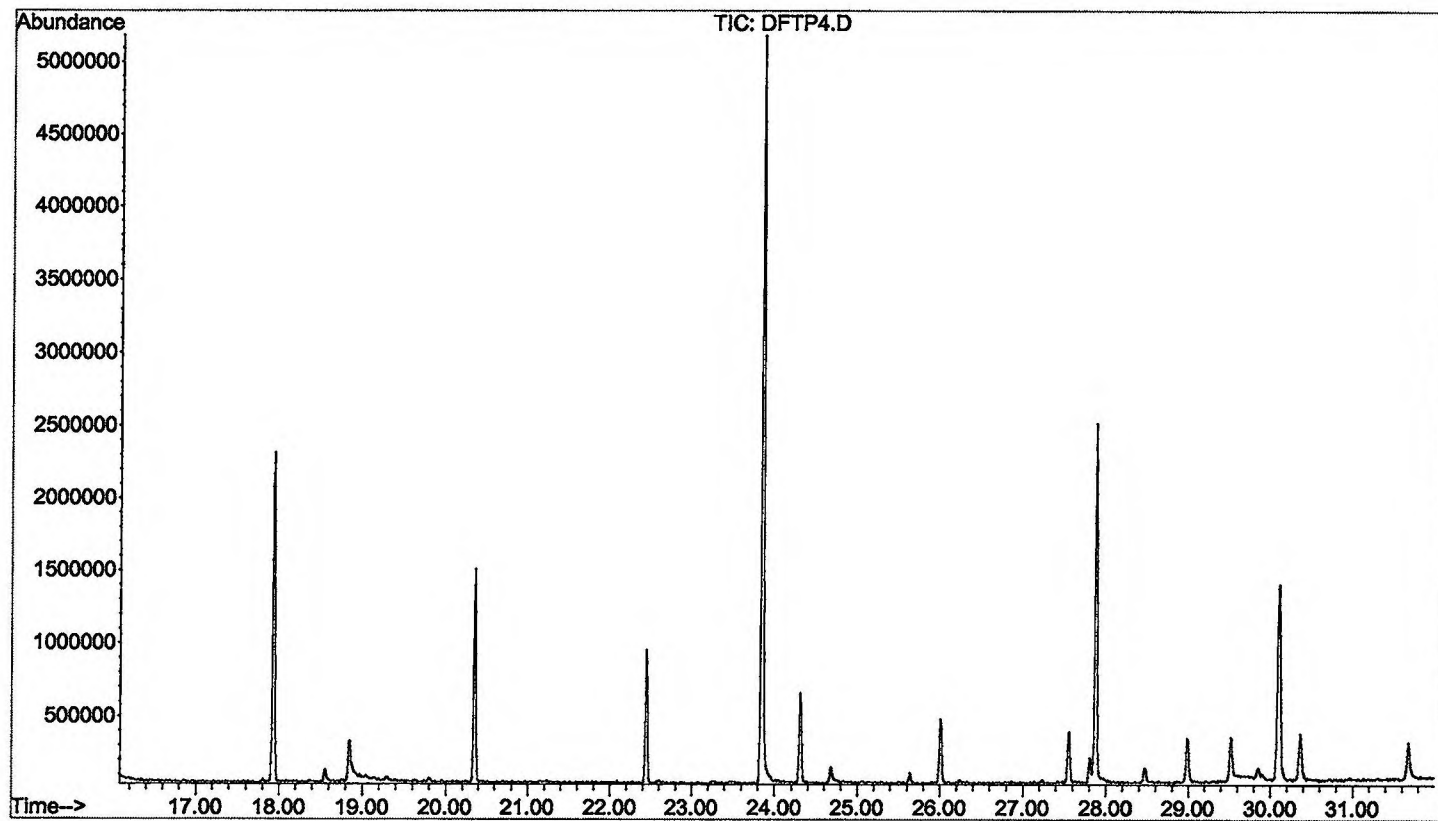
Method : C:\MSDCHEM\1\5973N\121401.M (RTE Integrator)
Title : Quantitation For Method 525.2

Spectrum Information: Scan 857

Target Mass	Rel. to Mass	Lower Limit%	Upper Limit%	Rel. Abn%	Raw Abn	Result Pass/Fail
51	198	10	80	22.7	165568	PASS
68	69	0.00	2	1.8	3887	PASS
69	198	0.00	100	30.0	219392	PASS
70	69	0.00	2	0.6	1373	PASS
127	198	10	80	47.5	347200	PASS
197	198	0.00	2	0.0	0	PASS
198	198	50	100	100.0	730752	PASS
199	198	5	9	6.6	47872	PASS
275	198	10	60	18.7	136768	PASS
365	198	1	100	2.0	14303	PASS
441	443	0.01	100	74.8	81360	PASS
442	198	50	100	80.2	586240	PASS
443	442	15	24	18.6	108776	PASS

DFTP4.D 121401.M Wed Jan 02 09:12:50 2002

File : C:\MSDCHEM\1\DATA\121901\DFTP4.D
 Operator : McLean
 Acquired : 20 Dec 2001 9:00 am using AcqMethod BIG3
 Instrument : Instrumen
 Sample Name:
 Misc Info :
 Vial Number: 14



Area Percent Report

Data File : C:\MSDCHEM\1\DATA\121901\DFTP4.D

Acq On : 20 Dec 2001 9:00 am

Sample :

Misc :

MS Integration Params: BENZO.P

Vial: 14

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	1297	1301	1305	rM 2	2406826	4105013	100.00%	82.502%
2	28.475	1364	1367	1371	rM 5	86148	178761	4.35%	3.593%
3	30.361	1571	1575	1580	rM	303919	691854	16.85%	13.905%

Sum of corrected areas: 4975628

Area Percent Report

Data File : C:\MSDCHEM\1\DATA\121901\DDT4.D

Acq On : 20 Dec 2001 9:39 am

Sample :

Misc :

MS Integration Params: BENZO.P

Vial: 15

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	28.983	1417	1423	1428	rM	451584	1030468	17.66%	15.010%
2	29.146	1433	1441	1448	rM	3011059	5834823	100.00%	84.990%

Sum of corrected areas: 6865291