

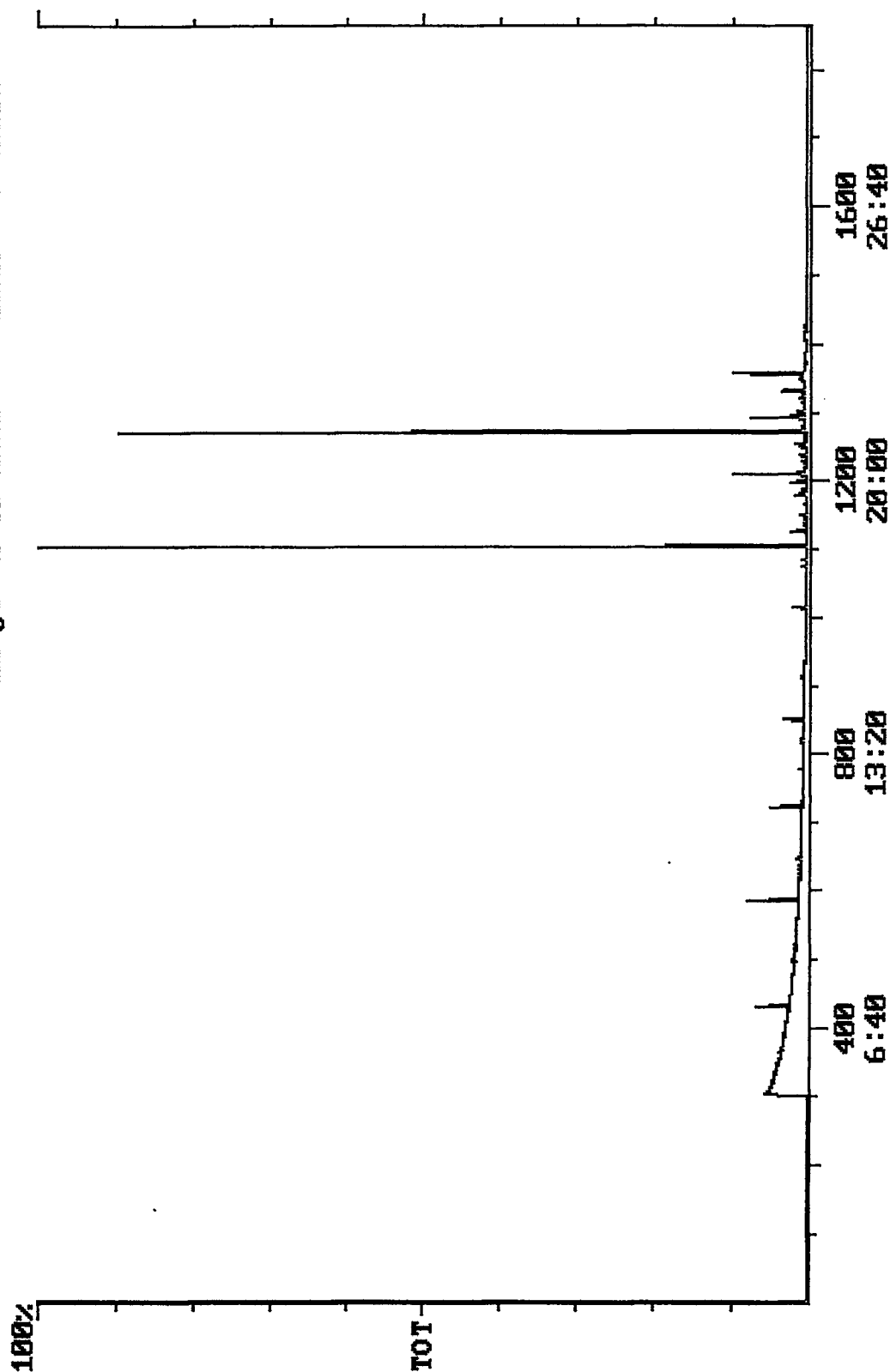
User: Bohk, Ted
Westchester Dept. of Labs & Research
Date: 11/30/2001 Time: 12:08:14

Sample: AD25041
Analysis: \$LP525 Organic Extractables
Units: ug
Started: 11/9/01 12:01 Ended: 11/9/01 12:01 Analyst: TB
Page: 1

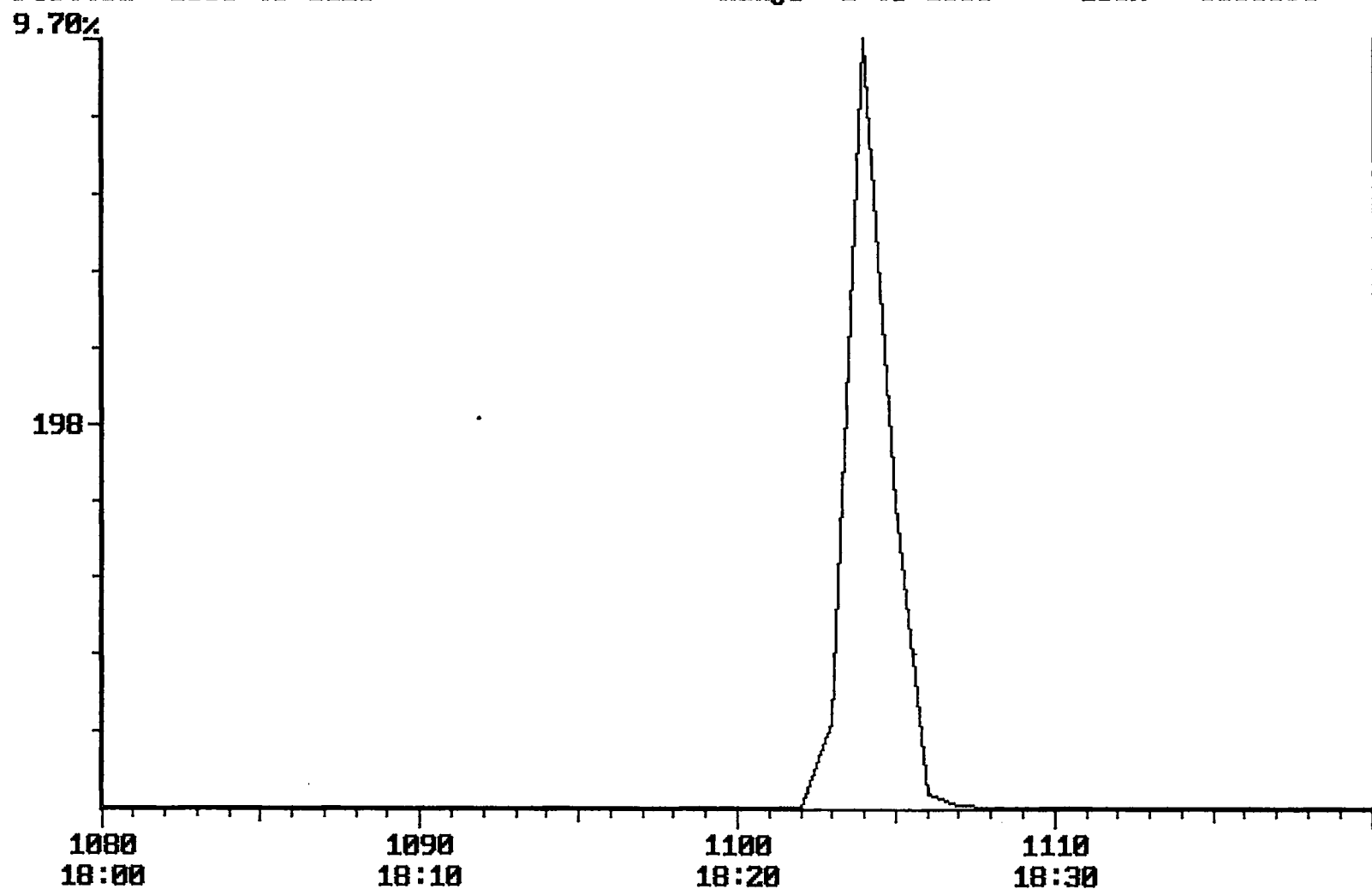
	Component Name	Viol	Result
1	DFTPP		passed
2	ENDRIN X		21
3	DDT		3.0

← flag.

Chromatogram Plot
 File: C:\DFTA1109 Date: Nov-09-1991 16:22:32
 Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN
 Scan No: 1 Retention Time: 0:01 RIC: 0 Mass Range: 0 - 0
 Plotted: 1 to 1860 Range: 1 to 1860 100% = 5568890



Chromatogram Plot File: C:\DFTA1109 Date: Nov-09-1991 16:22:32
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN
Scan No: 1080 Retention Time: 18:00 RIC: 23162 Mass Range: 47 - 86
Plotted: 1080 to 1120 Range: 1 to 1860 100% = 5568890



Mass List

File: C:\DFTA1109 Date: Nov-09-1991 16:22:32

Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN

Scan No: 1104 Retention Time: 18:24 RIC: 5568890 Mass Range: 47 - 444

Peaks: 169 Mass of Base Peak: 442 Int: 971385 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
47	2,710	0.28	84	3,614	0.37	125	3,313	0.34
49	14,156	1.46	85	5,421	0.56	127	164,457	16.93
50	118,674	12.22	86	11,746	1.21	128	15,662	1.61
51	350,602	36.09	91	3,915	0.40	129	84,939	8.74
52	12,349	1.27	92	3,313	0.34	130	10,240	1.05
55	4,518	0.47	93	38,855	4.00	132	2,710	0.28
56	9,036	0.93	94	3,614	0.37	134	3,313	0.34
57	12,650	1.30	98	34,337	3.53	135	7,530	0.78
61	5,421	0.56	99	13,855	1.43	136	4,819	0.50
62	4,819	0.50	101	8,132	0.84	137	2,710	0.28
63	10,542	1.09	103	5,120	0.53	141	15,060	1.55
67	3,614	0.37	104	6,927	0.71	142	3,915	0.40
68	5,722	0.59	105	5,722	0.59	143	4,819	0.50
69	240,361	24.74	106	3,313	0.34	147	5,722	0.59
70	1,807	0.19	107	79,518	8.19	148	14,759	1.52
73	3,614	0.37	108	9,638	0.99	149	4,216	0.43
74	35,240	3.63	109	4,518	0.47	153	5,120	0.53
75	30,120	3.10	110	147,590	15.19	154	4,819	0.50
76	15,662	1.61	111	17,469	1.80	155	7,228	0.74
77	259,939	26.76	112	3,313	0.34	156	8,734	0.90
78	15,361	1.58	116	4,216	0.43	157	3,012	0.31
79	30,722	3.16	117	93,072	9.58	158	2,710	0.28
80	20,783	2.14	118	9,036	0.93	160	3,614	0.37
81	18,674	1.92	122	6,325	0.65	161	5,722	0.59
82	4,819	0.50	123	8,132	0.84	162	2,710	0.28

Mass List

File: C:\DFTA1109 Date: Nov-09-1991 16:22:32

Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN

Scan No: 1104 Retention Time: 18:24 RIC: 5568890 Mass Range: 47 - 444

Peaks: 169 Mass of Base Peak: 442 Int: 971385 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
162	2,710	0.28	205	24,698	2.54	249	2,710	0.28
165	4,518	0.47	206	93,072	9.58	255	340,060	35.01
166	3,614	0.37	207	13,253	1.36	256	48,493	4.99
167	25,903	2.67	208	5,722	0.59	258	18,373	1.89
168	9,337	0.96	211	3,614	0.37	259	2,710	0.28
174	4,216	0.43	217	27,409	2.82	265	5,120	0.53
175	8,132	0.84	218	3,012	0.31	273	10,240	1.05
176	2,409	0.25	221	8,132	0.84	274	31,927	3.29
177	4,518	0.47	222	3,614	0.37	275	183,132	18.85
178	3,012	0.31	223	6,024	0.62	276	25,903	2.67
179	12,650	1.30	224	53,012	5.46	277	15,662	1.61
180	7,228	0.74	225	14,457	1.49	278	2,710	0.28
181	5,120	0.53	226	3,012	0.31	285	3,313	0.34
185	7,228	0.74	227	18,975	1.95	293	5,120	0.53
186	64,156	6.60	228	4,216	0.43	296	30,855	4.00
187	21,084	2.17	229	4,518	0.47	297	5,120	0.53
189	3,012	0.31	231	3,313	0.34	303	4,819	0.50
192	4,518	0.47	234	1,807	0.19	304	2,409	0.25
193	4,819	0.50	235	2,108	0.22	314	3,313	0.34
196	6,325	0.65	236	2,409	0.25	315	8,433	0.87
197	5,120	0.53	241	2,409	0.25	323	16,566	1.71
198	540,060	55.60	243	3,614	0.37	324	5,120	0.53
199	34,036	3.50	244	33,132	3.41	327	4,216	0.43
200	2,710	0.28	245	4,819	0.50	334	13,253	1.36
204	10,843	1.12	246	8,734	0.90	335	5,120	0.53

Mass List

File: C:\DFTA1109 Date: Nov-09-1991 16:22:32

Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN

Scan No: 1104 Retention Time: 18:24 RIC: 5568890 Mass Range: 47 - 444

Peaks: 169 Mass of Base Peak: 442 Int: 971385 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
162	2,710	0.28	205	24,698	2.54	249	2,710	0.28
165	4,518	0.47	206	93,072	9.58	255	340,060	35.01
166	3,614	0.37	207	13,253	1.36	256	48,493	4.99
167	25,903	2.67	208	5,722	0.59	258	18,373	1.89
168	9,337	0.96	211	3,614	0.37	259	2,710	0.28
174	4,216	0.43	217	27,409	2.82	265	5,120	0.53
175	8,132	0.84	218	3,012	0.31	273	10,240	1.05
176	2,409	0.25	221	8,132	0.84	274	31,927	3.29
177	4,518	0.47	222	3,614	0.37	275	183,132	18.85
178	3,012	0.31	223	6,024	0.62	276	25,903	2.67
179	12,650	1.30	224	53,012	5.46	277	15,662	1.61
180	7,228	0.74	225	14,457	1.49	278	2,710	0.28
181	5,120	0.53	226	3,012	0.31	285	3,313	0.34
185	7,228	0.74	227	18,975	1.95	293	5,120	0.53
186	64,156	6.60	228	4,216	0.43	296	38,855	4.00
187	21,084	2.17	229	4,518	0.47	297	5,120	0.53
189	3,012	0.31	231	3,313	0.34	303	4,819	0.50
192	4,518	0.47	234	1,807	0.19	304	2,409	0.25
193	4,819	0.50	235	2,108	0.22	314	3,313	0.34
196	6,325	0.65	236	2,409	0.25	315	8,433	0.87
197	5,120	0.53	241	2,409	0.25	323	16,566	1.71
198	540,060	55.60	243	3,614	0.37	324	5,120	0.53
199	34,036	3.50	244	33,132	3.41	327	4,216	0.43
200	2,710	0.28	245	4,819	0.50	334	13,253	1.36
204	10,843	1.12	246	8,734	0.90	335	5,120	0.53

Mass List

File: C:\DFTA1109 Date: Nov-09-1991 16:22:32

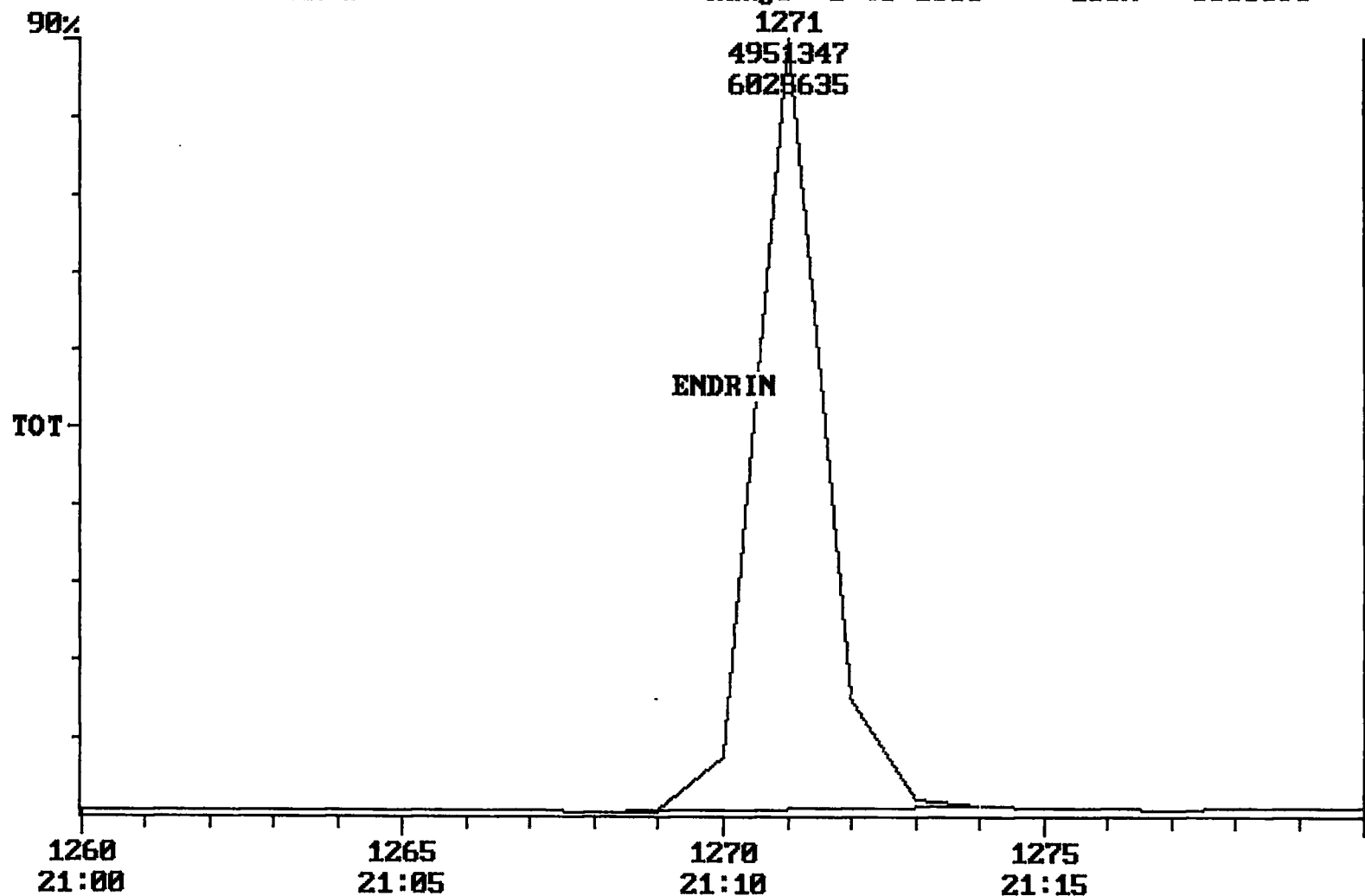
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN

Scan No: 1104 Retention Time: 18:24 RIC: 5568890 Mass Range: 47 - 444

Peaks: 169 Mass of Base Peak: 442 Int: 971385 Threshold: 0 %

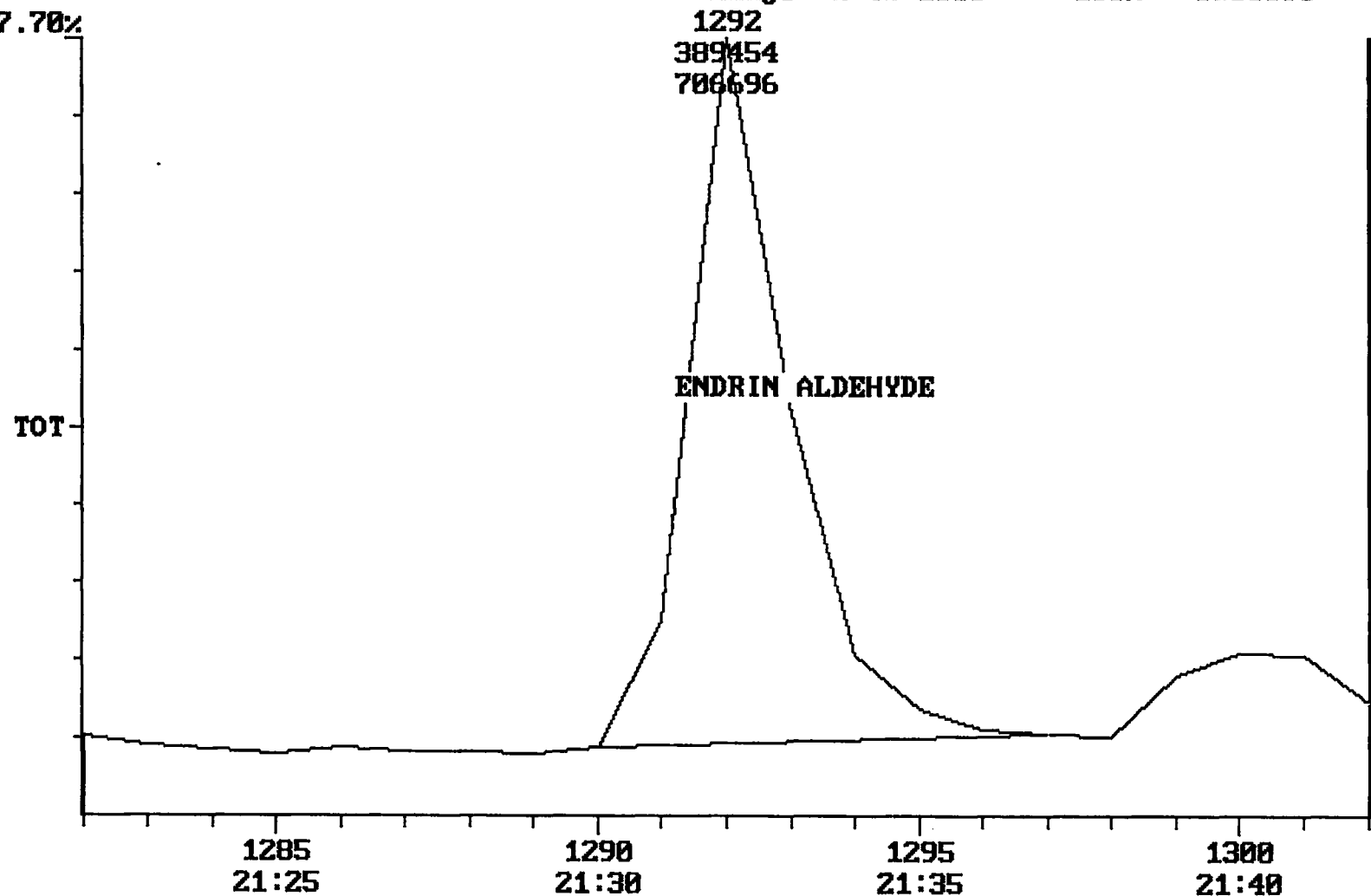
Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
197	5,120	0.53	241	2,409	0.25	323	16,566	1.71
198	540,060	55.60	243	3,614	0.37	324	5,120	0.53
199	34,036	3.50	244	33,132	3.41	327	4,216	0.43
200	2,710	0.28	245	4,819	0.50	334	13,253	1.36
204	10,843	1.12	246	8,734	0.90	335	5,120	0.53
205	24,698	2.54	249	2,710	0.28	341	2,108	0.22
206	93,072	9.58	255	340,060	35.01	346	4,819	0.50
207	13,253	1.36	256	48,493	4.99	352	5,722	0.59
208	5,722	0.59	258	18,373	1.89	353	6,325	0.65
211	3,614	0.37	259	2,710	0.28	354	11,144	1.15
217	27,409	2.82	265	5,120	0.53	365	43,975	4.53
218	3,012	0.31	273	10,240	1.05	366	9,939	1.02
221	8,132	0.84	274	31,927	3.29	372	14,156	1.46
222	3,614	0.37	275	183,132	18.85	373	6,325	0.65
223	6,024	0.62	276	25,903	2.67	402	4,518	0.47
224	53,012	5.46	277	15,662	1.61	403	8,433	0.87
225	14,457	1.49	278	2,710	0.28	404	3,614	0.37
226	3,012	0.31	285	3,313	0.34	421	4,216	0.43
227	18,975	1.95	293	5,120	0.53	422	6,927	0.71
228	4,216	0.43	296	38,855	4.00	423	42,168	4.34
229	4,518	0.47	297	5,120	0.53	424	8,734	0.90
231	3,313	0.34	303	4,819	0.50	441	138,855	14.29
234	1,807	0.19	304	2,409	0.25	442	971,385	100.00
235	2,108	0.22	314	3,313	0.34	443	162,349	16.71
236	2,409	0.25	315	8,433	0.87	444	12,349	1.27

Chromatogram Plot File: C:\DFTA1109 Date: Nov-09-1991 16:22:32
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN
Scan No: 1280 Retention Time: 21:20 RIC: 45198 Mass Range: 45 - 355
Plotted: 1260 to 1280 Range: 1 to 1860 100% = 5568890

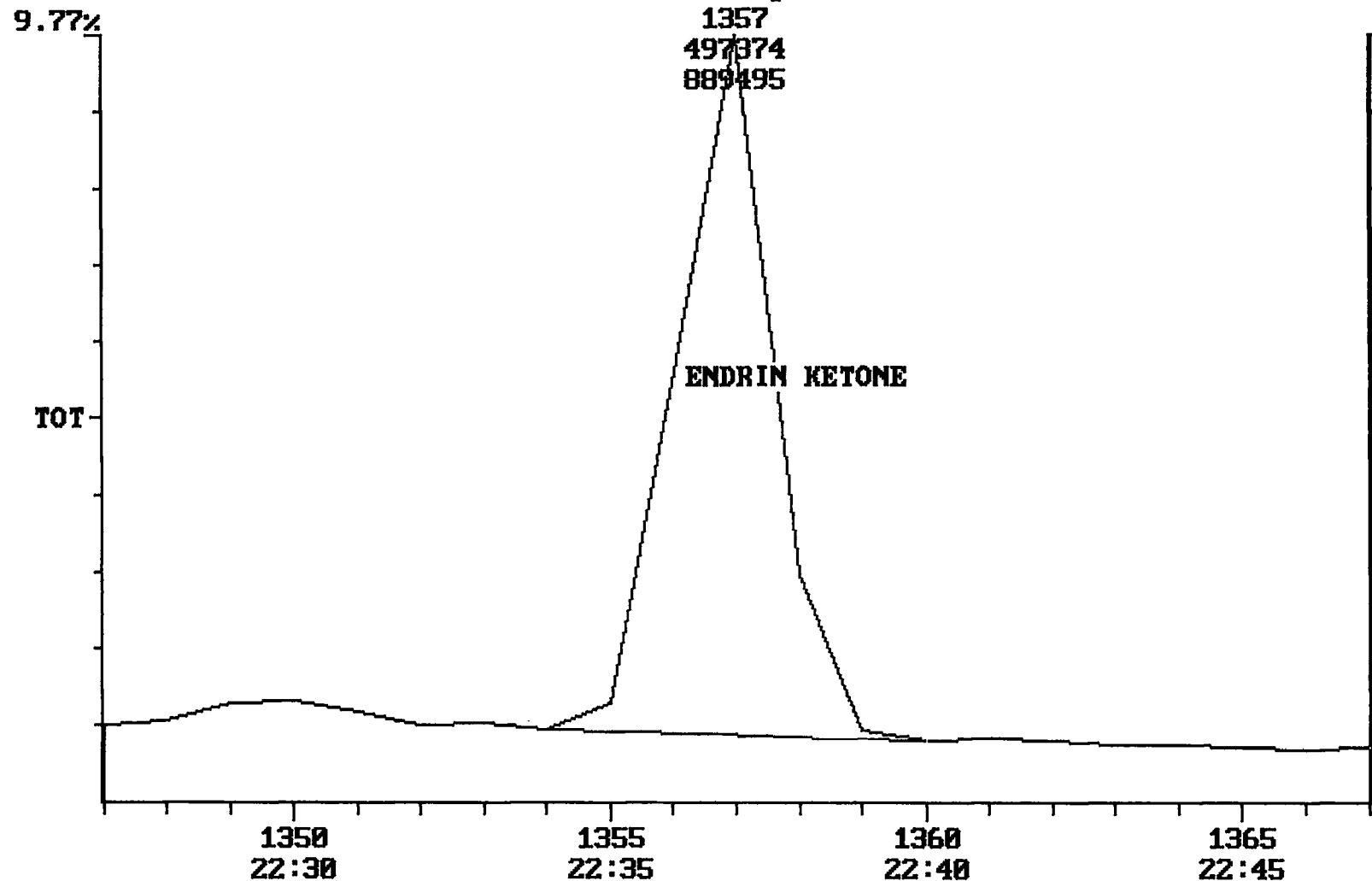


Breakdown = 20.9 %

Chromatogram Plot File: C:\DFTA1109 Date: Nov-09-1991 16:22:32
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN
Scan No: 1302 Retention Time: 21:42 RIC: 62169 Mass Range: 45 - 429
Plotted: 1282 to 1302 Range: 1 to 1860 100% = 5568890
7.70%



Chromatogram Plot File: C:\DFTA1109 Date: Nov-09-1991 16:22:32
 Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN
 Scan No: 1367 Retention Time: 22:47 RIC: 37521 Mass Range: 45 - 355
 Plotted: 1347 to 1367 Range: 1 to 1860 100% = 5568890



Chromatogram Plot

File: C:\DDTA1109

Date: Nov-09-1991 17:03:35

Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DDT

Scan No: 1

Retention Time: 0:01

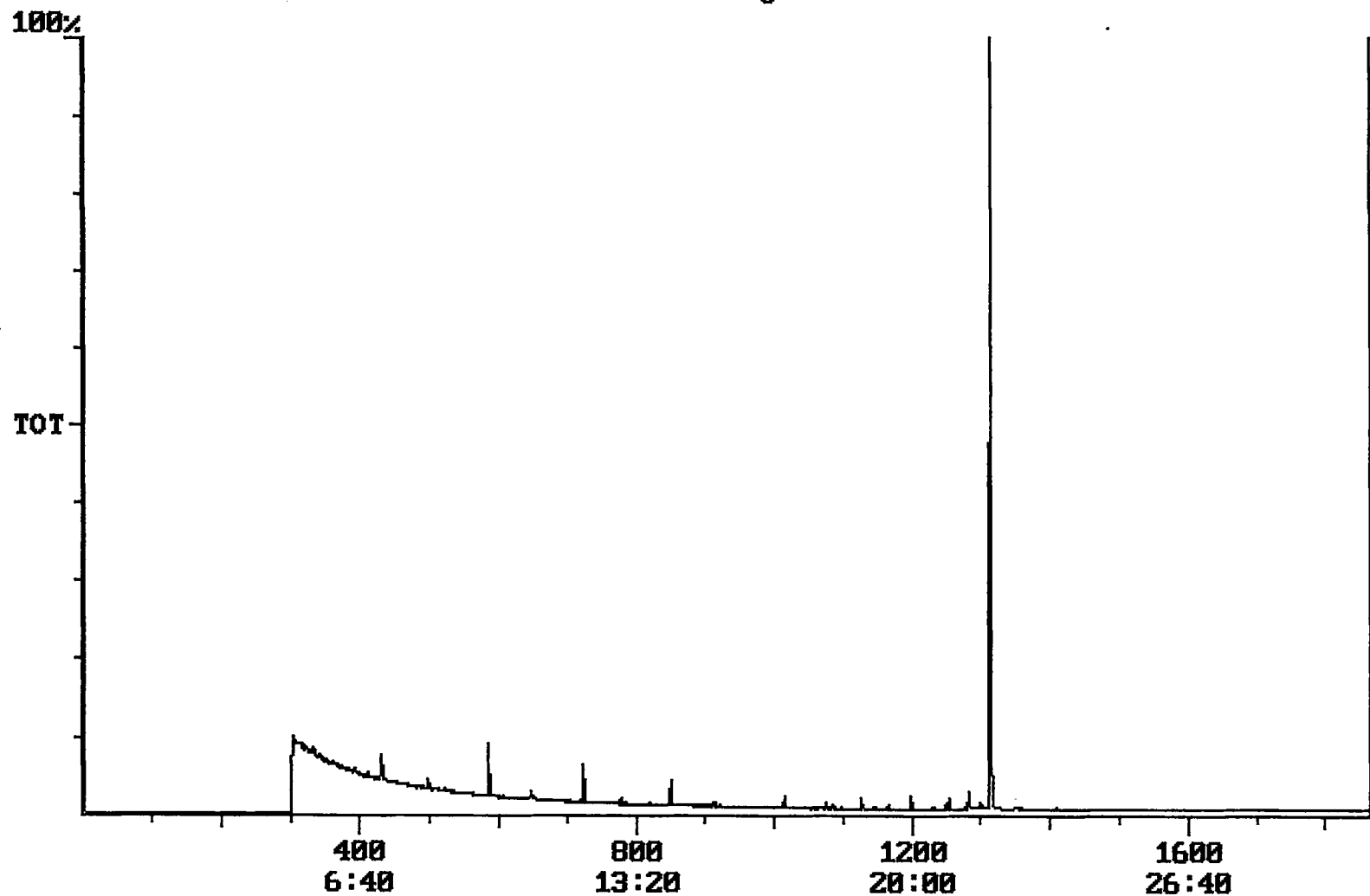
RIC: 0

Mass Range: 0 - 0

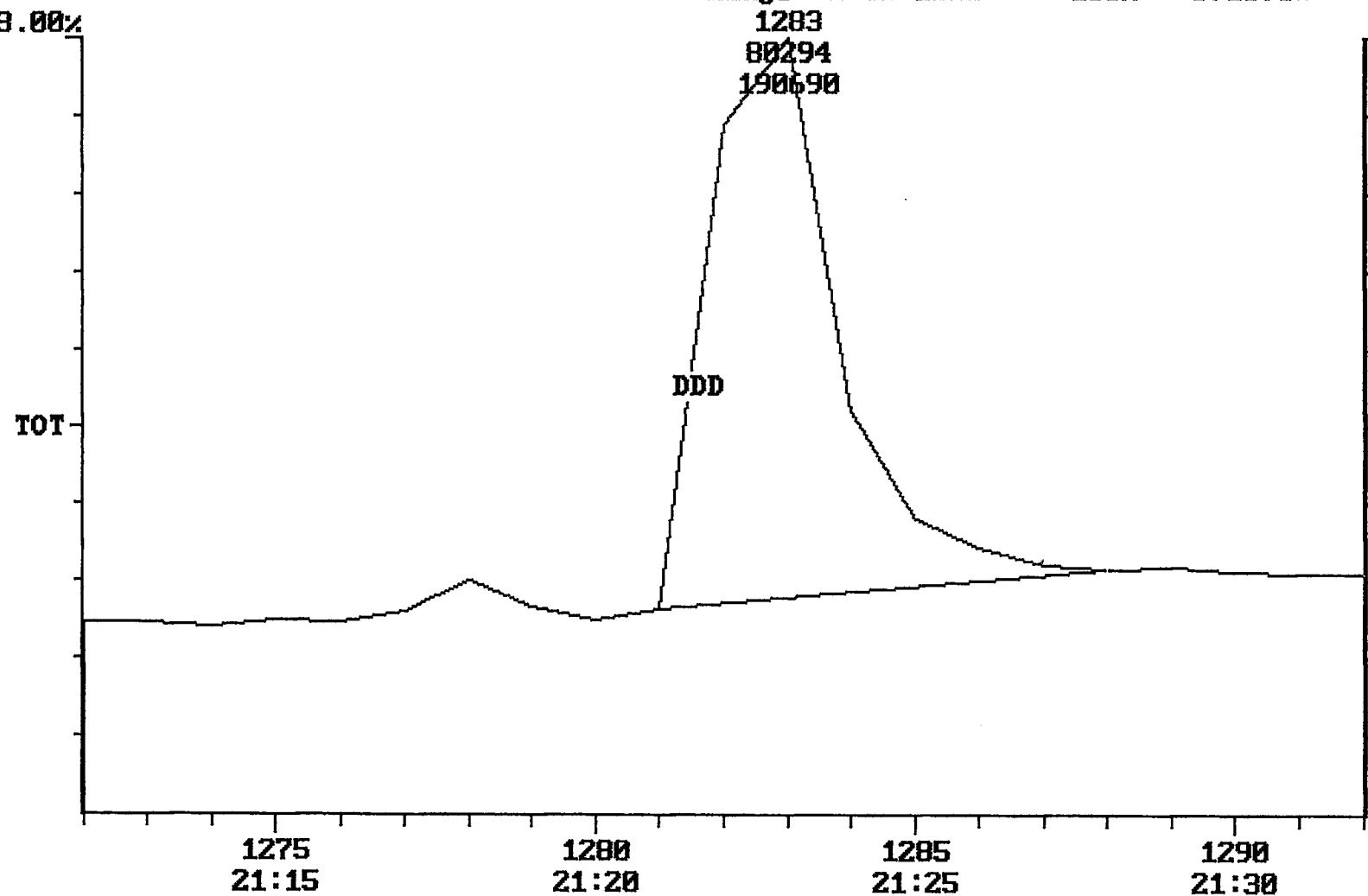
Plotted: 1 to 1860

Range: 1 to 1860

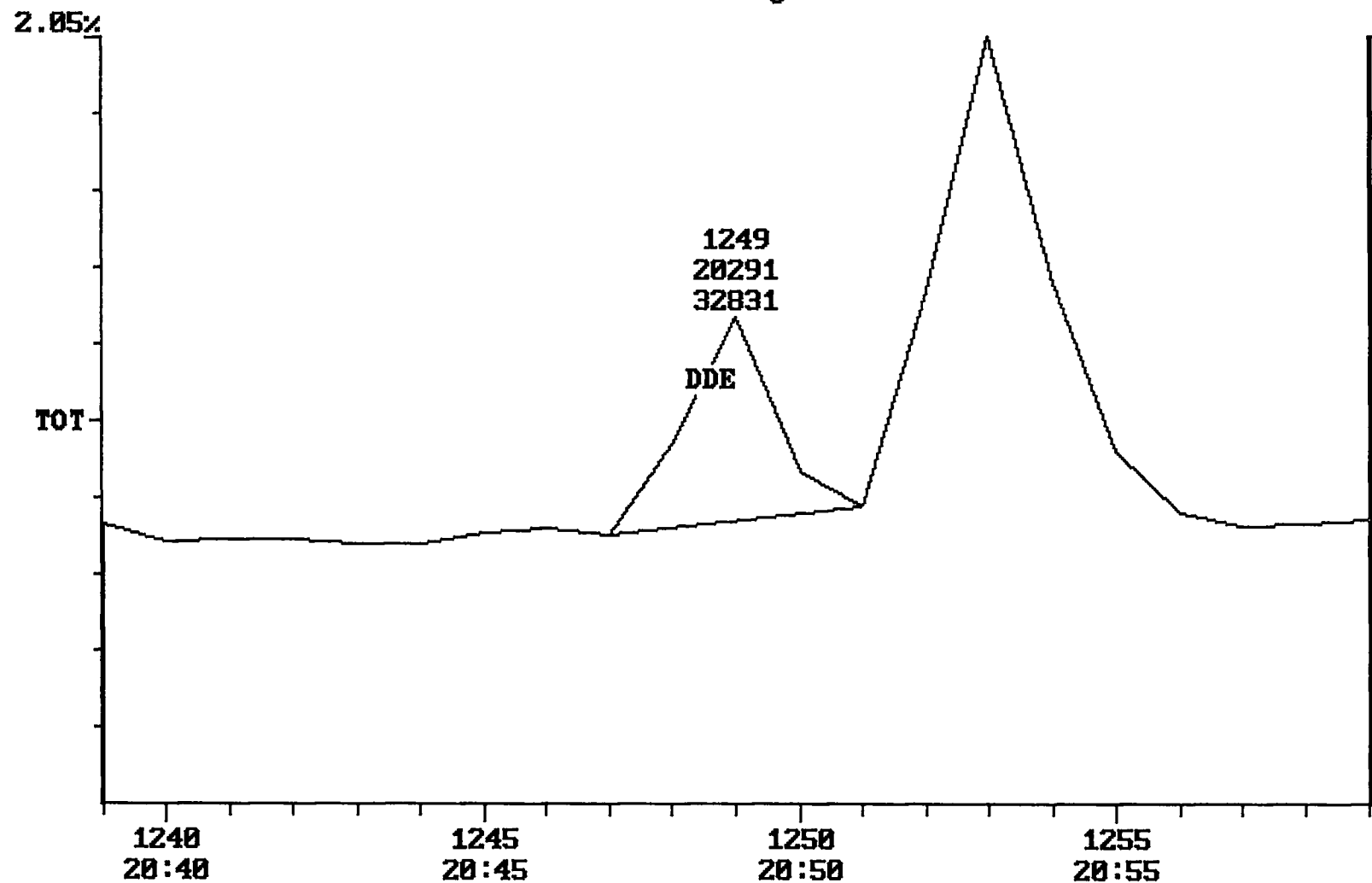
100% = 3710705



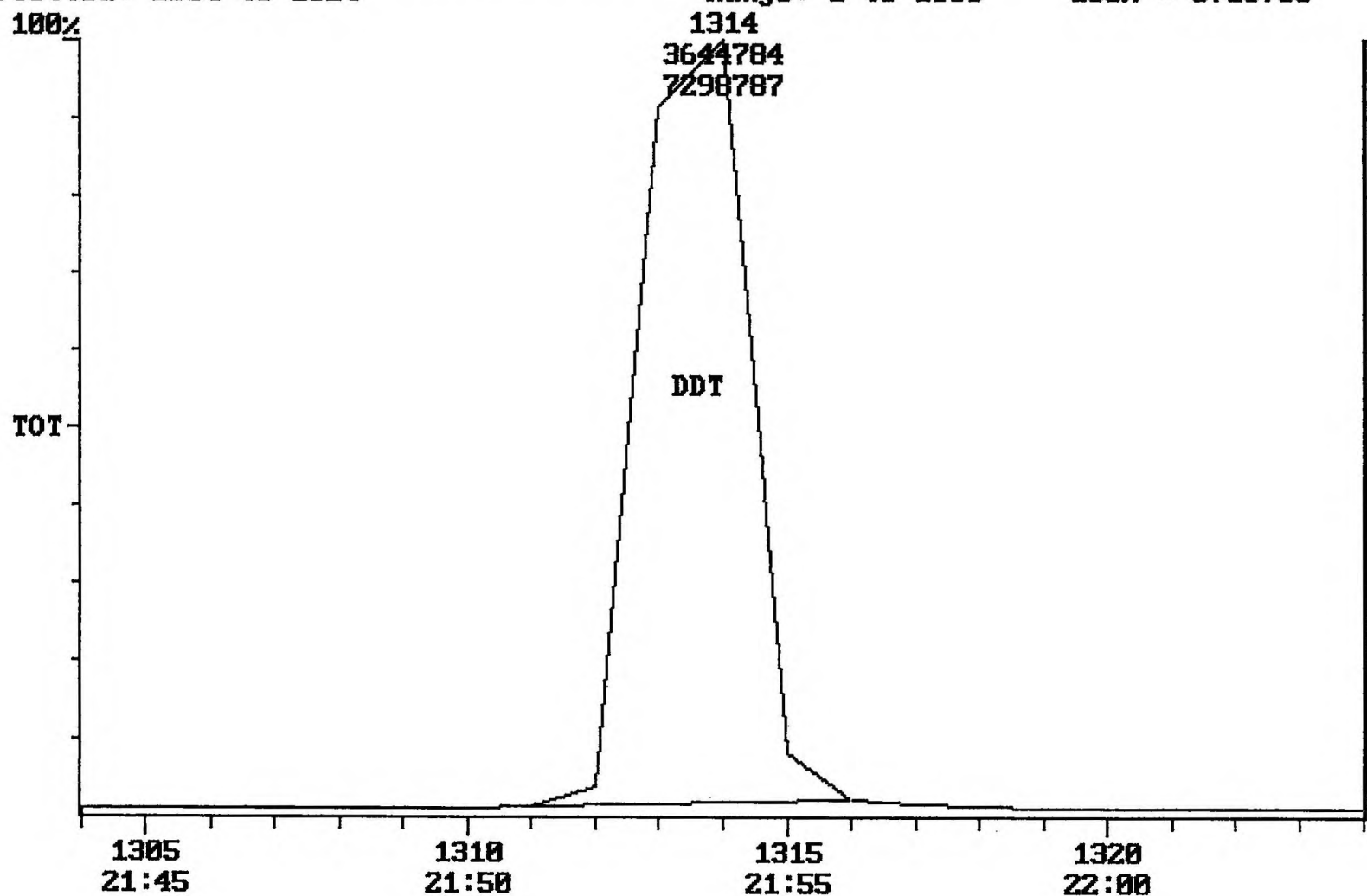
Chromatogram Plot File: C:\DDTA1109 Date: Nov-09-1991 17:03:35
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DDT
Scan No: 1292 Retention Time: 21:32 RIC: 34099 Mass Range: 47 - 237
Plotted: 1272 to 1292 Range: 1 to 1860 100% = 3710705
3.00%



Chromatogram Plot File: C:\DDTA1109 Date: Nov-09-1991 17:03:35
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DDT
Scan No: 1259 Retention Time: 20:59 RIC: 28111 Mass Range: 47 - 207
Plotted: 1239 to 1259 Range: 1 to 1860 100% = 3710705

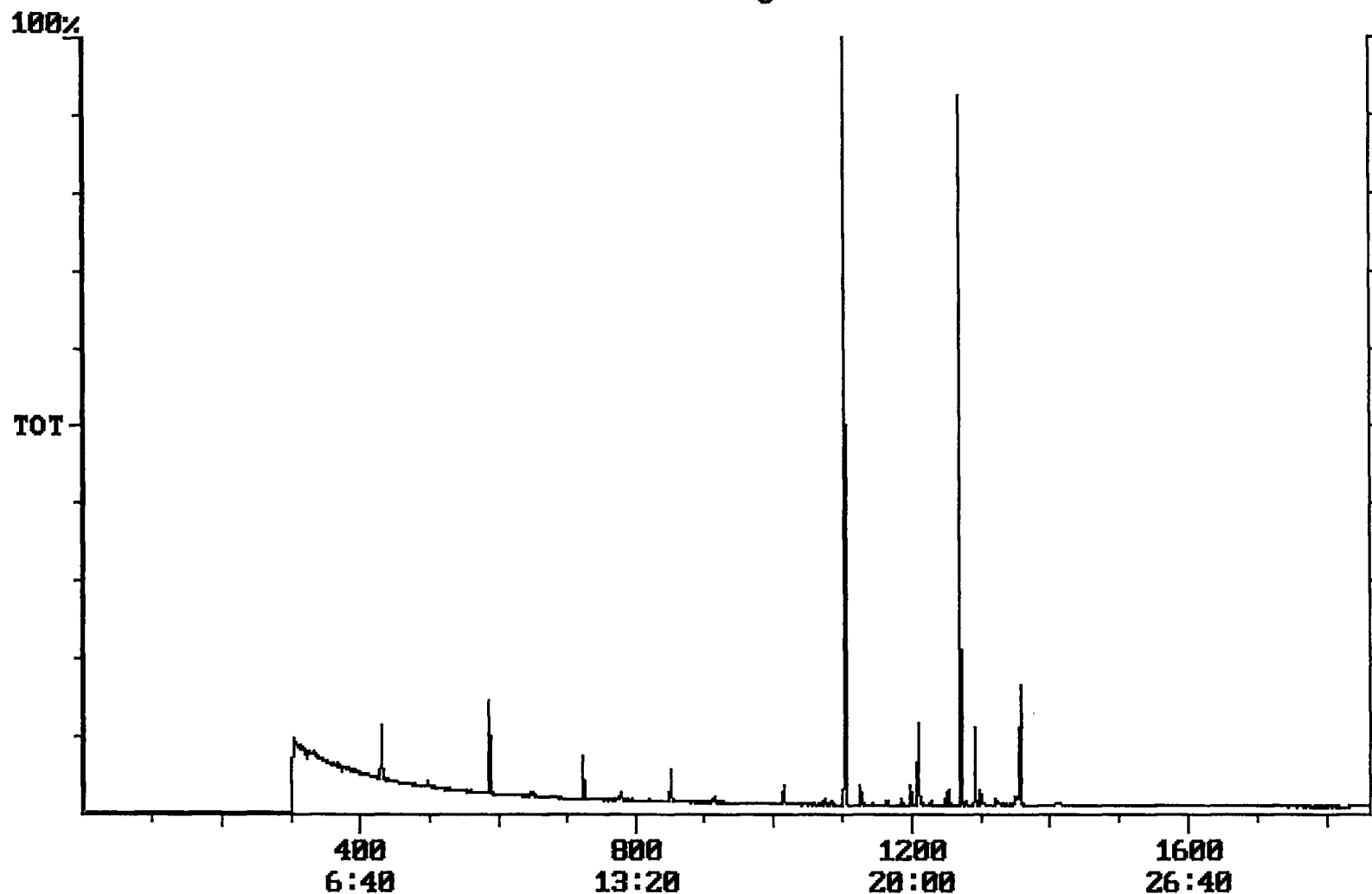


Chromatogram Plot File: C:\DDTA1109 Date: Nov-09-1991 17:03:35
 Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DDT
 Scan No: 1324 Retention Time: 22:04 RIC: 34198 Mass Range: 47 - 281
 Plotted: 1304 to 1324 Range: 1 to 1860 100% = 3710705

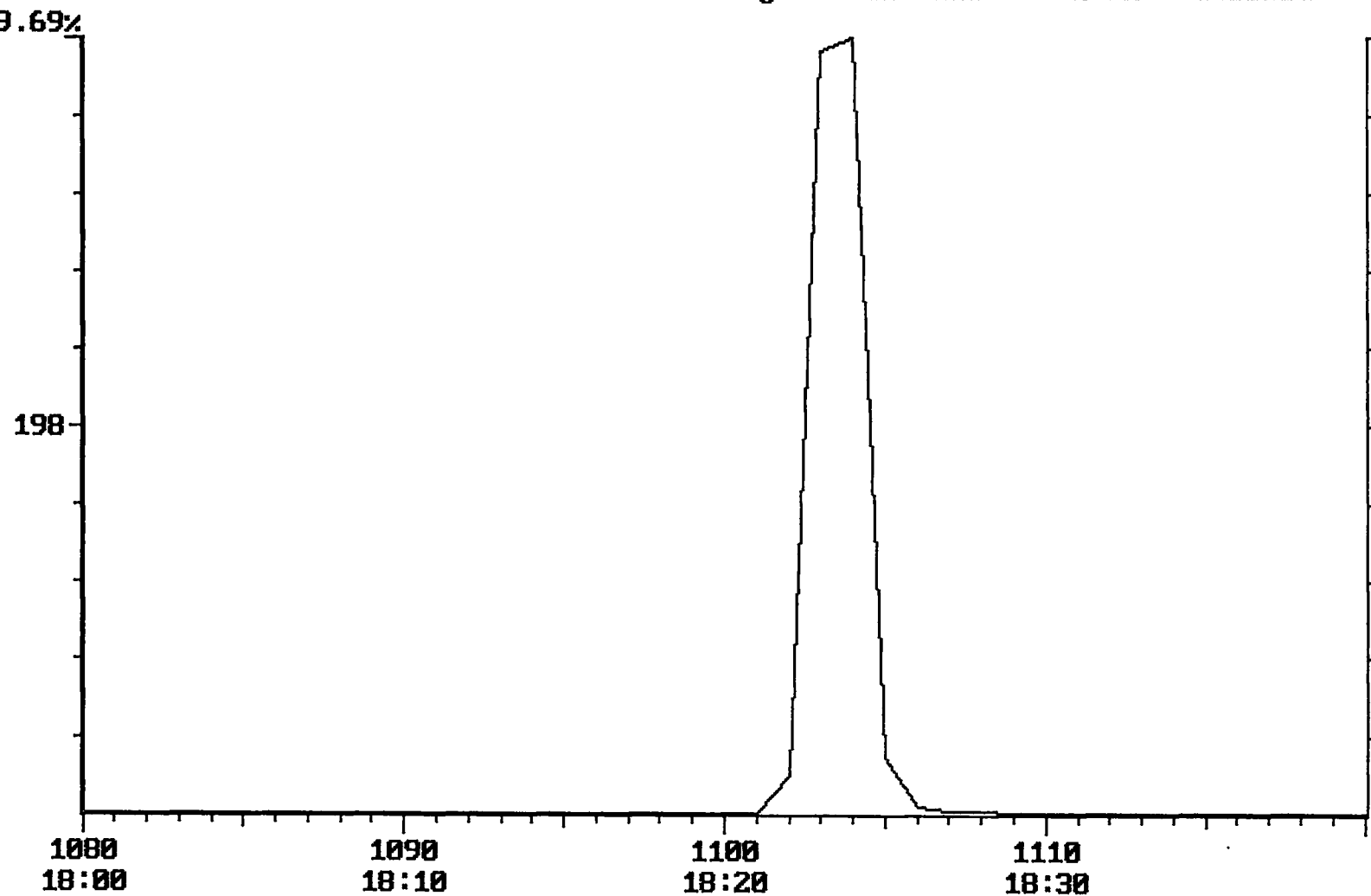


Breakdown = 3.0%

Chromatogram Plot File: D:\DFTB1109 Date: Nov-10-1991 04:35:50
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN
Scan No: 1 Retention Time: 0:01 RIC: 0 Mass Range: 0 - 0
Plotted: 1 to 1859 Range: 1 to 1859 100% = 4380084



Chromatogram Plot File: D:\DFTB1109 Date: Nov-10-1991 04:35:50
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN
Scan No: 1080 Retention Time: 18:00 RIC: 49512 Mass Range: 47 - 86
Plotted: 1080 to 1120 Range: 1 to 1859 100% = 4380084
9.69%



Mass List

File: D:\DFTB1109 Date: Nov-10-1991 04:35:50

Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN

Scan No: 1104 Retention Time: 18:24 RIC: 4137187 Mass Range: 47 - 444

Peaks: 163 Mass of Base Peak: 442 Int: 637740 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
47	5,288	0.83	84	6,971	1.09	124	2,163	0.34
48	2,403	0.38	85	2,884	0.45	125	3,365	0.53
49	19,230	3.02	86	10,096	1.58	127	134,615	21.11
50	90,865	14.25	91	4,326	0.68	128	11,778	1.85
51	271,875	42.63	92	2,163	0.34	129	56,009	8.78
52	11,057	1.73	93	29,326	4.60	130	5,288	0.83
55	4,807	0.75	94	2,884	0.45	134	2,884	0.45
56	8,173	1.28	96	2,884	0.45	135	8,413	1.32
57	10,817	1.70	98	29,086	4.56	141	12,259	1.92
61	3,605	0.57	99	15,384	2.41	142	3,365	0.53
62	3,846	0.60	101	6,250	0.98	146	3,125	0.49
63	11,538	1.81	103	7,451	1.17	147	6,971	1.09
68	2,644	0.41	104	5,048	0.79	148	9,855	1.55
69	200,240	31.40	105	5,769	0.90	149	3,125	0.49
73	3,605	0.57	106	1,923	0.30	153	2,644	0.41
74	30,288	4.75	107	59,375	9.31	155	5,048	0.79
75	30,769	4.82	108	10,336	1.62	156	6,730	1.06
76	9,855	1.55	109	3,365	0.53	158	2,403	0.38
77	203,125	31.85	110	108,173	16.96	160	2,884	0.45
78	10,096	1.58	111	12,019	1.88	161	4,567	0.72
79	23,076	3.62	116	3,605	0.57	165	4,567	0.72
80	13,701	2.15	117	74,519	11.68	166	3,125	0.49
81	14,903	2.34	118	3,605	0.57	167	20,192	3.17
82	3,846	0.60	122	5,048	0.79	168	8,653	1.36
83	3,125	0.49	123	6,730	1.06	173	2,644	0.41

Mass List

File: D:\DFTB1109 Date: Nov-10-1991 04:35:50

Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN

Scan No: 1104 Retention Time: 18:24 RIC: 4137187 Mass Range: 47 - 444

Peaks: 163 Mass of Base Peak: 442 Int: 637740 Threshold: 0 %

Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
174	3,365	0.53	217	23,798	3.73	275	124,278	19.49
175	4,326	0.68	218	4,567	0.72	276	13,942	2.19
177	3,605	0.57	221	9,134	1.43	277	13,701	2.15
179	9,855	1.55	222	2,884	0.45	293	3,125	0.49
180	8,173	1.28	223	5,048	0.79	294	1,923	0.30
181	2,644	0.41	224	39,182	6.14	296	28,846	4.52
185	9,615	1.51	225	10,576	1.66	297	3,605	0.57
186	54,086	8.48	227	24,519	3.84	303	4,086	0.64
187	11,778	1.85	228	1,923	0.30	304	2,884	0.45
189	3,365	0.53	229	4,567	0.72	315	4,807	0.75
191	2,403	0.38	235	1,923	0.30	316	2,403	0.38
192	3,125	0.49	236	2,163	0.34	323	13,942	2.19
193	4,807	0.75	243	3,125	0.49	324	2,403	0.38
196	8,173	1.28	244	29,086	4.56	327	3,846	0.60
197	3,605	0.57	245	3,846	0.60	334	9,615	1.51
198	424,519	66.57	246	8,173	1.28	335	3,605	0.57
199	23,798	3.73	249	2,163	0.34	346	2,403	0.38
203	2,644	0.41	255	236,778	37.13	352	2,403	0.38
204	9,855	1.55	256	33,894	5.31	353	2,884	0.45
205	17,548	2.75	257	3,365	0.53	354	6,971	1.09
206	72,355	11.35	258	11,778	1.85	355	2,644	0.41
207	10,336	1.62	259	3,125	0.49	364	3,125	0.49
208	3,846	0.60	265	4,086	0.64	365	23,076	3.62
211	3,125	0.49	273	7,211	1.13	366	4,567	0.72
216	1,442	0.23	274	23,557	3.69	371	1,923	0.30

Mass List

File: D:\DFTB1109 Date: Nov-10-1991 04:35:50

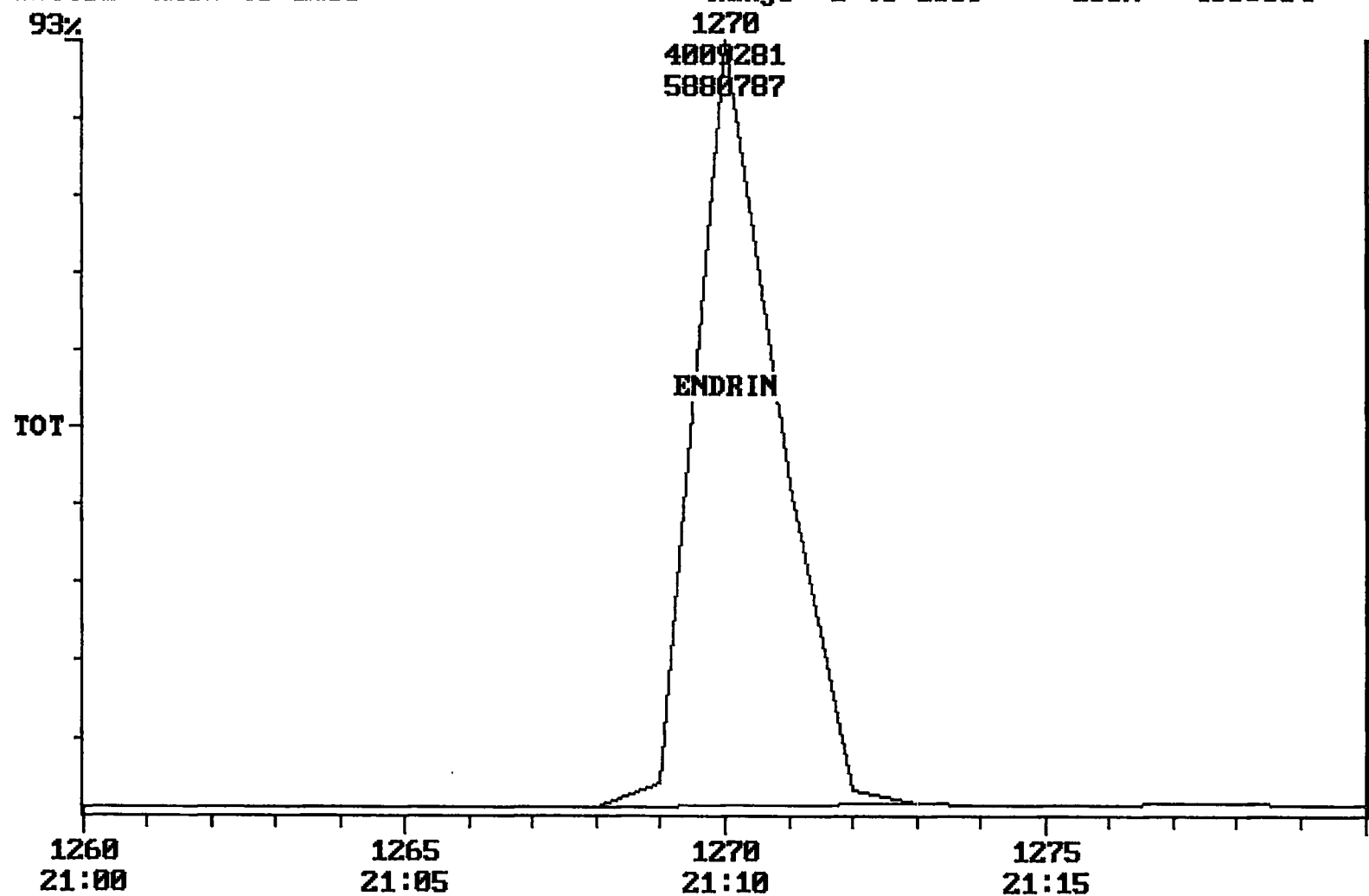
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN

Scan No: 1104 Retention Time: 18:24 RIC: 4137187 Mass Range: 47 - 444

Peaks: 163 Mass of Base Peak: 442 Int: 637740 Threshold: 0 %

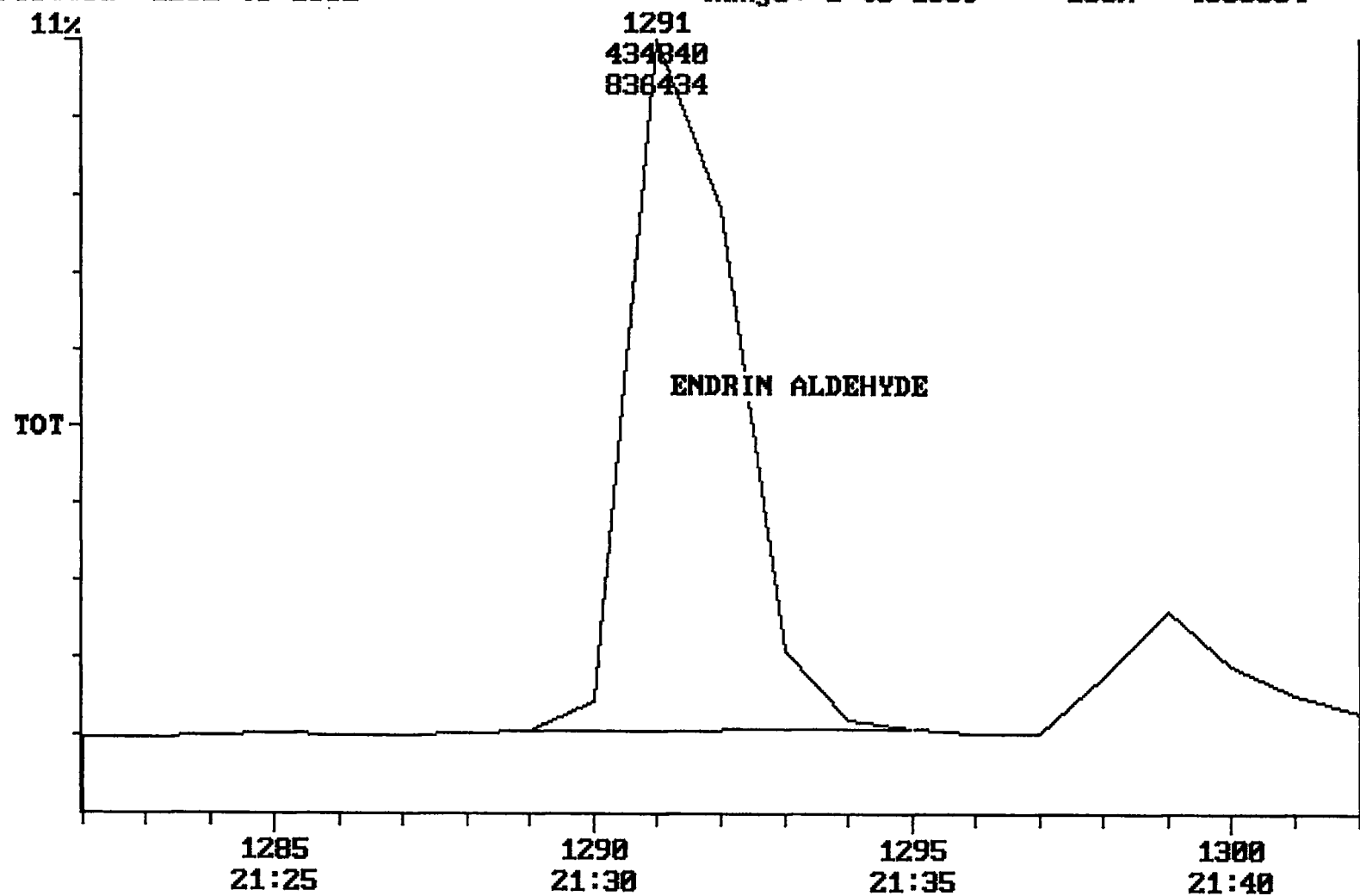
Mass	Intensity	% Base	Mass	Intensity	% Base	Mass	Intensity	% Base
196	8,173	1.28	244	29,086	4.56	327	3,846	0.60
197	3,605	0.57	245	3,846	0.60	334	9,615	1.51
198	424,519	66.57	246	8,173	1.28	335	3,605	0.57
199	23,798	3.73	249	2,163	0.34	346	2,403	0.38
203	2,644	0.41	255	236,778	37.13	352	2,403	0.38
204	9,855	1.55	256	33,894	5.31	353	2,884	0.45
205	17,548	2.75	257	3,365	0.53	354	6,971	1.09
206	72,355	11.35	258	11,778	1.85	355	2,644	0.41
207	10,336	1.62	259	3,125	0.49	364	3,125	0.49
208	3,846	0.60	265	4,086	0.64	365	23,076	3.62
211	3,125	0.49	273	7,211	1.13	366	4,567	0.72
216	1,442	0.23	274	23,557	3.69	371	1,923	0.30
217	23,798	3.73	275	124,278	19.49	372	9,855	1.55
218	4,567	0.72	276	13,942	2.19	373	2,403	0.38
221	9,134	1.43	277	13,701	2.15	383	1,923	0.30
222	2,884	0.45	293	3,125	0.49	402	2,163	0.34
223	5,048	0.79	294	1,923	0.30	403	4,086	0.64
224	39,182	6.14	296	28,846	4.52	421	4,086	0.64
225	10,576	1.66	297	3,605	0.57	422	2,884	0.45
227	24,519	3.84	303	4,086	0.64	423	29,567	4.64
228	1,923	0.30	304	2,884	0.45	424	3,605	0.57
229	4,567	0.72	315	4,007	0.75	441	90,865	14.25
235	1,923	0.30	316	2,403	0.38	442	637,740	100.00
236	2,163	0.34	323	13,942	2.19	443	115,384	18.09
243	3,125	0.49	324	2,403	0.38	444	4,086	0.64

Chromatogram Plot File: D:\DFTB1109 Date: Nov-10-1991 04:35:50
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN
Scan No: 1280 Retention Time: 21:20 RIC: 52278 Mass Range: 45 - 355
Plotted: 1260 to 1280 Range: 1 to 1859 100% = 4380084

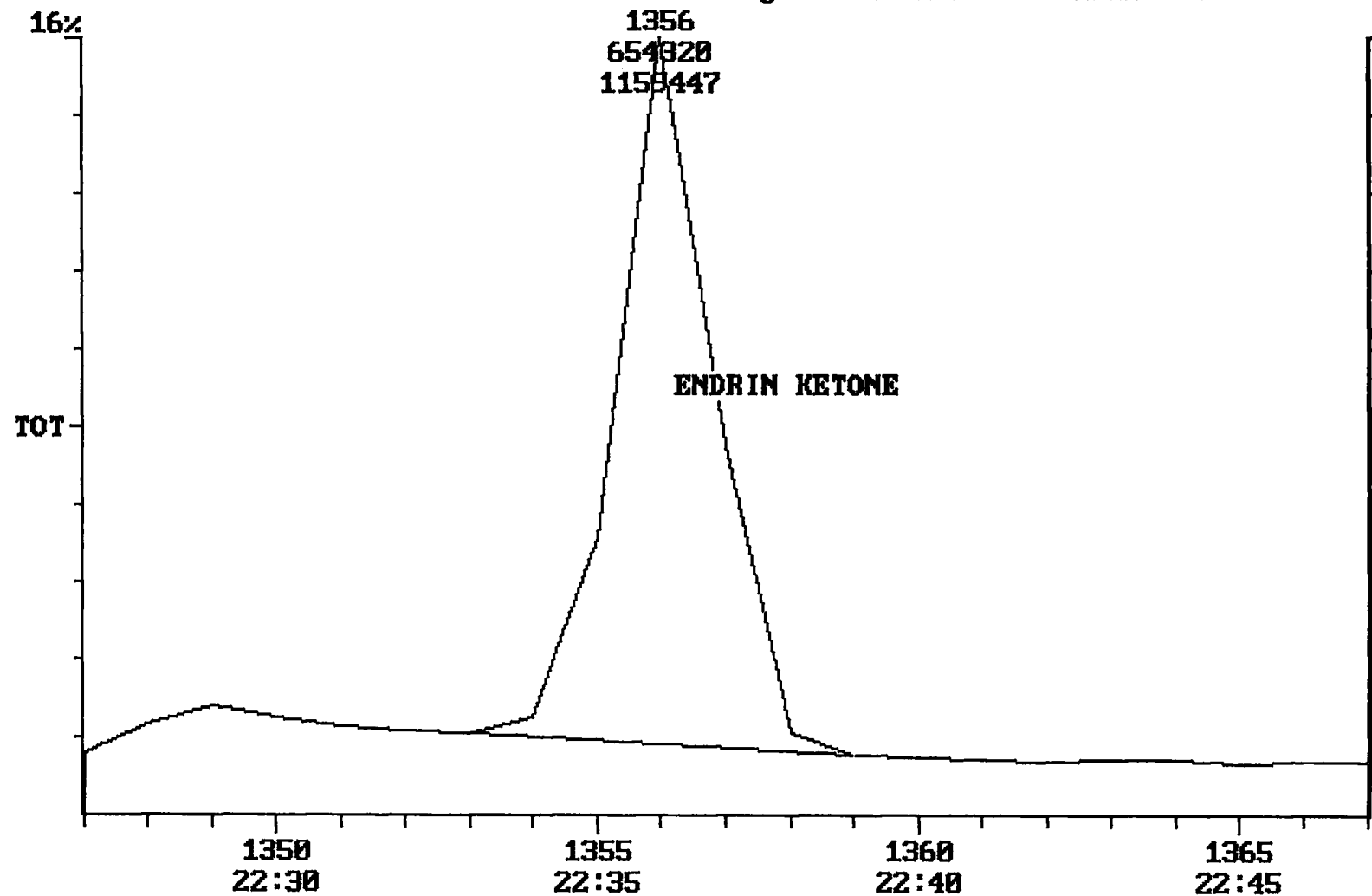


Breakdown = 25.3%

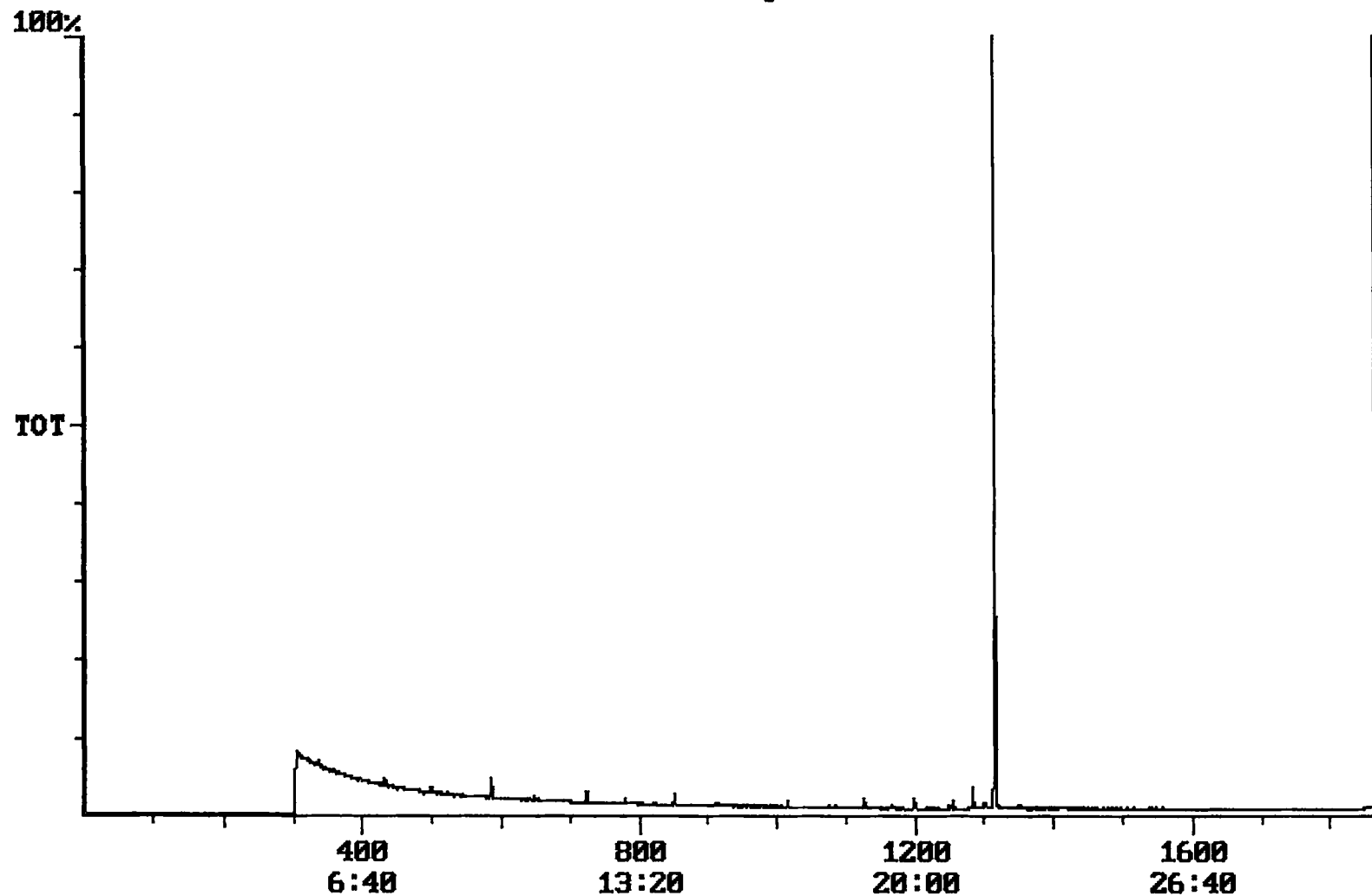
Chromatogram Plot File: D:\DFTB1109 Date: Nov-10-1991 04:35:50
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN
Scan No: 1302 Retention Time: 21:42 RIC: 62373 Mass Range: 45 - 429
Plotted: 1282 to 1302 Range: 1 to 1859 100% = 4380084



Chromatogram Plot File: D:\DFTB1109 Date: Nov-10-1991 04:35:50.
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DFTPP/ENDRIN
Scan No: 1367 Retention Time: 22:47 RIC: 48424 Mass Range: 45 - 281
Plotted: 1347 to 1367 Range: 1 to 1859 100% = 4380084



Chromatogram Plot File: D:\DDTB1109 Date: Nov-10-1991 05:10:13
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DDT
Scan No: 1 Retention Time: 0:01 RIC: 0 Mass Range: 0 - 0
Plotted: 1 to 1860 Range: 1 to 1860 100% = 4931445



Chromatogram Plot

File: D:\DDTB1109

Date: Nov-10-1991 05:10:13

Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DDT

Scan No: 1259

Retention Time: 20:59

RIC: 41882

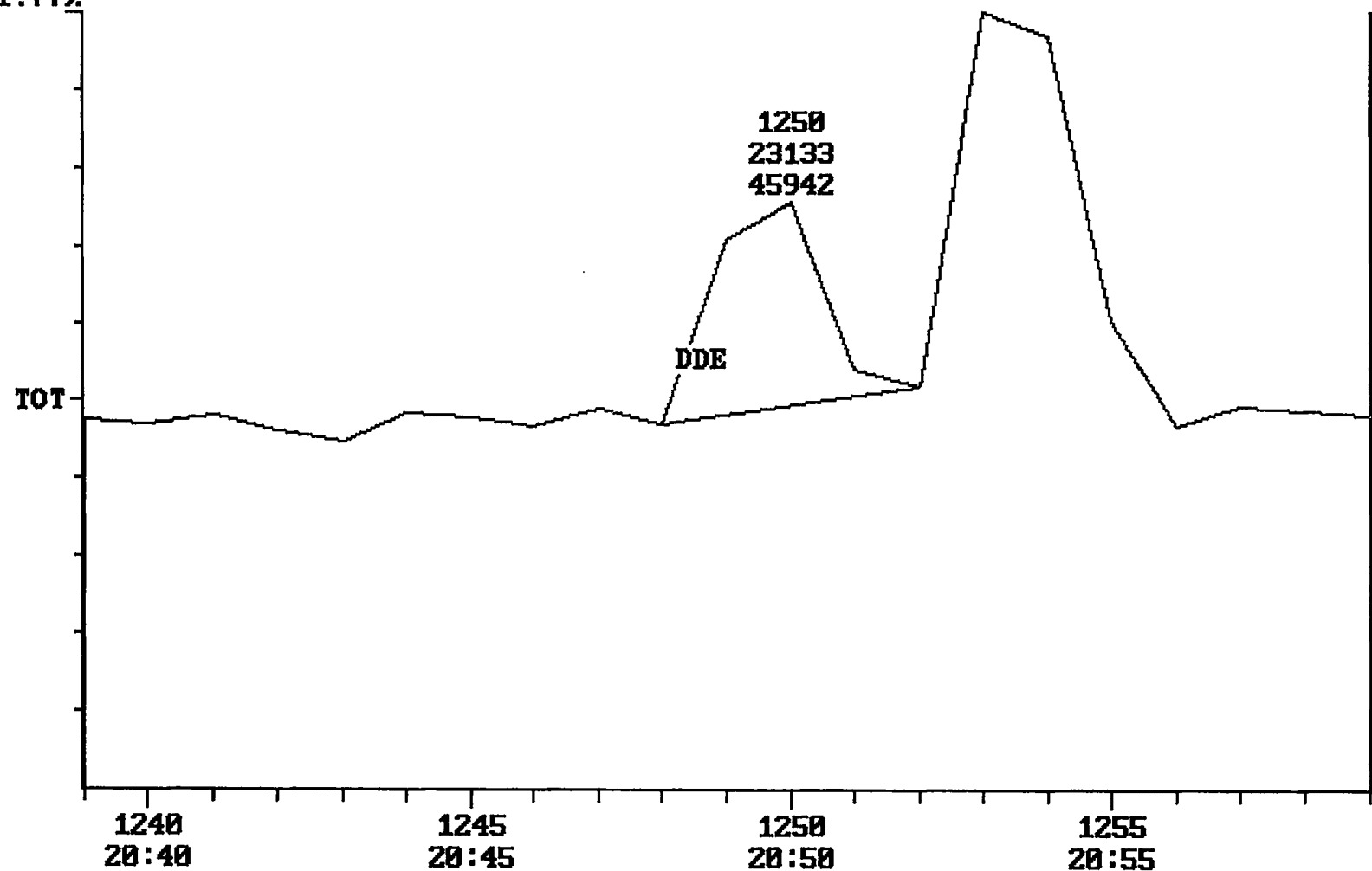
Mass Range: 47 - 207

Plotted: 1239 to 1259

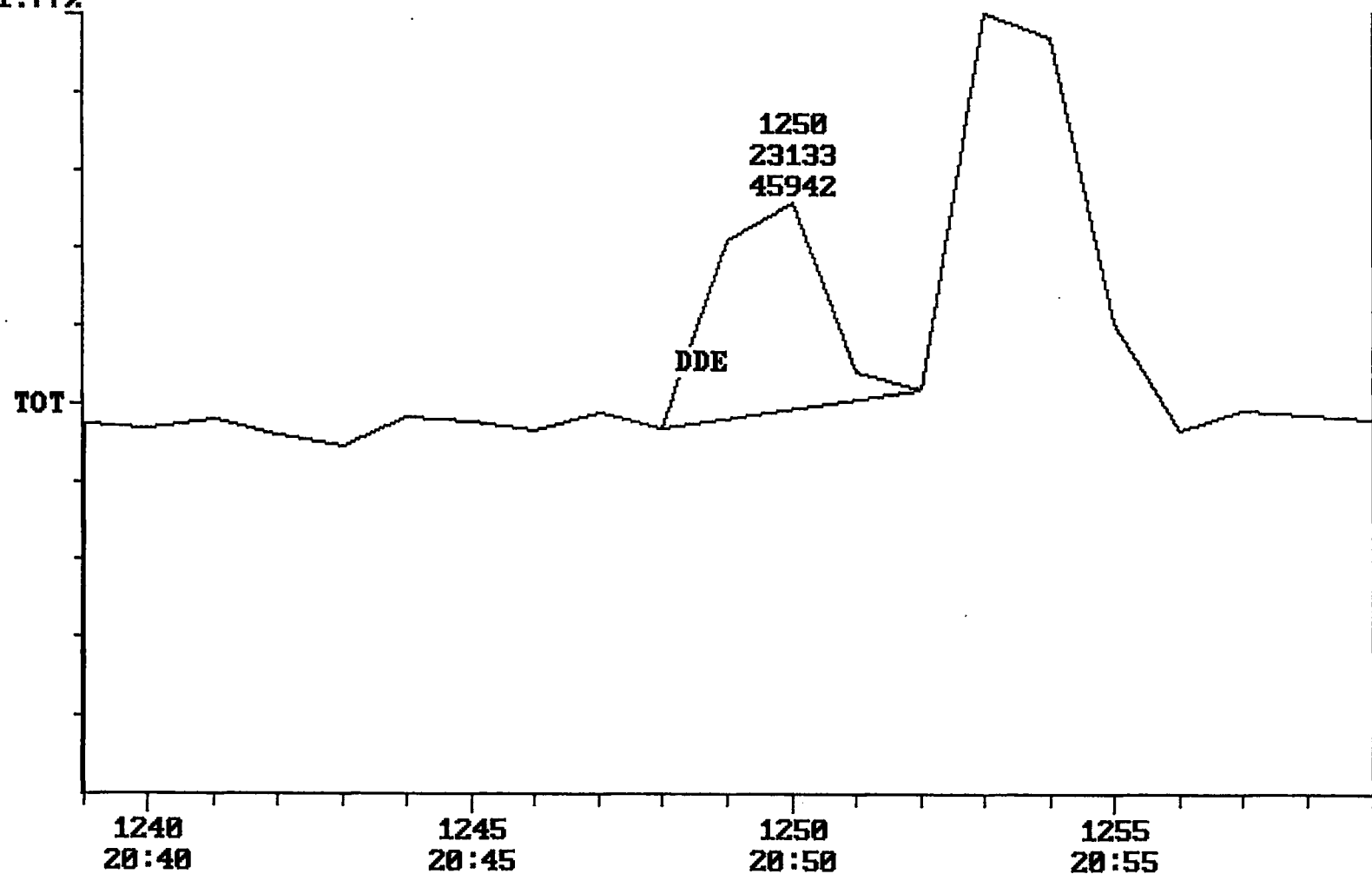
Range: 1 to 1860

100% = 4931445

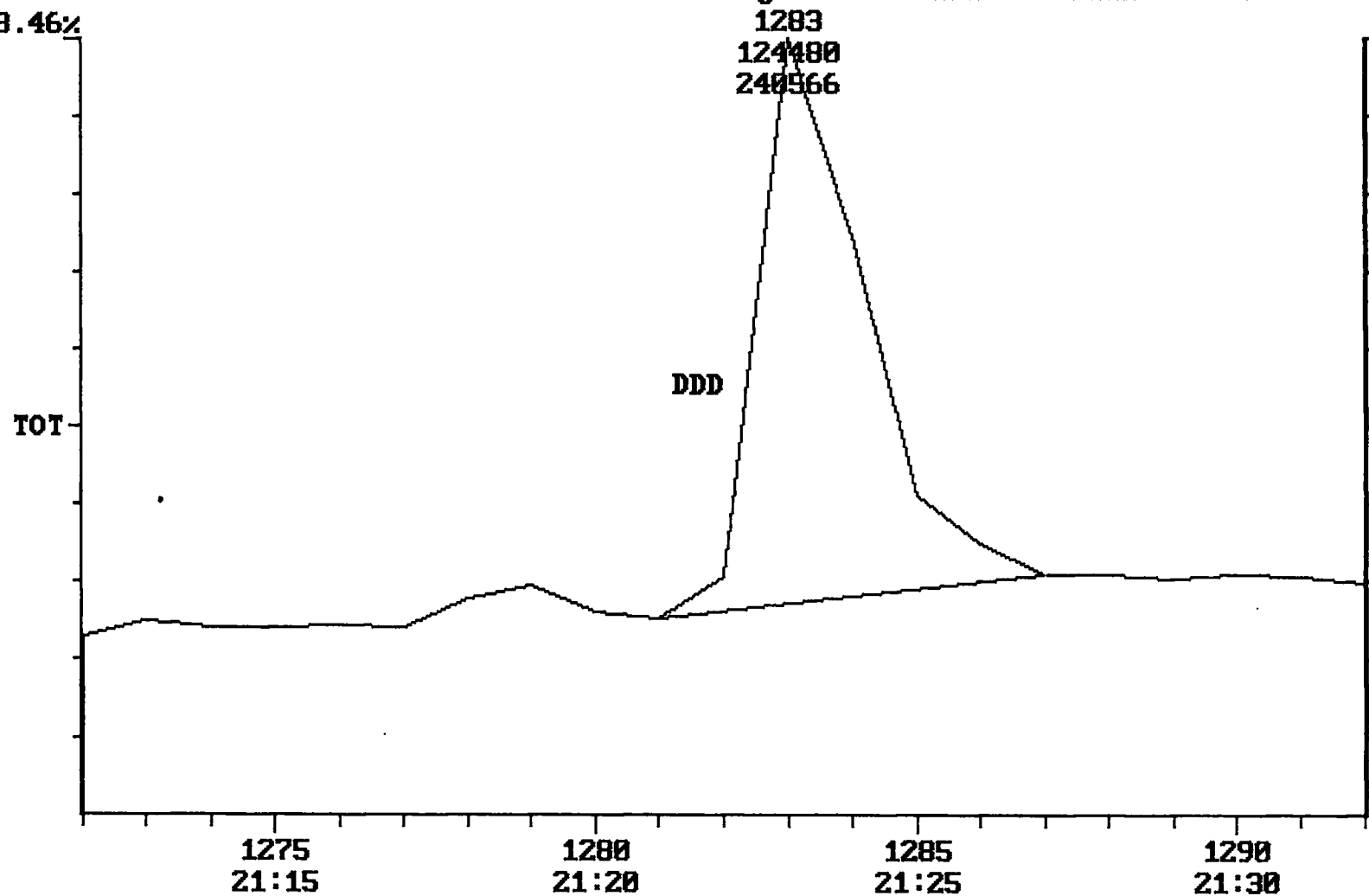
1.77%



Chromatogram Plot File: D:\DDTB1109 Date: Nov-10-1991 05:10:13
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DDT
Scan No: 1259 Retention Time: 20:59 RIC: 41882 Mass Range: 47 - 207
Plotted: 1239 to 1259 Range: 1 to 1860 100% = 4931445
1.77%



Chromatogram Plot File: D:\DDTB1109 Date: Nov-10-1991 05:10:13
Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DDT
Scan No: 1292 Retention Time: 21:32 RIC: 50691 Mass Range: 47 - 237
Plotted: 1272 to 1292 Range: 1 to 1860 100% = 4931445
3.46%



Chromatogram Plot

File: D:\DDTB1109

Date: Nov-10-1991 05:10:13

Comment: BOHLK; METHOD 525; INST 204; 11/09/01; DDT

Scan No: 1324

Retention Time: 22:04

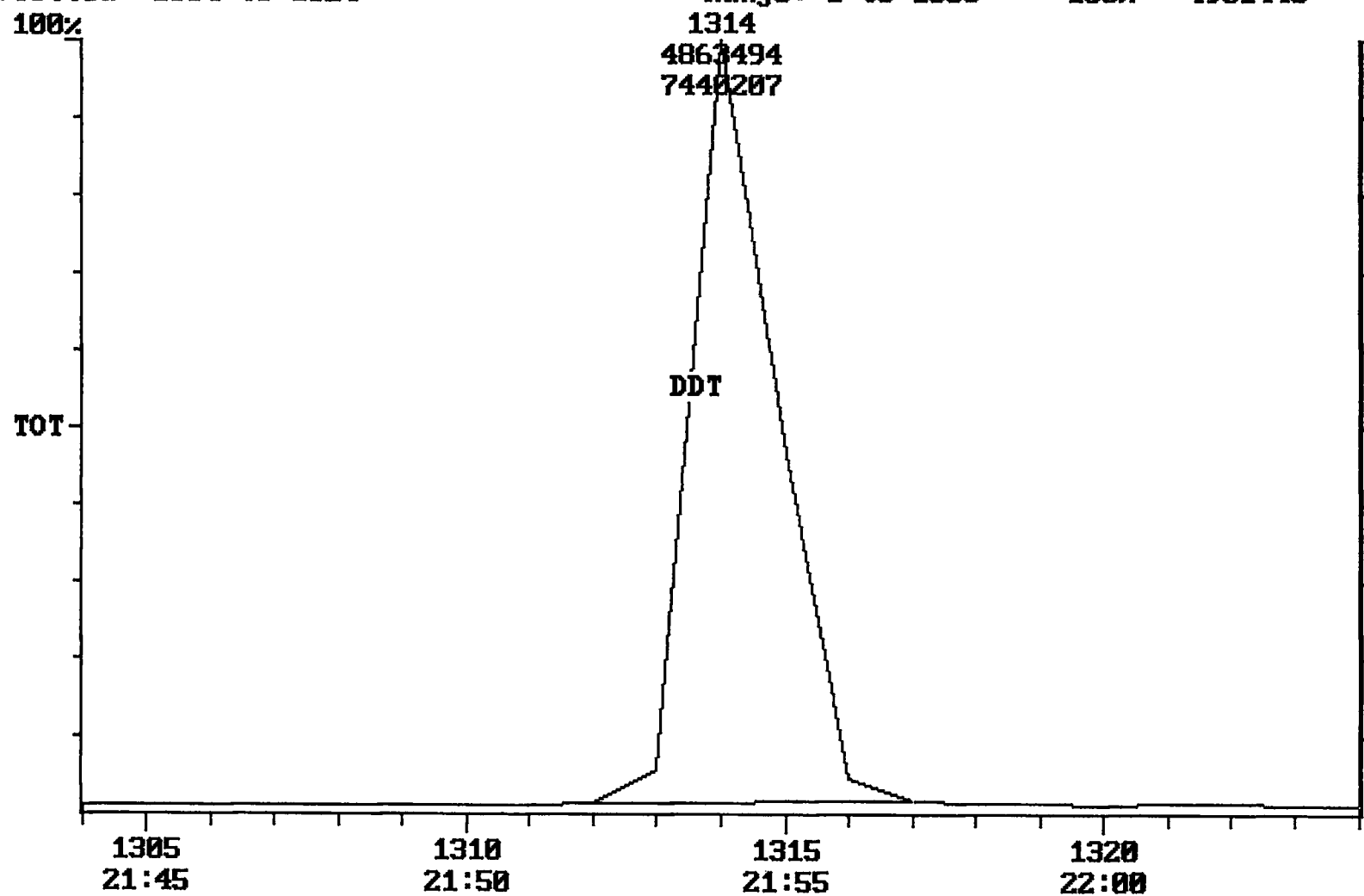
RIC: 50780

Mass Range: 47 - 281

Plotted: 1304 to 1324

Range: 1 to 1860

100% = 4931445

*Breakdown = 3.7 %*