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APPENDIX I

Table 21 Water quality of the rearing units at different stages of larvae rearing*

Larvae stage	salinity (ppt)	Temperature ($^{\circ}$ C)	pH	DO (ppm)	Ammonium (ppm)	Nitrate (ppm)
zoaea	25, 30	28-28.5	7.8-8.0	5.1-5.4	0.5-1.0	0-10
mysis	25, 30	27-28	7.7-8.0	5.2-5.4	0.5-1.0	10-25
postlarva	25, 30	26-28	7.7-8.2	5.1-5.4	0.5-1.0	10-25

* All parameters tested at Marine Research Unit. Ammonium and nitrate were assessed by test kits.

Table 22 Water quality for shrimp living

Water quality	Approximate range	Source
Temperature ($^{\circ}$ C)	25-30	Boyd and Tucker (1992)
	25-30	กรมประมง (2536)
Salinity (ppt)	23-25	Cheng and Laio (1986)
	15-25	กรมประมง (2534)
	15-30	Boyd and Tucker (1992)
pH	7-9	Boyd and Tucker (1992)
	7.5-8.5	ชลอ ลีสมสุวรรณ (2534)
	7.5-8.5	กรมประมง (2534)
DO (ppm)	\geq 3.5-อิมตัว	Boyd and Tucker (1992)
	5-7.5	กรมประมง (2534)
Ammonium (ppm)	0.4-2.0	Boyd and Tucker (1992)
	0.4-2.0	กรมประมง (2534)
Nitrate (ppm)	non toxic	Wetzel (1975)

APPENDIX II

STATISTICAL ANALYSIS

Raw data of length of post larva 15

OBS	TREAT	REP	SALINITY	LENGTH
1	1	1	25	12.0
2	1	1	25	11.8
3	1	1	25	9.9
4	1	1	25	13.6
5	1	1	25	11.4
6	1	1	25	12.1
7	1	1	25	10.8
8	1	1	25	11.5
9	1	1	25	11.0
10	1	1	25	11.4
11	1	1	25	10.9
12	1	1	25	12.3
13	1	1	25	11.4
14	1	1	25	10.3
15	1	1	25	11.2
16	1	1	25	11.8
17	1	1	25	11.1
18	1	1	25	10.2
19	1	1	25	12.2
20	1	1	25	8.5
21	1	2	25	12.1
22	1	2	25	12.2
23	1	2	25	10.1
24	1	2	25	9.6
25	1	2	25	11.6
26	1	2	25	8.5
27	1	2	25	11.7
28	1	2	25	12.1
29	1	2	25	12.8
30	1	2	25	11.7
31	1	2	25	10.2
32	1	2	25	10.2
33	1	2	25	12.2
34	1	2	25	14.6
35	1	2	25	11.5
36	1	2	25	10.2
37	1	2	25	12.9
38	1	3	25	12.4
39	1	3	25	10.7
40	1	3	25	11.7
41	1	3	25	10.3
42	1	3	25	12.7
43	1	3	25	10.4
44	1	3	25	11.7
45	1	3	25	12.0
46	1	3	25	8.4
47	1	3	25	14.1
48	1	3	25	12.1
49	1	3	25	12.2
50	1	3	25	11.4
51	1	3	25	10.5
52	1	3	25	10.0

53	1	3	25	11.6
54	1	3	25	10.8
55	1	3	25	12.3
56	1	3	25	11.5
57	1	3	25	14.5
58	1	1	30	10.1
59	1	1	30	11.6
60	1	1	30	12.9
61	1	1	30	12.5
62	1	1	30	9.7
63	1	1	30	12.0
64	1	1	30	12.7
65	1	1	30	11.3
66	1	1	30	10.2
67	1	1	30	11.1
68	1	1	30	11.5
69	1	1	30	11.9
70	1	1	30	11.2
71	1	1	30	11.6
72	1	1	30	11.1
73	1	1	30	11.9
74	1	1	30	10.9
75	1	1	30	12.5
76	1	1	30	10.6
77	1	1	30	11.6
78	1	2	30	11.6
79	1	2	30	10.4
80	1	2	30	10.8
81	1	2	30	12.1
82	1	2	30	11.2
83	1	2	30	10.7
84	1	2	30	11.1
85	1	2	30	10.5
86	1	2	30	11.7
87	1	2	30	10.5
88	1	2	30	10.7
89	1	2	30	11.2
90	1	2	30	12.6
91	1	2	30	9.5
92	1	2	30	10.2
93	1	2	30	12.1
94	1	2	30	11.3
95	1	2	30	10.7
96	1	2	30	11.5
97	1	2	30	10.7
98	1	3	30	12.1
99	1	3	30	9.6
100	1	3	30	9.8
101	1	3	30	9.3
102	1	3	30	9.9
103	1	3	30	10.4
104	1	3	30	11.4
105	1	3	30	9.9
106	1	3	30	11.1
107	1	3	30	11.0
108	1	3	30	8.8
109	1	3	30	8.9
110	1	3	30	10.8
111	1	3	30	12.0
112	1	3	30	11.5
113	1	3	30	11.1
114	1	3	30	9.0
115	1	3	30	11.6
116	1	3	30	12.7
117	1	3	30	11.6
118	2	1	25	10.5
119	2	1	25	10.9
120	2	1	25	11.0
121	2	1	25	11.2
122	2	1	25	9.7
123	2	1	25	10.1
124	2	1	25	11.6
125	2	1	25	8.9
126	2	1	25	9.8
127	2	1	25	10.3
128	2	1	25	10.8

129	2	1	25	11.3
130	2	1	25	11.2
131	2	1	25	11.5
132	2	1	25	9.6
133	2	1	25	10.6
134	2	1	25	11.5
135	2	1	25	12.3
136	2	2	25	10.9
137	2	2	25	10.8
138	2	2	25	12.1
139	2	2	25	12.1
140	2	2	25	11.5
141	2	2	25	12.5
142	2	2	25	11.7
143	2	2	25	11.2
144	2	2	25	12.4
145	2	2	25	10.9
146	2	2	25	11.9
147	2	2	25	11.9
148	2	2	25	13.2
149	2	2	25	10.9
150	2	3	25	0.0
151	2	1	30	11.1
152	2	1	30	10.4
153	2	1	30	10.1
154	2	1	30	9.9
155	2	1	30	11.5
156	2	1	30	9.7
157	2	1	30	9.1
158	2	1	30	11.2
159	2	1	30	8.6
160	2	1	30	11.3
161	2	1	30	11.6
162	2	1	30	11.4
163	2	1	30	12.2
164	2	1	30	10.2
165	2	1	30	11.2
166	2	1	30	12.3
167	2	1	30	11.5
168	2	1	30	10.2
169	2	1	30	10.7
170	2	1	30	11.9
171	2	2	30	10.7
172	2	2	30	10.4
173	2	2	30	12.7
174	2	2	30	10.5
175	2	2	30	10.5
176	2	2	30	10.8
177	2	2	30	11.5
178	2	2	30	11.3
179	2	2	30	11.4
180	2	2	30	10.7
181	2	2	30	10.6
182	2	2	30	12.9
183	2	2	30	10.4
184	2	2	30	10.2
185	2	2	30	10.6
186	2	2	30	11.3
187	2	2	30	11.0
188	2	2	30	9.6
189	2	2	30	11.1
190	2	2	30	11.2
191	2	3	30	10.1
192	2	3	30	10.9
193	2	3	30	10.9
194	2	3	30	11.2
195	2	3	30	10.6
196	2	3	30	10.4
197	2	3	30	10.9
198	2	3	30	11.3
199	2	3	30	9.3
200	2	3	30	11.2
201	2	3	30	11.6
202	2	3	30	11.2
203	2	3	30	10.5
204	2	3	30	10.8

205	2	3	30	11.4
206	2	3	30	11.2
207	2	3	30	11.7
208	2	3	30	9.7
209	2	3	30	9.8
210	2	3	30	11.5
211	3	1	25	11.1
212	3	1	25	11.7
213	3	1	25	8.9
214	3	1	25	11.2
215	3	1	25	12.4
216	3	1	25	11.3
217	3	1	25	12.0
218	3	1	25	10.3
219	3	1	25	10.7
220	3	1	25	10.4
221	3	1	25	10.3
222	3	1	25	11.0
223	3	1	25	11.2
224	3	1	25	12.7
225	3	1	25	11.8
226	3	1	25	10.9
227	3	2	25	10.9
228	3	2	25	11.6
229	3	2	25	11.6
230	3	2	25	9.9
231	3	2	25	12.4
232	3	2	25	11.2
233	3	2	25	11.5
234	3	2	25	10.2
235	3	2	25	11.3
236	3	2	25	9.8
237	3	2	25	12.0
238	3	2	25	11.1
239	3	2	25	10.8
240	3	2	25	11.2
241	3	2	25	10.2
242	3	2	25	10.1
243	3	2	25	9.7
244	3	2	25	9.4
245	3	2	25	11.1
246	3	2	25	11.3
247	3	3	25	9.9
248	3	3	25	10.0
249	3	3	25	10.5
250	3	3	25	10.2
251	3	3	25	11.4
252	3	3	25	8.6
253	3	3	25	11.8
254	3	3	25	9.2
255	3	3	25	10.3
256	3	3	25	8.6
257	3	3	25	8.0
258	3	3	25	9.7
259	3	3	25	8.9
260	3	3	25	8.6
261	3	3	25	9.9
262	3	3	25	12.0
263	3	1	30	12.1
264	3	1	30	11.2
265	3	1	30	10.7
266	3	1	30	11.3
267	3	1	30	10.9
268	3	1	30	11.7
269	3	1	30	10.8
270	3	1	30	11.8
271	3	1	30	12.3
272	3	1	30	10.9
273	3	1	30	11.6
274	3	1	30	12.5
275	3	1	30	11.1
276	3	1	30	13.3
277	3	2	30	10.1
278	3	2	30	10.7
279	3	2	30	11.9
280	3	2	30	11.3

281	3	2	30	8.7
282	3	2	30	11.2
283	3	2	30	10.4
284	3	2	30	10.5
285	3	2	30	10.7
286	3	2	30	11.1
287	3	2	30	10.9
288	3	2	30	11.2
289	3	2	30	10.7
290	3	2	30	10.5
291	3	2	30	10.7
292	3	2	30	10.8
293	3	3	30	11.2
294	3	3	30	12.0
295	3	3	30	11.6
296	3	3	30	11.5
297	3	3	30	11.5
298	3	3	30	12.9
299	3	3	30	10.7
300	3	3	30	10.7
301	3	3	30	11.9
302	3	3	30	10.8
303	3	3	30	11.1
304	3	3	30	11.0
305	3	3	30	10.6
306	3	3	30	10.2
307	3	3	30	12.0
308	3	3	30	10.6
309	3	3	30	11.6
310	3	3	30	10.3
311	3	3	30	11.2
312	3	3	30	10.9
313	4	1	25	10.5
314	4	1	25	11.2
315	4	1	25	9.1
316	4	1	25	11.3
317	4	1	25	9.8
318	4	1	25	10.3
319	4	1	25	10.6
320	4	1	25	10.7
321	4	1	25	11.3
322	4	1	25	10.5
323	4	1	25	11.1
324	4	1	25	10.1
325	4	1	25	10.8
326	4	2	25	11.0
327	4	2	25	10.3
328	4	2	25	9.7
329	4	2	25	12.1
330	4	2	25	12.1
331	4	2	25	11.9
332	4	2	25	10.6
333	4	2	25	10.9
334	4	2	25	11.8
335	4	2	25	9.6
336	4	2	25	10.5
337	4	2	25	10.4
338	4	2	25	11.4
339	4	2	25	10.7
340	4	2	25	10.7
341	4	2	25	10.9
342	4	2	25	12.6
343	4	2	25	10.9
344	4	2	25	12.1
345	4	2	25	11.5
346	4	3	25	12.8
347	4	3	25	14.5
348	4	3	25	11.2
349	4	3	25	10.9
350	4	3	25	10.6
351	4	3	25	11.5
352	4	3	25	11.6
353	4	3	25	10.6
354	4	3	25	11.5
355	4	3	25	10.5
356	4	3	25	11.3

357	4	3	25	11.0
358	4	3	25	11.4
359	4	3	25	10.2
360	4	3	25	11.5
361	4	3	25	10.4
362	4	3	25	11.1
363	4	3	25	12.6
364	4	3	25	11.0
365	4	3	25	11.0
366	4	1	30	13.0
367	4	1	30	11.8
368	4	1	30	9.5
369	4	1	30	11.7
370	4	1	30	11.7
371	4	1	30	12.2
372	4	1	30	10.8
373	4	1	30	12.8
374	4	1	30	12.4
375	4	1	30	12.2
376	4	1	30	13.6
377	4	1	30	10.7
378	4	1	30	9.1
379	4	1	30	12.1
380	4	1	30	11.4
381	4	1	30	10.1
382	4	1	30	12.1
383	4	1	30	8.9
384	4	1	30	11.4
385	4	1	30	9.9
386	4	2	30	10.0
387	4	2	30	9.2
388	4	2	30	8.3
389	4	2	30	8.8
390	4	2	30	9.1
391	4	2	30	10.8
392	4	2	30	8.8
393	4	2	30	10.0
394	4	2	30	9.8
395	4	2	30	10.1
396	4	2	30	10.0
397	4	2	30	10.8
398	4	2	30	10.3
399	4	2	30	9.2
400	4	2	30	10.8
401	4	2	30	8.9
402	4	2	30	11.6
403	4	2	30	8.8
404	4	2	30	11.1
405	4	2	30	9.3
406	4	3	30	10.7
407	4	3	30	10.5
408	4	3	30	11.2
409	4	3	30	9.7
410	4	3	30	11.4
411	4	3	30	10.5
412	4	3	30	10.2
413	4	3	30	10.6
414	4	3	30	11.0
415	4	3	30	11.3
416	4	3	30	9.5
417	4	3	30	10.8
418	4	3	30	9.6
419	4	3	30	10.6
420	4	3	30	9.6

ANOVA of length of postlarva 15 fed different diet at salinity 25,30 ppt.

General Linear Models Procedure

Dependent Variable: LENGTH

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	7	31.02156985	4.43165284	3.33	0.0019
Error	412	548.98840634	1.33249613		
Corrected Total	419	580.00997619			
	R-Square	C.V.	Root MSE	LENGTH Mean	
	0.053485	10.54167	1.154338	10.9502381	

Source	DF	Type I SS	Mean Square	F Value	Pr > F
TREAT	3	12.70960729	4.23653576	3.18	0.0240
SALINITY	1	1.02904656	1.02904656	0.77	0.3800
TREAT*SALINITY	3	17.28291600	5.76097200	4.32	0.0051

Source	DF	Type III SS	Mean Square	F Value	Pr > F
TREAT	3	12.84422681	4.28140894	3.21	0.0229
SALINITY	1	0.61158581	0.61158581	0.46	0.4985
TREAT*SALINITY	3	17.28291600	5.76097200	4.32	0.0051

General Linear Models Procedure

Dependent Variable: LENGTH

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	3	12.70960729	4.23653576	3.11	0.0264
Error	416	567.30036890	1.36370281		
Corrected Total	419	580.00997619			
	R-Square	C.V.	Root MSE	LENGTH Mean	
	0.021913	10.66440	1.167777	10.9502381	

Source	DF	Type I SS	Mean Square	F Value	Pr > F
TREAT	3	12.70960729	4.23653576	3.11	0.0264

Source	DF	Type III SS	Mean Square	F Value	Pr > F
TREAT	3	12.70960729	4.23653576	3.11	0.0264

Duncan's Multiple Range Test for variable: LENGTH
 Alpha= 0.05 df= 416 MSE= 1.363703
 Number of Means 2 3 4
 Critical Range 0.321 0.338 0.349
 Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	TREAT
A	11.226	117	1
B	10.889	102	3
B	10.844	93	2
B	10.801	108	4

General Linear Models Procedure

Dependent Variable: LENGTH

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	1	1.24484115	1.24484115	0.90	0.3436
Error	418	578.76513504	1.38460559		
Corrected Total	419	580.00997619			
	R-Square	C.V.	Root MSE	LENGTH Mean	
	0.002146	10.74582	1.176693	10.9502381	

Source	DF	Type I SS	Mean Square	F Value	Pr > F
SALINITY	1	1.24484115	1.24484115	0.90	0.3436

Source	DF	Type III SS	Mean Square	F Value	Pr > F
SALINITY	1	1.24484115	1.24484115	0.90	0.3436

Duncan's Multiple Range Test for variable: LENGTH
 Alpha= 0.05 df= 418 MSE= 1.384606
 Number of Means 2
 Critical Range 0.229
 Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	SALINITY
A	11.009	195	25
A	10.900	225	30

----- SALINITY=25 -----

General Linear Models Procedure
 Dependent Variable: LENGTH

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	3	18.45369436	6.15123145	3.42	0.0184
Error	191	343.74148513	1.79969364		
Corrected Total	194	362.19517949			

R-Square 0.050950 C.V. 12.18604 Root MSE 1.341527 LENGTH Mean 11.0087179

Source	DF	Type I SS	Mean Square	F Value	Pr > F
TREAT	3	18.45369436	6.15123145	3.42	0.0184
Source	DF	Type III SS	Mean Square	F Value	Pr > F
TREAT	3	18.45369436	6.15123145	3.42	0.0184

Duncan's Multiple Range Test for variable: LENGTH
 Alpha= 0.05 df= 191 MSE= 1.799694
 Number of Means 2 3 4
 Critical Range 0.552 0.581 0.599

Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	TREAT
A	11.419	57	1
B A	11.060	53	4
B	10.812	33	2
B	10.631	52	3

----- SALINITY=30 -----

General Linear Models Procedure
 Dependent Variable: LENGTH

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	3	11.32303434	3.77434478	4.06	0.0078
Error	221	205.24692121	0.92871910		
Corrected Total	224	216.56995556			

R-Square 0.052283 C.V. 8.841652 Root MSE 0.963701 LENGTH Mean 10.8995556

Source	DF	Type I SS	Mean Square	F Value	Pr > F
TREAT	3	11.32303434	3.77434478	4.06	0.0078
Source	DF	Type III SS	Mean Square	F Value	Pr > F
TREAT	3	11.32303434	3.77434478	4.06	0.0078

Duncan's Multiple Range Test for variable: LENGTH
 Alpha= 0.05 df= 221 MSE= 0.928719
 Number of Means 2 3 4
 Critical Range 0.365 0.381 0.393

Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	TREAT
A	11.158	50	3
A	11.042	60	1
B A	10.862	60	2
B	10.551	55	4

----- TREAT=1 -----

General Linear Models Procedure
 Dependent Variable: LENGTH

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	1	4.16847166	4.16847166	3.18	0.0770
Error	115	150.53460526	1.30899657		
Corrected Total	116	154.70307692			

R-Square 0.026945 C.V. 10.19197 Root MSE 1.144114 LENGTH Mean 11.2256410

Source	DF	Type I SS	Mean Square	F Value	Pr > F
SALINITY	1	4.16847166	4.16847166	3.18	0.0770
Source	DF	Type III SS	Mean Square	F Value	Pr > F
SALINITY	1	4.16847166	4.16847166	3.18	0.0770

Duncan's Multiple Range Test for variable: LENGTH
 Alpha= 0.05 df= 115 MSE= 1.308997
 Number of Means 2
 Critical Range 0.420

Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	SALINITY
A	11.419	57	25
A	11.042	60	30

----- TREAT=2 -----

General Linear Models Procedure

Dependent Variable: LENGTH

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	1	0.05226246	0.05226246	0.03	0.8737
Error	91	187.23698485	2.05754928		
Corrected Total	92	187.28924731			

	R-Square	C.V.	Root MSE	LENGTH Mean
	0.000279	13.22763	1.434416	10.8440860

Source	DF	Type I SS	Mean Square	F Value	Pr > F
SALINITY	1	0.05226246	0.05226246	0.03	0.8737

Source	DF	Type III SS	Mean Square	F Value	Pr > F
SALINITY	1	0.05226246	0.05226246	0.03	0.8737

Duncan's Multiple Range Test for variable: LENGTH

Alpha= 0.05 df= 91 MSE= 2.057549

Number of Means 2

Critical Range 0.618

Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	SALINITY
A	10.862	60	30
A	10.812	33	25

----- TREAT=3 -----

General Linear Models Procedure

Dependent Variable: LENGTH

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	1	7.08556802	7.08556802	7.86	0.0061
Error	100	90.13256923	0.90132569		
Corrected Total	101	97.21813725			

	R-Square	C.V.	Root MSE	LENGTH Mean
	0.072883	8.718550	0.949382	10.8892157

Dependent Variable: LENGTH

Source	DF	Type I SS	Mean Square	F Value	Pr > F
SALINITY	1	7.08556802	7.08556802	7.86	0.0061

Source	DF	Type III SS	Mean Square	F Value	Pr > F
SALINITY	1	7.08556802	7.08556802	7.86	0.0061

Duncan's Multiple Range Test for variable: LENGTH

Alpha= 0.05 df= 100 MSE= 0.901326

Number of Means 2

Critical Range 0.374

Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	SALINITY
A	11.158	50	30
B	10.631	52	25

----- TREAT=4 -----

General Linear Models Procedure

Dependent Variable: LENGTH

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	1	7.00566041	7.00566041	6.13	0.0149
Error	106	121.08424700	1.14230422		
Corrected Total	107	128.08990741			

	R-Square	C.V.	Root MSE	LENGTH Mean
	0.054693	9.895321	1.068786	10.8009259

Source

DF	Type I SS	Mean Square	F Value	Pr > F
1	7.00566041	7.00566041	6.13	0.0149

Source

DF	Type III SS	Mean Square	F Value	Pr > F
1	7.00566041	7.00566041	6.13	0.0149

Duncan's Multiple Range Test for variable: LENGTH

Alpha= 0.05 df= 106 MSE= 1.142304

Number of Means 2

Critical Range 0.409

Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	SALINITY
A	11.060	53	25
B	10.551	55	30

Raw data of survival of zoea stage

OBS	STAGE	TREAT	REPL	SALINITY	SURVIVE	PSURVIVE
1	ml	1	1	25	.	.
2	ml	1	2	25	.	.
3	ml	1	3	25	136	45.3333
4	ml	1	1	30	161	53.6667
5	ml	1	2	30	219	73.0000
6	ml	1	3	30	256	85.3333
7	ml	2	1	25	19	6.3333
8	ml	2	2	25	166	55.3333
9	ml	2	3	25	44	14.6667
10	ml	2	1	30	119	39.6667
11	ml	2	2	30	146	48.6667
12	ml	2	3	30	177	59.0000
13	ml	3	1	25	15	5.0000
14	ml	3	2	25	165	55.0000
15	ml	3	3	25	36	12.0000
16	ml	3	1	30	192	64.0000
17	ml	3	2	30	221	73.6667
18	ml	3	3	30	178	59.3333
19	ml	4	1	25	81	27.0000
20	ml	4	2	25	87	29.0000
21	ml	4	3	25	157	52.3333
22	ml	4	1	30	208	69.3333
23	ml	4	2	30	90	30.0000
24	ml	4	3	30	116	38.6667

General Linear Models Procedure

Dependent Variable: PSURVIVE

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	9	6730.510101	747.834456	2.15	0.1088
Error	12	4181.777778	348.481481		
Corrected Total	21	10912.287879			
	R-Square	C.V.	Root MSE	PSURVIVE Mean	
	0.616783	41.21999	18.66766	45.2878788	
Source	DF	Type I SS	Mean Square	F Value	Pr > F
REPL	2	713.272006	356.636003	1.02	0.3887
TREAT	3	1954.751323	651.583774	1.87	0.1885
SALINITY	1	3300.192308	3300.192308	9.47	0.0096
TREAT*SALINITY	3	762.294465	254.098155	0.73	0.5541
Source	DF	Type III SS	Mean Square	F Value	Pr > F
REPL	2	713.555556	356.777778	1.02	0.3886
TREAT	3	841.774522	280.591507	0.81	0.5148
SALINITY	1	2985.587596	2985.587596	8.57	0.0127
TREAT*SALINITY	3	762.294465	254.098155	0.73	0.5541

Duncan's Multiple Range Test for variable: PSURVIVE

Alpha= 0.05 df= 12 MSE= 348.4815

Number of Means 2

Critical Range 17.38

Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	SALINITY
A	57.861	12	30
B	30.200	10	25

General Linear Models Procedure

Duncan's Multiple Range Test for variable: PSURVIVE

Alpha= 0.05 df= 12 MSE= 348.4815

Number of Means 2 3 4

Critical Range 24.86 26.04 26.83

Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	TREAT
A	64.33	4	1
B	44.83	6	3
B	41.06	6	4
B	37.28	6	2

Raw data of survival of mysis stage

OBS	STAGE	TREAT	REPL	SALINITY	SURVIVE
1	p1	1	1	25	46
2	p1	1	2	25	50
3	p1	1	3	25	74
4	p1	1	1	30	42
5	p1	1	2	30	58
6	p1	1	3	30	58
7	p1	2	1	25	88
8	p1	2	2	25	47
9	p1	2	3	25	67
10	p1	2	1	30	80
11	p1	2	2	30	59
12	p1	2	3	30	56
13	p1	3	1	25	65
14	p1	3	2	25	55
15	p1	3	3	25	68
16	p1	3	1	30	81
17	p1	3	2	30	84
18	p1	3	3	30	60
19	p1	4	1	25	82
20	p1	4	2	25	73
21	p1	4	3	25	52
22	p1	4	1	30	63
23	p1	4	2	30	58
24	p1	4	3	30	53

General Linear Models Procedure

Dependent Variable: SURVIVE

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	9	1433.375000	159.263889	0.91	0.5454
Error	14	2459.583333	175.684524		
Corrected Total	23	3892.958333			
	R-Square	C.V.	Root MSE	SURVIVE Mean	
	0.368197	20.94210	13.25460	63.2916667	
Source	DF	Type I SS	Mean Square	F Value	Pr > F
REPL	2	311.0833333	155.5416667	0.89	0.4345
TREAT	3	680.4583333	226.8194444	1.29	0.3162
SALINITY	1	9.3750000	9.3750000	0.05	0.8207
TREAT*SALINITY	3	432.4583333	144.1527778	0.82	0.5039
Source	DF	Type III SS	Mean Square	F Value	Pr > F
REPL	2	311.0833333	155.5416667	0.89	0.4345
TREAT	3	680.4583333	226.8194444	1.29	0.3162
SALINITY	1	9.3750000	9.3750000	0.05	0.8207
TREAT*SALINITY	3	432.4583333	144.1527778	0.82	0.5039

Duncan's Multiple Range Test for variable: SURVIVE

Alpha= 0.05 df= 14 MSE= 175.6845

Number of Means 2
Critical Range 11.58

Duncan Grouping	Mean	N	SALINITY
A	63.917	12	25
A	62.667	12	30

Duncan's Multiple Range Test for variable: SURVIVE

Alpha= 0.05 df= 14 MSE= 175.6845

Number of Means 2 3 4
Critical Range 16.38 17.18 17.72

Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	TREAT
A	68.833	6	3
A	66.167	6	2
A	63.500	6	4
A	54.667	6	1

Raw data of survival of postlarva stage

OBS	STAGE	TREAT	REPL	SALINITY	SURVIVE	PSURVIVE
1	P15	1	1	25	31	62
2	P15	1	2	25	17	34
3	P15	1	3	25	40	80
4	P15	1	1	30	23	46
5	P15	1	2	30	22	44
6	P15	1	3	30	45	90
7	P15	2	1	25	18	36
8	P15	2	2	25	14	28
9	P15	2	3	25	.	.
10	P15	2	1	30	24	48
11	P15	2	2	30	26	52
12	P15	2	3	30	33	66
13	P15	3	1	25	18	36
14	P15	3	2	25	25	50
15	P15	3	3	25	16	32
16	P15	3	1	30	14	28
17	P15	3	2	30	16	32
18	P15	3	3	30	36	72
19	P15	4	1	25	13	26
20	P15	4	2	25	24	48
21	P15	4	3	25	21	42
22	P15	4	1	30	23	46
23	P15	4	2	30	26	52
24	P15	4	3	30	15	30

General Linear Models Procedure

Dependent Variable: PSURVIVE

Source	DF	Sum of Squares	Mean Square	F Value	Pr > F
Model	9	3203.004141	355.889349	1.38	0.2901
Error	13	3355.952381	258.150183		
Corrected Total	22	6558.956522			
	R-Square	C.V.	Root MSE	PSURVIVE Mean	
	0.488341	34.21687	16.06705	46.9565217	
Source	DF	Type I SS	Mean Square	F Value	Pr > F
REPL	2	1434.099379	717.049689	2.78	0.0990
TREAT	3	1327.560847	442.520282	1.71	0.2132
SALINITY	1	230.119826	230.119826	0.89	0.3623
TREAT*SALINITY	3	211.224090	70.408030	0.27	0.8440
Source	DF	Type III SS	Mean Square	F Value	Pr > F
REPL	2	1161.380952	580.690476	2.25	0.14
TREAT	3	1326.361345	442.120448	1.71	0.2135
SALINITY	1	276.269841	276.269841	1.07	0.3198
TREAT*SALINITY	3	211.224090	70.408030	0.27	0.8440

Duncan's Multiple Range Test for variable: PSURVIVE

Alpha= 0.05 df= 13 MSE= 258.1502

Number of Means 2

Critical Range 14.46

Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	SALINITY
A	50.500	12	30
A	43.091	11	25

Duncan's Multiple Range Test for variable: PSURVIVE

Alpha= 0.05 df= 13 MSE= 258.1502

Number of Means 2 3 4

Critical Range 20.50 21.48 22.15

Means with the same letter are not significantly different.

Duncan Grouping	Mean	N	TREAT
A	59.333	6	1
A	46.000	5	2
A	41.667	6	3
A	40.667	6	4

APPENDIX III

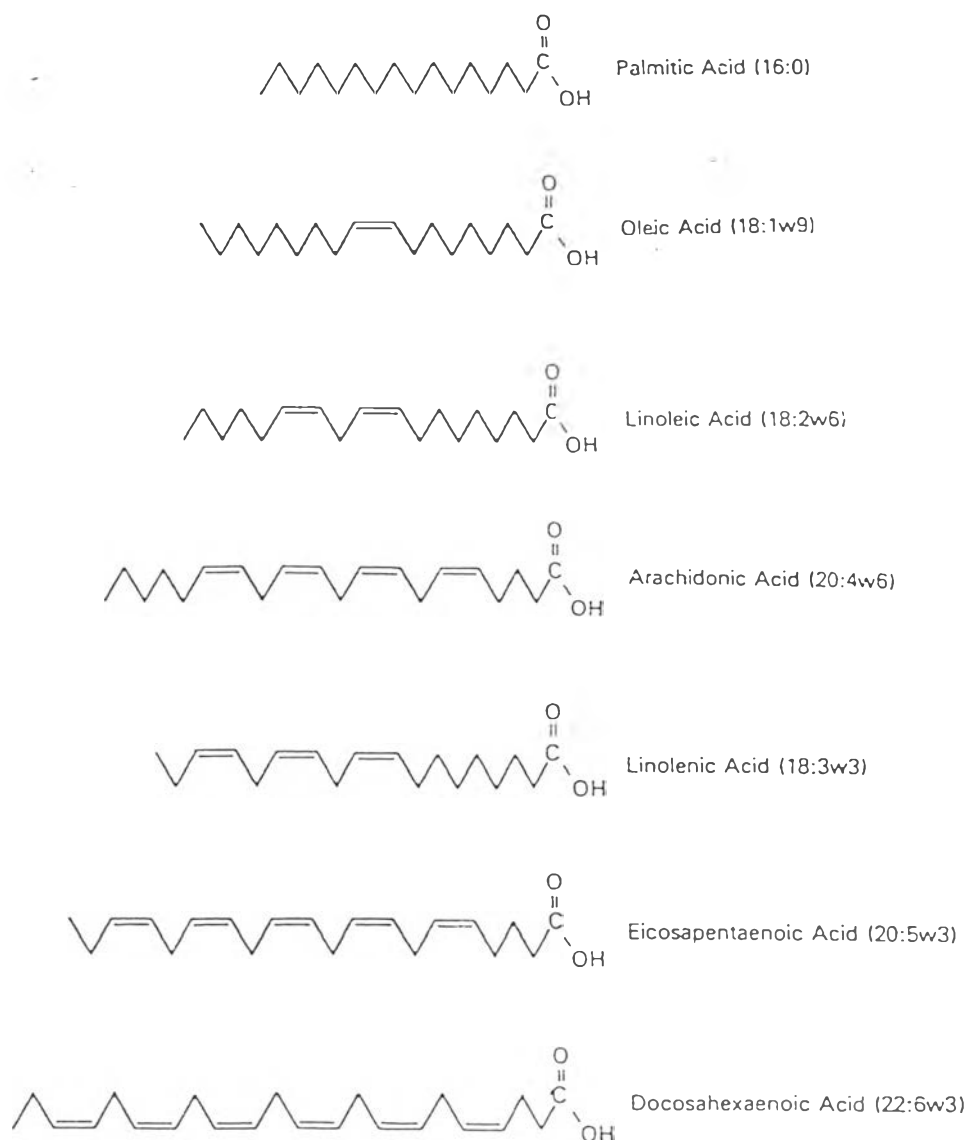


Figure 12 Structural formulas for omega6 (linoleic acid, 18:2 ω 6) and omega3 (alpha-linolenic acid, 18:3 ω 3) fatty acids. The first number (before the colon) give the number of carbon atoms in the molecule and the second give the number of double bonds. ω 6 indicate position of the first double bond in a given fatty acid molecule.

Table 23 Fatty acid composition of food fats (g fatty acid per 100g total fatty acid).

	C _{8:0}	C _{10:0}	C _{12:0}	C _{14:0}	C _{16:0}	C _{18:0}	C _{18:1}	C _{18:2}	C _{19:0}	C _{20:0}	C _{20:1}	C _{22:0}	C _{22:1}	C _{24:0}	C _{24:1}	C _{26:0}	C _{26:1}	C _{28:0}	C _{28:1}	C _{30:0}	C _{30:1}	C _{32:0}	C _{32:1}	C _{34:0}	C _{34:1}	C _{36:0}	C _{36:1}	References	
	Caprylic	Capric	Lauric	Myristic	Pentadecylic	Palmitic	Palmitoleic	Hexadecadienoic	Margaric	Stearic	Oleic	Linoleic	Linolonic	Stearidonic	Arachidic	Eicosenoic	Eicosadienoic	Arachidonic	Tinnodonic	Behenic	Erucic ^a Cetoleic ^b	Docosatetraenoic	Docosapentaenoic	Clupanodonic	Lignocenic	Sclacholeic			
<i>Plant lipids</i>																													
Almond oil	0	0	0	trace	-	7.0	0.6	-	0.1	1.2	67.3	23.7	0.2	-	0	-	-	-	-	0	0	-	-	-	-	-	0	-	21
Cocoa butter	-	0	trace	0.1	-	27.6	0.4	-	trace	33.9	34.0	3.1	trace	-	0.9	-	-	-	-	0	0	-	-	-	-	-	-	-	21
Coconut oil	10.1	6.6	45.8	18.4	-	8.0	trace	-	-	2.4	6.0	1.7	0	-	-	-	-	-	-	0	0	-	-	-	-	-	-	-	21
Corn oil	0	0	0	trace	-	11.5	0.1	-	trace	2.0	28.3	56.3	1.0	-	0.6	-	-	-	-	-	0	0	-	-	-	-	-	-	21
Cottonseed oil	0	trace	0.1	0.6	-	22.6	0.5	-	trace	2.6	18.1	52.3	1.7	-	0.5	-	-	-	-	0.2	0.7	-	-	-	-	-	-	-	21
Oliveseed oil	-	0	trace	0.04	-	6.7	0.1	-	-	3.1	12.1	77.3	0.7	-	0	-	-	-	-	0	-	-	-	-	-	-	-	-	21
Linseed oil	-	0	trace	trace	-	6.0	0.1	-	trace	5.1	21.1	14.9	53.0	-	trace	-	-	-	-	trace	0	-	-	-	-	-	-	-	22
Margarine**	2.2-5.0	1.9-3.7	14.2-29.3	6.1-11.4	-	9.4-22.1	-	-	-	3.3-8.0	14.3-38.1	10.3-29.3	0-2.5	-	-	-	-	-	-	0.4	44.4	-	-	-	-	-	0.3	2.7	21
Mustard-seed oil	0	0	trace	0.05	-	2.9	0.2	-	-	1.0	19.4	9.7	21.4	-	0.7	-	-	-	-	-	-	-	-	-	-	-	-	-	24
Olive oil, European†	0	0	0	trace	-	11.8	0.9	-	trace	2.8	74.5	8.7	1.0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	21
-, Tunisian	-	0	0	0	-	16.8	1.9	-	trace	2.5	58.9	18.8	0.9	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	21
Palm-kernel oil	4.9	3.9	48.7	16.0	-	7.6	0	-	-	1.9	14.2	2.6	0	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	21
Palm oil	-	-	trace	1.5	-	45.1	trace	-	trace	4.8	36.8	10.2	0.5	-	0.5	-	0	-	-	trace	0.2	-	-	-	-	-	-	-	21
Peanut oil	-	0	trace	trace	-	10.6	0.1	-	0.1	3.5	49.5	28.8	1.3	-	1.8	-	-	-	-	3.1	0.1	-	-	-	-	-	-	1.6	21
Rapeseed oil	0	0	0	0.05	-	3.6	0.2	-	trace	1.4	15.5	13.9	19.0	-	0.8	7.5	0.5	-	-	0.5	45.5†	-	-	-	-	-	0.5	-	21
Safflower oil	0	0	trace	0.1	-	7.5	0.1	-	trace	2.6	12.7	75.5	0.8	-	0.7	-	-	-	-	0.2	0.1	-	-	-	-	-	-	-	21
Sesame oil	-	0	trace	trace	-	8.9	0.3	-	trace	6.0	40.7	41.7	1.7	-	0.8	-	-	-	-	0	0	-	-	-	-	-	-	-	21
Soybean oil	0	0	0	0.1	-	10.7	0.1	-	0.1	3.8	23.0	52.4	8.9	-	0.6	-	-	-	-	0.4	trace	-	-	-	-	-	-	-	21
Sunflower-seed oil	-	0	trace	0.1	-	6.6	0.1	-	trace	4.3	22.4	65.2	0.3	-	0.4	-	-	-	-	0.7	0	-	-	-	-	-	-	0	21
Walnut oil	-	0	0	0.02	-	7.0	0.1	-	trace	2.1	18.7	59.8	13.3	-	trace	-	-	-	-	0	0	-	-	-	-	-	-	-	21
<i>Animal lipids</i>																													
Cow's milk	1.0	2.5	2.9	10.1	-	24.7	3.3	-	-	9.5	22.5	4.0	2.8	-	trace	-	-	-	-	-	-	-	-	-	-	-	-	-	23
Eggs	-	-	-	0.6	-	25.5	4.5	-	-	9.9	45.4	13.0	1.7	-	-	-	-	-	-	-	-	-	-	-	-	-	-	-	23
<i>Fish</i>																													
Cod	-	-	-	1.8	-	0.5	33.4	2.4	0.9	4.0	11.8	1.2	0.8	1.2	-	1.6	-	3.2	12.4	-	0.7	-	0.6	21.9	-	-	-	-	23
Cod-liver oil	-	-	-	2.8	-	0.4	10.7	6.9	1.2	3.7	23.9	1.5	0.9	2.6	-	8.8	0.5	1.0	8.0	-	5.3	1.3	14.3	-	-	-	-	0.5	23
Eel	-	-	-	3.7	0.1	0.2	12.8	11.8	0.9	3.2	25.7	1.0	0.4	1.6	trace	6.7	0.2	0.2	12.7	trace	3.8	0.2	1.5	6.9	trace	1.0	-	-	23
Herring	-	-	-	5.9	1.6	0.6	14.0	12.4	0.7	1.4	27.8	1.0	trace	-	-	28.3	-	0.5	0.8	trace	trace	trace	0.7	0.5	trace	0.9	-	-	24
Mackerel	-	-	-	7.6	-	0.4	18.3	8.3	0.5	2.2	16.9	1.6	0.6	2.8	-	9.4	-	0.4	8.6	-	11.6	trace	1.3	7.6	-	-	0.8	-	24
Salmon	-	-	-	4.9	-	0.5	28.2	5.3	1.0	3.9	19.3	1.1	1.3	3.4	-	3.1	-	3.9	7.1	-	2.8	-	1.2	10.8	-	-	0.8	-	24
Trout	-	-	-	15.1	-	trace	15.3	5.6	1.5	3.8	27.2	1.5	0.9	-	trace	8.0	-	0.5	3.4	trace	10.5	-	0.7	2.5	trace	0.7	-	-	24
Tunny	-	-	-	2.1	-	0.8	11.9	8.2	1.5	4.1	19.8	4.6	5.2	1.5	-	3.0	0.6	2.2	5.0	-	1.3	0.6	2.6	19.6	-	-	-	-	24
Beef	-	-	-	4.5	-	0.6	22.1	2.8	0.8	6.1	21.7	2.1	1.2	0.5	-	2.0	-	3.0	13.2	-	trace	-	1.5	17.3	-	-	-	-	24
Chicken	-	-	-	3.2	0.6	26.9	6.3	-	1.2	13.0	42.0	2.0	1.3	-	-	trace	-	1.0	trace	-	-	trace	-	-	-	-	-	-	24
Duck	-	-	-	1.3	trace	26.7	7.2	-	trace	7.1	39.8	13.5	0.7	-	-	0.6	-	0.7	trace	-	-	trace	-	1.0	-	-	-	-	24
Lamb	-	-	-	0.6	trace	22.8	4.4	-	trace	5.5	52.8	12.1	0.6	-	-	trace	-	trace	trace	-	-	trace	-	-	-	-	-	-	24
Pork	-	-	-	5.4	0.6	24.2	1.3	-	1.0	20.9	38.2	2.5	2.5	-	-	trace	-	0	trace	-	-	trace	-	-	-	-	-	-	24
Turkey	-	-	-	1.6	trace	27.1	3.4	-	trace	13.8	43.8	7.4	0.9	-	-	0.7	-	trace	trace	-	-	trace	-	-	-	-	-	-	24

* In plant lipids C_{18:3} + C_{18:1}. † New strains contain less erucic acid. ** Values based on 9 Swiss samples; for American brands, see WEINBAUER et al.

Table 24 Content of ω 3 Fatty acids and other Fat Component in Selected Fish.

Fish	Fatty acids							Cholesterol
	Total fat	Total saturated	Total mono-unsaturated	Total poly-unsaturated	18:3	20:5	22:6	
Anchovy,								
European	4.8	1.3	1.2	1.6	—	0.5	0.9	—
Bass, striped	2.3	0.5	0.7	0.8	Tr	0.2	0.6	80
Bluefish	6.5	1.4	2.9	1.6	—	0.4	0.8	59
Carp	5.6	1.1	2.3	1.4	0.3	0.2	0.1	67
Catfish, brown								
bullhead	2.7	0.6	1.0	0.8	0.1	0.2	0.2	75
Catfish, channel	4.3	1.0	1.6	1.0	Tr	0.1	0.2	58
Cod, Atlantic	0.7	0.1	0.1	0.3	Tr	0.1	0.2	43
Croaker, Atlantic	3.2	1.1	1.2	0.5	Tr	0.1	0.1	61
Flounder,								
unspecified	1.0	0.2	0.3	0.3	Tr	0.1	0.1	46
Grouper, red	0.8	0.2	0.1	0.2	—	Tr	0.2	—
Haddock	0.7	0.1	0.1	0.2	Tr	0.1	0.1	63
Halibut,								
Greenland	13.8	2.4	8.4	1.4	Tr	0.5	0.4	46
Pacific	2.3	0.3	0.8	0.7	0.1	0.1	0.3	32
Herring, Pacific	13.9	3.3	6.9	2.4	0.1	1.0	0.7	77
Herring, Round	4.4	1.3	0.8	1.5	0.1	0.4	0.8	28
Mackerel, king	13.0	2.5	5.9	3.2	—	1.0	1.2	53
Mullet, striped	3.7	1.2	1.1	1.1	0.1	0.3	0.2	49
Ocean perch	1.6	0.3	0.6	0.5	Tr	0.1	0.1	42
Plaice, European	1.5	0.3	0.5	0.4	Tr	0.1	0.1	70
Pollock	1.0	0.1	0.1	0.5	—	0.1	0.4	71
Pompano, Florida	9.5	3.5	2.6	1.1	—	0.2	0.4	50
Salmon, Chinook	10.4	2.5	4.5	2.1	0.1	0.8	0.6	—
Salmon, pink	3.4	0.6	0.9	1.4	Tr	0.4	0.6	—
Snapper, red	1.2	0.2	0.2	0.4	Tr	Tr	0.2	—
Sole, European	1.2	0.3	0.4	0.2	Tr	Tr	0.1	50
Swordfish	2.1	0.6	0.8	0.2	—	0.1	0.1	39
Trout, rainbow	3.4	0.6	1.0	1.2	0.1	0.1	0.4	57
Tuna, albacore	4.9	1.2	1.2	1.8	0.2	0.3	1.0	54
Tuna, unspecified	2.5	0.9	0.6	0.5	—	0.1	0.4	—

* Values are given as g/100 g edible portion, raw except for cholesterol, which is given as mg. Dash (—) denotes lack of reliable data for nutrient known to be present. Tr, trace (less than 0.05 g/100 g of food).

Adapted from the U.S. Department of Agriculture Provisional Table on the Content of Omega-3 Fatty Acids and Other Fat Components in Seafoods, as presented by Simopoulos et al.

Table 25 Effects of ω 3 Fatty acids on Factor Involved in the pathophysiology of Atherosclerosis and Inflammation.

Factor	Function	Effect of ω 3 fatty acid
Arachidonic acid	Eicosanoid precursor; aggregates platelets; stimulates white blood cells	↓
Thromboxane	Platelet aggregation; vasoconstriction; increase of intracellular Ca^{++}	↓
Prostacyclin ($PGI_{2\beta}$)	Prevent platelet aggregation; vasodilation; increase cAMP	↑
Leukotriene (LTB_4)	Neutrophil chemoattractant; increase of intracellular Ca^{++}	↓
Tissue plasminogen activator	Increase endogenous fibrinolysis	↑
Fibrinogen	Blood clotting factor	↓
Red cell deformability	Decreases tendency to thrombosis and improves oxygen delivery to tissues	↑
Platelet activating factor (PAF)	Activates platelets and white blood cells	↓
Platelet-derived growth factor (PDGF)	Chemoattractant and mitogen for smooth muscles and macrophages	↓
Oxygen-free radicals	Cellular damage; enhance LDL uptake via scavenger pathway; stimulate arachidonic acid metabolism	↓
Lipid hydroperoxides	Stimulate eicosanoid formation	↓
Interleukin 1 and tumor necrosis factor	Stimulate neutrophil O_2 free radical formation; stimulate lymphocyte proliferation; stimulate PAF; express intercellular adhesion molecule-1 on endothelial cells; inhibit plasminogen activator, thus, procoagulants	↓
Endothelial-derived relaxation factor (EDRF)	Reduces arterial vasoconstrictor response	↑
VLDL	Related to LDL and HDL level	↓
HDL	Decreases the risk for coronary heart disease	↑
Lp(a)	Lipoprotein (a) is a genetically determined protein that has atherogenic and thrombogenic properties	↓
Triglycerides and chylomicrons	Contribute to postprandial lipemia	↓

Adapted from Weber, P.C. and Leaf, A., *World Rev. Nutr. Diet.* 66, 218, 1991.

APPENDIX IV

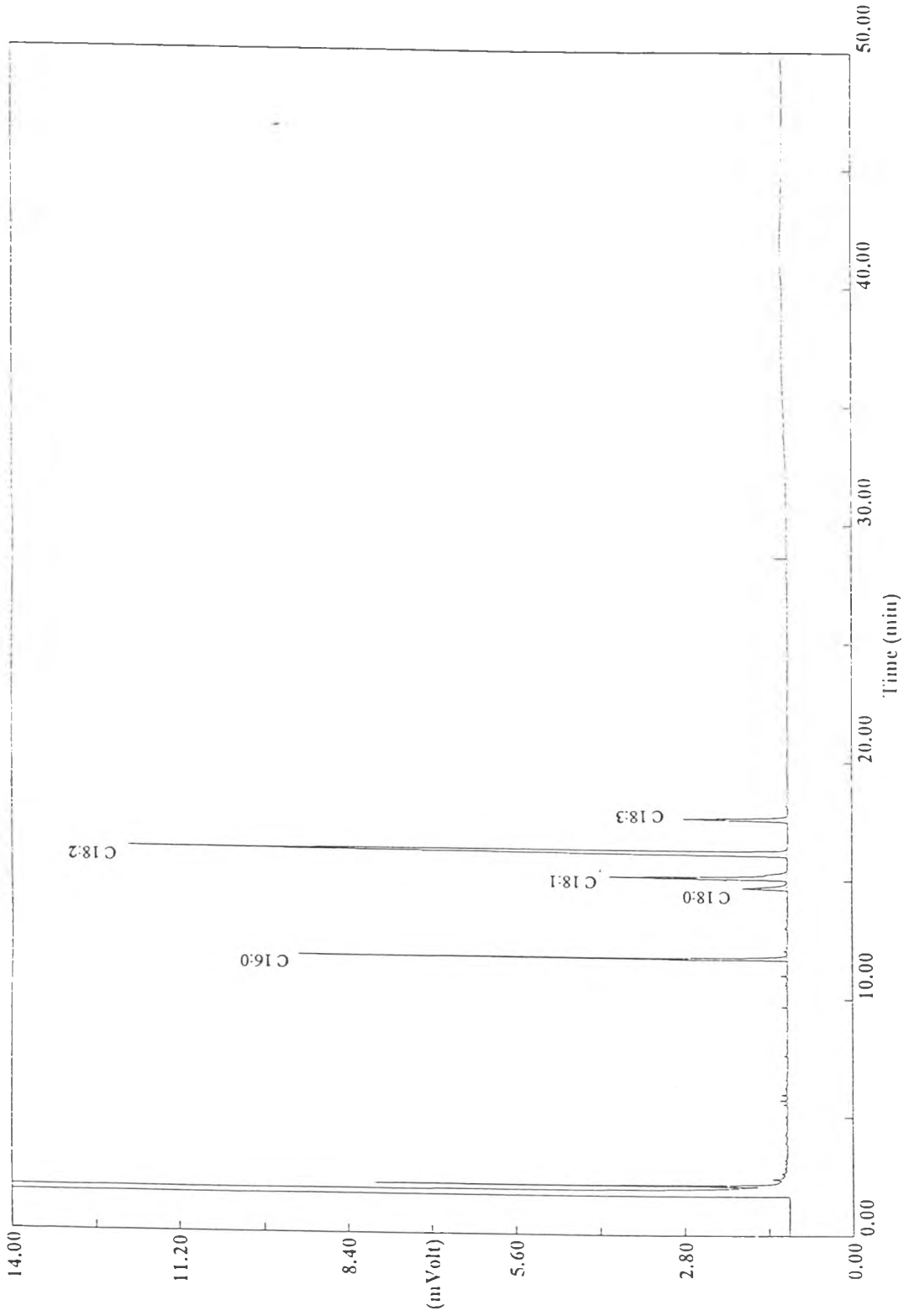


Figure 13 Gas-liquid chromatogram of fatty acid of total lecithin derived from soybean .

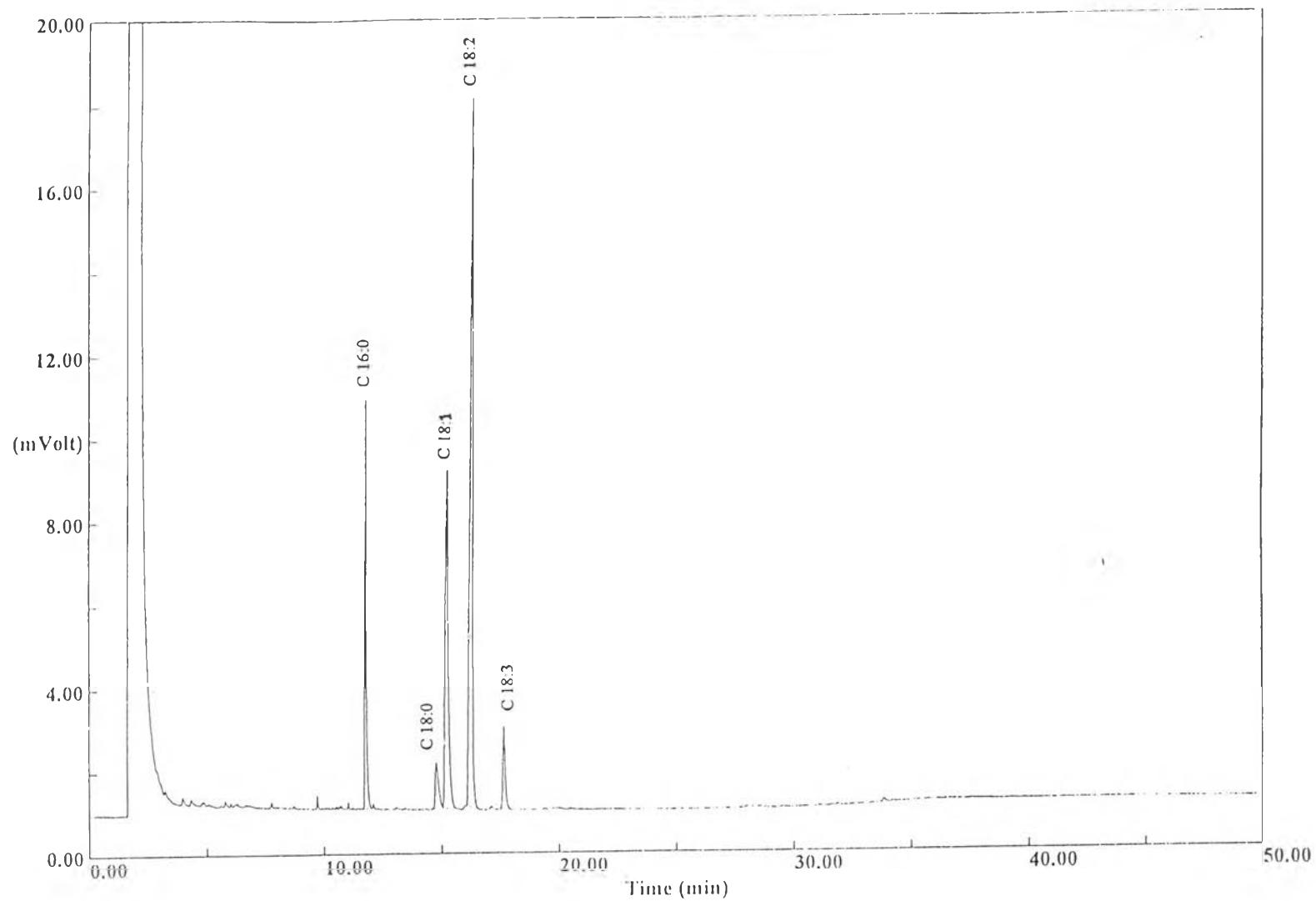


Figure 14 Gas-liquid chromatogram of fatty acid of triglycerides fraction of lecithin derived from soybean.

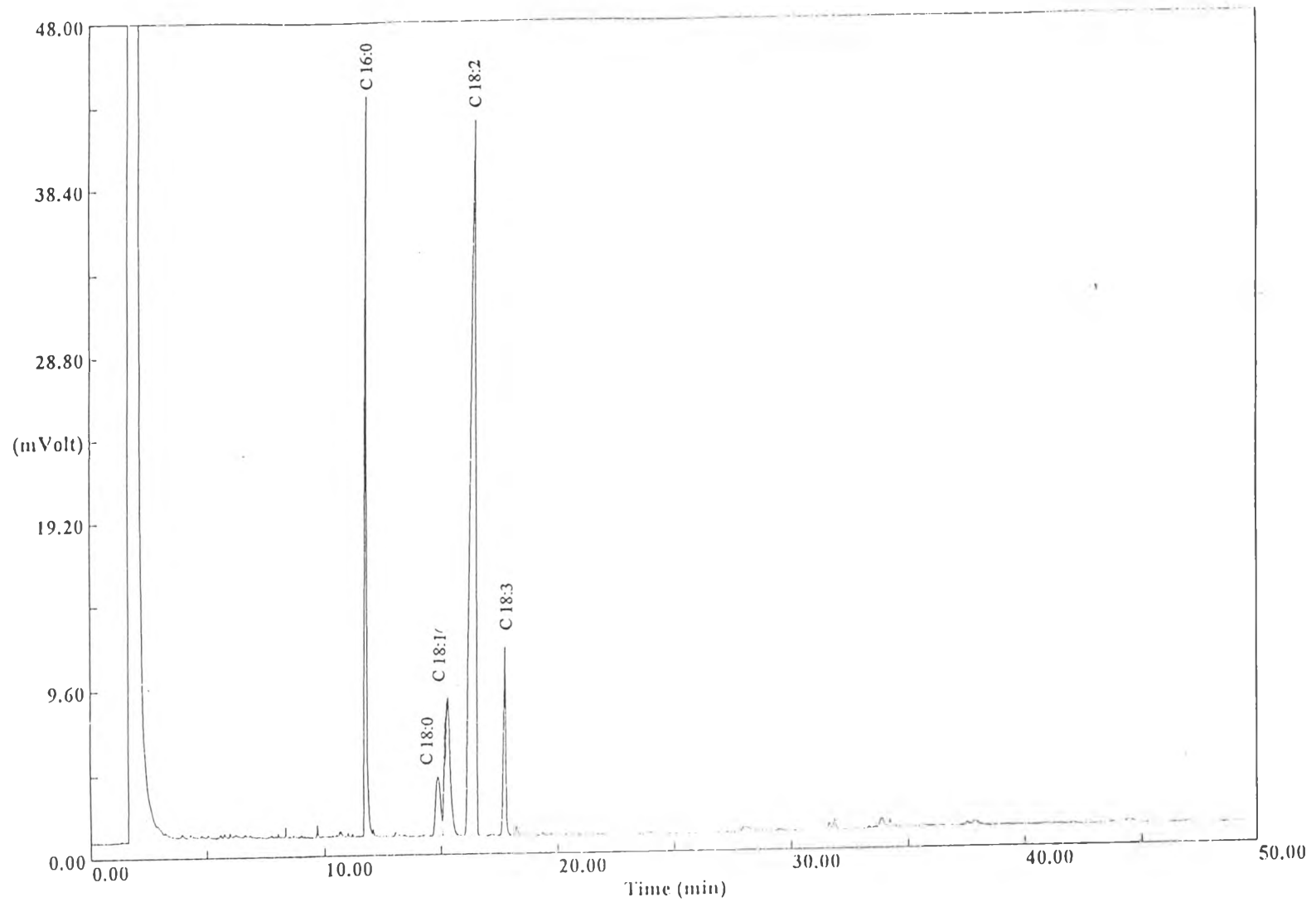


Figure 15 Gas-liquid chromatogram of fatty acid of phospholipid fraction of lecithin derived from soybean.

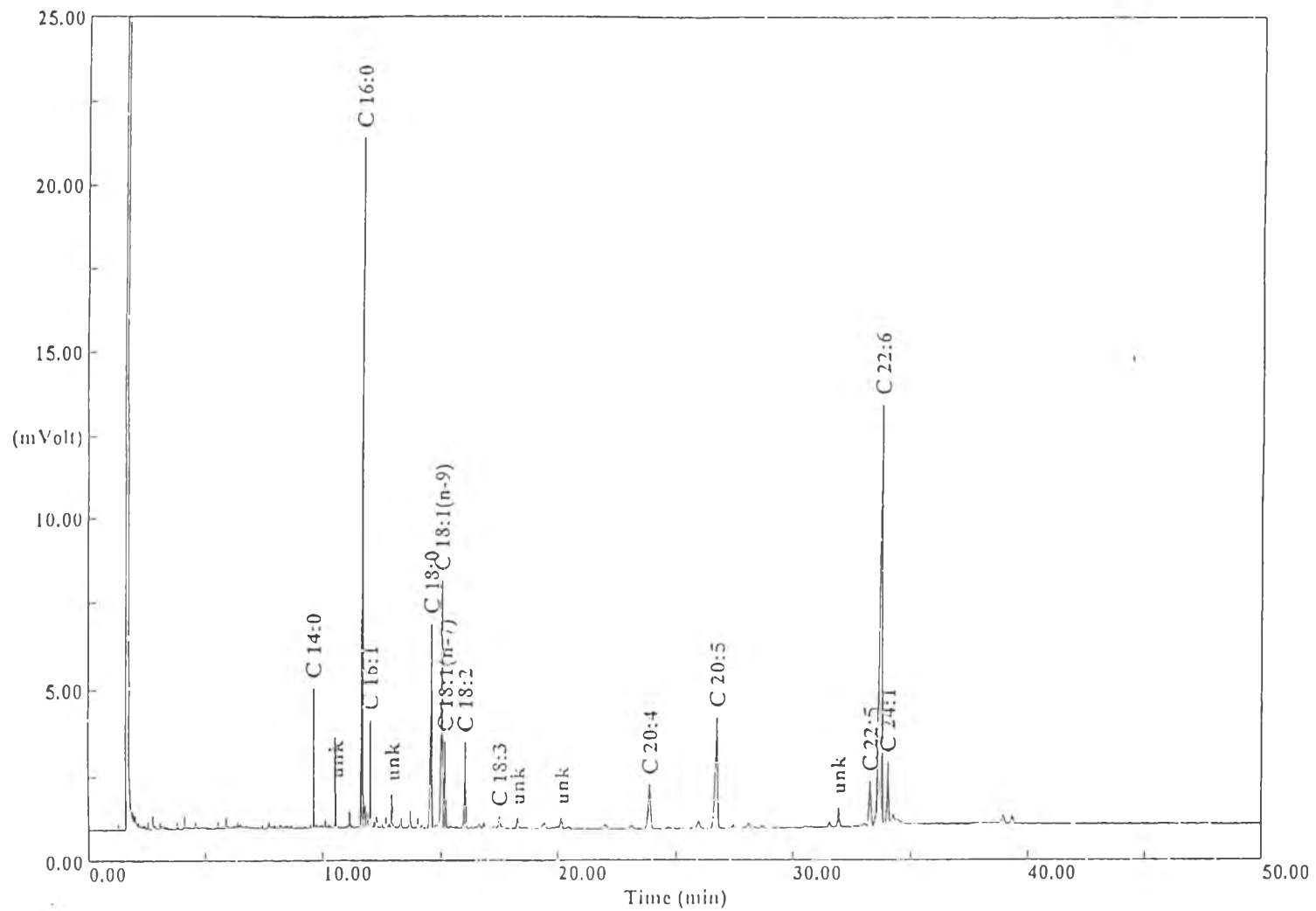


Figure 16 Gas-liquid chromatogram of fatty acid of total lecithin derived from Danish fish meal .

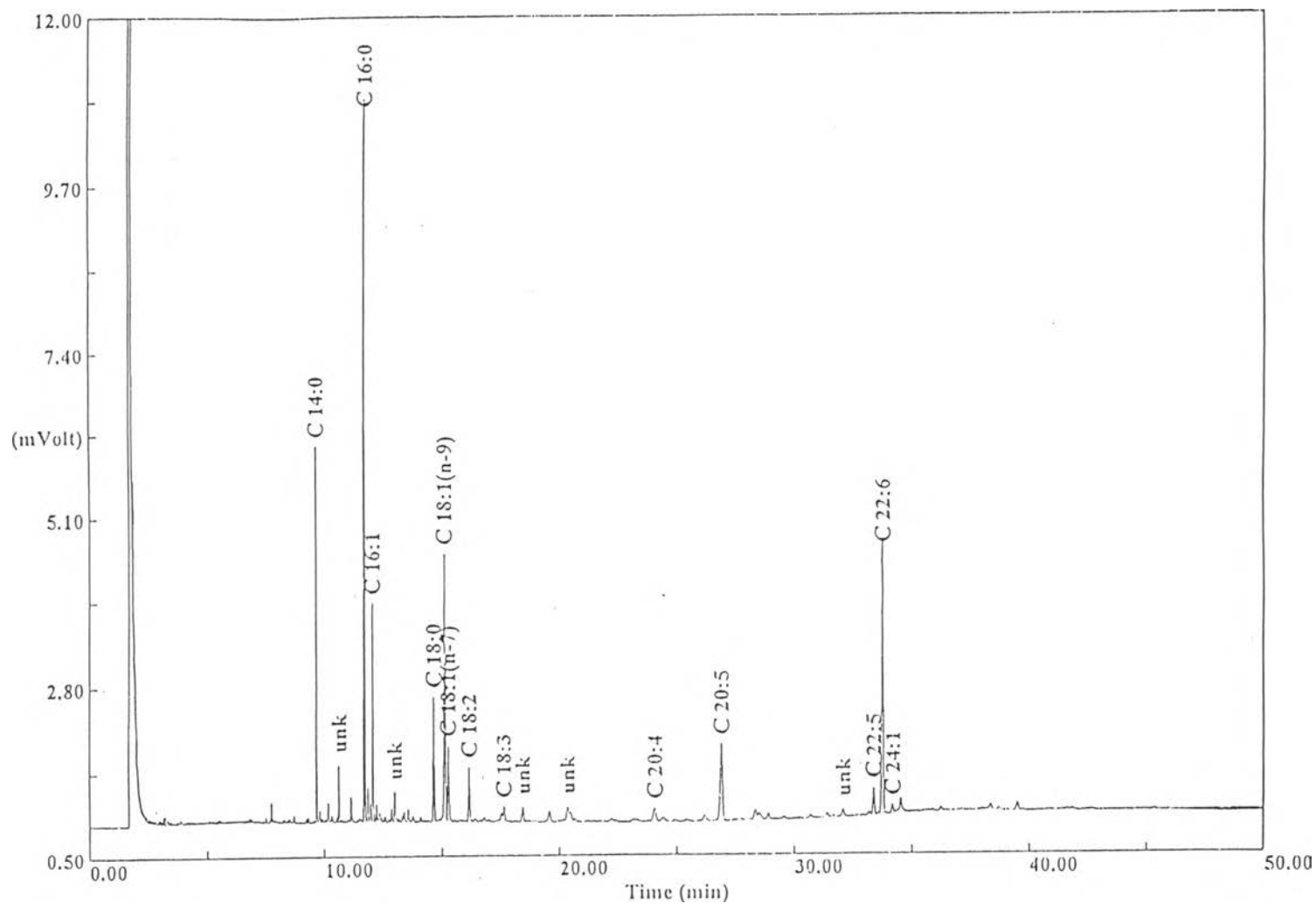


Figure 17 Gas-liquid chromatogram of fatty acid of triglyceride fraction of lecithin derived from Danish fish meal .

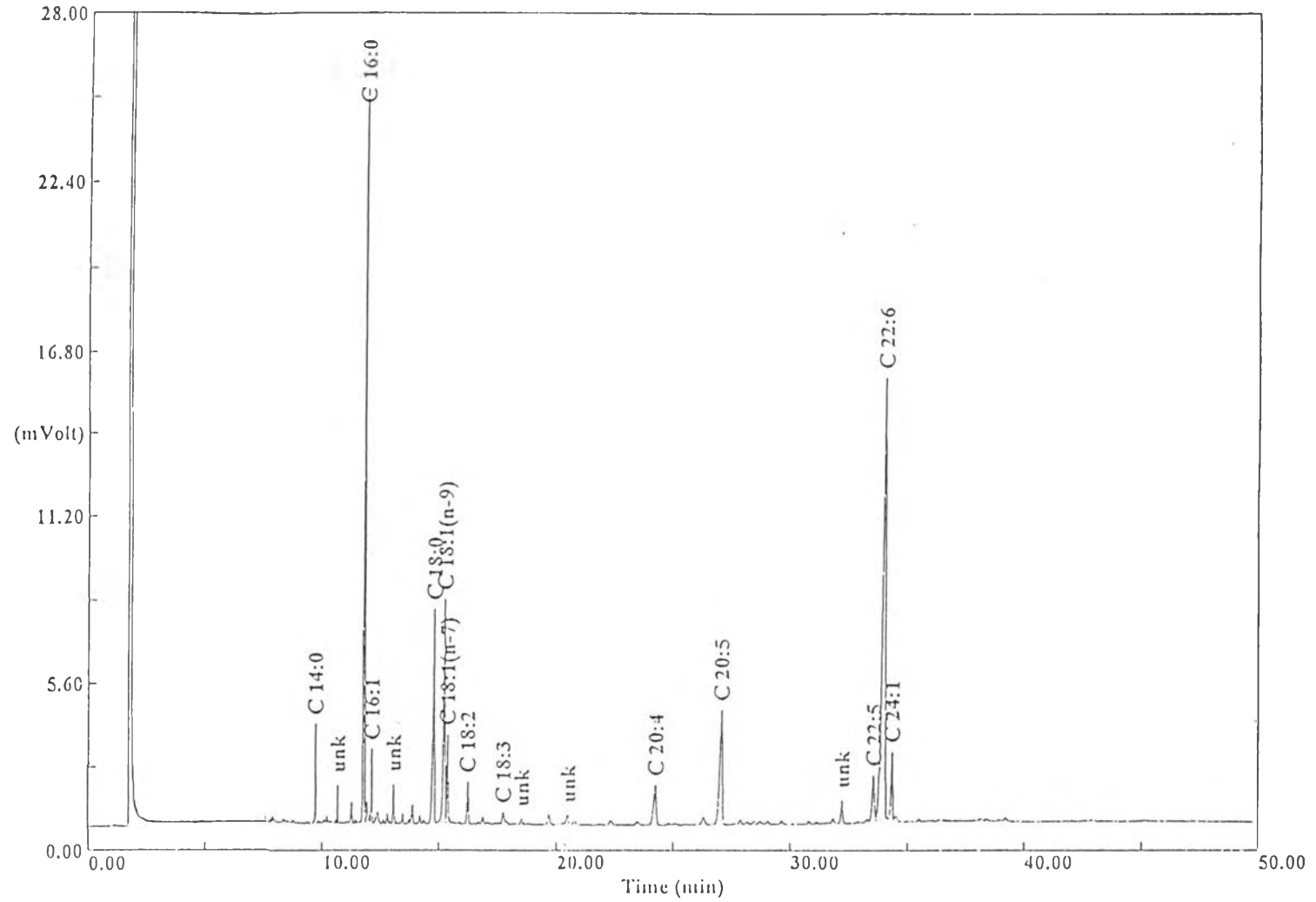


Figure 18 Gas-liquid chromatogram of fatty acid of phospholipid fraction of lecithin derived from Danish fish meal .

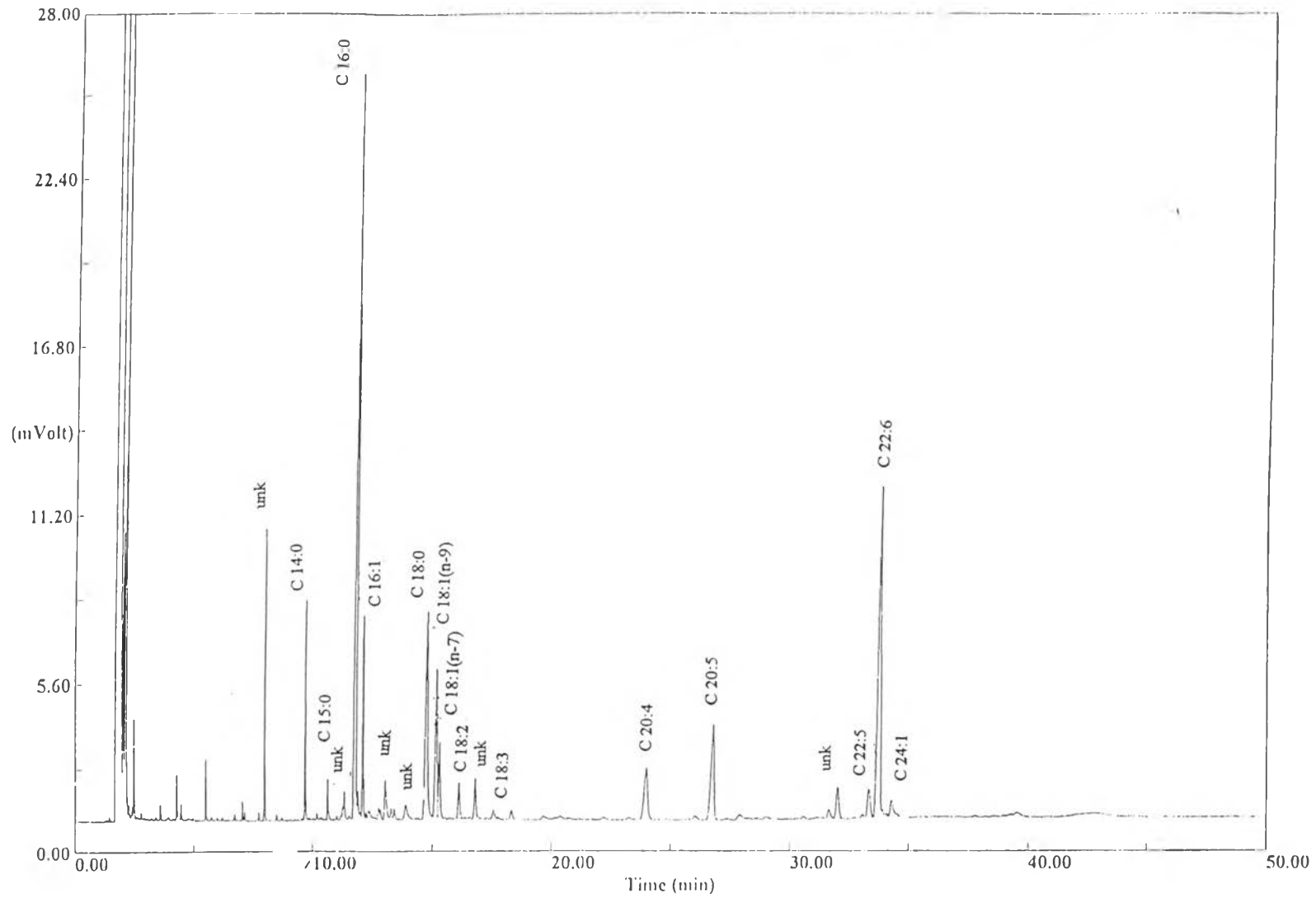


Figure 19 Gas-liquid chromatogram of fatty acid of total lecithin derived from local fish meal.

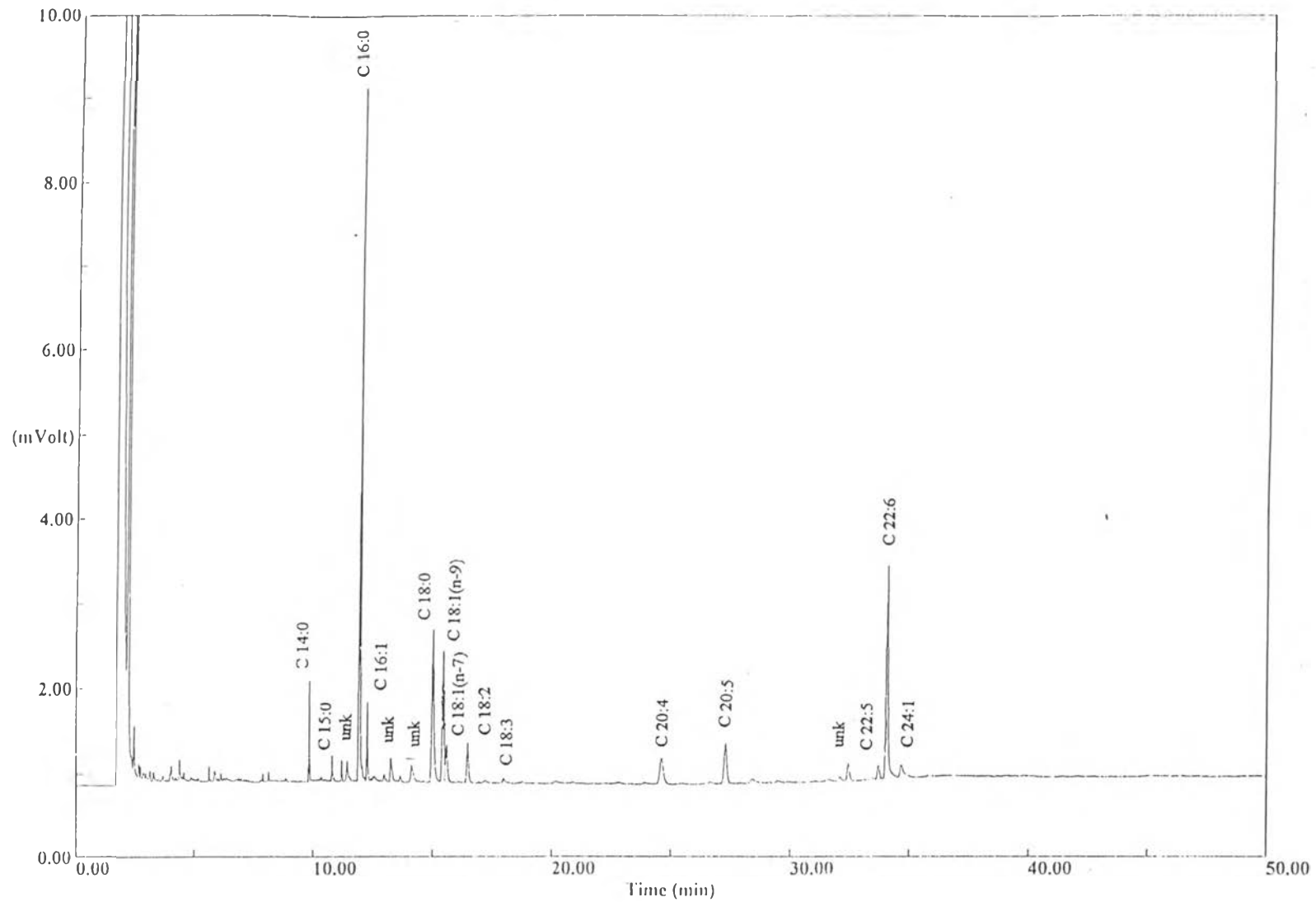


Figure 20 Gas-liquid chromatogram of fatty acid of triglyceride fraction of lecithin derived from local fish meal.

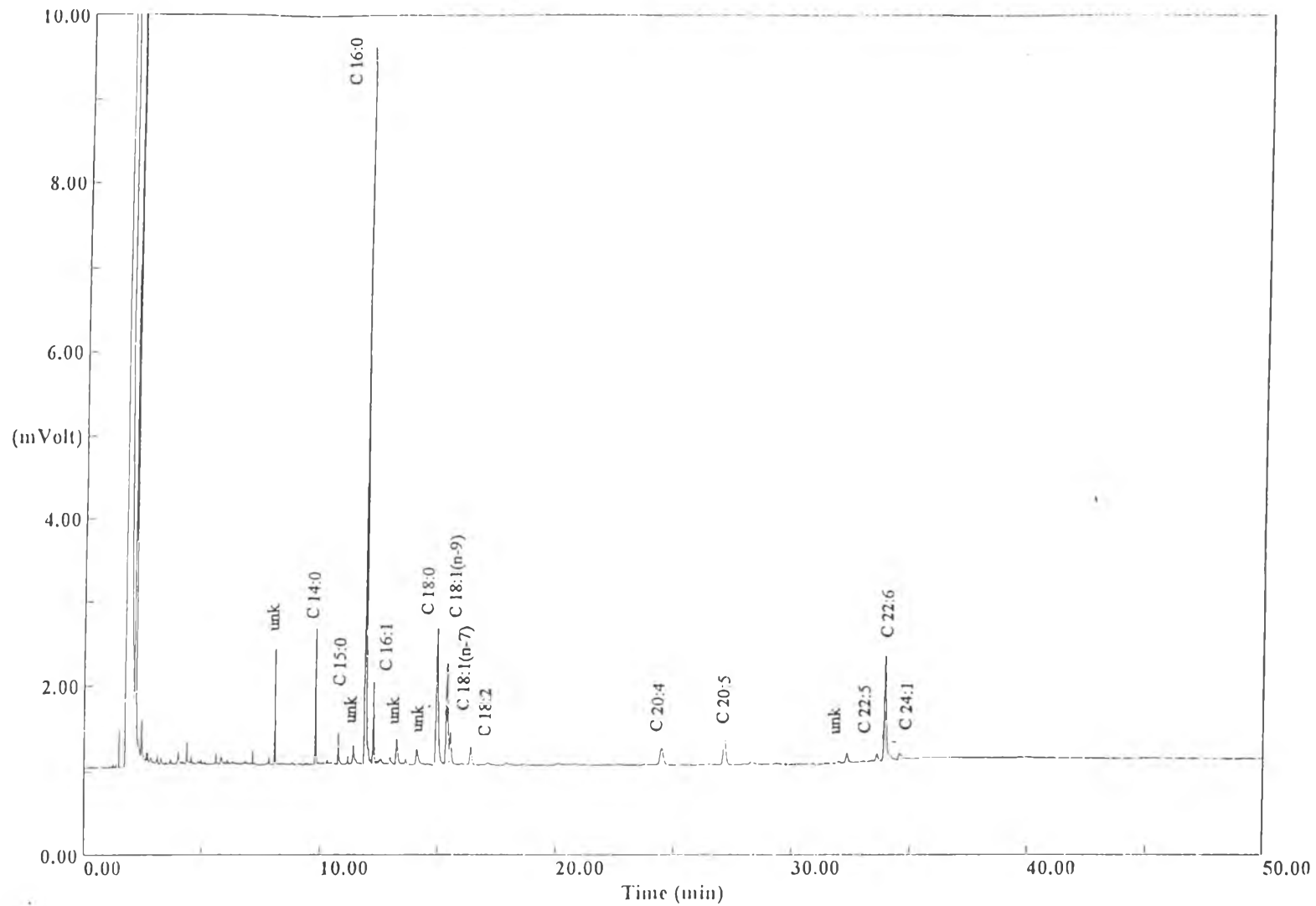


Figure 21 Gas-liquid chromatogram of fatty acid of phospholipid fraction of lecithin derived from local fish meal.

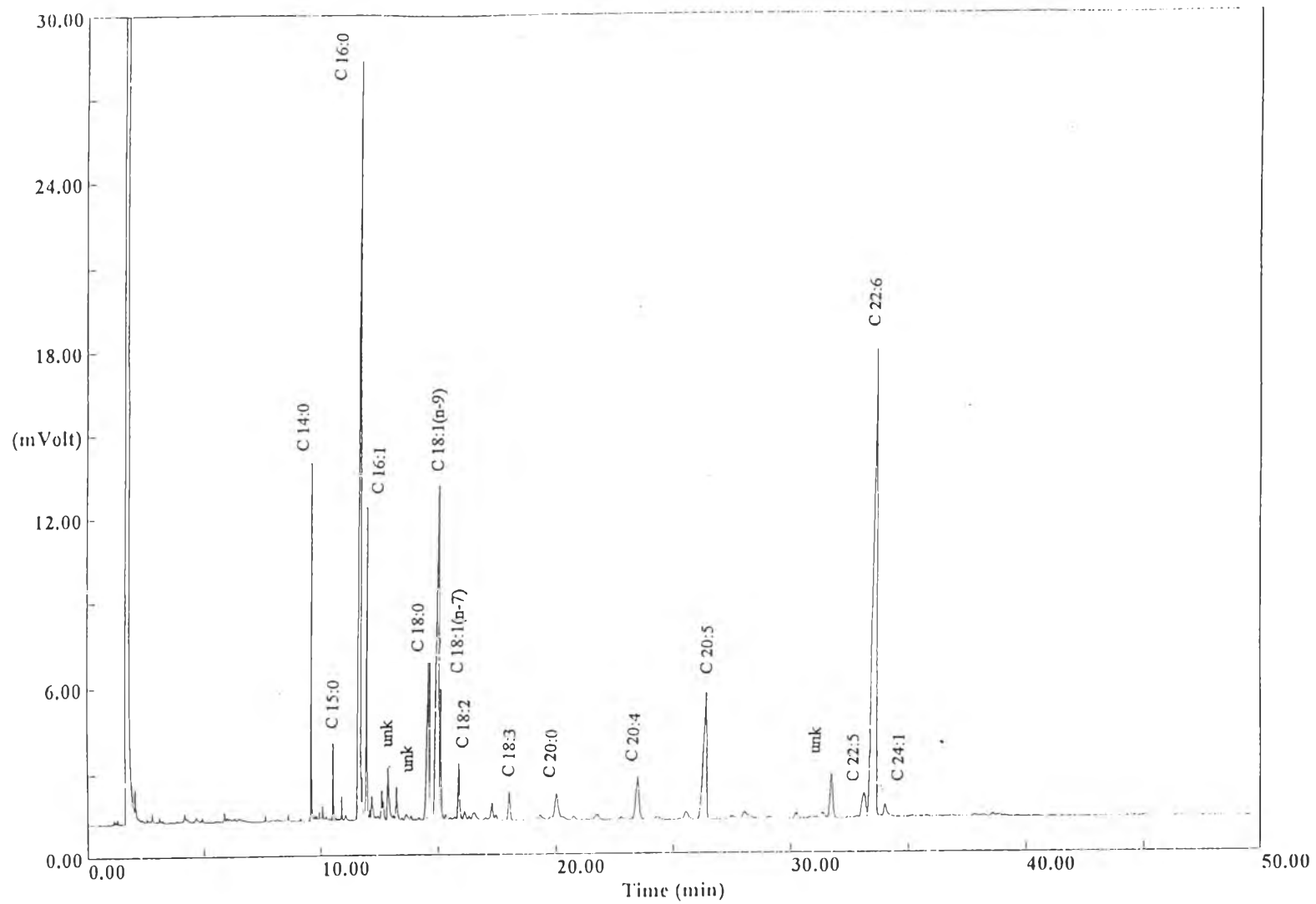


Figure 22 Gas-liquid chromatogram of fatty acid derived from fish oil.

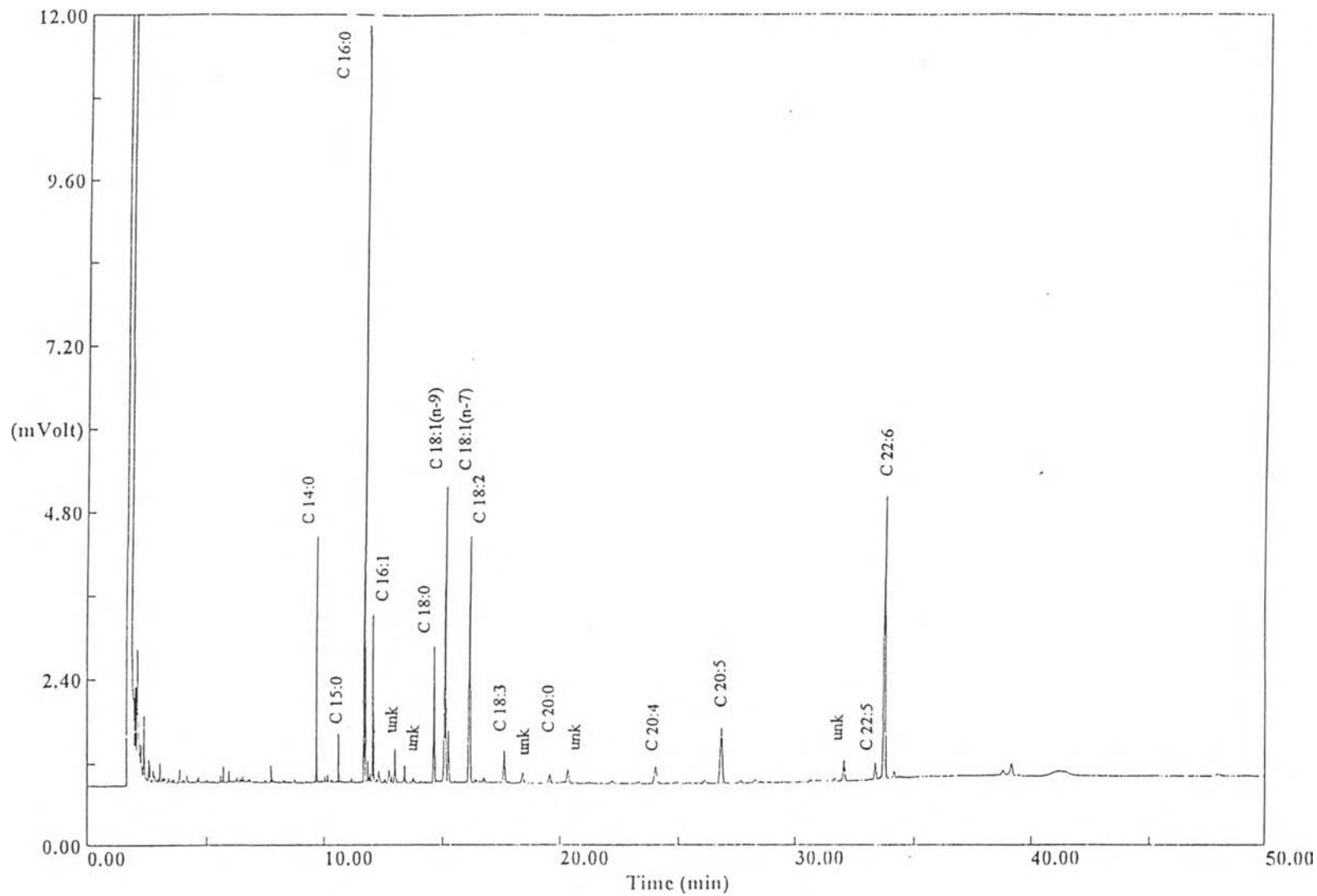


Figure 23 Gas-liquid chromatogram of fatty acid of Soybean lecithin - added diet (SAD) .

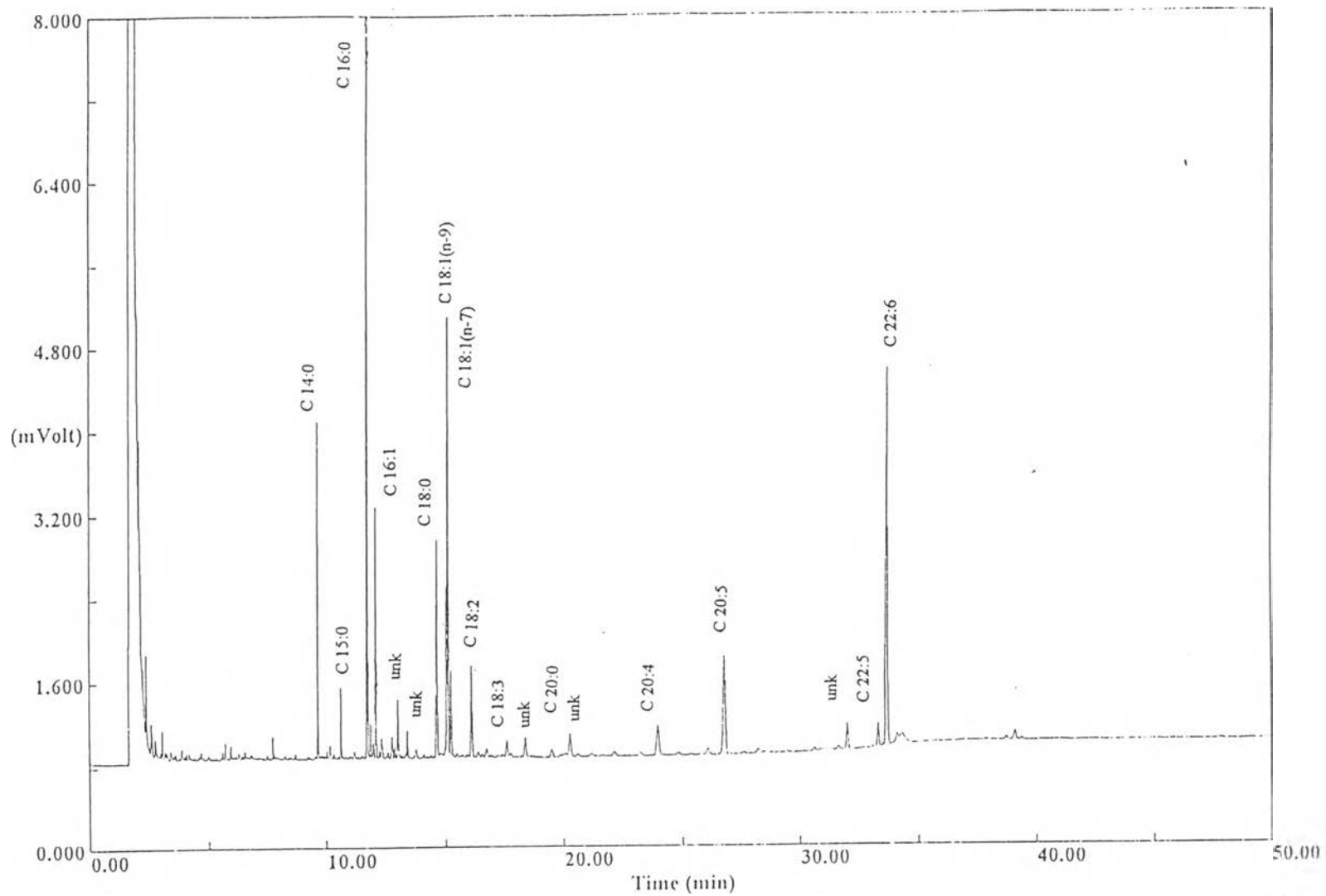


Figure 24 Gas-liquid chromatogram of fatty acid of Danish fish meal lecithin - added diet (DAD).

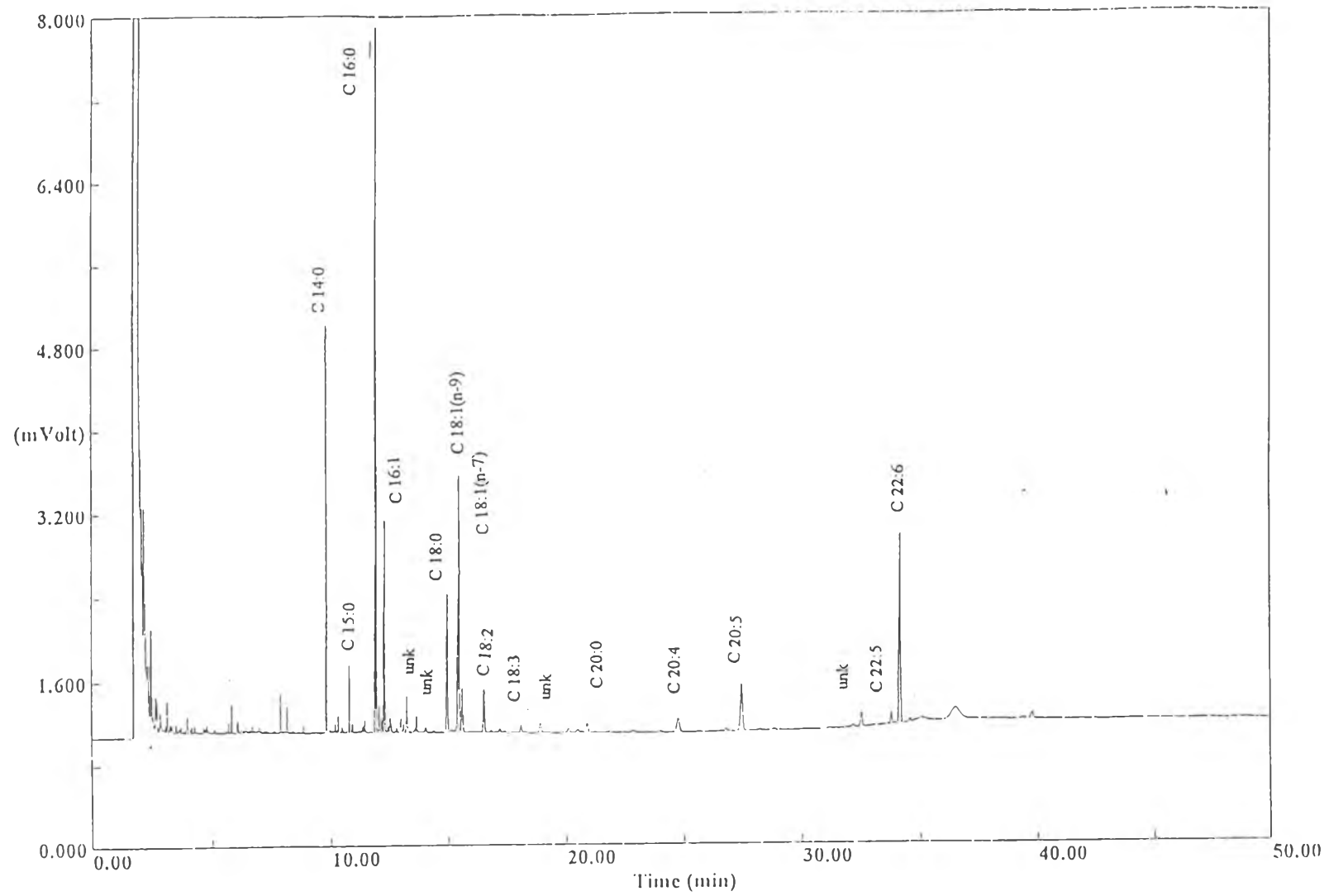


Figure 25 Gas-liquid chromatogram of fatty acid of local fish meal lecithin - added diet (LAD).

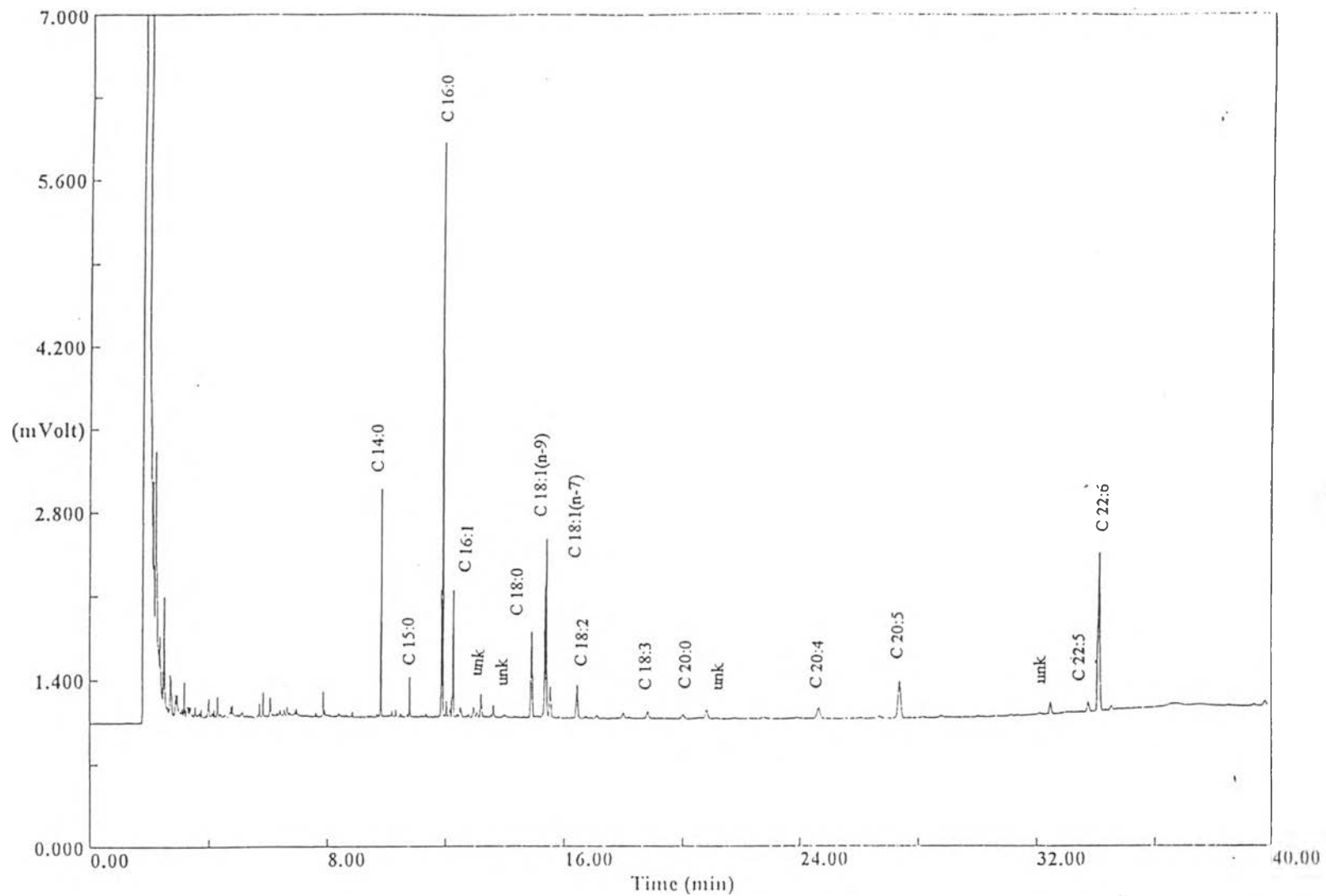


Figure 26 Gas-liquid chromatogram of fatty acid of Free - lecithin added diet (CD).

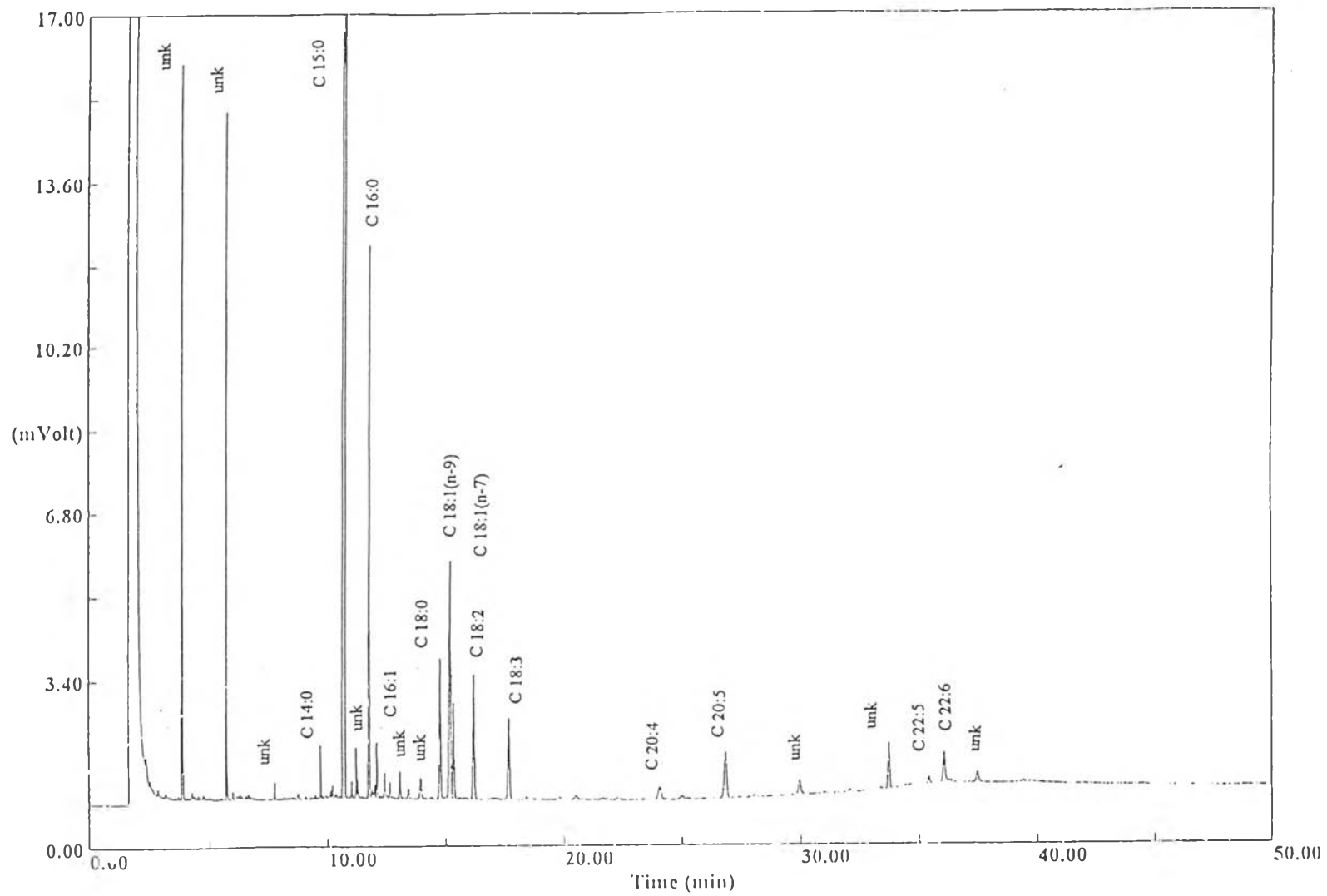


Figure 27 Gas-liquid chromatogram of fatty acid of shrimp fed SAD .

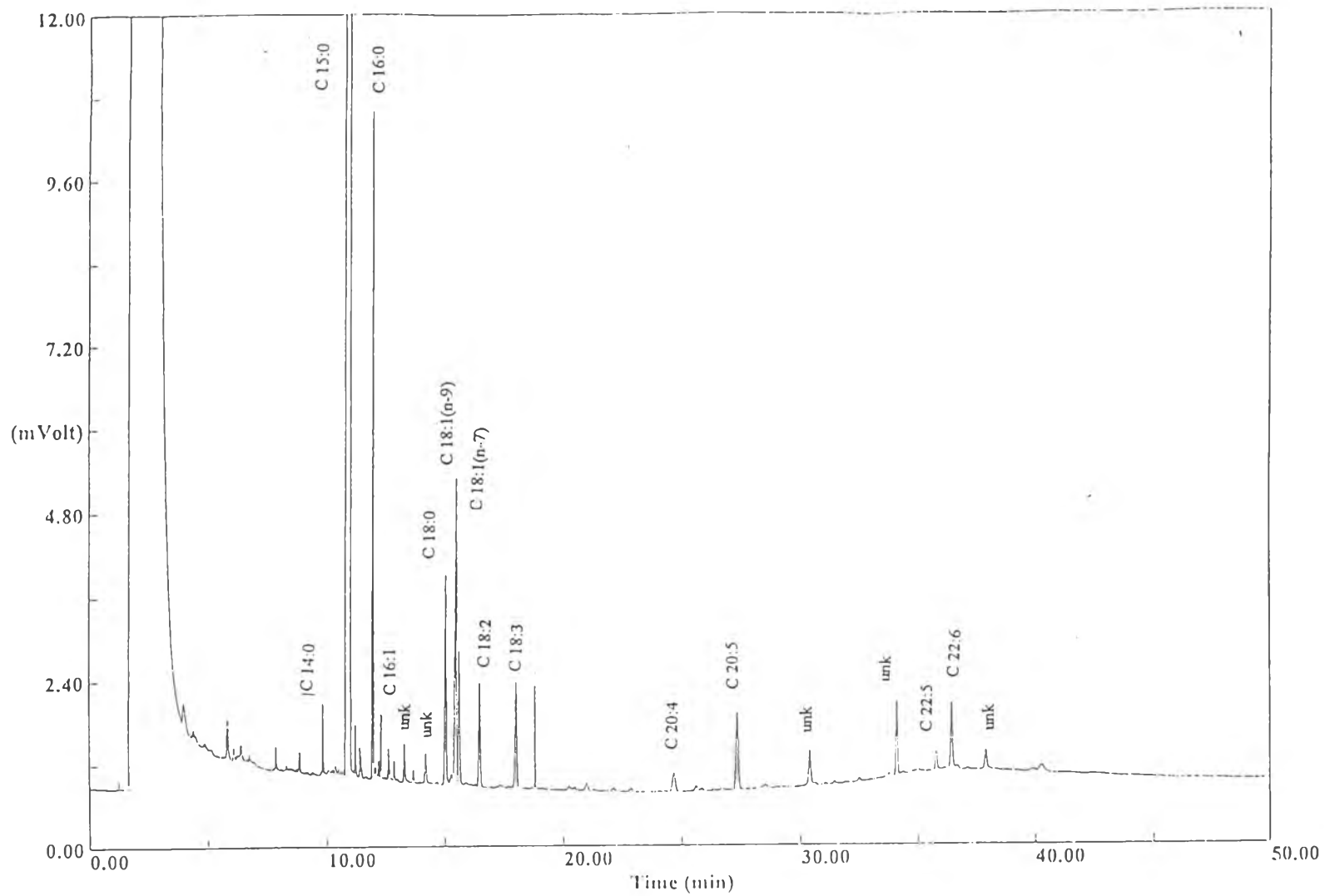


Figure 28 Gas-liquid chromatogram of fatty acid of shrimp fed DAD.

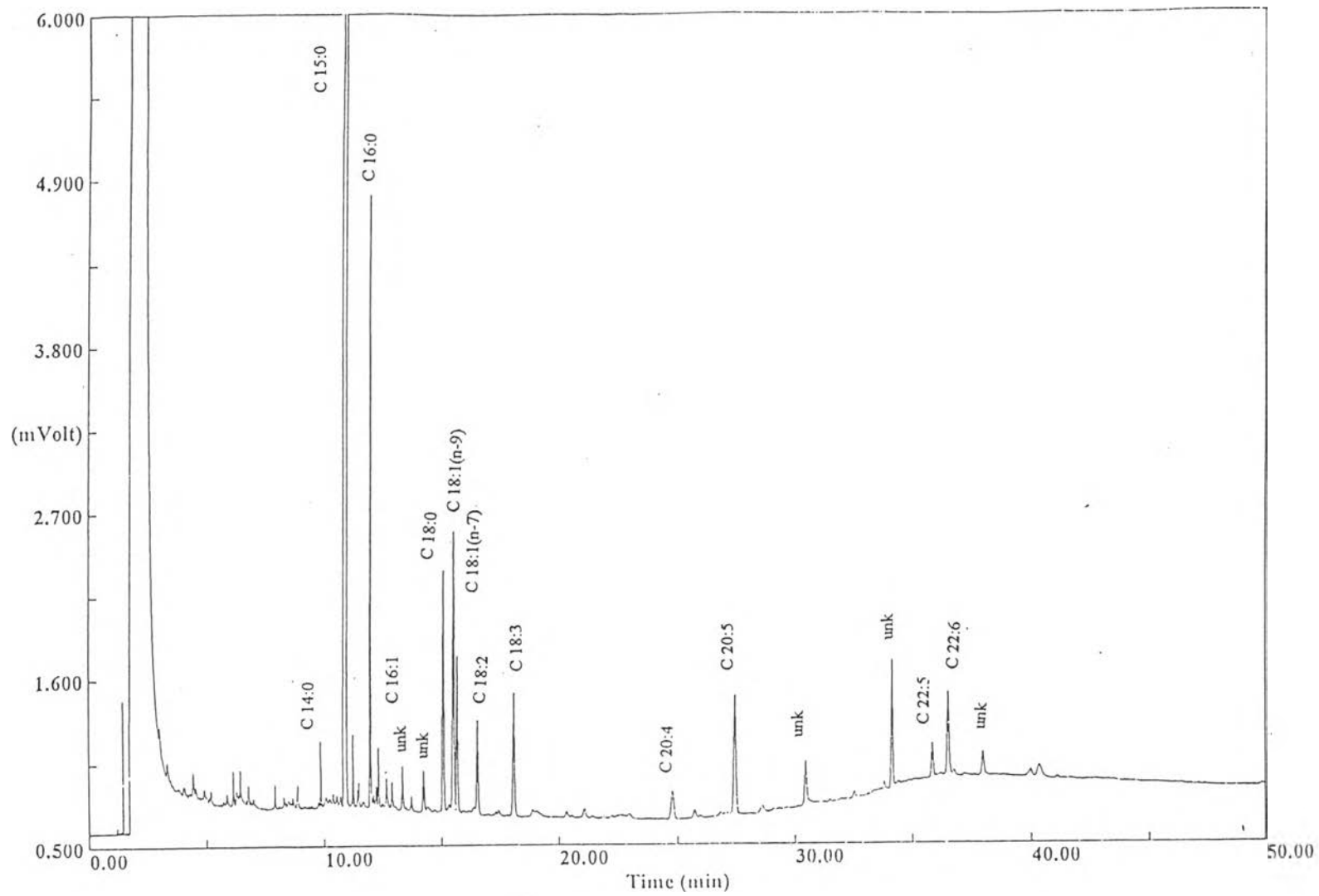


Figure 29 Gas-liquid chromatogram of fatty acid of shrimp fed LAD.

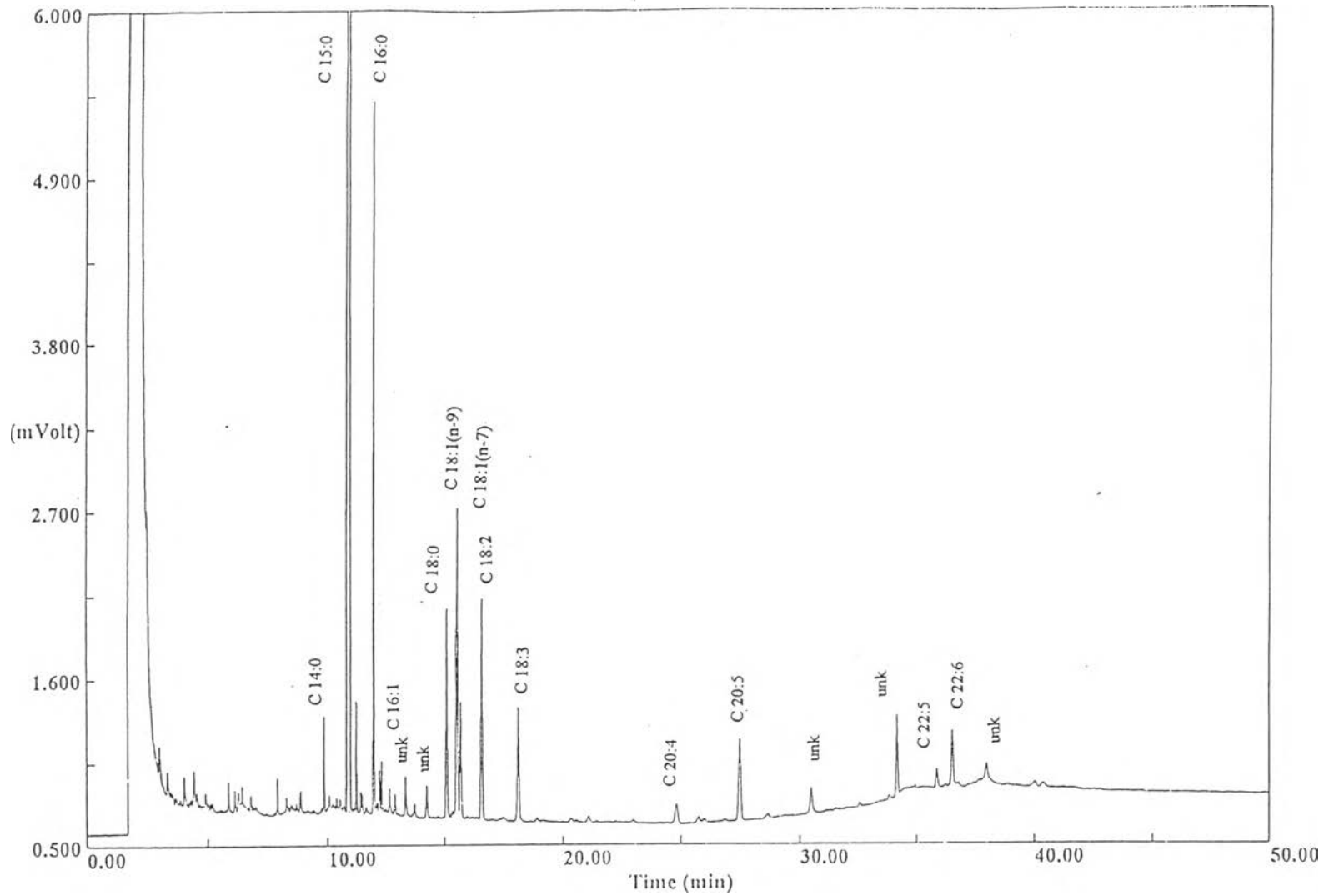


Figure 30 Gas-liquid chromatogram of fatty acid of shrimp fed CD.

BIOGRAPHY

Miss Pimporn Innopakun was born on January 22, 1973 in Saraburi Province, Thailand. She graduated with a Bachelor degree in Biology, Faculty of Science, Chulalongkorn University in 1995. She has studied for Master degree in Biotechnology Programme at Chulalongkorn University since 1995.

