

## Changes in the Haemolymph Metabolites, Protein, Carbohydrate, and Free Amino Acid of *Galleria mellonella* L. Parasitized by the Pupal Parasitoid, *Brachymeria lasus* Walker

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*Galleria mellonella* L. was parasitized by the pupal parasitoid, *Brachymeria lasus* Walker in the laboratory. Haemolymph total protein and carbohydrate concentrations from the hosts parasitized by *B. lasus* decreased within 1st day following parasitization. However, haemolymph total free amino acids, glycine, proline, arginine, isoleucine, phenylalanine, leucine, valine, glutamic acid, histidine, serine, and tyrosine increased and free amino acids decreased were threonine, aspartic acid, methionine, cysteine, alanine, and lysine.

**KEY WORDS:** Parasitization, Haemolymph, *Brachymeria lasus* Walker, *Galleria mellonella* L. Protein, Carbohydrate, Free amino acids

Hymenopterous endoparasitoids were known to cause many physiological and biochemical changes in the host insect, an action Vinson and Iwantsch (1980) termed "host regulation." These changes included alternations in growth, food utilization and development rate of host that are thought to benefit the developing parasitoid. Changes that have been reported in the host haemolymph after parasitization include changes in specific gravity, freezing point depression, dry weight, hormone titers, plus differences in amino acid, carbohydrate, lipid, and protein concentrations and in the number of electrophoretically separable proteins (Baldwin and House, 1952; Corbet, 1968; Barras *et al.*, 1969; Dahlman, 1970; Brewer *et al.*, 1973; Thompson and Barlow, 1974; Dahlman, 1975; Vinson and Iwantsch, 1980; Dahlman and Green, jr, 1981; Beckage and Riddiford, 1982; Thompson, 1982).

*Brachymeria lasus* was best known as a solitary

endoparasitoid of fall webworm pupae, *Hyphantria cunea* in Korea, and this insect was polyphagous. Currently, there was interest in *B. lasus* as a possible biocontrol agent for gypsy moth, *Lymantria dispar* and fall webworm, *Hyphantria cunea*.

After *B. lasus* parasitizes the host insect, greater wax moth, *Galleria mellonella*, we examined the changes in haemolymph metabolites—protein, carbohydrate, and free amino acid concentrations of host for studies on in vitro culture of the parasitoid.

### Materials and Methods

*Galleria mellonella* pupae were reared at 32°C in darkness, and maintained according to the methods of Dutky *et al.* (1962). Wax moth pupae were removed from their cocoons for exposure to the parasitoids.

Wax moth pupae were collected daily for experiments in which pupal age was required, and were used to rear *B. lasus* for colony maintenance.

The parasitoids were reared in 9-cm diam petri

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dish at  $25 \pm 1^\circ\text{C}$  with 16L/8D photoregion and fed a 10% honey water solution soaked into cotton dental wicks.

Wax moth pupae (0-day-old pupae) were exposed to *B. lausus* in 9 cm diam petri dish at  $25 \pm 1^\circ\text{C}$  and then the pupae parasitized kept in clear petri dish. The pupae (0-day-old pupae) unparasitized also were kept in the same condition.

Haemolymph samples were obtained by puncturing near the head of pupae with fine needle and being collected into eppendorf tubes in ice with capillary tubes at 24-h intervals after pupae were given control treatment or treatment with the parasitoids. Haemolymph from several pupae was pooled to obtain a minimum volume of  $70 \mu\text{l}$ , but usually the volume of the sample was  $70\text{--}100 \mu\text{l}$ . To collect the haemolymph of parasitized host pupae, the egg and larva of *B. lausus* in the host pupae was identified by dissecting host pupae in Ringer's solution (128 mM NaCl, 1.8 mM  $\text{CaCl}_2$ , 1.3 mM KCl; pH 7.4) under stereo microscope.

A small amounts of phenylthiourea (PTU) was added to the appropriate amount of haemolymph to prevent melanization and this haemolymph was centrifuged at 10,000 g for 10 min at  $4^\circ\text{C}$  to remove haemocytes and tissue debris and supernatant was taken and stored at  $-20^\circ\text{C}$  for use.

Total protein concentration was determined by the Folin and Ciocateau's Phenol Reagent with bovine serum albumin as standard (Lowry *et al.*, 1951), and total carbohydrate concentration also was determined by Anthrone methods with glucose as standard (Joseph H. Roe, 1954).

Free amino acids were estimated by Pico-Tag amino acid analysis system (Waters Assoc) with Pierce H (17 species of free amino acid contained in  $2.5 \mu\text{mol/ml}$ ) as standard (Brian *et al.*, 1984). The  $50 \mu\text{l}$  of haemolymph sample and 10 volumes of diluent standard were placed in sample tubes and dried under vacuum. Samples were dried down again after adding  $10 \mu\text{l}$  of ethanol-water-triethylamine (TEA) (2:2:1) to each sample tubes. The derivatization was made fresh daily and consisted of ethanol-TEA-water-phenylisothiocyanate (PITC) (7:1:1:1). The phenylthiocarbonyl (PTC) derivative of free amino acids was formed by adding  $20 \mu\text{l}$  of ethanol-TEA-water-phenylisothiocyanate to the dried

samples and then in the vacuum vials for 20 min at room temperature. Ethanol-TEA-water-phenylisothiocyanate was then removed under using the Workstation. The column used was C18 reverse-phase column with  $4 \mu\text{m}$  packing ( $150 \text{ mm} \times 3.9 \text{ id}$ ). Each samples ( $10 \mu\text{l}$ ) were injected after adding  $250 \mu\text{l}$  of sample diluent to each sample tubes and then dissolving samples. The solvent system consisted of two eluents: (A) an aqueous buffer and (B) 60% acetonitril in water. The typical buffer was 0.14 M sodium acetate containing 0.5 ml/TEA and titrated to pH 6.35 with glacial acetic acid. A gradient which was run for the separation was a linear increase from 0% B to 46% B in 10 min, followed by 46% B to 100% B in the next 4 min.

## Results

The concentrations of haemolymph protein and carbohydrate in wax moth pupae parasitized by *B. lausus* has significantly lower levels than that of unparasitized hosts during the entire experimental periods (Figs. 1 and 2).

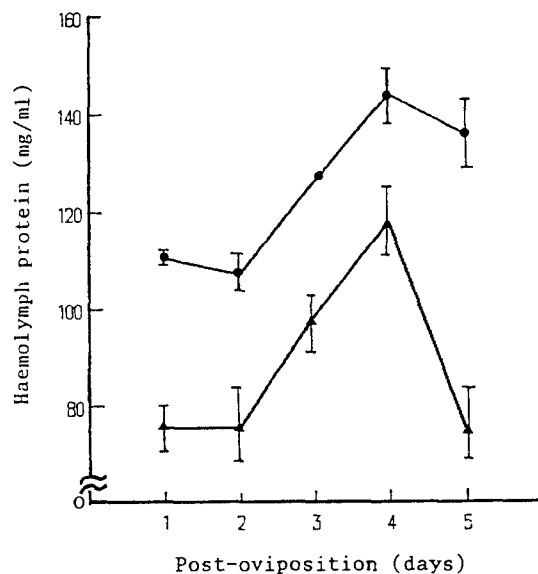


Fig. 1. The haemolymph total protein concentrations of *Galleria mellonella* unparasitized and parasitized by *Brachymeria lausus*, as 0-day-old pupae. Each value represents the mean of several individuals. ▲: Parasitized, ●: Unparasitized

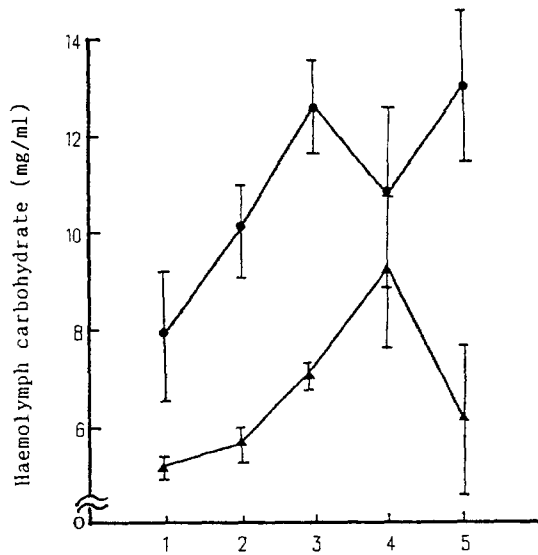


Fig. 2. The haemolymph total carbohydrate concentrations of *Galleria mellonella* unparasitized and parasitized by *Brachymeria lasus*, 0-day-old pupae. Each value represents the haemolymph of several individuals. ▲: Parasitized, ●: Unparasitized

Quantitatively, total free amino acid concentration in wax moth pupae was approximately 3200  $\mu\text{g/ml}$  haemolymph in all the experimental periods, and was dominated by threonine, methionine, lysine, glutamic acid, alanine, proline, tyrosine, and arginine, together with lesser amounts of histidine, valine, leucine, and serine, which varied among experimental periods. Glycine, isoleucine, phenylalanine, aspartic acid, and cysteine were minor constituents in all the experimental periods and were present at variable concentrations all less than 100  $\mu\text{g/ml}$  haemolymph (Tables, 1 and 2).

Parasitization resulted in a increase in total free amino acid concentration except 1st day of the experimental periods (Fig. 3) and a modification in free amino acid compositions. Increased free amino acids were glycine, proline, arginine, isoleucine, phenylalanine, leucine, valine, glutamic acid, histidine, serine, and tyrosine, and decreased free amino acids were threonine, aspartic acid, methionine, cysteine, alanine, and lysine (Tables. 1 and 2).

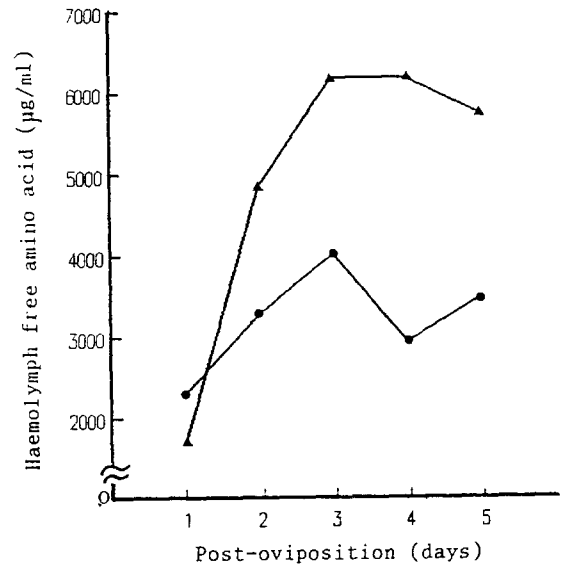


Fig. 3. The haemolymph total free amino acid concentrations of *Galleria mellonella* unparasitized and parasitized by *Brachymeria lasus*, 0-day-old pupae. Each value represents the haemolymph of several individuals. ▲: Parasitized, ●: Unparasitized

## Discussion

Various results on the concentration of haemolymph protein and carbohydrate have been obtained from studies with hymenopterous insect parasite. Dahlman (1975) reported that haemolymph trehalose concentrations decreased in *Manduca sexta* parasitized by *Apanteles congregatus*. In contrast, the concentration in *Heliothis virescens* was unaltered after infection by *Cardiophiles nigriceps* (Dahlman and Vinson, 1975). Dahlman and Vinson (1975), however, reported a marked elevation in the haemolymph concentration of trehalose in *H. virescens* parasitized by *Microplitis croceipes*. Similar results were reported for *Trichoplusia ni* parasitized by *Hyposoter exiguae* (Thompson, 1982).

The effects of the gregarous insect parasite, *Apanteles congregatus* on the total haemolymph protein level in *Manduca sexta* parasitized as first instar larvae was examined by Dahlman (1969). Early fourth and fifth instar host insects had levels similar to uninfected individuals. Smilowitz and Smith (1977) demonstrated alternations in

**Table 1.** Haemolymph concentrations essential free amino acids in *Galleria mellonella* unparasited and parasitized by *Brachymeria lasus*, as 0-day-old pupae.

Amino acid concentrations ( $\mu\text{g/ml}$ )	Post-oviposition (days)										Change R.C.(%)*		
	1 day	Change (%)	2 day	Change (%)	3 day	Change (%)	4 day	Change (%)	5 day	Change (%)		Average	Change (%)
Histidine	U** 208.021	-55	128.540	+107	254.661	-32	106.665	+244	234.559	+60	186.489	+37	9.8
	P*** 92.795		266.695		173.644		367.050		374.557		254.948		9.9
Arginine	U 408.203	-53	120.912	+321	240.091	+119	170.398	+334	201.497	+305	228.220	+150	12.1
	P 191.577		580.979		525.461		740.104		816.276		570.880		22.3
Threonine	U 34.714	+82	513.902	-71	507.650	-38	231.500	+19	341.153	-33	325.784	-35	17.2
	P 63.172		150.655		313.132		275.927		261.187		212.815		8.3
Valine	U 104.509	-8	184.945	+54	258.963	+90	208.803	+86	221.817	+74	195.824	+68	10.4
	P 95.594		284.345		490.873		388.587		386.438		329.159		12.8
Methionine	U 109.509	+6	396.249	-13	337.551	+7	379.522	-59	219.553	-40	288.477	-23	15.3
	P 116.501		342.844		360.847		153.634		130.848		220.935		8.6
Isoleucine	U 29.826	+84	74.344	+97	127.901	+127	100.728	+128	108.733	+128	88.306	+120	4.7
	P 54.840		146.748		290.011		230.093		248.222		193.844		7.2
Leucine	U 50.840	+71	124.268	+82	268.686	+82	197.858	+91	242.686	+66	176.868	+79	9.4
	P 87.119		226.249		488.786		377.831		403.459		316.689		12.3
Phenylalanine	U 36.312	+44	69.219	+50	131.811	+81	87.324	+94	108.625	+103	86.658	+81	4.6
	P 52.144		103.868		238.214		169.342		220.967		156.907		6.1
Lysine	U 186.022	-25	348.042	+19	393.072	-38	276.467	+32	373.279	+4	315.376	-2	16.7
	P 139.953		413.093		242.452		365.936		387.659		309.819		12.1
Total	U 1168.041	-23	1960.412	+28	2520.385	+24	1759.295	+74	2051.902	+57	1892.003	+36	
	P 893.000		2512.476		3123.420		2068.504		3229.613		2566.003		

\*Relative concentration, \*\*Unparasitized, \*\*\*Parasitized.  $\text{Change (\%)} = \frac{(\text{Parasitized} - \text{Unparasitized})}{\text{Unparasitized}} \times 100$

Table 2. Haemolymph concentrations of nonessential free amino acids in *Galleria mellonella* unparasitized and parasitized by *Brachymeria lasus*, as O-day-old pupae.

Amino acid concentrations ( $\mu\text{g/ml}$ )	Post-oviposition (days)											
	1 day	Change (%)	2 day	Change (%)	3 day	Change (%)	4 day	Change (%)	5 day	Change (%)	Average	Change R.C.(%)*
Aspartic acid	U** 3.913	-75	5.546	-78	73.920	+109	41.797	-68	123.912	-89	49.818	3.8
	P*** 0.986		1.196		154.269		13.347		13.869		36.733	1.5
Glutamic acid	U 128.767	-42	207.792	-25	401.941	+56	217.179	+8	338.461	+47	258.828	19.8
	P 74.587		155.328		628.017		452.357		497.208		361.499	14.5
Serine	U 53.993	-20	159.404	-19	140.422	+28	82.406	+70	112.781	+29	109.801	8.4
	P 43.210		129.404		180.244		140.500		145.327		127.796	5.1
Glycine	U 209.665	-13	72.439	+547	71.457	+682	46.931	+868	95.208	+1203	95.208	+440
	P 182.371		487.998		558.817		731.194		611.712		514.418	20.6
Alanine	U 200.740	-68	398.355	-20	311.801	-18	244.097	+124	229.058	+82	276.810	21.2
	P 64.359		318.005		254.184		302.090		416.562		271.040	10.8
Proline	U 419.599	-22	269.669	+280	192.739	+366	215.948	+466	227.724	+420	265.142	+251
	P 326.150		1024.767		897.868		1222.754		1185.012		931.310	37.2
Tyrosine	U 99.193	-3	164.869	+20	273.050	+25	281.149	-15	345.139	-2	232.680	+4
	P 96.626		198.325		342.554		238.321		339.860		243.137	9.7
Cysteine	U 10.138	+28	36.623	-54	23.301	+56	16.435	-36	16.416	-41	20.583	-16
	P 13.017		16.817		36.249		10.600		9.740		17.285	0.7
Total	U 1126.008	-29	1314.697	+77	1488.631	+105	1174.558	+165	1440.421	+54	1308.870	+91
	P 801.306		2332.136		3052.202		3111.163		2219.290		2503.218	

\*Relative concentration, \*\*Unparasitized, \*\*\*Parasitized. Change (%) =  $\frac{\text{Parasitized} - \text{Unparasitized}}{\text{Unparasitized}} \times 100$

haemolymph proteins of *Pieris rapae* parasitized as second instar larvae by *Apanteles glometatus*. Protein concentrations in both uninfected and infected hosts followed a cyclical pattern during host development, but parasitized larvae had somewhat higher concentrations during the late fourth and early fifth instars and significantly lower levels than uninfected insects during the remainder of larval development. The latter decline was coincident with fat body destruction. Thompson (1982) also demonstrated the effects of parasitization by *H. exiguae* on the total haemolymph protein concentrations of fourth instar *T. ni*. Parasitized larvae had significantly elevated haemolymph concentration immediately following parasitization. The higher concentrations were maintained until approx. 4 days post-infection at which time the concentration in unparasitized larvae surpassed that of infected individuals.

Investigations on alternations in electrophoretic protein patterns were performed in *Heliothis virescens* parasitized by *Cardiochiles migriceps* (Vinson and Barras, 1970) and in *Heliothis zea* parasitized by *Microplitis croceipes* (Barras *et al.*, 1972). In both studies it was concluded that the developing parasite utilized or destroyed certain haemolymph proteins while causing a shift in free amino acids to the bound state or the secretions of other proteins into the haemolymph.

In investigations such as above, the concentrations of haemolymph carbohydrate and protein in the host insects parasitized by larval parasitoids show mainly a tendency to increase. However, the concentrations of haemolymph carbohydrate and protein in the wax moth pupae parasitized by *B. lasus* show a tendency to decrease (Figs. 1 and 2). It is suggested that such decrease takes place because the host wax moth pupae parasitized is not feeding.

Amino acids may be of considerable importance in the nutritional and metabolic adaptations of insect parasite. Corbet (1968) reported that *N. canescens* spend a considerable portion of its larval life as a first instar larva and completes most of its larval development during the late last larval stadium of the host. The composition of the host haemolymph changed dramatically during this period. It was suggested that such changes after feeding behavior of the first instar larva, in-

creasing the rate of nutrient uptake and inducing growth. Among the change occurring in the host haemolymph was a sharp increase in free amino acid concentration.

In similar investigations, Barras *et al.* (1969) reported that following parasitization of *H. virescens* by *C. migriceps*, the concentration of certain free amino acid decreased while their levels increased in the peptide and protein fractions. The result, however, was not consistent with all the amino acids. Fisher and Ganesalingam (1970) demonstrated altered amino acid levels in *Anagasta kuehniella* one week after parasitization by *Devorgilla canescens*. Phenylalanine and leucine were absent in parasitized hosts while two unknown amino acids, which were absent in uninfected hosts, were reported at high levels in parasitized. Thompson (1986) also reported that following parasitization of *Trichoplusia ni* by *Hyposoter exiguae*, the concentration of total haemolymph glucogenic amino acid decreased.

*In vitro* studies demonstrated that amino acid alone can fulfill the energy requirements for complete larval development of a number of hymenopterous parasitoids (Thompson, 1976, 1983; Yazgan, 1972). The parasitoid *B. lasus* grew well *in vitro* on media in which nitrogen was supplied as a mixture of protein and free amino acids. Larvae failed to develop properly, however, on media containing protein alone. Studies on amino acid metabolism of *Exeristes roborator* further indicated the importance of amino acids for growth and development of insect parasitoids. A comparison of the rate of incorporation of carbohydrate precursor into amino acid with the synthesis rate of protein from amino acid indicated that the protein synthesized by amino acid synthesis. Thus, developing larvae require a source of amino acid for maximal growth and development, and a low rate of amino acid synthesis may represent an adaptation of the parasite, which feeds on host haemolymph, a rich source of amino acids (Thompson, 1982).

In this experiment, therefore, the concentration of increased haemolymph free amino acids in the wax moth pupae parasitized by *B. lasus* is thought to benefit to the developing parasitoid because of requiring a lot of free amino acid for maximal growth and development of the parasitoid (Fig. 3,

Tables. 1 and 2). But, it is not known yet whether the concentration of increased haemolymph free amino acids is due to the fat body destruction or haemolymph proteolysis in the host pupae.

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무늬수중다리좀벌(*Brachymeria lasus Walker*)에 의해 기생된 꿀벌부채명나방(*Galleria mellonella L.*)번데기의 혈림프 대사물질에서 단백질, 탄수화물, 유리아미노산의 변화  
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꿀벌부채명나방이 무늬수중다리좀벌에 의해 기생 되었을때 이 기주의 혈림프에서 대사물질의 변화 생겼다. 이 기생벌에 의해 기생된 기주와 기생되지않은 기주의 혈림프에서 총단백질과 총탄수화물의 농도를 비교하여보면, 기생된 기주에서 전체적으로 뚜렷이 감소하는 경향을 나타내었다. 그러나, 총유리아미노산은 기생된 후 첫날을 제외하고 기생된 기주에서 전체적으로 뚜렷이 증가하는 경향을 나타내었다. 그리고 유리아미노산 중에서, glycine, proline, arginine, isoleucine, phenylalanine, leucine, valine, glutamic acid, histidine, serine, tyrosine은 뚜렷이 증가하는 경향을 나타내었고, threonine, aspartic acid, methionine, cysteine, alanine, lysine는 약간의 감소하는 경향을 나타내었다.



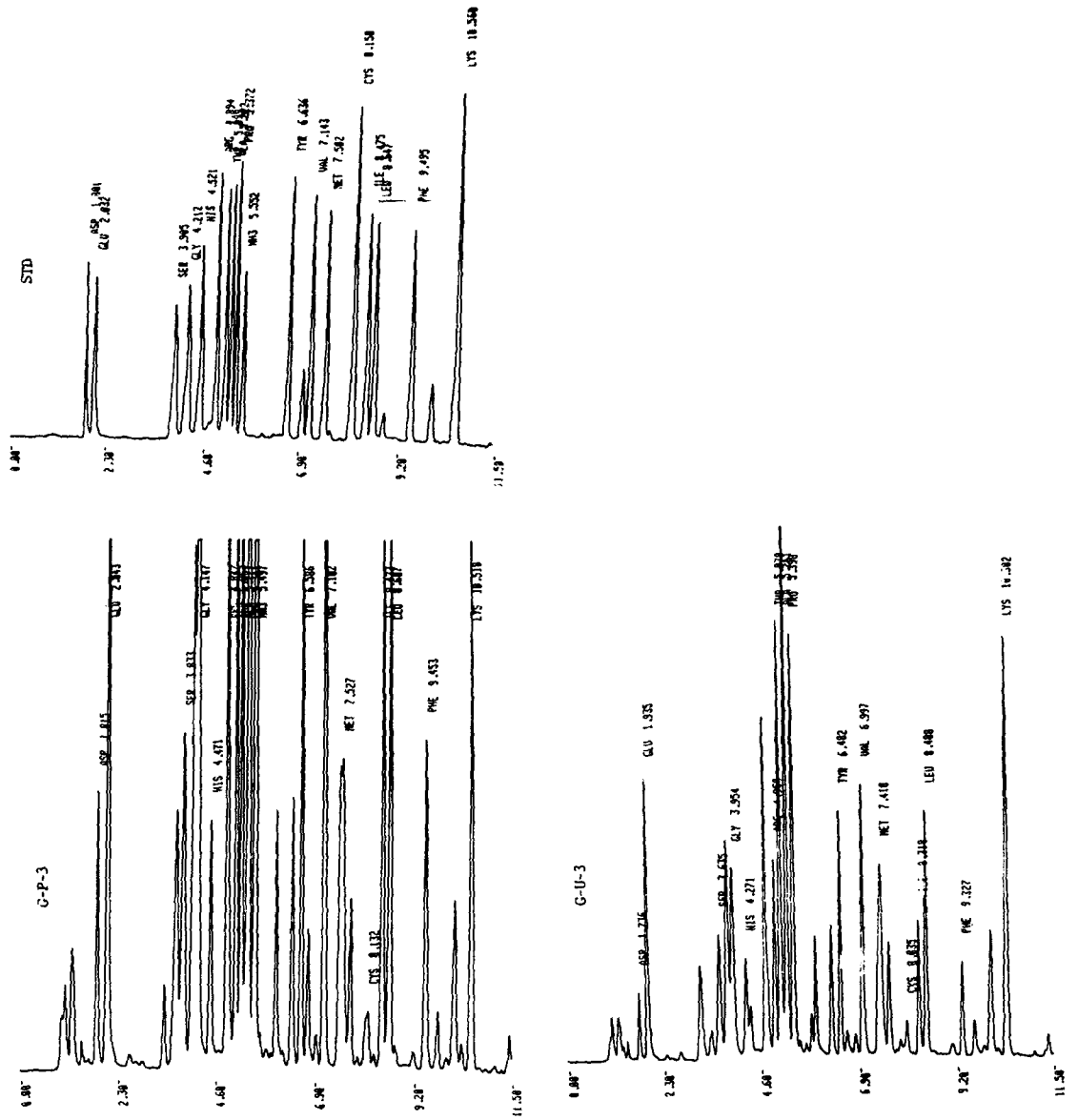


Plate. Chromatograms of free amino acids. STD: Standard, G: *Galleria mellonella*, P: Parasitized, U: Unparasitized, 3: Post-oviposition (days).