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Detection of glutathione S-transferase and activity levels of the esterases associated with insecticide resistance in field collected mosquito *Culex pipiens* (Diptera: Culicidae) from Egypt

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Abstract

Chemical insecticides are still the main part of vector mosquitoes control program. The effectiveness is eventually reduces as mosquitoes become resistant to them with repeated use. Therefore, understanding the mechanisms of resistance will be required. The present study was carried out to evaluate the potential biochemical mechanism in contributing toward insecticide resistance in Culex pipiens field populations in Egypt. Natural populations of Culex pipiens were collected from Giza, Beheira, Sohag and Monufia Governorates. The present finding showed difference in enzyme activity in Culex pipiens natural populations compared with laboratory colony. Monufia populations show the highest level of beta and alpha esterases (9.2 and 5.7 times higher activity than S-lab respectively). The most significant values of GST were in Sohag and Giza populations (3.8 and 2.4 times higher activity than S-lab respectively). A significant increase in specific carboxylesterases was detected only in Beheira populations (1.8 times higher activity than S-lab). Mechanisms of resistance are discussed in relation to biochemical assay results.

Keywords: Culex pipiens, resistance, esterases, GST, Egypt

1. Introduction

Mosquitoes are the most important vectors of human and animal diseases. In Egypt, the mosquito Culex pipiens is a main vector of lymphatic filariasis (Harb et al. 1993) [17]. It is also capable of transmitting Rift Valley fever Virus (Hoogstraal et al.1979) [22] and West Nile Virus (Taylor et al. 1956 and Soluman et al. 2010) [53, 49]. Its control is essential to prevent disease transmission.

The use of insecticides on adult and immature stages is efficient strategy for mosquito - borne disease control. Mosquitoes control big problem with regard to developing insecticide resistance. The most common mechanisms of resistance are the increase in metabolic detoxification of insecticides before it reaches its target site and changes in sensitivity of the target site (Hemingway et al. 1986 and Price 1991) ^[20, 37].

Three major enzyme families, esterases, glutathione Stransferase (GST) and Mixed Function Oxidases (MFO) such as cytochrome P450 oxidases are mainly involved in insecticides metabolism. Each of these enzymes catalyzes a wide range of detoxification reaction (Raymond et el. 1998; Scott 1999; Wheelock et al. 2005; Ranson and Hemingway 2005) [43, 46, 56, 40]. Measuring the activity of these enzymes in natural population is an important step in monitoring insecticide resistance. Therefore, the aim of the present work was to evaluate the levels of detoxification enzymes (non specific esterases, glutathione S – transferase and specific carboxylesterases) in natural populations and laboratory

2. Materials and Methods Mosquito samples

Natural populations of Culex pipiens were collected as larval stage from breeding sites of four Governorates, Giza, Beheira, Sohag and Monufia. Laboratory strain was obtained from Namru (S-lab).

Biochemical assay

Batches of 50 third instar larvae were homogenized in 1.5ml of ice-cold phosphate buffer, the homogenates were centrifuged at 4000 rpm for 30 min at 4°C using cooling centrifuge. The supernatant was used as enzyme samples. Total proteins were determined by the method of Bradford (1976) [6]. Protein reagent was prepared by dissolving 100mg of Coomassie Brilliant blue G-250 in 50 ml 95% ethanol. To this solution 100ml 85% (W/V) phosphoric acid were added. The resulting solution was diluted to a final volume of 1 liter. Sample solution (50µl) or for preparation of standard curve 50µl of serial concentrations containing 10 to 100µg bovine serum albumin (standard protein) were pipetted into test tubes. The volume in the test tube was adjusted to 1 ml with phosphate buffer (0.1 M, PH 6.6). Five millimeters of protein reagent were added to test tube and the contents were mixed either by inversion or vortexing. The absorbance at 595 nm was measured after 2 min. and before 1 hr. against blank prepare from 1 ml of phosphate buffer and 5 ml protein reagent

Nonspecific esterases (alpha and beta esterases) were determined according to Van Asperen (1962) [54] using αnaphthyl acetate or β-naphthyl acetate as substrates, respectively. The reaction mixture consisted of 5ml substrate solution $(3x10^{-4} \text{ M} \alpha \text{ or } \beta\text{-naphthyl acetate}, 1\%$ acetone and 0.1 M phosphate buffer, PH 7) and 20µl of larval homogenate. The mixture was incubated for exactly 15 min at 27°C, then 1ml of diazoblue color reagent (prepared by mixing 2 parts of 1% diazoblue B and 5 parts of 5% sodium lauryl sulphate) was added. The developed color was read at 600 or 555 nm for α- and β-naphthol produced from hydrolysis of the substrate, respectively. α-and β-naphthol standard curves were prepared by dissolving 20 mg α- or βnaphthol in 100ml phosphate buffer, PH7 (stock solution). Ten milliliters of stock solution were diluted up to 100ml by the buffer. Aliquots of 0.1, 0.2, 0.4, 0.8 and 1.6 ml of diluted solution (equal to 2, 4, 8, 16 and $32\mu g$ naphthol) were pipetted into test tubes and were completed to 5ml by phosphate buffer. One milliliter of diazoblue reagent was added and the developed color was measured as mentioned before.

Glutathione S-transferase (GST) catalyzed the conjugation of reduced glutathione (GSH) with 1-chloro 2, 4-dinitrobenzene (CDNB) via the –SH group of glutathione. The conjugate, S-(2, 4-dinitro-phenyl)-L-glutathione could be detected as described by the method of Habig *et al.* (1974) [15]. The reaction mixture consisted of 1ml of the potassium salt of phosphate buffer (PH 6.5), 100µl of GSH and 200 µl of larval homogenate. The reaction started by the addition of 25µl of the substrate CDNB solution. The concentration of both GSH and CDNB was adjusted to be 5mM and 1mM, respectively. Enzyme and reagents were incubated at 30°c for 5 min. The increment in absorbance at 340 nm was recorded against blank containing everything except the enzyme to determine the nanomole substrate conjugated / min. / larvae using a molar extinction coefficient of 9.6 /mM /cm.

Carboxylesterase activity was measured according to the method described by simpson $et\ al.\ (1964)^{[48]}$, using methyl n butyrate (MeB) as substrate. The reaction mixture contained 200µl enzyme solution, 0.5ml 0.067 M phosphate buffer (PH7) and 0.5ml MeB (4mM). The test tubes were incubated at 37°c for exactly 30 min. 1ml of alkaline hydroxylamine (equal volume of 2M hydroxylamine chloride and 3.5 M NaOH) was added to the test tubes. Then 0.5 ml of HCL (1 part of conc. HCL and 2parts of Δ H2O) was added. The mixture was shaken vigorously and allowed to stand for 2 min. 0.5ml of ferric chloride solution (0.9M Fecl3 in 0.1M Hcl) was added and mixed well. The decrease in MeB resulting from hydrolysis by carboxylesterases was read at 515nm.

Statistical analysis

One - way ANOVA revealed that the mean of all tested enzyme activities (P<0.001). The data were subjected to analysis of variance (ANOVA) with at least three repeats (Duncan 1955)^[14].

3. Results

In non-specific esterases (NSE) activity by using α or β naphthyl acetate, Monufia populations showed the highest level of beta and alpha esterases (9.2 and 5.7 times higher activity than S-lab strain respectively). Giza and Sohag populations exhibited a significant increase in α -esterases (1.7 and 1.3 times higher activity than S-lab strain respectively) and non-significant with β -esterases. Populations of Beheira did not show any significant elevation of alpha and beta esterases (Tables and Fig. 1 and 2).

Table 1: Alpha esterases activity in 3rd instar larvae of *Culex pipiens* in comparison with laboratory strain.

Natural populations	Alpha esterases (μg α – naphthol/min/mg protein)	
populations	Mean ± SD	Enzyme level
Giza	44 ± 3.2^{b}	1.7
Beheira	14 ± 1.07^{d}	0.5
Sohag	33 ± 1.05^{bc}	1.3
Monufia	149 ± 12.99^{a}	5.7
laboratory strain	26 ± 1.08^{cd}	0.0

Mean values followed by different litters are significantly different

 $Enzyme\ level = \frac{Enzymatic\ activity\ in\ field\ populations}{Enzymatic\ activity\ in\ laboratory\ colony}$

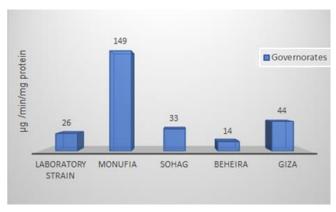


Fig 1: Alpha esterases activity for larvae of *Culex pipiens* populations.

Table 2: Beta esterases activity in 3rd instar larvae of *Culex pipiens* in comparison with laboratory strain.

Natural populations	Beta esterases $(\mu g \beta - naphthol/min/mg protein)$	
populations	Mean ± SD	Enzyme level
Giza	24.4 ± 1.6^{b}	1.5
Beheira	12.04 ± 1.5 b	0.8
Sohag	15.3 ± 0.87^{b}	0.96
Monufia	147 ± 13.6^{a}	9.2
laboratory strain	15.9 ± 1.04^{b}	0.0



Fig 2: Beta esterases activity for larvae of *Culex pipiens* populations.

In regard to glutathione S-transferases (GST), all populations revealed a significant increase as compared to S-lab except in Behrira populations did not show significant elevation. Sohag, Giza and Monufia populations showed 3.8, 2.4 and 1.4 times higher activity than S-lab strain respectively (Table and Fig. 3).

Table 3: Glutathione S-transferase activity in 3rd instar larvae of *Culex pipiens* in comparison with laboratory strain.

Natural	GST activity (µmol substrate conjucated/min/mg protein)		
populations	Mean ± SD	Enzyme level	
Giza	621 ± 72 ^b	2.4	
Beheira	245 ± 14^{d}	0.96	
Sohag	968 ± 21 ^a	3.8	
Monufia	361 ± 14^{c}	1.4	
laboratory strain	255 ± 8 ^d	0.0	

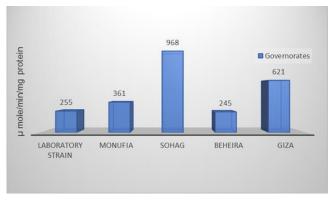


Fig 3: GST activity for larvae of Culex pipiens populations.

The activity level of specific carboxylesterases (CarE) was significantly increased in Beheira populations, 1.8 times higher than S-lab strain. Populations of Giza, Sohog and Monufia were nearly in the same range, the enzyme levels were 1.1, 1 and 0.98 respectively and did not show any significant differences from one to another compared to S-lab strain (Table and Fig. 4).

Table 4: Carboxylesterases activity in 3rd instar larvae of *Culex pipiens* in comparison with laboratory strain.

Natural populations	Carboxylesterases (μgMeb/min/mgprotein)		
	Mean ± SD	Enzyme level	
Giza	131 ± 6.2^{c}	1.1	
Beheira	218± 11.02b	1.8	
Sohag	120± 4 ^c	1	
Monufia	$114 \pm 3.6^{\circ}$	0.98	
laboratory strain	$120 \pm 4.5^{\circ}$	0.0	

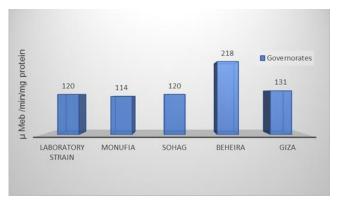


Fig 4: Carboxylesterases activity for larvae of *Culex pipiens* populations.

4. Discussion

Many organophosphate (OP) insecticides including, temephos, chlorpyrifos, pirimiphos-methyl and fenthion are most widely used in *Culex pipiens* control in Egypt, they are effective as larvicide for mosquitoes. In order to understand the resistance of *C. pipiens* to OP insecticides, We evaluated the levels of detoxification enzymes (non-specific esterases, glutathione S-transferase and specific carboxylesterases) in natural populations.

The major mechanism of OP insecticide resistance in *Culex* mosquitoes involves the elevation of one or more esterases. The esterases which preferentially hydrolyses α and β -naphthyl acetate are called A and B esterases respectively. Esterases A and B are coded by two closely linked gene Est-3 and Est-2 respectively (Pasteur *et al.* 1981) [³⁴].

The current study indicated that, the highest level of β and α -esterases in Monufia populations. The association between the activity of α and β esterases has been documented previously such as A2 and B2, A4-B4, A5-B5 and A8-B8 (Rivet *et al.*1993; Wirth *et al.* 1995; Qiao *et al.* 1998 and Wirth 1998) [44, 58, 38, 59]. A3-B3 has been described in *CX. Tarsalis* (Pasteur 1987) [36]. The occurrence of incidence might be due to co-amplification of two esterase genes (Est-2 and Est3 loci) which commonly found in organophosphate resistance (Rooker *et al.*1996 and Hemingway *et al.* 2004) [45,21]

Many previous studies observed the association between the alpha and beta esterases involved in the OP resistance in several Culex species. Resistance to fenthion in C. pipiens judged by its hydrolysis of alpha and beta-naphthyl acetates (Stone and Brown 1969) [51]. Two sets of over-produced esterases (A2-B2 and A4-B4), known to be involved in resistance to temephos in C. pipiens from Tunisia (Ben Cheikh and Pasteur 1993) [5]. In Martinique esterases A2 and B2 are known to confer a 100 fold resistance to temephos in C. p. quinquefasciatus (Yebakima et al. 1995) [60]. The A5 plus B5 demonstrated a significant level of temephos resistant in C.pipiens from Italy (Severini 1997) [47] and Cyprus (Wirth and Georghiou 1996) [57]. A moderate level of temephos resistance was observed in CX. quinquefasciatus from Barazil and biochemical profile of detoxification enzymes showed increased activity for α-esterase and βesterase in 30 and 17% respectively in strain resistance to temephos compared to susceptible colony (Amorim et al 2013)^[3]. The increase of detoxification by esterases was also involved in primiphos methyl resistance, overproduced esterases A and B among the Tunisia populations of C. pipiens resistance (Tabbabi et al 2017) [52].

The present investigation has also revealed that Giza and Sohag populations exhibited a significant increase in alpha esterase and non-significant with beta esterase. Similar observation by Pasteur et al 1981[35] who mentioned that ethanol formulating from chlorpyrifos hydrolysis could be demonstrated from esterase A but not for esterse B. Chlopyrifos resistance in *C. quinquefasciatus* from Californa and C. pipiens from France associated with a highly active detoxifying esterases encoded by the Est-3A gene (Raymond et al. 1986 and Raymond et al 1987) [41, 42]. In Egypt resistant strain of Pectinophora gossypiella to chlorpyrifos showed significant elevation in the alpha-esterase (Abo-El Seoud et al. 2005) [2]. Also the level of non-specific esterases(αnaphthol) significantly increase in both field populations and chlorpyrifos resistant strain of C. pipiens compared with laboratory colony in Egypt (Abd El-Samie and Abd El-Basset 2012) [1].

Glutathione S-trasferases (GST) enzymes are particularly important in the metabolism of OP insecticides, in which either the alkyl or aryl substituents on phosphorus can be cleaved. In O-dealkylation (methyl or ethyl groups) the GSH is conjugation with the alkyl portion of OP insecticides, while in the O-dearylation the GSH reacts with leaving group (Wei *et al.* 2001) ^[55]. Many workers have reported the involvement of GST in OP insecticides (Hayes *et al.* 2005 and Che-Mendoza *et al.* 2009) ^[18, 8].

The present investigation with regared to GST, all populations revealed a significant increase as compared to S-lab except in Beheira populations which did not show significant elevation. Results confirmed by previous studies, GST mediated detoxification in the chlorpyrifos resistance in

Cx quinquefasciatus (Liu et al. 2005) [29]. Also Daaboub et al. (2017) [11] showed that the increased detoxification by Est (and or GST) had a minor role in the chlorpyrifos resistance. The increased detoxification by Est (and or GST) was also involved in pirimphos methy resistance in C. pipiens of field populations from Tanisia (Tabbabi et al 2017) [52]. Although glutation greatly increased the demethylation of methyl termed O-dealkylation, or specifically demethylation, deethylation and involved the removal of only one of the alkyl groups, thus destroying the toxicity of fenthion in C. pipiens strain (Stone 1969) [50]. In CX quinquefasciatus, GST activity increased in temephos at low dose, whereas decreased significantly at high concentration (Muthusamy and Shivakumar 2015) [30]. Amorim et al. (2013) [3] suggested that the activity of GST was not significant between moderate level of resistance to temephos strain and susceptible colony of CX.quinquefasciatus. Similar observation for resistance strain of Ae. Aegypti to temephos showed increased for α and β esterases whereas the activity of GST was not significantly different from susceptible strain (Dhang et al. 2008; Muthusamy and Shirakumar 2015) [12, 31].

The most commonly observed change that has been linked to resistance development was the increase in the activity of carboxylesterases (Hemingway and Karunaratne 1998) [19]. Carboxylesterases were able to detoxify insecticides by catalyzing the hydrolysis of carboxyl ester, carboxy amide ester and carboxy thioesters and also certain phosphoric acid esters (Heymann and Jakoby 1980) [22]. For *Culex* mosquito this enzyme was referred to as A2 because of its preferential hydrolysis of α - over β - naphthyl acetat (Katterman *et al.* 1992) [27].

Our study detected the activity level of specific carboxylesterase, which was significantly increased only in Beheira populations. Many previous studies observed that *C. pipiens, C. quinquefasciatus* and other Culicidae, their OP-resistance was due to esterase isozymes which can break down OP compounds by phosphatas-type hydrolysis (Pasteur and Georghiou 1980) [33]. The alkaline phosphatases activites might be used as biomarkers in chlorpyrifos resistance in *C. pipiens* in Egypt (Abd El-Samie and Abd El- Basset 2012) [11]. Thionase activity was more important in the mechanism of resistance strain of *CX. P. fatigans* resistance to fenthion (Stone and Brown1969) [51]. Biochemical assay suggested a role for increased carboxylesterases activities in conferring temephos resistance in *Ae. Aegypti* larvae in Thailand (Pardin *et al.* 2014) [32].

The juvenods (JH) (such as methoprene and pyriproxyfen) with the benzoylphenyl urea chitin synthesis inhibitor (diflobenzuron) introduced a group of insecticides termed insect growth regulators (IGR). This IGR and Bti (*Bucillus thuringiesis*) are used as larvecidal in mosquitoes control in Egypt.

Specific and non specific esterases have been shown to hydrolyze JH to JH –acid in several tissues of different insects (Yu and Terriere 1978) [61]. The most abundant primary metabolite of JH in *Culex* and *Aedes* larvae was the hydroxyl ester (Quistad *et al.* 1975) [39]. Juvenile hormones analogues such as mothoprene have been identified as that possess an α and β unsaturated methyl ester at one end of the molecule and an epoxide at other. The methyl ester of JH is hydrolyzed by a JH –specific esterases (JHE) and the epoxid by JH epoxid hydrolas (JHEh) (Kamita and Hammock 2010) [26]. Hooper (1976) [24] concluded that esterases hydrolysing malathion and α - NA were similar to each other and were

distinct from those hydrolyzing JH. Downer *et al* (1975) ^[13] mentioned that the juvenoid methoprene could depress the activity of a non – specific esterase in *Aedes aegypti*.

On the other hand Ae. aegypt populations with resistance to the organophosphate temephos were less susceptible to pyriproxyfen, indicating a possible cross resistance in mosquitoes between these two insecticides families (Andrighetti et al. 2008) [4]. In Malaysia, Aedes larvae tolerance against temephos was reported and exhibited moderate resistance toward pyriproxyfen (Chen 2005) [9] but susceptible to diflobenzuron (Lau et al. 2015) [28]. Also Braga et al. (2005) [7] indicated a potential cross resistance when larvae were exposed to temephos and methoprene. Resistance to methoprene associated with activation of the esterases and mixed function oxidases were responsible for the metabolic resistance to conventional chemical insecticides, including OP insecticide. C.pipiens selected for mothoprene showed cross resistance to other juvenoids but not to diflubenzuron (Ishaay 2009) [25].

The effectiveness of Bti as a larvicide is an agent that does not cross resistance with motheprene (Cornel *et al.* 2002) ^[10]. The enzymatic activities of GST and carboxylesterases could be affected by Bti treatment. The activity of GST and carboxylesterases were increased after Bti treatment, but recovered to normal level quickly (Han *et al.* 2015) ^[16].

5. Conclusion

Our studies suggested that the difference in enzyme activity in *C.pipiens* natural populations could pose a threat for future control strategies. In addition, the early detection of resistance is essential for the successful implementation of insecticide resistance management strategies. These data will help monitoring and resistance management as a part of any program of vector control.

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