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Mini Review

The Karyotype of the Forensically Important Flesh Flies (Diptera: Sarcophagidae), Mini Review

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Abstract

Maggots of flesh flies were known as one of the main players in depletion of the earthily decomposed remains. Sarcophagidae colonized the carrion few minutes after death, and were known as the initial colonizers in several research papers. In consequence, they could be used in the minimum Post-Mortem Intervals (PMImini) estimation. Also, some sarcophagid species proved their capability in detecting toxins/ drugs in carcasses in the entomotoxicology field. The difficulty to identify sarcophagid females and larvae hindered their role as entomological evidence. Cytogenetics would aid in the identification of the immatures and their sex; the differentiation between adult species; and/ or the categorization of the taxa. Sarcophagidae received little attention concerning their karyotype measurements, especially in Egypt where there is only one unpublished thesis (El-Bassiony 2001). This mini review listed the previous studies and findings on sarcophagid's karyotypes, especially the solitary data from Egypt.

Keywords: Entomotoxicology; Forensic entomology; Wohlfahrtia; Sarcophaga

Introduction

The study of succession pattern of insects on carrion was the most important application of forensic entomology. Family Sarcophagidae enclosed over 2500 species [1] which fed on carrion including human bodies. Additionally, sarcophagid flies have considerable forensic importance to estimate PMImini [2,3]. While flies belonging to family

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Calliphoridae were known as the first insect's colonizers on carrion in most previous studies [4], other studies described Sarcophagidae as the initial and dominant colonizers on carrion in Egypt [5]. Calliphoridae was considered a key group for forensic entomologists due to the extensive biological, taxonomical, cytotaxonomical, and behavioral data, in contrary to Sarcophagidae which faced the lack of corresponding studies. The sarcophagid's identification was difficult and perplexing, as they were cryptic and/or morphologically similar.

Main Body

The diploid number of chromosomes in most sarcophagid species, was previously noted as 2n=12 [6-15]. However, 2n=19-20 and 2n=14 have been recorded for Pseudosarcophaga affinis [9], and Helicophagella melanura [16], respectively. The karyotypes of four sarcophagid flies, Sarcophaga surcoufi (Villeneuve), Sarcophaga dux (Thomson), Sarcophaga aegyptiaca (Salem), and Wohlfahrtia nuba (Wiedemann), were studied from three locations in Egypt [14], and W. nuba has reduced 2n=10 number of chromosomes (Table 1, Figure 1). This case has never been reported before in Sarcophagidae, except for Peckia (Squamatodes) ingens [17], but this case was found in many calyptrate Diptera such as Muscina stabulans (Muscidae), Haematobia irritans (Muscidae) [18], Stomoxys calcitrans (Muscidae) [19], and Cephalopina titillator (Oestridae) [20]. Differences in both chromosome size and number can result from fissions and fusions [21]. Chromosomal fusion, especially those which decreased the chromosome number, was explained lately as a trend of karyotype evolution in many organisms [22]. Also, evolutionary turnover can convert autosomes to sex chromosomes and revert them back through the species diverge [23,24].

Several authors described the differences between the designations of autosomal pairs and sex pair in several Sarcophagidae (Table 1) [9,10,13,14]. The sex pair showed XX designation in females, and XY in males [25]. Three metacentric and two submetacentric autosomal pairs were observed in Pattonella intermutans (Sarcophagidae), and also the X and Y chromosomes were described as subtelocentric and submetacentric respectively [13]. In the four Egyptian species (Figure 1), both S. surcoufi and S. dux have four metacentric pairs of chromosomes and two submetacentric chromosomes [14]. On the other hand, S. aegyptiaca chromosomes were all metacentric except for the third autosome. In the same study, the autosomes of W. nuba have metacentric designation, while X and Y chromosomes were submetacentric and telocentric, respectively [14]. While the autosomal morphology of Sarcophagidae was more or less similar in the different species (Table 1), the sex chromosomes were heterochromosomes and species specific [10]. The sex chromosomes were greatly varied in size and morphology among species within several genera. Nine sarcophagid flies have acrocentric sex chromosomes [9], three Parasarcophaga species have metacentric and telocentric ones [12], two Parasarcophaga [12], and nine Boettcherisca species [15] have small dot-like chromosomes. The large verity in the size and morphology of the sex chromosomes might be a result of heterochromatin accumulation or deletion, where the changes in the amount and distribution of heterochromatin resulted mainly in the evolution of the karyotype of a species [12,26].

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Species	Chromosome number							
	I	п	ш	IV	v	x	Y	TCL (µm)
Parasarcophaga argyrostoma [11] Arm ratio Designation	1.15 m	1.81 sm	1.82 sm	1.17 m	1.24 m	NA t	NA (15)* t	NA
Parasarcophaga argyrostoma [12] length Arm ratio Designation	1.19 m	1.82 sm	1.89 sm	1.18 m	1.33 m	4.9±0.24 NA NA	3.2(7~10)* NA NA	48.8±2.80
Parasarcophaga misera [12] length Arm ratio Designation	1.67 m	1.97 sm	2.01 sm	1.31 m	1.44 m	0.9±0.06 NA NA	0.5(5~10)* NA NA	48.9 ±3.11
Parasarcophaga albiceps [12] length Arm ratio Designation	1.17 m	1.19 m	1.09 m	1.16 m	1.63 m	1.2±0.01 NA NA	NA NA NA	51.5 ±1.27
Parasarcophaga ruficornis [12] length Arm ratio Designation	1.44 m	1.39 m	1.57 m	1.24 m	1.19 m	8.9±0.55 NA NA	6.3(5~10)* NA NA	64.3 ±1.19
Parasarcophaga knabi [12] length Arm ratio Designation	1.18 m	1.20 m	1.17 m	1.15 m	1.19 m	13.3±0.53 1.28 m	9.7(10)* 1.15 m	48.9 ±2.10
Pattonella intermutans [13] length Arm ratio Designation	5.3 1.23 m	4.5 1.91 sm	3.9 1.45 m	3.8 1.75 sm	2.9 1.16 m	7.7 3.32 st	3.4(10)* 2.1 sm	NA
Sarcophaga surcoufi [14] Length ± S.E. Arm ratio Designation	6.06± 0.17 1.67 m	5.68 ± 0.16 1.15 m	5.31 ± 0.15 1.74 sm	5.01 ± 0.17 1.83 sm	4.66 ± 0.14 1.15 m	5.83±0.21 1.17 m	4.53 ± 0.19(23~50)* 1.28 m	26.70 ± 0.74
Sarcophaga dux [14] length ± S.E. Arm ratio Designation	5.00 ± 0.19 1.45 m	4.77 ± 0.16 1.22 m	4.49 ± 0.15 1.74 sm	4.27±0.12 1.75 sm	4.01±0.12 1.23 m	5.07 ± 0.20 1.33 m	$\begin{array}{c} 3.77 \pm 0.14 (25 {\sim} 50)^{*} \\ 1.50 \\ m \end{array}$	22.52 ± 0.72
Sarcophaga aegyptiaca [14] Length ± S.E. Arm ratio Designation	5.39 ± 0.20 1.52 m	4.91 ± 0.20 1.26 m	4.63 ± 0.18 1.79 sm	4.30 ± 0.17 1.65 m	4.08 ± 0.17 1.28 m	4.82 ± 0.20 1.47 m	3.80 ± 0.14(21~50)* 1.39 m	23.24 ± 0.90
Wohlfahrtia nuba [14] Length ± S.E. Arm ratio Designation	$\begin{array}{c} 4.17 \pm 0.12 \\ 1.57 \\ m \end{array}$	3.76 ± 0.09 1.59 m	3.50 ± 0.06 1.32 m	3.22 ± 0.10 1.39 m		4.81±0.07 2.06 sm	$3.58 \pm 0.07(23 \sim 50)^{*}$	14.63 ± 0.25

Table 1: Mean length (µm), arm ratio and designation of chromosomes of somepublished and unpublished data of sarcophagid flies' karyotypes [1-14].

TCL=mean total haploid autosomal complement length,

NA=not applicable,

m=metacentric chromosome,

sm=submetacentric chromosome,

st=subtelocentric, and t= telocentric chromosome.

*Number of Y chromosomes measured~ to the total number of chromosomes measured, in each species, in brackets.

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Figure 1: Neuroganglial chromosomes of four sarcophagid species [14].

- A: Female and B: male Sarcophaga surcoufi,
- C: Female
- D: male Sarcophaga dux,
- E: Female
- F: male Sarcophaga aegyptiaca

G: Female and H: male Wohlfahrtia nuba. Arrows show X and Y chromosomes. Bar=5 $\mu m.$

The previous studies on genome size evaluated the phylogenetic relations in the taxonomy field [27-29]; estimated the completeness of genomes [30]; and helped in assessing genetic linkage maps [30]. Genome size was usually considered constant within species, and could vary widely between closely related species [22]. Limited intraspecific variation was an assumption in measurement and comparison of genome sizes. The inter- and intraspecific genome size variation might be due to differences in heterochromatin content and the amount of repetitive DNA in the autosomal complement [31]. The average Total haploid Complement Lengths (TCL) of species belonging to the genera *Pseudosarcophaga*, *Sarcophaga*, *Kellymyia* and *Protodexia* were recorded in 1953 [9]. Also, the karyotype measurements of *S. surcoufi* exhibit significantly longer chromosome lengths, and longer TCL than *S. dux*, *S. aegyptiaca* and *Wohlfahrtia nuba* [14].

Conclusion

The genetics analyses could provide cue for the chromosome's evolution in sarcophagid flies, and could supply information which is of both forensic significance and evolutionary interest.

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