

Systemic orchestration of cell size throughout the body: Influence of sex and rapamycin exposure in *Drosophila melanogaster*

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Supplementary material

Developmental time

To obtain a group of the same age for histological analysis, vials of developing flies from the second generation were checked daily. This procedure also allowed us to count the days from the day the parental flies were placed in the vial to the appearance of the first adults in the vial, which served as a rough measure of the duration of development of each isoline under a given treatment, without distinguishing between sexes (N=28). Similar to thorax length and cell sizes (see Methods of the main work), developmental times (days) were analysed with a general linear mixed model (GLMM) that included treatment (rapamycin vs. control) as a fixed factor and isoline as a random factor. To increase homogeneity of variance, developmental times were rank-transformed for the analysis. Results of the analysis are shown in Supplementary tables (Table 1S).

Histological procedures and cell size measurements

We used a stereomicroscope (Olympus SZX12, Olympus, Tokyo, Japan) to dissect flies and measure thorax length. We used a light microscope (Eclipse 80i, Nikon, Tokyo, Japan), camera

(Axio Cam MRc5, Zeiss, Oberkochen, Germany) and ZEN software (ver. 2011, ZEISS, Oberkochen, Germany) to image the wings, legs, ommatidia, flight muscles (with bright-field microscopy) and Malpighian tubules (with phase contrast objectives).

Flies were dissected using a microtome knife and forceps. After dissection, wings were preserved at -20°C in Eppendorf tubes. For measurements, wings were mounted on microscopy slides using ST Ultra and CV Ultra (Leica Biosystems, Nussloch, Germany) and digitalized under 20× objective magnification. According to Dobzhansky [1], each trichome represents one wing epidermal cell. Thus, we calculated the mean wing epidermal cell size (μm^2) from the ratio of defined wing area and the number of corresponding trichomes. Based on previous methods [2,3], we counted trichomes on the dorsal wing blade in a 0.031-mm² circle placed between the cubital and distal veins with the help of two macros embedded in ImageJ software (National Institutes of Health (NIH), Bethesda, MD, USA).

Heads were preserved with 1 ml of 20% methanol (POCH, Gliwice, Poland). Ommatidia, which form compound eyes in *Drosophila*, are composed of a fixed number of cells [4]. Thus, based on previous studies [5-7], we used their size as a proxy of cell size. To conduct the measurements, both eyes were covered with a transparent 5SecondFix UV glue and dried with a UV lamp (Ontel Products, Fairfield, USA). Such prepared replicas were removed from the eyes and flattened on a slide by incising in three places. Next, a drop of glycerine (POCH) was added, and the slide was covered with a coverslip. On the top of each slide, fishing weights (100 g) were placed for 24 h to allow flattening of replicas, and then the coverslip edges were covered with colourless nail polish (Vipera, Kraków, Poland). Prepared slides were digitalized under 10× objective magnification. We estimated the mean ommatidium size (μm^2) using automatized methods [5] integrated into ImageJ (NIH).

Legs were preserved with 1 ml of 70% ethanol (Lineal Chemicals, Warszawa, Poland). For epidermal cell measurements, legs were placed individually on microscopic glass slides to obtain photos of the 8th row of the basitarsus under 20× objective magnification. Using ZEN software (Zeiss), we measured the distance (μm) between the first and last bristles of the 8th row (based on Hannah-Alava nomenclature [8]) and counted the total number of bristles. According to Held [9], the bristle number in a row and cell number along the basitarsus are coordinated as are the bristle interval and cell diameter. Therefore, to calculate the proximal cell size (μm), the measured distance was divided by the bristle number in the row. For consistency with other cell types, linear measures of leg cells were squared (μm^2).

During dissection, the Malpighian tubules were taken with two entomological pins (Paradox, Kraków, Poland) from the severed abdomen on the microscopic glass with a drop of PBSTX (mixture of phosphate-buffered saline (PBS; Sigma Aldrich, Saint Louis, USA) and Triton X-100 (POCH)). After preparation, slides were covered with a coverslip, and Malpighian tubules were digitalized under 20× magnification with a phase contrast objective. We estimated the proxy of cell size forming Malpighian tubules (μm^2) from the density of cell nuclei/nucleoli in a known area of the tubule. We outlined the area of Malpighian tubules using ImageJ software with a LiveWire Plugin (NIH) and counted the number of cell nuclei/nucleoli in this area. We carried out measurements in five to ten photos for each individual.

Dissected thoraxes were used to measure dorsal longitudinal muscle cells, a subset of indirect flight muscle cells. According to the modified methodology of Barbosa et al. [10], to fix this tissue, thoraxes were placed in Duboscq–Brasil's solution for 2 h. Within this time, Eppendorf tubes with thoraxes were placed in a beaker with water standing on a hot plate (50 °C) for 10 minutes. Next, thoraxes were washed in PBSTX, dehydrated in 70%, 80%, 90% and 96% ethanol (Linegal Chemicals), butanol (Chempur, Piekary Śląskie, Poland) and isopropanol (Leica), cleared in pure Clearene (Leica), then in a mixture of Clearene (Leica) and Paraplast Plus (Leica) and finally embedded in Paraplast Plus (Leica). Thoraxes were cut with a rotary microtome (Hyrax M55, Zeiss) into 6- μm thick cross-sections. The slides were deparaffinized in Clearene (Leica), rehydrated in isopropanol (Leica) and 96%, 90%, 80%, 70% and 50% ethanol (Linegal) and distilled water, and stained with Ehrlich haematoxylin (Carl Roth) and then with Gömöri trichrome (BioOptica, Milano, Italy). Then, the slides were washed in 0.2% acetic acid (POCH), dehydrated in 96% ethanol (Linegal), cleared in a mixture of phenol (Chempur) and xylene (POCH) and then in pure xylene (POCH), and embedded in CV Mount (Leica). We analysed muscle tissue under 20× objective magnification and photographed one bundle from each of the six pairs of muscle bundles. Using ImageJ (NIH), we outlined the area of individual muscle cells (fibres) (μm^2).

Overall, each isoline and sex was represented by two individual flies per each treatment. Each fly was characterized by the mean cell size in each tissue type. The means were calculated from the following number of cell-size related structures: thorax – from 26 to 71 muscle cells, wings – from 143 to 273 trichomes, eyes - from 88 to 548 ommatidia, legs – from 11 to 16 bristles, Malpighian tubules – from 78 to 338 nuclei/nucleoli.

References

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Supplementary tables

Table S1

Comparison of development time in *Drosophila melanogaster* fed a diet with or without rapamycin with GLMM. Developmental times (days) were rank-transformed before analysis. The results show that rapamycin supplementation delayed development by ~12%. The percentage effect size was estimated based on the model predictions for each treatment group, after back-transforming the ranks to units of development time (days).

Fixed effects	Coefficients	t	P
Intercept (control)	9.214	5.879	<0.0001
Treatment (rapamycin)	10.571	5.518	<0.0001
Random effects			
Variance estimates			
Isoline (intercept)		8.70	
Residual		25.69	

Table S2

GLMM comparisons of adult *Drosophila melanogaster* after feeding diets with and without rapamycin. Models include thorax length (mm) as a continuous covariate. N indicates number of flies.

	Cell size: thorax muscle (μm^2)		Cell size: wings (μm^2)		Cell size: ommatidia (μm^2)		Cell size: legs (μm^2)		Cell size: Malpighian tubules (μm^2)	
	N=111		N=112		N=109		N=112		N=56	
Fixed effects	t	P	t	P	t	P	t	P	t	P
Intercept (control female)	1.0	0.3367	2.2	0.0336	8.4	<0.0001	0.7	0.5027	-1.0	0.3406
Thorax length	1.9	0.0582	8.4	<0.0001	5.3	<0.0001	3.3	0.0015	2.9	0.0057
Treatment (rapamycin)	-2.0	0.0508	-1.0	0.3097	-4.6	<0.0001	0.4	0.7126	-4.9	<0.0001
Sex (male)	-1.0	0.3126	-5.7	<0.0001	-1.9	0.0586	-0.2	0.8383		
Random effects										
variance estimations										
Isoline (intercept)	213.6		153.3		45.4		1901		0	
Residual	2520.6		58.8		68.6		2412		25251	