



# THE USE OF DROSOPHILA MELANOGASTER AS A MODEL ORGANISM TO STUDY THE EFFECT OF INNATE IMMUNITY ON METABOLISM

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## Abstract

Apart from its traditional role in disease control, our study provides evidence that suggest a role of the JAK/STAT innate immune pathway in regulating metabolic homeostasis. Owing to the importance of this “immune-metabolic alignment” in dictating a state of health or disease, a proper mechanistic understanding of this alignment is crucial in opening up for promising therapeutic approaches against a broad range of chronic, metabolic, and inflammatory disorders like obesity, diabetes, and inflammatory bowel diseases. Since the JAK/STAT pathway is evolutionary conserved between invertebrates and vertebrates, our potential findings in the fruit fly serves as a platform for further immune-metabolic translational studies in more complex mammalian systems including humans.

## OBJECTIVES

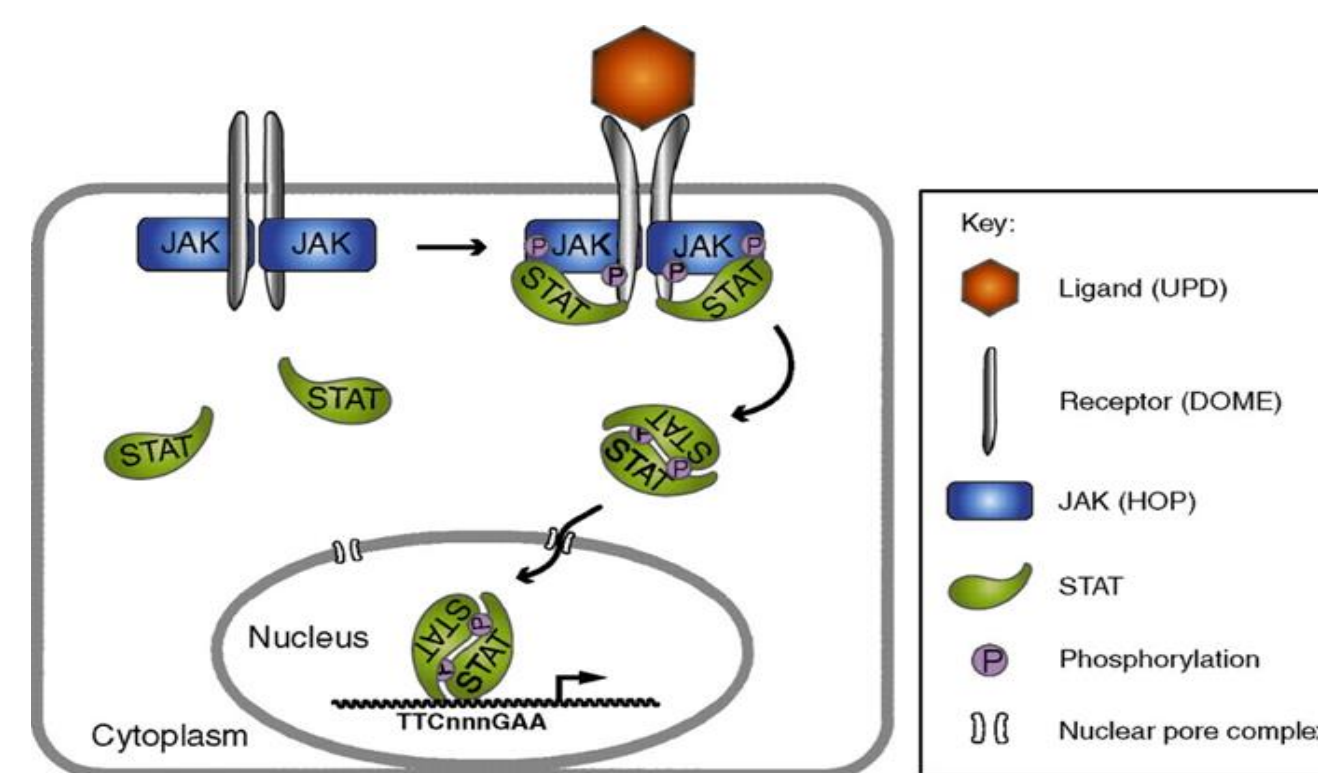
- Compare the body weight of *dome*<sup>G0282</sup> and *stat92E*<sup>EY10528</sup> mutant flies to *yw* control flies.
- Score for normal lipid storage and distribution in the fat body and gut of *yw*, *dome*<sup>G0282</sup>, and *stat92E*<sup>EY10528</sup> flies.
- Assess systemic glucose and triacylglyceride levels in *yw*, *dome*<sup>G0282</sup>, and *stat92E*<sup>EY10528</sup> flies.
- Detect relative gene expression of different metabolic peptide hormones (*Tk*, *AstC*, *DH31*) in *yw*, *dome*<sup>G0282</sup>, and *stat92E*<sup>EY10528</sup> flies.

## Methodology

- *Drosophila melanogaster* husbandry and maintenance
- Organ dissection and fluorescence microscope.
- Nutritional assays (Glucose and Triacylglyceride ).
- Relative gene expression of metabolic peptide hormone
- *Acetobacter* and *Lactobacillus* sp. colony forming unit (CFU) assay.
- Quantification and Statistical Analysis.

## RESULTS

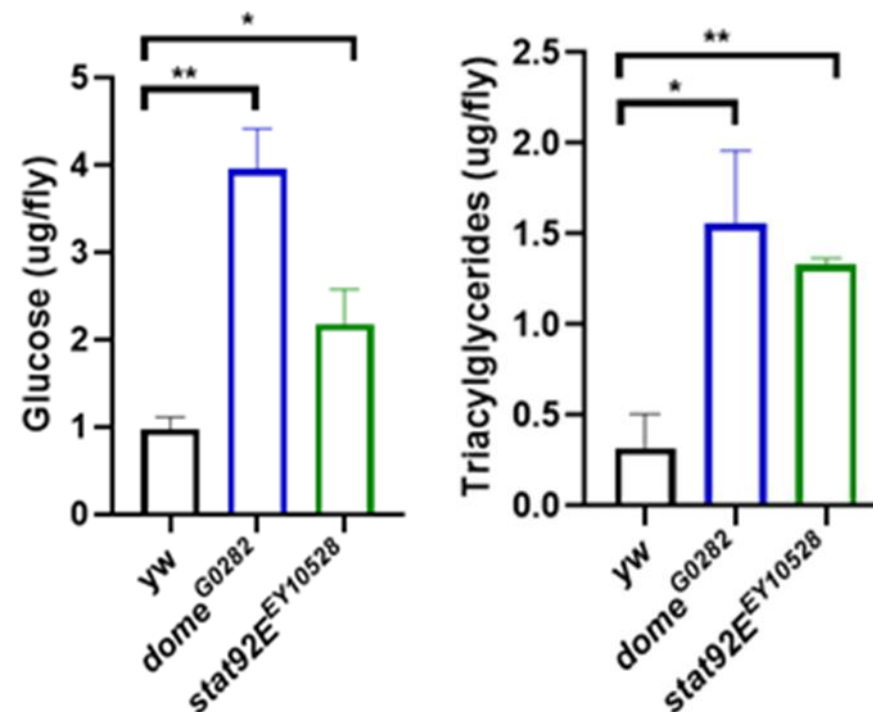
### The JAK/STAT Signaling Pathway



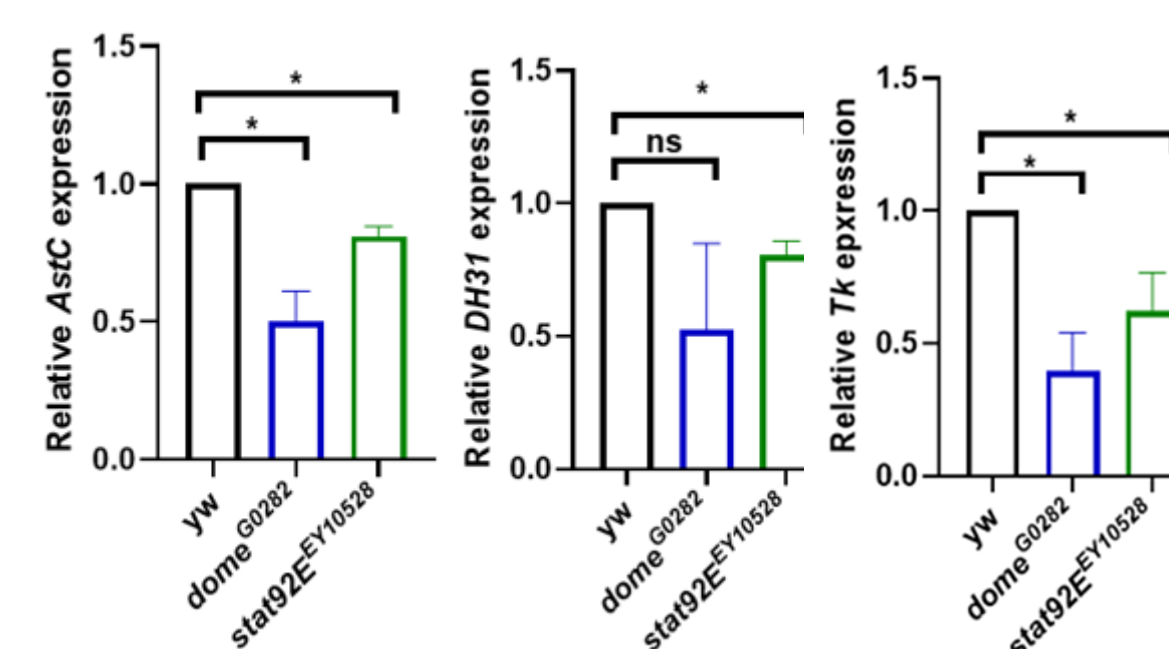
### Gain of body weight in JAK/STAT pathway mutants



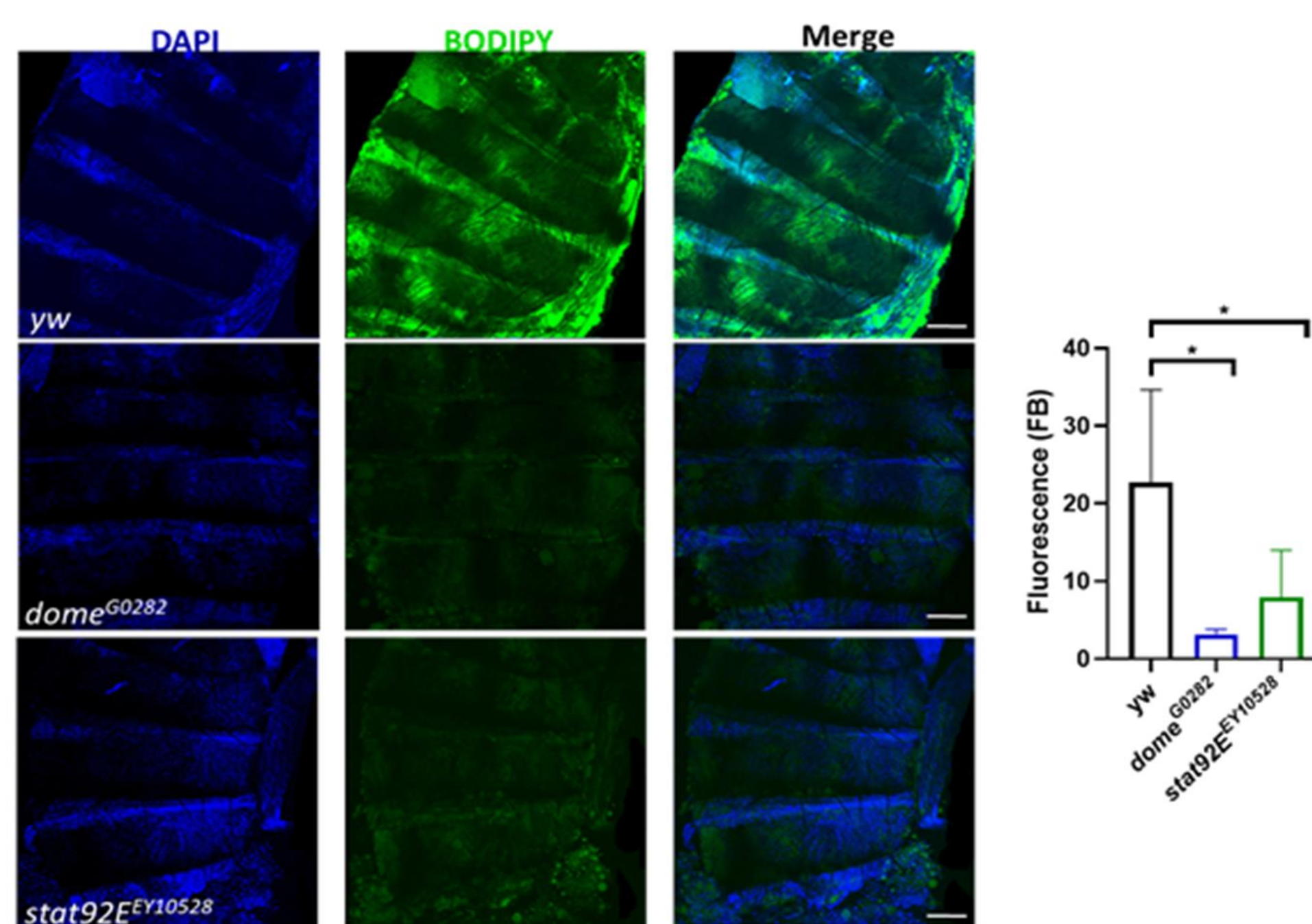
### Elevation in glucose and triacylglyceride levels in JAK/STAT Pathway mutants



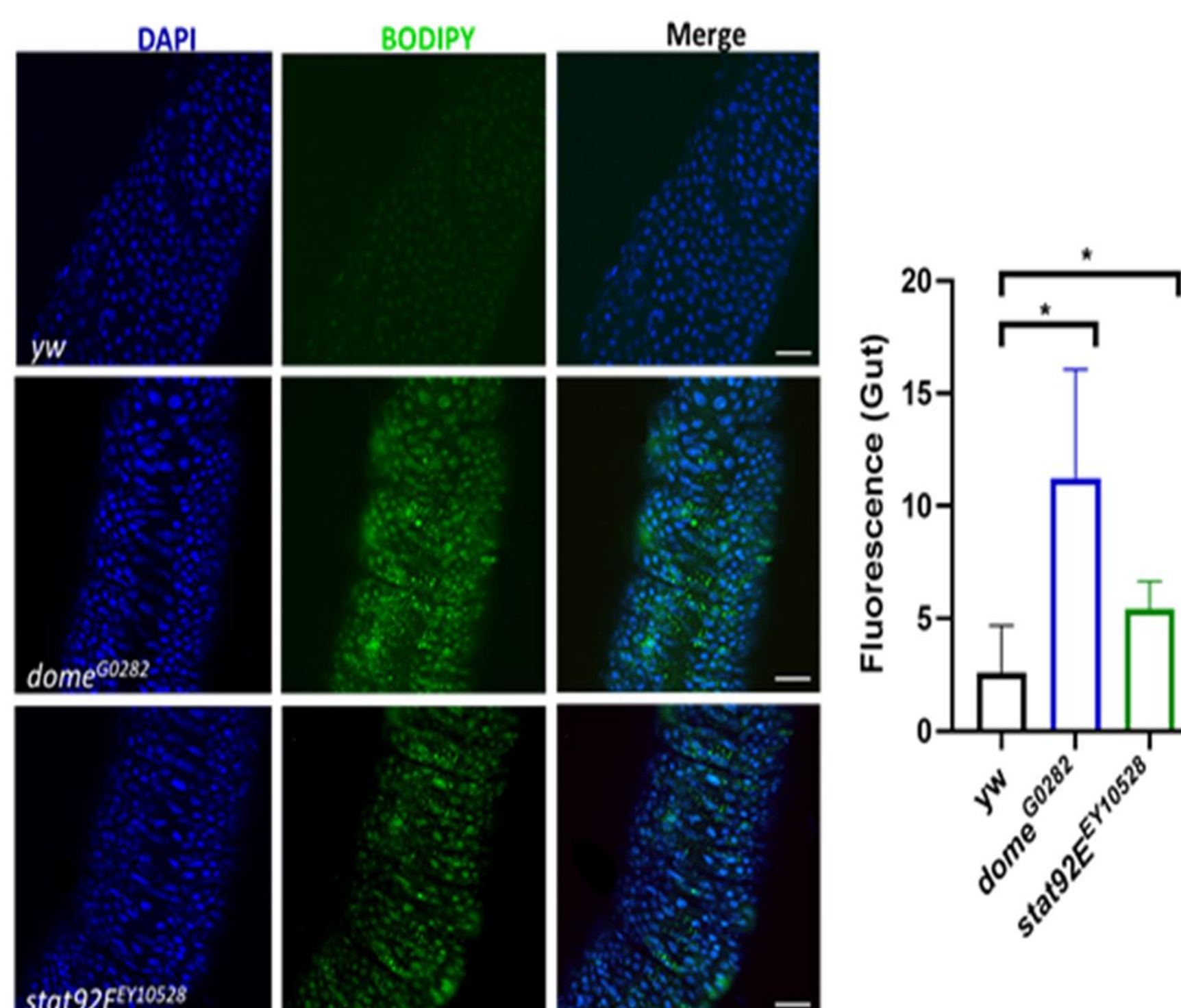
### Down-regulation of metabolic peptide hormones expression in JAK/STAT Pathway mutants



### Lipid depletion from the fat body of JAK/STAT pathway mutants



### Irregular lipid accumulation in the gut of JAK/STAT mutants



## Conclusion

Blocking JAK/STAT signaling alters the metabolic profile of mutant flies. Both *dome*<sup>G0282</sup> and *stat92E*<sup>EY10528</sup> mutants had increased body weight, lipid deprivation in their fat body (normal lipid storage organ), irregular accumulation of lipid droplets in the gut, significant elevation of glucose and triacylglyceride levels, and significant decrease in the relative gene expression of different peptide hormones (*Tk*, *AstC*, and *DH 31*) known to regulate metabolic homeostasis in flies.

## ACKNOWLEDGMENT

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