

## Overexpression of grappa encoding a histone methyltransferase enhances stress resistance in *Drosophila*

Submitted by Olivier List on Mon, 10/19/2015 - 10:17

Titre	Overexpression of grappa encoding a histone methyltransferase enhances stress resistance in <i>Drosophila</i>
Type de publication	Article de revue
Auteur	List, Olivier [1], Togawa, Toru [2], Tsuda, Manabu [3], Matsuo, Takashi [4], Elard, Loic [5], Aigaki, Toshiro [6]
Pays	Su�de
Editeur	Mendelska s�llskapet
Ville	Lund
Type	Article scientifique dans une revue � comit� de lecture
Ann�e	2009
Langue	Anglais
Date	Jan-02-2009
Num�ro	1
Pagination	19-28
Volume	146
Titre de la revue	Hereditas
ISSN	0018-0661

R sum  en anglais

Histone deacetylases, such as silent information regulator 2 (Sir2) and Rpd3 are involved in chromatin silencing and implicated in lifespan determination in several organisms. The yeast Dot1 gene encoding a histone methyltransferase affects localization of silencing proteins including Sir2, and plays an essential role in the repair of damaged DNA. However, it is not known whether an alteration of a histone methyltransferase activity influences lifespan or stress resistance, which is often associated with extended lifespan. Here we investigated whether the *Drosophilagrappa* (*gpp*) gene, a Dot1 homolog influences lifespan and stress resistance using transgenic flies overexpressing *gpp* and those bearing a partial loss-of-function mutation. Overexpression of *gpp* throughout the adult stage did not extend the lifespan, but significantly enhanced resistances when they were kept on medium containing 1% H<sub>2</sub>O<sub>2</sub>, or those with poor nutrients. As well, *gpp*-overexpressing flies were behaviourally more active than control flies. We investigated whether *gpp* overexpression induced anti-oxidant genes, Catalase, Sod, Sod2, GstD2, *dhd*, TrxT and Trx-2. However, none of these genes was induced. A partial loss-of-function mutations in *gpp* dramatically reduced the lifespan under oxidative and caloric stresses. Taken together, these results demonstrated that *gpp* is required for normal lifespan and stress resistance, and that its overexpression increases stress resistance in *Drosophila*, without obvious induction of representative anti-oxidant genes.

URL de la notice <http://okina.univ-angers.fr/publications/ua14120> [7]

DOI 10.1111/j.1601-5223.2008.02080.x [8]

## Liens

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Publié sur *Okina* (<http://okina.univ-angers.fr>)