

Suppression of him-14(it44ts) by a transgene insertion expressing GFP::COSA-1

Chloé Girard, Chantal C. Akerib, Anne M. Villeneuve

▶ To cite this version:

Chloé Girard, Chantal C. Akerib, Anne M. Villeneuve. Suppression of him-14(it44ts) by a transgene insertion expressing GFP::COSA-1. microPublication Biology, 2021, 2021, pp.000430. 10.17912/micropub.biology.000430. hal-03852730

HAL Id: hal-03852730

https://hal.science/hal-03852730

Submitted on 15 Nov 2022

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers. L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Suppression of *him-14(it44ts)* by a transgene insertion expressing GFP::COSA-1

Chloé Girard^{1,2}, Chantal C Akerib¹ and Anne M Villeneuve^{1§}

Abstract

Meiotic crossover formation requires the activity of multiple pro-crossover factors, including the MutSy complex and the cyclin-related protein COSA-1, that become concentrated together at the sites of crossover recombination intermediates. Here we show that a transgene insertion expressing GFP::COSA-1 can suppress the crossover deficit caused by a partial reduction in MutSy function. Our data, combined with previous findings, support a model in which COSA-1 promotes crossover formation, at least in part, through positive regulation of MutSy function.

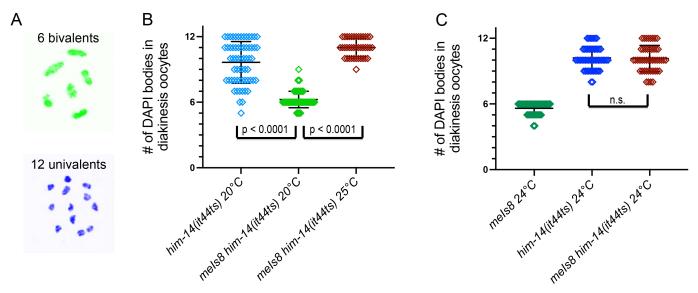


Figure 1. Suppression of the him-14(it44ts) crossover deficit at semi-permissive temperature by the GFP::COSA-1expressing transgene insertion meIs8: A) Example oocyte karyotypes (modified from Zalevsky et al. 1999) illustrating a normal wild-type karyotype (top) with all 6 pairs of homologous chromosomes connected by chiasmata (6 bivalents), and a karyotype from a recombination-deficient mutant (bottom) with 12 achiasmate chromosomes (12 univalents). B and C) Graphs quantifying numbers of resolvable DAPI-stained bodies in diakinesis oocyte nuclei; each diamond represents a nucleus, and mean and standard deviation are indicated. him-14(it44ts) is a temperature-sensitive mutation affecting the HIM-14/MSH-4 subunit of the MutSy complex; meIs8 is a transgene insertion expressing a GFP::COSA-1 fusion protein. Fixation and DAPI staining were conducted as in Bessler et al. 2007, and numbers of DAPI bodies were counted in oocyte nuclei in the -1 to -3 positions. For the 20°C data, worms raised at the semi-permissive temperature of 20°C were fixed at 24 h post L4; for the 24° and 25°C data, worms were raised at 20°C until the L4 stage, then were shifted to 24° or 25°C for 24 h before fixation. (Note that this assay tends to underrepresent the actual numbers of DAPI bodies present, as some chromosomes may be too close together to be resolved unambiguously.) Genotypes and numbers of nuclei scored: B) him-14(it44ts) unc-4 20°C, n = 62; meIs8 him-14(it44ts) unc-4 20°C, n = 65; meIs8 him-14(it44ts) unc-4 25°C, n = 40; C) meIs8 24C°, n = 63; him-14(it44ts) 24° C, n = 53; $meIs8\ him-14(it44ts)\ 24C^{\circ}$, n = 47. p-values are for two-sided Mann Whitney tests. Because of the temperaturesensitive nature of him-14(it44ts), all plotted data for each graph are derived from cohorts of worms raised, fixed and stained in parallel in the same experiment. The finding that meIs8 suppresses the him-14(it44ts) CO deficit at 20°C has been replicated in at least four independent experiments. The findings that meIs8 him-14(it44ts) unc-4 is severely defective in CO formation at 25°C and that suppression of him-14(it44ts) by meIs8 is abrogated at 24C° have each been replicated in at least two independent experiments.

Description

¹Departments of Developmental Biology and Genetics, Stanford University School of Medicine, Stanford, CA 94305, U.S.A.

²current affiliation: Université Paris-Saclay, CEA, CNRS, Institute for Integrative Biology of the Cell (I2BC), 91198, Gif-sur-Yvette, France

[§]To whom correspondence should be addressed: annev@stanford.edu



8/24/2021 - Open Access

Formation of crossovers (COs) during meiosis depends on the coordinated activities of multiple pro-CO factors, including the MutSγ complex and COSA-1, that become concentrated at the sites of CO-designated recombination intermediates in an interdependent manner (Yokoo *et al.* 2012; Woglar and Villeneuve 2018). The MutSγ complex is a heterodimer comprising the conserved proteins MSH-5 and MSH-4 (aka HIM-14, encoded by the *him-14* gene) (Zalevsky *et al.* 1999; Kelly *et al.* 2000). COSA-1 is a member of the cyclin superfamily and is hypothesized to function as part of a cyclin-dependent protein kinase (CDK) complex (Yokoo *et al.* 2012). The MutSγ complex was suggested to be a potential target for this hypothesized COSA-1 dependent CDK activity, as MSH-5 has multiple potential CDK phosphorylation sites and can be phosphorylated *in vitro* by a mammalian CDK. These and other data led to a model in which COSA-1-dependent phosphorylation boosts the activity of MutSγ as part of a positive feedback loop that concentrates pro-CO factors at CO sites (Yokoo *et al.* 2012).

In the course of constructing strains for experiments using the temperature-sensitive missense mutation him-14(it44ts), we obtained additional evidence consistent with a role for COSA-1 as a positive regulator of MutSy function. Specifically, we found that the presence of mels8, a transgene insertion expressing a functional GFP::COSA-1 fusion protein (Yokoo et al. 2012), resulted in significant rescue of the CO deficit caused by the him-14(it44ts) mutation at 20°C, a semi-permissive temperature at which MutSy activity is reduced but not eliminated. We evaluated the success or failure of CO formation by examining DAPI-stained bodies present in oocytes at diakinesis, the last stage of meiotic prophase. Wild-type oocytes contain six DAPI bodies, reflecting six connected homolog pairs (bivalents) harboring CO-based connections (chiasmata), whereas up to 12 DAPI bodies can be resolved in mutants lacking COs, reflecting unconnected individual chromosomes (univalents). In contrast to the high frequency of univalents observed in him-14(it44ts) mutant oocytes at the semi-permissive temperature of 20°C, six bivalents (indicative of successful CO formation for all homolog pairs) were observed in most meIs8 him-14(it44ts) oocytes at 20°C. This finding suggests that an elevated level of COSA-1 protein, due to the presence of the transgene insertion expressing GFP::COSA-1, can increase the likelihood of CO formation in the context of reduced MutSy activity. However, suppression of the him-14(it44ts) CO defect does not occur at the more restrictive temperatures of 24°C or 25°C where activity of MutSy is severely impaired. We note that six DAPI bodies are consistently observed at 24°C in diakinesis oocytes of AV818 worms in which mels8 is the only source of COSA-1 activity (mean = 6.0 ± 0.0 , n = 72), indicating that functional GFP::COSA-1 is expressed from meIs8 at 24°C. Together our data suggest that the presence of meIs8 does not bypass the requirement for MutSy activity to form COs, but instead suppresses him-14(it44ts) by augmenting the residual MutSy activity present at semi-permissive temperature. Our data are consistent with a model in which COSA-1 promotes CO formation, at least in part, through positive regulation of MutSy.

Reagents

Strain	Genotype	Available from
KK323	him-14(it44ts) unc-4(e120)/mnC1 II	AV lab
AV893	meIs8[unc-119(+) pie-1promoter::gfp::cosa-1] him-14(it44ts) unc-4(e120) II	AV lab
AV630	meIs8[unc-119(+) pie-1promoter::gfp::cosa-1] II	CGC
AV902	him-14(it44ts) II	AV lab
AV956	meIs8[unc-119(+) pie-1promoter::gfp::cosa-1] him-14(it44ts) II	AV lab
AV818	meIs8[unc-119(+) pie-1promoter::gfp::cosa-1] cosa-1(tm3298) II	AV lab

References

Bessler JB, Reddy KC, Hayashi M, Hodgkin J, Villeneuve AM. 2007. A role for *Caenorhabditis elegans* chromatin-associated protein HIM-17 in the proliferation vs. meiotic entry decision. Genetics 175: 2029-37. PMID: 17237503.

Kelly KO, Dernburg AF, Stanfield GM, Villeneuve AM. 2000. *Caenorhabditis elegans* msh-5 is required for both normal and radiation-induced meiotic crossing over but not for completion of meiosis. Genetics 156: 617-30. PMID: 11014811.

Woglar A, Villeneuve AM. 2018. Dynamic Architecture of DNA Repair Complexes and the Synaptonemal Complex at Sites of Meiotic Recombination. Cell 173: 1678-1691.e16. PMID: 29754818.

Yokoo R, Zawadzki KA, Nabeshima K, Drake M, Arur S, Villeneuve AM. 2012. COSA-1 reveals robust homeostasis and separable licensing and reinforcement steps governing meiotic crossovers. Cell 149: 75-87. PMID: 22464324.

Zalevsky J, MacQueen AJ, Duffy JB, Kemphues KJ, Villeneuve AM. 1999. Crossing over during *Caenorhabditis elegans* meiosis requires a conserved MutS-based pathway that is partially dispensable in budding yeast. Genetics 153: 1271-83.



8/24/2021 - Open Access

PMID: 10545458.

Funding: This work was supported by NIH grants R01GM067268 and R35GM126964 to AMV.

Author Contributions: Chloé Girard: Conceptualization, Investigation. Chantal C Akerib: Investigation. Anne M Villeneuve: Conceptualization, Formal analysis, Visualization, Writing - original draft.

Reviewed By: Anonymous

History: Received July 15, 2021 Revision received July 31, 2021 Accepted August 16, 2021 Published August 24, 2021

Copyright: © 2021 by the authors. This is an open-access article distributed under the terms of the Creative Commons Attribution 4.0 International (CC BY 4.0) License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Citation: Girard, C; Akerib, CC; Villeneuve, AM (2021). Suppression of *him-14(it44ts)* by a transgene insertion expressing GFP::COSA-1. microPublication Biology. https://doi.org/10.17912/micropub.biology.000430