



Figure S1 STAT92E enhances repeat-mediated variegation. Eye pigmentation phenotypes of adult flies of indicated genotypes raised at 25°C. The light orange background was due to the presence of *mini-w*⁺ in the *UAS-Stat92E*⁺

transgene. Note the deceased pigmentation when UAS- $Stat92E^+$ was expressed by the basal level of hsp70-Gal4 transgene (which does not carry $mini-w^+$).



Figure S2 Loss of Stat92E⁺ does not affect HP1 expression in imaginal discs. A 3^{rd} instar wing imaginal disc bearing $Stat92E^{-/-}$

cells (marked by lack of GFP) is shown for HP1 levels (stained by anti-HP1, red).



Figure S3 Long-term loss of STAT92E reduced HP1. A *Stat92E* homozygous loss of function cell (marked by GFP) was induced during

embryogenesis and examined for HP1 (red) in late $3^{\rm rd}$ instar larval salivary gland.



Figure S4 Specificity of anti-STAT92E and anti-pSTAT92E. (a) A 3rd instar wing imaginal disc bearing *Stat92E^{-/-}* cells (marked by lack of GFP) was stained by anti-STAT92E (red). Note that the red signals closely matches STAT92E protein levels, and that no signal is detected in *Stat92E^{-/-}* cells. (b) S2 cells were transfected with Hop-V5 and/or STAT92E-V5. Total cell lysates were subjected to SDS-PAGE and transferred to nitrocellulose

membrane and immunoblotted sequentially with anti-pSTAT92E, anti-V5, and anti-STAT92E, respectively. Note that anti-pSTAT92E does not cross-react with non-phospho-STAT92E (lack of signals in lane 1, 2, left panel), and that anti-STAT92E does not detect other proteins (right panel). Extra bands on the right panel were due to incomplete stripping. Anit-V5 detected non-specific bands (bands across all 3 lanes in middle panel).



Figure S5 HP1 and STAT92E distribution in hop^{GOF} mutant larvae. Salivary glands isolated from $hop^{GOF/+}$ ($hop^{Tum-l/+}$) larvae were stained with anti-HP1 (red) and anti-STAT92E (green). (a) In glands isolated from $hop^{Tum-l/+}$ larvae raised at 29°C, both HP1 and STAT92E appear diffused, lacking prominent heterochromatin foci. (b) In $hop^{Tum-l/+}$ larvae raised at 25°C, a moderately nonpermissive temperature¹, a fraction of the nuclei (32%; n=64/198 nuclei) exhibited nearly normal patterns of HP1 foci (left panel), whereas in nearly all of these nuclei (95%; n=61/64 nuclei), STAT92E appeared dispersed and did not colocalize with HP1 (middle panel). These results suggest that STAT92E disperses in response to Hop^{Tum-I} phosphorylation, and this process precedes changes in HP1 subnuclear localization.



Figure S6 Effects of H2O2/vanadate and CHX on STAT92E phosphorylation and histone H3 K9 methylation. S2 cells were cultured with or without CHX for 1 h, and then stimulated with $H_2O_2/vanadate$ for indicated times **(a)**, or

incubated in the presence of CHX for indicated times (b). Cell extracts were blotted sequentially with anti-pSTAT92E and anti-STAT92E, or with anti-H3mK9, respectively. Membrane was stripped between blots.



Figure S7 Full-length protein gels. Note that Fig. 2f used a lower percentage gel and longer run, which resulted in a greater

separation phospho-STAT92E from STAT92E bands (compared with Fig. 5b).

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References:

1. Hanratty, W. P. & Dearolf, C. R. The Drosophila Tumorous-lethal hematopoietic oncogene is a dominant mutation in the hopscotch locus. Mol Gen Genet 238, 33-7 (1993).