## A Balance Between Euchromatic (JIL-1) and Heterochromatic [SU(VAR)2-5 and SU(VAR)3-9] Factors Regulates Position-Effect Variegation in *Drosophila*

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**ABSTRACT** In this study, we show that the haplo-enhancer effect of *JIL-1* has the ability to counterbalance the haplo-suppressor effect of both *Su(var)3-9* and *Su(var)2-5* on position-effect variegation, providing evidence that a finely tuned balance between the levels of JIL-1 and the major heterochromatin components contributes to the regulation of gene expression.

HE essential JIL-1 histone H3S10 kinase (Jin et al. 1999; Wang et al. 2001) is a major regulator of chromatin structure (Deng et al. 2005, 2008) that functions to maintain euchromatic domains while counteracting heterochromatization and gene silencing (Ebert et al. 2004; Lerach et al. 2006; Zhang et al. 2006; Bao et al. 2007). In the absence of the JIL-1 kinase, the major heterochromatin markers H3K9me2, HP1a [Su(var)2-5], and Su(var)3-7 spread to ectopic locations on the chromosome arms (Zhang et al. 2006; Deng et al. 2007, 2010). These observations suggested a model for a dynamic balance between euchromatin and heterochromatin (Ebert et al. 2004; Zhang et al. 2006; Deng et al. 2010), where, as can be monitored in position-effect variegation (PEV) arrangements, the boundary between these two states is determined by antagonistic functions of a euchromatic regulator (JIL-1) and the major determinants of heterochromatin assembly, e.g., Su(var)3-9, HP1a, and Su(var)3-7 (for review see Weiler and Wakimoto 1995; Girton and Johansen 2008). In support of this model, Deng et al. (2010) recently showed that Su(var)3-7 and JIL-1 loss-of-function mutations have an antagonistic and counterbalancing effect on gene expression using PEV assays; however, potential dynamic interactions between JIL-1 and the other two heterochromatin genes, Su(var)3-9 and Su(var)2-5, were not

addressed in this study. Interestingly, in other genetic interaction assays monitoring the lethality as well as the chromosome morphology defects associated with the null JIL-1 phenotype, only a reduction in the dose of the Su(var)3-9 gene (Zhang et al. 2006; Deng et al. 2007) rescued both phenotypes. In contrast, in the same assays a reduction of *Su(var)3-7* rescued the lethality, but not the chromosome defects (Deng et al. 2010), and no genetic interactions were detectable between JIL-1 and Su(var)2-5 (Deng et al. 2007). Thus, these findings indicate that while Su(var)3-9 activity may be a major factor in the lethality and chromatinstructure perturbations associated with loss of the JIL-1 histone H3S10 kinase, these effects are likely to be uncoupled from HP1a and, to a lesser degree, from Su(var)3-7. This raises the question of whether JIL-1 dynamically interacts with the two other heterochromatin genes, Su(var)2-5 and Su(var)3-9, in regulating gene expression, as it does with Su (var)3-7.

To answer this question, we explored the effect of various combinations of loss-of-function alleles of *JIL-1* and *Su(var) 3-9* or *Su(var)2-5* on PEV caused by the *P*-element insertion line *118E-10* (Wallrath and Elgin 1995; Wallrath *et al.* 1996). Insertion of this *P* element (*P[hsp26-pt, hsp70-w]*) into euchromatic sites results in a uniform red-eye phenotype whereas insertion into a known heterochromatin region of the fourth chromosome results in a variegating eye phenotype (Cryderman *et al.* 1998; Bao *et al.* 2007) (Figures 1 and 2). It has been demonstrated that loss-of-function *JIL-1* alleles can act as haplo-enhancers of PEV, resulting in increased silencing of gene expression (Deng *et al.* 2010), whereas loci for structural components of heterochromatin

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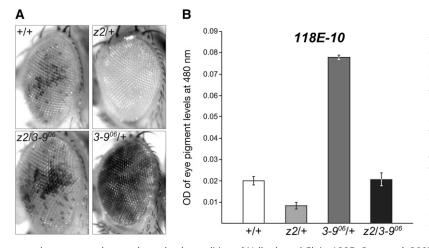
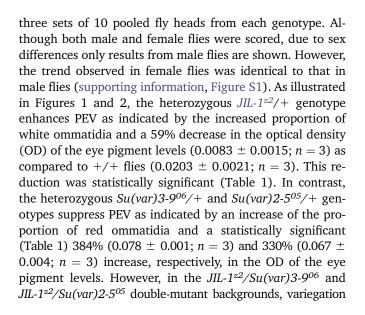
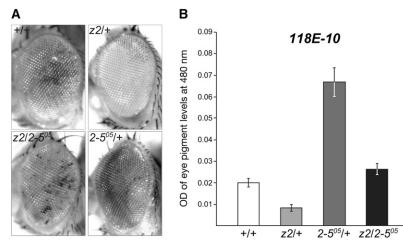


Figure 1 Counterbalancing effect of JIL-1 and Su(var)3-9 loss-of-function alleles on the PEV of the P-element insertion line 118E-10. (A) Examples of the degree of PEV in the eyes of wild-type JIL-1 and Su(var)3-9 (+/+), JIL-1z2/+ (z2/+), Su(var)3-9<sup>06</sup>/+ (3-9<sup>06</sup>/+), and JIL-1<sup>z2</sup>/Su(var)3-9<sup>06</sup> (z2/3-906) flies in a 118E-10/+ background. All images are from male flies. (B) Histograms of the amount of eye pigment in +/+, JIL-1<sup>22</sup>/+ (z2/+), Su(var)3-9<sup>06</sup>/+ (3-9<sup>06</sup>/+), and JIL-1<sup>z2</sup>/Su(var)3-9<sup>06</sup> (z2/3-9<sup>06</sup>) male flies heterozygous for 118E-10. Fly stocks were maintained according to standard protocols (Roberts 1998). Oregon-R was used for wild-type preparations. The JIL-1z2 allele is described in Wang et al. (2001) and in Zhang et al. (2003). The Su (var)3-906 and Su(var)2-505 alleles are described in Schotta et al. (2002) and in Eissenberg et al. (1992). The P-element P[hsp26-pt, hsp70-w] insertion line 118E-10 was the generous gift of L. Wallrath. The hsp70 promoter is leaky and promotes sufficient expression to generate a variegated

eye phenotype under non-heat-shock conditions (Wallrath and Elgin 1995; Bao *et al.* 2007). PEV assays were performed as previously described in Lerach *et al.* (2006) and in Bao *et al.* (2007). In short, various combinations of *JIL-1, Su(var)3-9*, or *Su(var)2-5* alleles were introduced into the *118E-10* or  $w^{n4}$  PEV arrangements by standard crossing. To quantify the variegated phenotype, adult flies were collected from the respective crosses at eclosion, aged 6 days at 25°, frozen in liquid nitrogen, and stored at  $-80^{\circ}$ C until assayed. The pigment assays were performed essentially as in Kavi and Birchler (2009) using three sets of 10 fly heads of each genotype collected from males and females. For each sample, the heads from the 10 flies were homogenized in 200 µl of methanol with 0.1% hydrochloric acid and centrifuged, and the OD of the supernatant was spectrophotometrically measured at a wavelength of 480 nm.

such as Su(var)3-9, Su(var)2-5, and Su(var)3-7 act as strong haplo-suppressors (Eissenberg et al. 1990; Reuter et al. 1990; Tschiersch et al. 1994). In the experiments, the transgenic reporter line 118E-10 was crossed into JIL-1<sup>z2</sup>/+, Su(var)3- $9^{06}/+$ , and Su(var)2- $5^{05}/+$  mutant backgrounds as well as into JIL-1z2/Su(var)3-906 and JIL-1z2/Su(var)2-505 doublemutant backgrounds. The JIL-1z2 allele is a true null allele (Wang et al. 2001; Zhang et al. 2003), the loss-of-function Su (var)3-9<sup>06</sup> allele is due to a DNA insertion (Schotta et al. 2002), and the  $Su(var)2-5^{05}$  loss-of-function allele is associated with a frameshift resulting in a nonsense peptide containing only the first 10 amino acids of HP1a (Eissenberg et al. 1992). Thus, to test whether the heterozygous  $JIL-1^{z2}$ allele could counterbalance the suppression of the  $Su(var)3-9^{06}$ or Su(var)2-505 loss-of-function alleles of the PEV of 118E-10, we compared the eye pigment levels of the various genotypes (Figures 1 and 2 and Table 1). Pigment assays were performed essentially as in Kavi and Birchler (2009) using





**Figure 2** Counterbalancing effect of *JIL-1* and *Su(var)2-5* loss-of-function alleles on the PEV of the *P*-element insertion line *118E-10*. (A) Examples of the degree of PEV in the eyes of wild-type *JIL-1* and *Su(var)2-5* (+/+), *JIL-1<sup>22</sup>/+* (*z2/+*), *Su(var)2-5<sup>05</sup>/+* (*2-5<sup>05</sup>/+*), and *JIL-1<sup>22</sup>/Su(var) 2-5<sup>05</sup>* (*z2/2-5<sup>05</sup>*) files in a *118E-10/+* background. All images are from male flies. (B) Histograms of the levels of eye pigment in +/+, *JIL-1<sup>22</sup>/+* (*z2/+)*, *Su(var)2-5<sup>05</sup>* (*z2/2-5<sup>05</sup>/+*), and *JIL-1<sup>22</sup>/Su(var)2-5<sup>05</sup>* (*z2/2-5<sup>05</sup>*) male flies heterozygous for *118E-10*.

Table 1	Statistical	comparison of	eye pigment ass	ays

Genotype <sup>a</sup>	JIL-1 <sup>z2</sup> /+	3-9 <sup>06</sup> /+	JIL-1 <sup>z2</sup> /3-9 <sup>06</sup>
118E-10/+ +/+ JIL-1 <sup>22</sup> /+ 3-9 <sup>06</sup> /+	P < 0.005 	P < 0.0001 P < 0.0001 —	P > 0.8 P < 0.005 P < 0.0001
Genotype <sup>a</sup>	JIL-1 <sup>z2</sup> /+	2-5 <sup>05</sup> /+	JIL-1 <sup>z2</sup> /2-5 <sup>05</sup>
118E-10/+ +/+ JIL-1 <sup>z2</sup> /+ 2-5 <sup>05</sup> /+	P < 0.005	P < 0.0001 P < 0.0001 —	P > 0.06 P < 0.005 P < 0.0005
Genotype	JIL-1 <sup>z2</sup> /+	<i>3-9<sup>06</sup>/+</i>	JIL-1 <sup>z2</sup> /3-9 <sup>06</sup>
W <sup>m4</sup> /Y +/+ JIL-1 <sup>z2</sup> /+ 3-9 <sup>06</sup> /+	P < 0.005 	P < 0.0001 P < 0.0001 	P < 0.0001 P < 0.0001 P < 0.001
Genotype	JIL-1 <sup>z2</sup> /+	2-5 <sup>05</sup> /+	JIL-1 <sup>z2</sup> /2-5 <sup>05</sup>
W <sup>m4</sup> /Y +/+ JIL-1 <sup>22</sup> /+ 2-5 <sup>05</sup> /+	P < 0.005 	P < 0.0001 P < 0.0001 —	P < 0.0001 P < 0.0001 P < 0.001

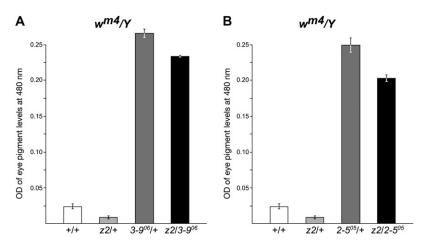
For each genotype, the average pigment level from three sets of measurements from 10 pooled fly heads were compared using a two-tailed Student's *t*-test. <sup>a</sup> Only male flies were scored.

of the proportion of red ommatidia was intermediate, and the eye pigment levels ( $0.0207 \pm 0.0031$  and  $0.0263 \pm 0.0035$ , respectively; n = 3) were statistically indistinguishable from genotypes with +/+ levels of JIL-1, Su(var)3-9, and Su(var)2-5 proteins (Figures 1 and 2 and Table 1).

To test whether a heterozygous *JIL-1* null allele also could counterbalance the suppression of the *Su(var)3-9<sup>06</sup>* or *Su(var)2-5<sup>05</sup>* alleles of the PEV of  $w^{m4}$ , we performed experiments similar to those described above for *118E-10*. As illustrated in Figure 3 for male flies, the heterozygous *JIL-1<sup>z2</sup>/+* genotype enhances PEV as indicated by a 67% decrease in the OD of the eye pigment levels (0.008 ± 0.002; n = 3) as compared to +/+ flies (0.024 ± 0.003, n = 3). This reduction was statistically significant (Table 1).

In contrast, the heterozygous  $Su(var)3-9^{06}/+$  and Su(var)2- $5^{05}/+$  genotypes suppress PEV as indicated by a statistically significant (Table 1) 1115% (0.2677  $\pm$  0.0061; n = 3) and 1023% (0.2454  $\pm$  0.0103; n = 3) increase, respectively, in the OD of the eye pigment levels. However, in the JIL- $1^{z2}$ / Su(var)3-9<sup>06</sup> and JIL-1<sup>z2</sup>/Su(var)2-5<sup>05</sup> double-mutant backgrounds, the eye pigment levels (0.2337  $\pm$  0.0011 and 0.2043  $\pm$  0.0037, respectively; n = 3) were significantly reduced (Table 1) by 13% and 17% as compared to heterozygous  $Su(var)3-9^{06}/+$  and  $Su(var)2-5^{05}/+$  genotypes, indicating that a heterozygous JIL-1 null allele has the abilty to counterbalance the suppression of the  $Su(var)3-9^{06}$  or  $Su(var)2-5^{05}$  alleles of the PEV of  $w^{m4}$ . However, it should be noted that it has been demonstrated that JIL-1 can act both as an enhancer and as a suppressor of  $w^{m4}$  PEV, depending on the precise levels of JIL-1 (Lerach et al. 2006; Deng et al. 2010). Thus, the genetic interactions between JIL-1 and the Su(var)3-9 and Su(var)2-5 alleles in regulating the PEV of  $w^{m4}$  are likely to be more complex than in the case of 118E-10 where reduced levels of JIL-1 always act as an enhancer (Bao et al. 2007). In females where the enhancer effect of the heterozygous JIL-1<sup>z2</sup> allele is less pronounced than in males, a statistically significant counterbalancing effect was detected only in flies of the JIL-1<sup>z2</sup>/Su (var)2-5<sup>05</sup> genotype (Figure S2).

These results demonstrate that the haplo-enhancer effect of *JIL-1* has the ability to counterbalance the haplo-suppressor effect of both Su(var)3-9 and Su(var)2-5 on the PEV of two different alleles. In previous experiments, a genetic interaction between *JIL-1* and Su(var)2-5 was not detected (Deng *et al.* 2007). However, the assays used to probe for interactions were viability and rescue of polytene chromosome morphology. As indicated by the experiments presented here, these parameters are likely to be independent of and separate from the mechanisms contributing to epigenetic regulation of PEV and gene silencing. Consequently, the present experiments, taken together with those of Deng *et al.* (2010) using a *JIL-1* null allele, provide strong evidence that a finely tuned balance between the levels of JIL-1 and all of the major heterochromatin components Su(var)3-9, HP1a, and



**Figure 3** Counterbalancing effect of *JIL-1* with *Su(var)3-9* and *Su(var)2-5* loss-of-function alleles on the PEV of  $w^{rn4}$ . (A) Histograms of the levels of eye pigment in +/+, *JIL-1<sup>z2</sup>/+* (*z2/+*), *Su(var)3-9<sup>06</sup>/+* (3-9<sup>06</sup>/+), and *JIL-1<sup>z2</sup>/Su(var)3-9<sup>06</sup>* (*z2/* 3-9<sup>06</sup>)  $w^{rn4}$  male flies. (B) Histograms of the levels of eye pigment in +/+, *JIL-1<sup>z2</sup>/+* (*z2/+*), *Su(var)2-5<sup>05</sup>/+* (2-5<sup>05</sup>/+), and *JIL-1<sup>z2</sup>/Su(var)2-5<sup>05</sup>* (*z2/2-5<sup>05</sup>*)  $w^{rn4}$  male flies.

Su(var)3-7 contributes to the regulation of PEV and gene expression.

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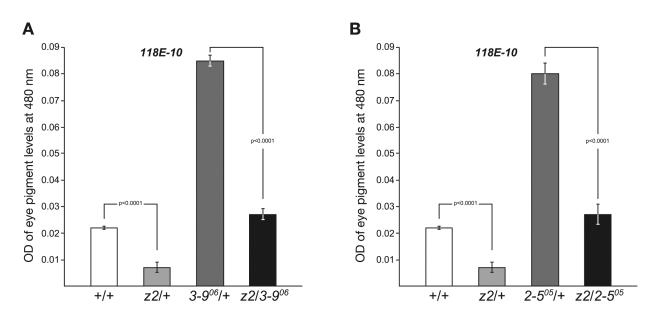
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## A Balance Between Euchromatic (JIL-1) and Heterochromatic [SU(VAR)2-5 and SU(VAR)3-9] Factors Regulates Position-Effect Variegation in Drosophila

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**Figure S1** Counterbalancing effect of *JIL-1* with *Su(var)3-9* and *Su(var)2-5* loss-of-function alleles on PEV of *118E-10* in female flies. (A) Histograms of the amount of eye pigment in +/+, *JIL-1<sup>22</sup>/+* (*z2/+*), *Su(var)3-9<sup>06</sup>/+* (*3-9<sup>06</sup>/+*), and *JIL-1<sup>22</sup>/Su(var)3-9<sup>06</sup>* (*z2/3-9<sup>06</sup>*) female flies heterozygous for *118E-10*. (B) Histograms of the levels of eye pigment in +/+, *JIL-1<sup>22</sup>/+* (*z2/+*), *Su(var)2-5<sup>05</sup>/+* (*2-5<sup>05</sup>/+*), and *JIL-1<sup>22</sup>/Su(var)2-5<sup>05</sup>/2/2-5<sup>05</sup>* (*z2/2-5<sup>05</sup>/*) female flies heterozygous for *118E-10*. The average pigment level from three sets of measurements from 10 pooled flyheads were compared using a two-tailed Student's t-test.

