Tubby-tagged balancers for the Drosophila X and second chromosomes

Ramona Lattao,¹ Silvia Bonaccorsi,¹ Xiao Guan,² Steven A. Wasserman² and Maurizio Gatti^{1,*}

¹Dipartimento di Biologia e Biotecnologie "Charles Darwin" and Istituto di Biologia e Patologia Molecolari del CNR; Sapienza, Università di Roma; Rome, Italy; ²Section of Cell and Developmental Biology; University of California; La Jolla, CA USA

Key words: balancer chromosomes, Tubby, Drosophila melanogaster

We generated *FM7a* and *CyO* balancer chromosomes bearing a *Tubby*¹ (*Tb*¹) dominant transgene. Flies heterozygous for these *FM7a* and *CyO* derivatives exhibit a phenotype undistinguishable from that elicited by the *Tb*¹ mutation associated with the *TM6B* balancer. We tested two of these *Tb*-bearing balancers (*FM7-TbA* and *CyO-TbA*) for more than 30 generations and found that the *Tb*¹ transgene they carry is stable. Thus, these new *Tb*-tagged balancers are particularly useful for balancing lethal mutations and distinguish homozygous mutant larvae from their heterozygous siblings.

One of the great advantages of Drosophila melanogaster as model organism is the availability of balancer chromosomes. These chromosomes suppress recombination with their homologues, allowing the maintenance of lethal and sterile mutants as balanced heterozygotes. All balancers carry dominant markers that are visible in adult flies, but only a subset have markers that unambiguously distinguish homozygous mutant larvae from their heterozygous siblings. The latter balancers include those that express high levels of the GFP protein under the indirect control, via the UAS/GAL4 system, of either the Kruppel (Kr),¹ or hsp70,² promoter. In addition, there are direct-drive balancers that express GFP under the control of the actin A5c promoter³ or YFP under the control of Deformed (Dfd HZ2.7rev) or glass (GMR) enhancer elements.⁴ Each of the GFP- or YFP-expressing balancers has specific advantages, but all share a common drawback: These balancers require the use of a dissecting microscope equipped with an UV light source, which for reliable fluorescence detection is preferably used in the dark.

The *TM6B* balancer⁵ carries the *Tb*¹ dominant mutation, which results in squat larvae and pupae. We have been using this balancer for many years to unambiguously distinguish homozygous mitotic mutants dying at late larval stages from their heterozygous siblings (reviewed in ref. 6). This balancer proved particularly useful when mutant larvae are rare and one has to examine several vials (or bottles) to find third instar larvae suitable for dissection and cytological analysis. To generate new tools for easy detection of larvae homozygous for lethal mutations on the X or the second chromosome, we decided to generate *Tb*-marked versions of *FM7a*⁷ and *CyO*,⁸ respectively.

We have previously demonstrated that the Tubby phenotype results from the deletion in the Tb^{t} chromosome of DNA encoding amino acids 167–190 of the TwdlA cuticle protein.⁹ A genomic fragment spanning the *TwdlA^{Tb1}* transcription unit, as well as 1

kb of DNA upstream, was inserted into a modified polylinker sequence in the pCaSpeR vector. Using standard methods, we coinjected DNA with a $\Delta 2$ -3 transposase source into $w^{1/8}$ embryos, selected for w^* eye color, and generated balanced stocks. All such stocks (n = 10) exhibited the squat larval and pupal phenotype characteristic of the Tb' mutation. One such stock, referred to hereafter as $P\{Tb'\}/CyO$, was used as the starting point for mobilizing Tb' onto X and 2^{nd} chromosome balancers.

To generate $FM7a-P\{Tb^{1}\}$ we crossed FM7a; $P\{Tb^{1}\}/CyO$; TMS/+ males to w/w females (TMS is a $\Delta 2$ -3-bearing third chromosome balancer expressing the P-element transposase described in ref. 10); from approximately 1,000 progeny we recovered two FM7 chromosomes that co-segregated with Tb (we propose to name these balancers FM7-TbA and FM7-TbB). To generate a second chromosome balancer with a $P\{Tb^l\}$ insertion we used a CyO balancer bearing the additional markers S and bw^{I} (designated as CyO-Sbw in FlyBase). We crossed w/w; $P\{Tb^{I}\}/CyO$ -Sbw; TMS/+ females to w; CyO/Sco males and recovered three $C_{\gamma}O$ -Sbw-P{Tb} chromosomes from approximately 1,000 progeny (we propose to name these balancers CyO-TbA, CyO-TbB and CyO-TbC). To assess the utility of these FM7a and CyO Tb-bearing balancers, we compared their Tb phenotype with the Tb' mutant phenotype associated with TM6B.5 To quantify the squat phenotype elicited by the balancers we measured the axial ratio¹¹ of larvae and pupae (AR, length/width) heterozygous for each balancer. It has been previously shown that the AR does not depend on larval and pupal size and provides a reliable measure of the Tb phenotype.9 As shown in Figure 1, the ARs observed in FM7-TbA, FM7-TbB, CyO-TbA, CyO-TbB, CyO-TbC and TM6B heterozygotes are fully comparable and significantly different from those of wild-type or non-Tb-bearing larvae and pupae. We thus conclude that the FM7a and CyO Tb-tagged chromosomes are highly suitable to distinguish larvae and pupae

^{*}Correspondence to: Maurizio Gatti; Email: maurizio.gatti@uniroma1.it Submitted: 05/31/11; Revised: 07/06/11; Accepted: 07/07/11 DOI:

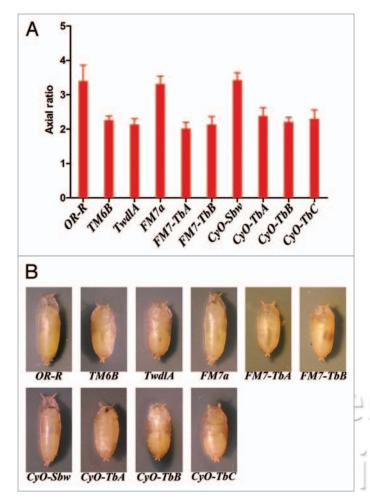


Figure 1. Quantitation of the squat phenotype of pupae heterozygous for different *Tb*-bearing chromosomes or balancers; *OR-R* is a wild-type Oregon R stock; *TwdlA* contains a single copy of the *TwdlA*^{Tb1} transgene; *FM7a* and *CyO-Sbw* are the balancer chromosomes used to construct the *Tb*-bearing derivatives indicated as *FM7-TbA*, *FM7-TbB*, *CyO-TbA*, *CyO-TbB* and *CyO-TbC*. (A) Axial ratios (ARs; length/width, ±standard deviation) determined from digital photographs of at least 40 pupae of each genotype (B) examples of pupae heterozygous for different *Tb* mutations and transgenes.

References

- Casso D, Ramirez-Weber FA, Kornberg TB. GFPtagged balancer chromosomes for *Drosophila melanogaster*. Mech Develop 1999; 88:229-32; PMID: 10534621.
- Rudolph T, Lu B, Westphal T, Szidonya J, Eissenberg J, Reuter G. New type of *CyO* and *TM3* green balancers. Dros Inf Serv 1999; 82:99-100.
- Reichhart JM, Ferrandon D. Green balancers. Dros Inf Serv 1998; 81:201-2.
- Le T, Liang Z, Patel H, Yu MH, Sivasubramaniam G, Slovitt M, et al. A new family of Drosophila balancer chromosomes with a *w dfd*-GMR yellow fluorescent protein marker. Genetics 2006; 174:2255-7; PMID: 17057238; DOI: 10.1534/genetics.106.063461.
- 5. Craymer L. *TM6B*: Third Multiple Six, B structure. Dros Inf Serv 1984; 60:234.
- Gatti M, Baker BS. Genes controlling essential cell cycle functions in *Drosophila melanogaster*. Genes Dev 1989; 3:438-53; PMID: 2498166.
- Merriam JR. FM7: First multiple seven. Dros Inf Serv 1968; 43:64.
- Oster II. A new crossing-over suppressor in chromosome 2 effective in the presence of heterologous inversions. Dros Inf Serv 1956; 30:145.
- Guan X, Middlebrooks BW, Alexander S, Wasserman SA. Mutation of TweedleD, a member of an unconventional cuticle protein family, alters body shape in Drosophila. Proc Natl Acad Sci USA 2006; 103:16794-9; PMID: 17075064; DOI:10.1073/ pnas.0607616103.

that bear the balancers from those that are homozygous for the balanced chromosome.

We generated the two FM7a and the three CyO Tb-bearing balancers in October 2009 and March 2010, respectively. Within two months from the time of their generation, one balancer of each type (FM7-TbA and CyO-TbA) was used to rebalance 40 chromosomes bearing different lethal mutations associated with either the X or the second chromosome. These stocks were reexamined in March 2011 (after more than 30 generations) and none of them had lost the Tb marker from the balancer. We thus conclude that the Tb' transgene is very stable and that the FM7-TbA and CyO-TbA balancers are suitable for the establishment and long-term maintenance of balanced stocks.

Using inverse PCR (an EcoRI fragment was ligated and amplified using the 5'-CAGCTCCATAGTTATAGCCGC and CGTTAAGTGGATGTCTTCTTG-3' primers), we also determined the insertion sites of the Tb' transgenes within the *FM7-TbA* and *CyO-TbA* balancers. The insertion in *FM7-TbA* mapped at 17C in the intergenic space between *CG15047* and *beadex*; accordingly, this balancer is homozygous-viable. The Tb'transgene of *CyO-TbA* was inserted into the *Cytochrome P450 reductase* (*Cpr*) gene in region 26C3. We did not determine whether this insertion inactivates the *Cpr* gene.

In summary, we have generated Tb-marked versions of FM7a and CyO. These chromosomes carry a stable Tb^{1} transgene that has the same expressivity as the original Tb^{\prime} mutation. We believe that the use of the FM7-TbA and CyO-TbA balancers will facilitate a number of experimental strategies, allowing researchers to readily distinguish homozygous larvae and pupae from their heterozygous siblings simply by looking through the vial or the bottle used to grow the flies. In addition, the availability of balancers with different dominant larval markers such as Tb or GFP will allow construction of stocks carrying two lethal mutations balanced on Tb- and GFP-bearing balancers, respectively. This will be particularly helpful for selection and analysis of single and double mutants from the same culture. The FM7-TbA and CyO-TbA balancers can be obtained from the Bloomington Drosophila Stock Center at Indiana University (http://flystocks. bio.indiana.edu).

- Lindsley DL, Zimm. In: The Genome of *Drosophila* melanogaster. Academic Press, Inc. 1992; 1-1133.
- Letsou A, Alexander S, Orth K, Wasserman SA. Genetic and molecular characterization of tube, a Drosophila gene maternally required for embryonic dorsoventral polarity. Proc Natl Acad Sci USA 1991; 88:810-4; PMID: 1899484.