

FLPe functions in zebrafish embryos

Andrew C. Wong · Bruce W. Draper ·
Alison L. Van Eenennaam

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Abstract To assay the efficiency of the FLP/FRT site-specific recombination system in *Danio rerio*, a construct consisting of a muscle-specific promoter driving EGFP flanked by FRT sites was developed. FLPe capped RNA was microinjected into transgenic single cell stage zebrafish embryos obtained by crossing hemizygous transgenic males with wild-type females. By 48 h post fertilization (hpf), the proportion of embryos displaying green fluorescence following FLPe RNA microinjection was significantly lower (7.7%; $P < 0.001$) than would be expected from a cross in the absence of the recombinase (50%). Embryos that retained fluorescence displayed marked mosaicism. Inheritance of the excised transgene in non-fluorescent, transgenic embryos was verified by PCR analysis and FLPe-mediated recombination was confirmed by DNA sequencing. Sperm derived from confirmed transgenic males in these experiments was used to fertilize wild-type eggs to determine whether germline excision of the transgene had occurred. Clutches sired by FLPe-microinjected males contained 0–4% fluorescent embryos. Transgenic males that were phenotypically wild-type produced no fluorescent

progeny, demonstrating complete excision of the transgene from their germline. FLPe microinjected males that retained some fluorescent muscle expression produced a small proportion of fluorescent offspring, suggesting that in mosaic males not all germline cells had undergone FLPe-mediated transgene excision. Our results show that FLPe, which is derived from *Saccharomyces cerevisiae*, is an efficient recombinase in zebrafish maintained at 28.5°C.

Keywords FLP · FLPe · Recombinase · Zebrafish

Abbreviations

EGFP	Enhanced green fluorescent protein
FLPe	Enhanced FLP recombinase
FRT	FLP recognition target
hpf	Hours post fertilization

Introduction

Already established as an important developmental model, zebrafish (*Danio rerio*) are emerging as a powerful genetic model as well. Zebrafish possess certain characteristics that make them ideally suited for developmental and genetic studies, namely short generation time, large number of offspring, and optical clarity of externally fertilized embryos. Embryogenesis is finished in about 24 h, organogenesis is completed in about 5 days, and sexual maturity is

A. C. Wong · A. L. Van Eenennaam (✉)
Department of Animal Science, University of California,
Davis, CA 95616, USA
e-mail: alvaneenennaam@ucdavis.edu

B. W. Draper
Department of Molecular and Cellular Biology,
University of California, Davis, CA 95616, USA

reached at 3 months. A single female can provide hundreds of embryos at a time, either by natural crosses or by in vitro fertilization. Eggs are externally fertilized, develop synchronously outside of the mother and remain optically transparent up to 72 hpf, allowing for easy visualization of internal processes as well as the expression of fluorescently tagged transgenes and other reporter constructs.

Twelve years after the first successful in vivo demonstration of Cre/loxP in transgenic mice (Lakso et al. 1992; Orban et al. 1992), the Cre/loxP site-specific recombination system was introduced into zebrafish (Dong and Stuart 2004). The extensive use of Cre recombinase in mice has been proven to be a powerful tool for the study of mammalian genetics. Cre (cyclization recombination) is a 38-kDa protein from bacteriophage P1 that can recombine target DNA recognition sequences called loxP (locus of crossing over [x] in P1) sites. The loxP site is a 34-bp DNA sequence that consists of two 13-bp inverted repeats, which act as binding sites for Cre, flanking an asymmetrical 8-bp core region. Depending on the orientation and location of the recombinase recognition sites, Cre can catalyze excision, integration, inversion, or translocation of DNA segments. The biological role of Cre/loxP in the circular plasmid form of bacteriophage P1 is to resolve DNA dimers into plasmid monomers for equal partitioning to

daughter cells during cell division (Austin et al. 1981; Sternberg and Hamilton 1981). Gene targeting in mouse embryonic stem cells can add recombinase recognition sites to specific locations in the genome, enabling a variety of sophisticated genetic manipulations, including chromosomal inversions, site-specific transgene insertion, conditional transgenesis and conditional gene inactivation (reviewed in Nagy 2000). A database has been designed recently to keep track of the over 500 different Cre-expressing transgenic mice (Nagy et al. 2009).

Although the use of Cre recombinase in zebrafish is still in its infancy when compared to the mouse, sophisticated implementation of the Cre/loxP system, such as site-directed gene integration at mutant lox sites (Liu et al. 2007) and heat shock-inducible gene activation (Langenau et al. 2005; Thummel et al. 2005; Feng et al. 2007; Hans et al. 2009), have been recently reported. To further enhance the zebrafish genetic toolkit, the addition of a second site-specific recombination system, such as FLP/FRT, would be beneficial and allow for more elegant experimentation. FLP (flipase) is a 43-kDa protein from the 2-μm circle of *Saccharomyces cerevisiae*. Like Cre, FLP can recombine 34-bp DNA recognition sequences called FRT (FLP recognition target) sites, consisting of two imperfect 13-bp inverted repeats flanking an asymmetrical 8-bp core region (Fig. 1b). The biological role of FLP/FRT in

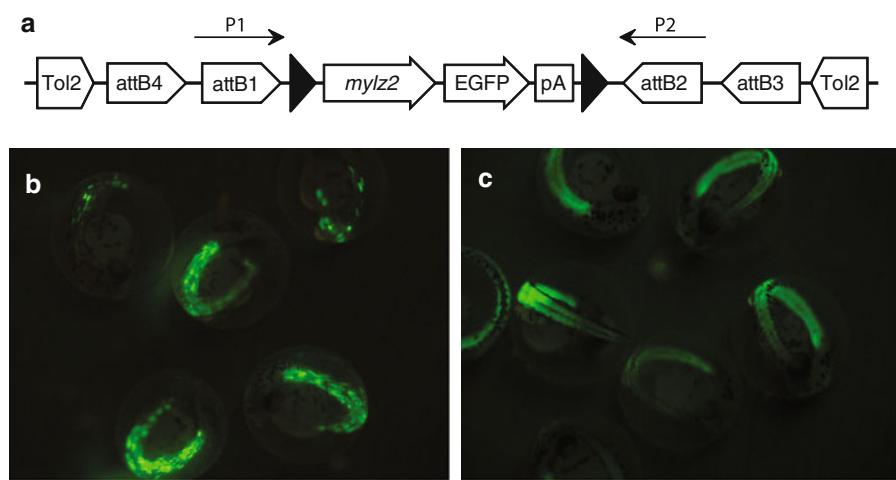


Fig. 1 **a** Schematic drawing of FRT *mylz2*-EGFP construct for assaying FLP-mediated site-specific recombination. Black triangles represent FRT sites. The Tol2Kit (Kwan et al. 2007), which utilizes Gateway cloning (Invitrogen), was used to create this construct. Tol2 sites at either end of the construct facilitated zebrafish transgenesis. **b** Founder zebrafish embryos

microinjected with the construct, FRT *mylz2*-EGFP, displayed mosaicism by 48 hpf. **c** Hemizygous F₁ embryos derived from mosaic founder fish that genotyped positive for the transgene in their sperm displayed even distribution of EGFP expression throughout

S. cerevisiae is to maintain a high copy number of the 2-μm circle by inverting half of the plasmid during replication such that a large multimeric plasmid is created which is later resolved into individual monomers (Futcher 1986). An in vitro study by Buchholz et al. (1996) determined that the most favorable temperature for FLP activity was near 30°C, consistent with the yeast origin of the FLP/FRT site-specific recombination system. An enhanced version of FLP recombinase, FLPe, was developed by cycling mutagenesis to improve its thermostability so as to prevent its inactivation at higher temperatures (Buchholz et al. 1998). Although predominantly used in *Drosophila* for mitotic recombination and gene activation (reviewed in Bischof and Basler 2008), the FLP/FRT recombination system has also been used in mice, both separately (Ludwig et al. 1996; Dymecki 1996; Rodriguez et al. 2000; Kanki et al. 2006) and in conjunction with Cre/loxP (Meyers et al. 1998; Turakainen et al. 2009; Yamamoto et al. 2009). To assay the efficiency of the FLP/FRT site-specific recombination system in *Danio rerio*, FLPe capped RNA was microinjected into transgenic zebrafish embryos carrying an FRT-flanked EGFP expression cassette. Injected fish were observed for muscle-specific GFP expression, and outcrossed to wild-type zebrafish to assess germline excision efficiency.

Materials and methods

Zebrafish care

The *AB line of wild-type zebrafish were raised and spawned as previously described (Westerfield 2000). Zebrafish were maintained at a water temperature of 28.5°C in an air temperature controlled room with a 14 h light/10 h dark photoperiod. Zebrafish were fed a combination of paramecia, *Artemia* nauplii (Argent Laboratories, Redmond, WA) and commercial fish food twice daily.

DNA transgene construction

mylz2-EGFP (Xu et al. 1999), was amplified from pMLC2f-EGFP1 with primers containing FRT sites (underlined): 5'- ggggacaagtttgtacaaaaaaaggcaggctgaagttcctattcttagaaagtataggacttcattcgccacagaggaagtgacacc-3' and 5'- ggggaccacttttacaaagactgggtgaagt-

cctatactttcttagagaataggaaacttctttatttgtgaaatttgtgatgcata-3'. The primers also contained *attB* sites (italicized) such that the PCR product would be compatible with the middle entry clone of the Tol2Kit (Kwan et al. 2007). A 5' entry clone and a 3' entry clone were combined in an LR reaction with the FRT-flanked *mylz2*-EGFP middle entry clone and a pDestTol2pA2 destination vector (Kwan et al. 2007) to produce FRT *mylz2*-EGFP (Fig. 1a).

In vitro synthesis of Tol2 RNA

Tol2 transposase RNA was generated from pCS2FA-transposase (Kwan et al. 2007). The plasmid was linearized with *NotI* and capped RNA was synthesized using the mMESSAGE mMACHINE SP6 Kit (Ambion, Austin, TX). RNA was purified with an RNeasy Mini Kit (Qiagen, Valencia, CA) and LiCl precipitated.

Microinjection and production of a stable transgenic zebrafish line

Tol2 transposon-mediated transgenesis methods (Kwan et al. 2007) were used to produce stable lines of transgenic zebrafish. Immediately prior to microinjection, FRT *mylz2*-EGFP plasmid DNA and Tol2 transposase RNA were combined to a final concentration of 5 ng/μl each. The DNA/RNA mixture was microinjected into the blastodisc of single cell-stage wild-type zebrafish embryo. Injected embryos were then moved to a 28.5°C incubator and screened for EGFP by 48 hpf. Germline positive transgenic fish were identified as those which produced EGFP positive T₁ embryos.

Fluorescent microscopy

Microinjected zebrafish were anesthetized in tricaine/MS-222 and observed under a MZ16F stereomicroscope (Leica, Bannockburn, IL) with a GFP2 filter set. Photographs were obtained with a 12-megapixel DFC 500 digital camera (Leica).

In vitro synthesis of FLPe RNA and microinjection

FLPe RNA was generated from pCSFLPe (Werdien et al. 2001). The plasmid was linearized with *NotI*

and capped RNA was synthesized using the mMES-SAGE mMACHINE SP6 Kit (Ambion). RNA was purified with an RNeasy Mini Kit (Qiagen) and LiCl precipitated. FLPe RNA was quantified and diluted down to 75 ng/μl before being microinjected into single cell transgenic zebrafish embryos. T₂ single-cell embryos from multiple clutches derived from outcrossing hemizygous transgenic T₁ male zebrafish were microinjected with approximately 2 nl of FLPe capped RNA over the course of several days. Fish that had inherited the transgene, as evidenced by a fluorescent phenotype or by the presence of recombinase-excision footprint, were raised to sexual maturity and sperm from males was genotyped and used to fertilize eggs from wild-type females to determine whether excision of the transgene had occurred in the germline.

PCR analysis and DNA sequencing

FLPe excision primers, 5'-CTCGGAATGTCAATAC ACCTTGCAGAG-3' and 5'-GTTGATGCCGGTGA ACGTGCAAA-3', were used to verify site-specific excision from FLPe RNA-injected transgenic zebrafish. Genomic DNA from finclips of FLPe RNA-injected zebrafish was used as the template for PCR. PCR products were gel purified with a QIAquick Gel Extraction Kit (Qiagen) and cloned into a pCR2.1-TOPO vector (Invitrogen). Colonies were grown in liquid LB with the appropriate antibiotic and plasmid DNA was obtained with a QIAprep Spin Miniprep Kit (Qiagen). DNA sequence was determined on a 3730 Capillary Electrophoresis Genetic Analyzer (Applied Biosystems, Foster City, CA).

Results and discussion

Production of a stable transgenic zebrafish line containing FRT *mylz2*-EGFP

To create a stable transgenic line of zebrafish containing FRT *mylz2*-EGFP (Fig. 1a), 716 wild-type zebrafish eggs were microinjected with the DNA construct/transposase RNA mix. Of the 240 embryos that survived past day five, 39 were mosaic for EGFP expression (Fig. 1b). At sexual maturity, sperm from the 27 EGFP mosaic zebrafish males was obtained and a PCR screen for the transgene was performed to

identify male transgenic germline positive founders. One male genotyped positive for the transgene (data not shown) and was used for in vitro fertilization of wild-type zebrafish eggs. Of the 387 wild-type zebrafish eggs that were in vitro fertilized with the PCR-positive sperm, 78 T₁ embryos were hemizygous for EGFP (Fig. 1c).

An advantage of using the Tol2 transposon-mediated transgenesis method is that transgene insertions are always single copy (Kawakami 2005; Kwan et al. 2007) unlike the concatemers that are common with traditional microinjection methods. A 1:1 ratio of GFP+ to GFP- fish (30:32, 12:9, 10:11, 89:95, 29:24, 20:20, 28:27, 8:8) was observed when transgenic T₁ offspring were outcrossed to wild-type zebrafish and suggested that the transgene was segregating as a single locus.

Site-specific excision from FLPe RNA microinjection

Hemizygous transgenic T₁ male zebrafish were outcrossed to wild-type female zebrafish to produce T₂ embryos for FLPe RNA microinjection. A clutch of embryos obtained by outcrossing a hemizygous transgenic fish would be expected to yield 50% transgenic fish (EGFP+) and 50% wild-type fish (EGFP-). Survival and proportion of fluorescent embryos did not differ between clutches in the same treatment group ($P > 0.05$; data not shown) and so data for all clutches in each treatment group were pooled. At 48 h after FLPe RNA injection, only 7.7% (17/220) embryos showed any level of muscle-specific EGFP expression (Fig. 2a, b). The remaining 203 embryos appeared wild-type (Table 1). A control hemizygous transgenic outcross that was not microinjected with FLPe RNA produced a clutch with an expected 1:1 ratio of EGFP + to EGFP- fish (Fig. 2c, d; Table 1).

To confirm FLPe-mediated excision of the EGFP transgene from FLPe RNA-injected embryos, PCR was performed on genomic DNA obtained from finclips of injected zebrafish with primers P1 and P2 (Fig. 1a). Site-specific recombination between the FRT sites was expected to produce an 803-bp fragment. As shown in Fig. 2e, finclips collected from eight FLPe RNA-injected transgenic zebrafish with a mosaic GFP phenotype produced an 800-bp fragment indicative of successful FLPe-mediated excision. A 3.7-kb intact fragment was not observed in these

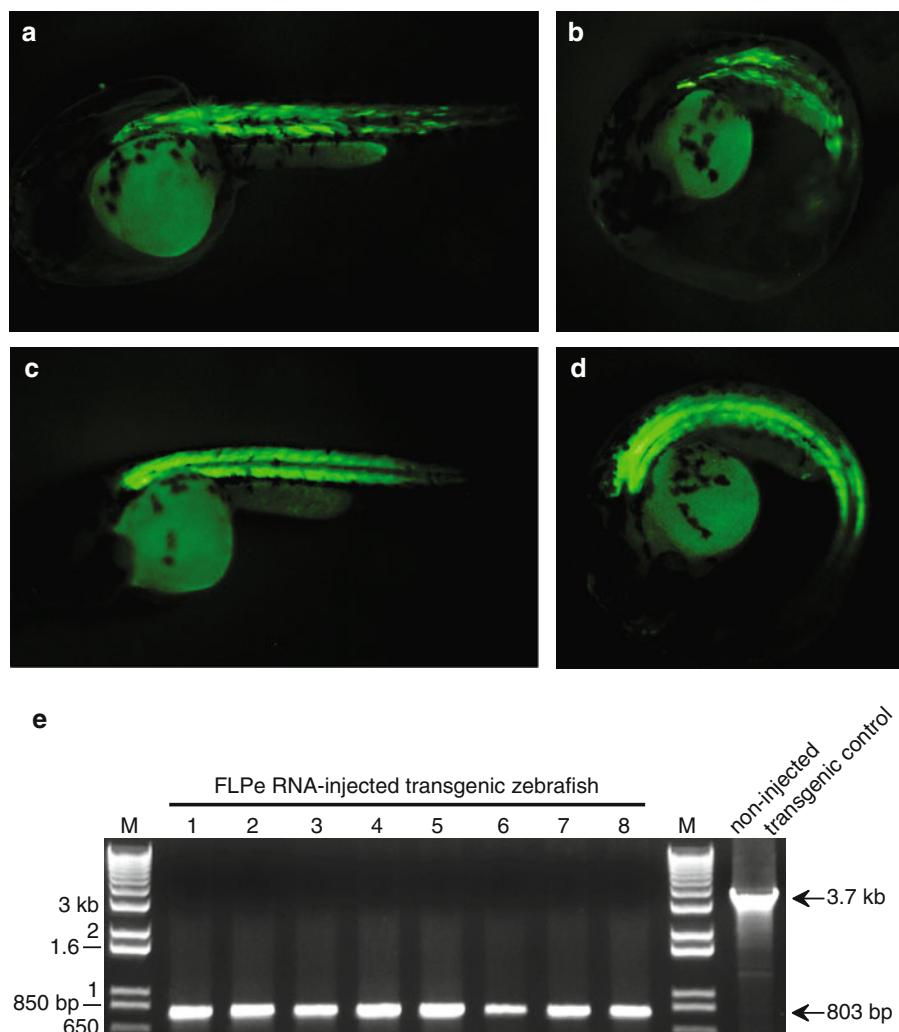


Fig. 2 Mosaic EGFP expression after FLPe recombinase RNA microinjection. Forty-eight hours post fertilization, mosaic expression of fluorescence can be observed in FLPe RNA-injected transgenic zebrafish embryos (**a, b**). Individual muscle cells that have not had the FRT-flanked transgene excised can be seen fluorescing green. In contrast, non-injected transgenic controls (**c, d**), show evenly distributed EGFP expression throughout the muscles of the zebrafish. **e** PCR confirmation of recombinase-mediated site-specific recombination. Genomic

DNA from finclips of FLPe RNA-injected transgenic zebrafish was used as a DNA template for PCR to confirm excision of the FRT-flanked EGFP transgene. In FLPe-injected zebrafish (1–8), site-specific recombination produced an 803-bp fragment, indicative of excision. In the non-injected transgenic control, an unexcised 3.7-kb band is expected. Excision products were cloned and sequenced to verify precise FLPe-mediated site-specific recombination in zebrafish

Table 1 Proportion of fish displaying muscle-specific fluorescence following microinjection of FLPe-capped RNA into single cell embryos derived from outcrossing hemizygous FRT *mylz2*-EGFP transgenic fish with wild-type fish

Recombinase microinjected	EGFP+ ^a	EGFP−	Total	χ^2	P
FLPe	17 (7.7%)	203 (92.3%)	220	157.25	<0.001
Control	28 (49.1%)	29 (50.9%)	57	0.017	>0.995

^a Includes fish that displayed any evidence of EGFP expression

Table 2 Segregation ratios observed in the offspring derived from outcrossing confirmed FRT *mylz2*-EGFP transgenic fish that had been microinjected with FLPe-capped RNA at the single cell stage

Recombinase microinjected	Parental phenotype	Offspring phenotype		Total	χ^2	<i>P</i>
		EGFP+	EGFP-			
FLPe	Mosaic muscle-specific fluorescence	34 (4.04%)	806 (95.96%)	842	711.5	<0.001
FLPe	No fluorescence	0 (0%)	339 (100%)	339	339	<0.001

samples possibly due to a reduction in the amount of non-recombined template available for amplification as well as preferential amplification of the shorter PCR product. In non-injected transgenic zebrafish controls, a 3.7-kb intact fragment was produced with primers P1 and P2 (Fig. 1a). To verify that the excised PCR fragments were indeed due to FLPe-mediated site-specific recombination, PCR products were sequenced and were found to contain a single 34-bp FRT footprint (data not shown).

To ascertain whether excision had occurred in the germline, FLPe RNA-injected T₂ transgenic fish were raised to sexual maturity and outcrossed to wild-type zebrafish. If excision had occurred in the germline, a deviation from the 1:1 ratio of transgenic to wild-type offspring would be expected. Only 4.0% (34/842) of embryos derived from outcrosses of FLPe RNA-injected fish that exhibited a mosaic phenotype appeared EGFP+ by 48 hpf (Table 2). When multiple FLPe RNA-injected transgenic fish that lacked a fluorescent phenotype (EGFP-) were outcrossed to wild-type zebrafish, 0.0% (0/339) of the embryos were positive for EGFP expression (Table 2) suggesting that these fish had successfully excised the transgene from the germ cell lineage.

During zebrafish development, primordial germ cells can be specified as those containing *vasa*, a molecular marker for germline cells. *Vasa* transcript can be found as early as the two-cell stage and the number of *vasa*-containing cells (4) remains constant up to the 1,000-cell stage (3 hpf) (Raz 2002). However, somites, which eventually differentiate into sclerotome (vertebrae), dermatome (dermis), and myotome (skeletal muscle), do not begin to appear until the segmentation stage (10–24 hpf) (Kimmel et al. 1995). The fact that none of the FLPe RNA-injected transgenic fish that lacked a fluorescent phenotype were germ cell-mosaic suggests that FLPe excision frequently occurred at a very early stage, thereby excising the transgene from all cell lineages.

One other study has examined FLPe site-specific recombination in zebrafish (Boniface et al. 2009). In that study transgenic zebrafish expressing Cre or FLPo (a mouse codon-optimized version of FLPe) under the control of a heat shock promoter were crossed to a recombinase reporter line and inversion efficiencies were recorded. The authors reported that following an hour long heat shock (38°C) induction, Cre expression inverted 82% of the germline reporter construct whereas FLPo expression inverted only 31% of the germline reporter construct. Given FLP activity is optimal near 30°C (Buchholz et al. 1996), it would be interesting to compare the efficiency of these two recombinases, and that of wild-type FLP in zebrafish maintained in ambient water temperatures of 28.5°C.

In conclusion, we found that the FLP/FRT site-specific recombination system from *S. cerevisiae* was very active in zebrafish maintained at a water temperature of 28.5°C. FLPe-mediated excision occurred rapidly following RNA microinjection into single-cell embryos, and frequently removed the FRT-flanked transgene from the germ cell lineage of transgenic fish. Used in conjunction with Cre/loxP, FLP/FRT will allow zebrafish researchers to design more elegant experiments similar to those already seen in mice (Kondo et al. 2006) and plants (Djukanovic et al. 2008).

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