

Expression and Visualization of Red Fluorescent Protein (RFP) in *Neurospora crassa*Michael Freitag[#] and Eric U. Selker

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We report the expression of *Discosoma* red fluorescent protein (RFP) and RFP fusion proteins in *Neurospora crassa*. RFP was expressed under the control of the *Neurospora ccg-1* promoter in transformants with single copies integrated at the *his-3* locus by gene targeting. Because this RFP gene, *tdimer2(12)*, contains a 677 bp direct tandem repeat of dsRed, RFP constructs underwent RIP at high frequency in *rid*⁺ strains. Fusion proteins of RFP to the amino terminus of *Neurospora* heterochromatin protein 1 (HP1) were localized to heterochromatic foci in *Neurospora* nuclei, consistent with prior findings with carboxy-terminal HP1-GFP fusion proteins.

Heterologous expression of jellyfish Green Fluorescent Protein (GFP) has become an indispensable tool for biological studies. We and others have reported successful expression of GFP in *Neurospora* (Freitag *et al.*, 2001; 2004a; 2004b; Fuchs *et al.*, 2002). Many applications of live cell imaging require simultaneous observation of more than one protein, however, prompting this study. Some pairs of fluorescent fusion proteins (e.g., green and red, or cyan and yellow) are typically observed simultaneously, because their excitation and emission spectra are well separated, but multi-color imaging is feasible with appropriate hardware and software. Here we describe the development of *Neurospora* plasmids for the expression of fusion proteins with an RFP gene that was derived from dsRed (or drFP583), first cloned from a *Discosoma* coral species (Matz *et al.*, 1999).

We began by testing transcriptional fusions of *Neurospora* promoters to an enhanced version of dsRed, *tdimer2(12)* (Campbell *et al.*, 2002). We chose this version because it matures more quickly and is brighter than dsRed, properties achieved by engineering and fusing two dsRed genes and separating them by a 42 bp spacer (which encodes the peptide linker “GHGTGSTGSGSSGT”), thus generating an almost perfect 677 bp direct repeat (Campbell *et al.*, 2002). We expected that *Neurospora* transformants with this gene would undergo RIP at high frequencies when crossed in wild type (i.e., *rid*⁺) strains.

The *tdimer2(12)* gene was excised from pDNT and pDCT (generously provided by Melissa Rolls, Univ. of Oregon) by digestion with *Apa* I and *Xba* I and inserted into pBM61 (Margolin *et al.*, 1997) to yield pMF331 and pMF332, respectively (Fig. 1). In pMF331, a plasmid to generate amino-terminal RFP fusions, a consensus Kozak translation initiation sequence precedes the 1391 bp *tdimer2(12)* gene, which is followed by a 12 bp glycine linker (“GSGG”) and a 63 bp region with multiple rare cloning sites (Fig. 1A). The *Apa* I and *Eco* RI sites can accept promoters for the genes to be tested. In pMF332, a plasmid to express carboxy-terminal RFP fusions, the multiple cloning site is followed by the glycine linker and the *tdimer2(12)* gene (Fig. 1B). To test amino-terminal RFP fusions in the pMF331 plasmid, the *Neurospora ccg-1* promoter from pMF272 (Freitag *et al.*, 2004a) was amplified by PCR with *Pfu* polymerase and inserted between the *Apa* I and *Eco* RI sites of pMF331 to yield pMF334 (Fig. 1C). The three plasmids are available from the Fungal Genetic Stock Center and their complete sequences have been submitted to the GenBank database (accession numbers: pMF331, DQ250997; pMF332, DQ250998; pMF334, DQ250999).

We transformed a *N. crassa his-3* strain (N623; FGSC# 6103) and *rid his-3* strain (N2240; FGSC# 9014) with pMF334 by electroporation (Margolin *et al.*, 1997). Transformants were selected on Vogel’s minimal medium with sorbose to induce colonial growth. Under these conditions, expression of RFP was detected with an Olympus SZX12 fluorescence stereo microscope in macroconidia and hyphae after three to five days in heterokaryotic prototrophic transformants. Fluorescent strains were picked to slants with minimal sucrose medium; fluorescence in aerial hyphae was detectable after overnight incubation at 32 C.

For conventional epifluorescence microscopy, *Neurospora* conidia were spotted on microscope slides in droplets containing minimal medium with 1.5 % sucrose; hyphae were examined on agar blocks (Freitag *et al.*, 2004b). We used a Zeiss Axioplan 2 microscope equipped with UV and FITC filter sets and images were captured with an Orca II digital camera (Hamamatsu, San Jose, CA) and Openlab software (Improvision, Coventry, UK). Images were processed with Photoshop software (version 4.0; Adobe, San Jose, CA). RFP accumulated in macroconidia, allowing visualization in germ tubes and young hyphae after 3 - 12 hours (Fig. 2).

We expected to observe high RIP frequencies in crosses of primary transformants because of the tandem duplication in *tdimer2(12)*. Indeed, 99 of 100 His⁺ progeny of one N623-derived transformant crossed to a *his-3* (N625; FGSC# 6525) strain were RFP⁺. Crosses of N2240-derived *rid* RFP⁺ transformants to a *rid his-3* strain (N2257; FGSC# 9015) yielded only RFP⁺ His⁺ progeny, as

FIGURE 1 FREITAG AND SELKER RFP EXPRESSION IN NEUROSPORA

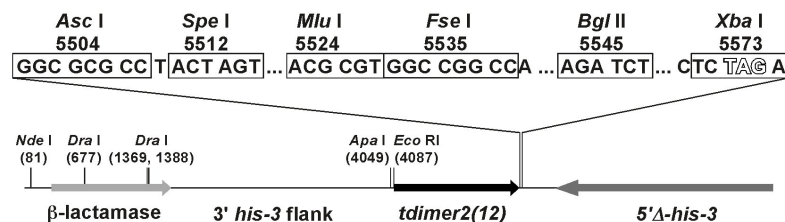
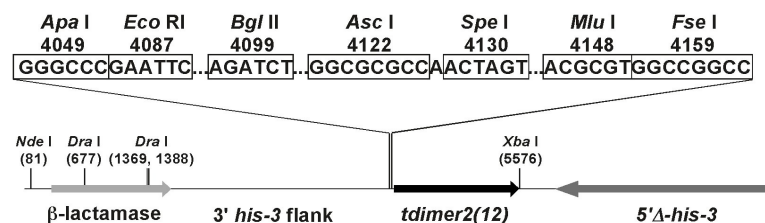
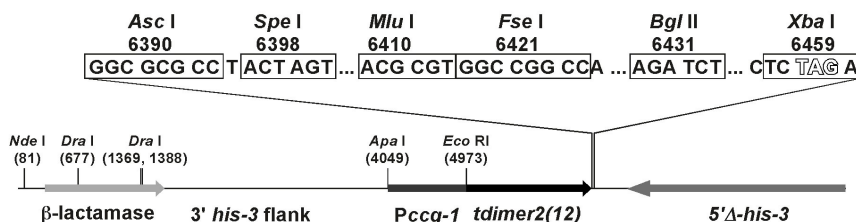
A pMF331 (N-terminal *tdimerRedN*) 8291 bp**B** pMF332 (C-terminal *tdimerRedC*) 8294 bp**C** pMF334 (*Pccg-1-tdimerRedN*) 9177 bp

Figure 1. Partial maps of pMF331, pMF332 and pMF334. (A) Amino-terminal fusions to the direct tandem repeat *tdimer2(12)* gene can be constructed in pMF331. The unique *Apa* I and *Eco* RI cloning sites allow insertion of promoter elements. (B) Carboxy-terminal fusions can be generated in pMF332. (C) For cytosolic RFP expression, pMF334 was constructed by inserting the *Neurospora ccg-1* promoter previously used in GFP constructs (Freitag *et al.*, 2004a; 2004b) into the *Apa* I and *Eco* RI sites of pMF331. Plasmid landmarks are illustrated as follows: β -lactamase gene for ampicillin resistance; 5' Δ -*his-3*, truncated *Neurospora his-3* allele for gene targeting; 3'-*his-3* flank, downstream flank of the *his-3* gene for gene targeting. The multiple cloning sites are shown above the construct schematics. Plasmids are available from the Fungal Genetic Stock Center; their complete sequences have been submitted to the GenBank database (accession numbers: pMF331, DQ250997; pMF332, DQ250998; pMF334, DQ250999).

expected because RIP is abolished in homozygous *rid* crosses (Freitag *et al.*, 2002). Subsequent crosses of homokaryotic *rid his-3⁺::tdimer2(12)* RFP⁺ strains with *rid⁺* strains yielded no RFP⁺ progeny (0/100). This confirmed our previous finding that *rid* shows little or no effect on RIP frequency in heterozygous crosses (Freitag *et al.*, 2002). We note that this RFP construct could serve as a sensitive locus to screen for RIP-deficient mutants. When RFP⁺ progeny are desired, however, we suggest that investigators introduce RFP constructs in *rid* strains to avoid RIP.

To test simultaneous live cell imaging of GFP- and RFP-tagged versions of the same protein, we generated a translational fusion of RFP with *Neurospora* heterochromatin protein 1 (HP1). We previously showed that carboxy-terminal GFP fusions of the HP1 gene (*hpo*) localized to heterochromatin and complemented DNA methylation defects and growth phenotypes of *hpo* null mutants (Freitag *et al.*, 2004b). We wanted to verify that neither the position nor the type of tag interfere with HP1 localization. We inserted an *Asc*I - *Bam*HI segment of the *hpo* coding region between the *Asc*I and *Bgl*II sites of pMF334 to generate a *tdimer2(120-hpo)* fusion (pMF344). We targeted this construct to *his-3* by electroporation of N623 and N2240 strains. RFP-HP1 was targeted to heterochromatin (Fig. 3, RFP), as expected from previous results with the HP1-GFP fusion protein (Freitag *et al.*, 2004b). To observe co-localization of RFP- and GFP-tagged HP1, we generated heterokaryons by growing RFP-HP1 transformants with an HP1-GFP strain (N2540; Freitag *et al.*, 2004b). Unforced heterokaryons were usually established overnight. Conidia with both RFP and GFP fluorescence were re-grown and observed after overnight growth at 32 C on Petri plates with sucrose as the carbon source. GFP- and RFP-tagged HP1 fusion proteins were localized to heterochromatic regions and their distribution was largely congruent (Fig. 3). These results indicate that neither the location of the tag (i.e., amino- vs. carboxy-terminal) nor the type of tag (coral RFP vs. jellyfish GFP) altered the expected localization of *Neurospora* HP1.

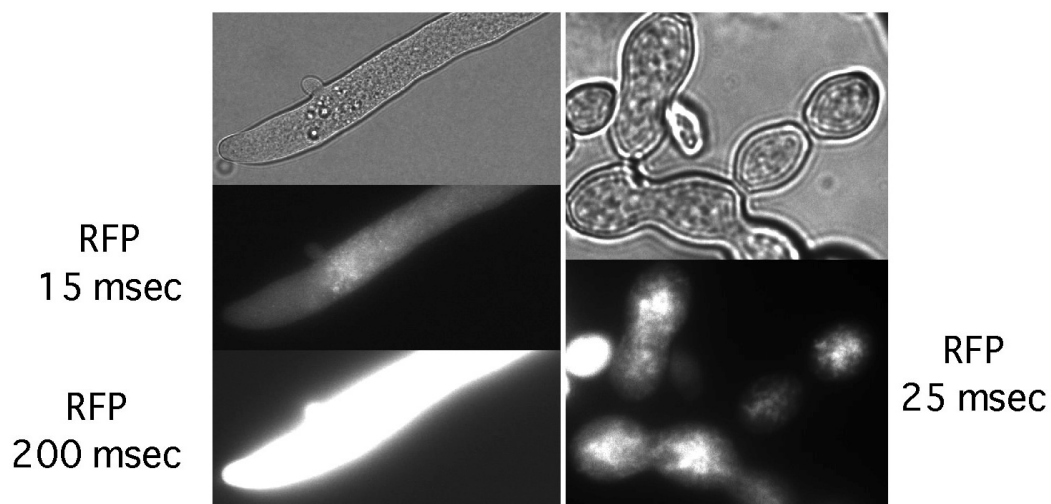


Figure 2. Expression of cytosolic RFP in *N. crassa*. *Neurospora* hyphae (left) and conidia (right) were examined by epifluorescence microscopy (see text). Even heterokaryotic transformants generated by insertion of pMF334 into the *his-3* locus gave strong fluorescence (exposure times at 100% lamp power and 70% gain are shown). We noticed that under these conditions RFP bleached faster than GFP constructs (Freitag *et al.*, 2004a; 2004b).

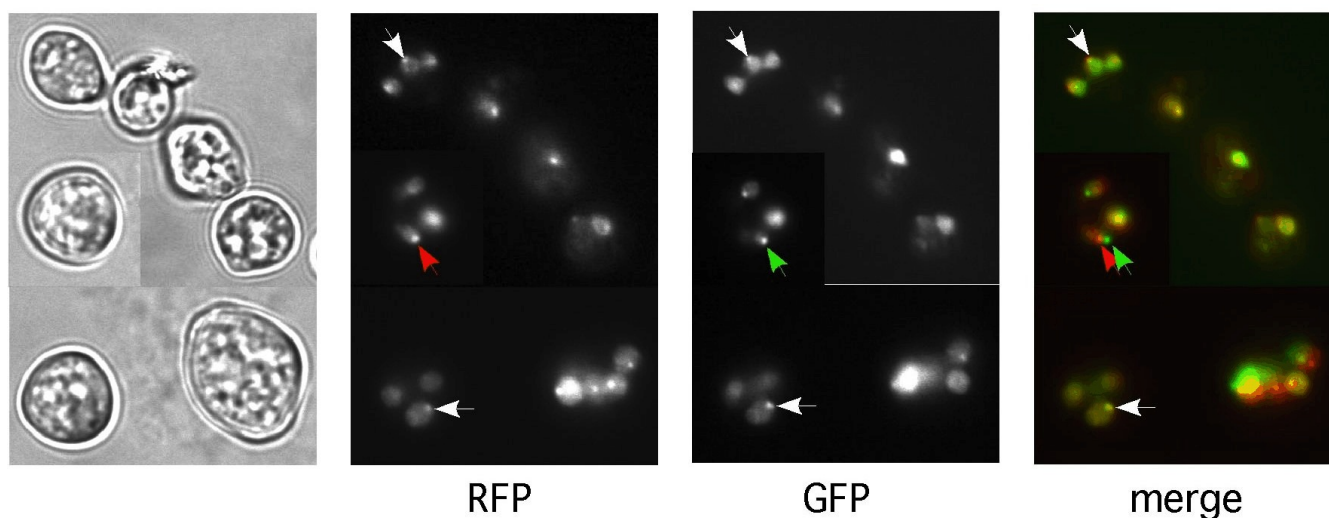


Figure 3. Colocalization of amino-terminal RFP-HP1 and carboxy-terminal HP1-GFP fusion proteins in heterochromatic regions. Conidia were examined by epifluorescence microscopy (see text). Heterochromatic foci in nuclei were detected in heterokaryotic strains expressing RFP-HP1 (RFP) and HP1-GFP (GFP), and overlapped with foci detected by staining with the DNA dye Hoechst 33258 (data not shown). The outline of nuclei is visible in the RFP and GFP panels of the lower right conidium with four nuclei. RFP-HP1 and HP1-GFP exhibit largely congruent distribution (see white arrows in the RFP, GFP and merged panels). Because of rapid nuclear movement, foci do not always align perfectly (red and green arrows in the RFP, GFP and merged panels). Panels are composites of three images.

In conclusion, expression of RFP in *Neurospora* can be achieved by using a strong fungal promoter. The cytosolic RFP expression profile was as expected from previous studies with the *cgc-1* promoter (Freitag *et al.*, 2004a). We generated a set of RFP plasmids that allow the expression of amino- or carboxy-terminal fusion proteins under the control of the *cgc-1* or other (e.g., endogenous) promoters. All three plasmids are designed for targeting to the *Neurospora his-3* locus but can also be used in co-transformation

experiments with other selectable markers (M. Freitag and E.U. Selker, unpublished results). In combination with our GFP constructs (Freitag *et al.*, 2004a; 2004b), these RFP constructs should prove useful for studies in which two different fluorescent tags are required.

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