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To verify the functionality of the DsRed encoded by pDsRed-SKL, we transformed a *S. macrospora* wildtype strain (Culture collection, Department for General and Molecular Botany, Ruhr-University of Bochum) either with plasmid pRHN1 or pDsRed-SKL (Fig. 2A and 2B).

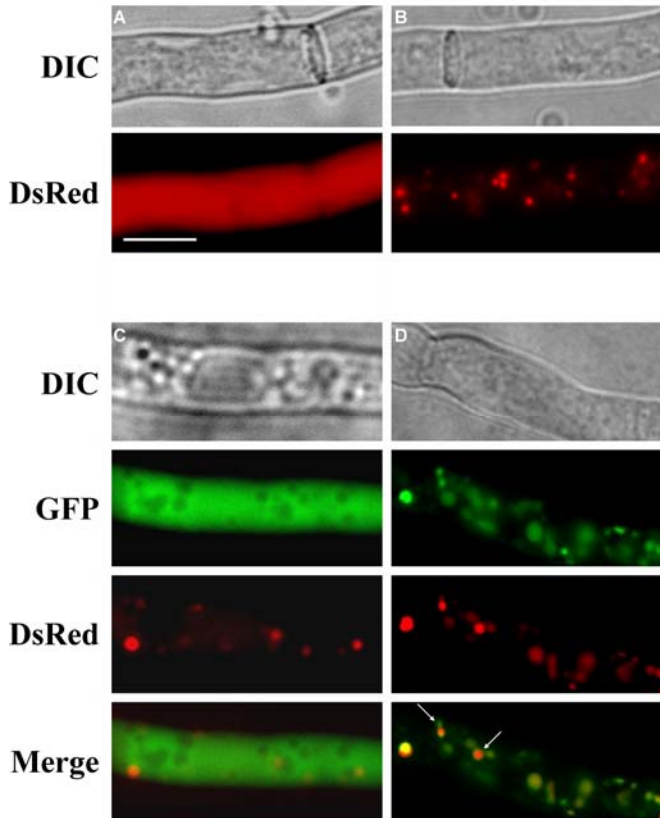


Figure 2. Expression of cytosolic and peroxisomal targeted DsRed and GFP in *S. macrospora*. (A) Hyphae of a strain transformed with pRHN1. (B) Hyphae of a strain transformed with pDsRed-SKL. Expression of DsRed-SKL leads to DsRed import into peroxisomes, visible as red dots in the cytoplasm. (C) Co-transformation with pIG1783 and pDsRed-SKL results in green fluorescent hyphae with red colored peroxisomes. (D) Parallel expression of SKL-tagged GFP and SKL-tagged DsRed. Both tagged proteins are not imported into the same exact organelle at the same rate, resulting in green or red dots after merging (see white arrows). Scale bar, 10 μ m. A color reproduction of the figure is available with the on-line version of this article (<http://www.fgsc.net/>).

Transformants were selected on CMS-medium (0.15 % (w/v) KH_2PO_4 ; 0.05 % (w/v) KCl; 0.05 % (w/v) MgSO_4 ; 0.37 % (w/v) NH_4Cl ; 1 % (w/v) glucose; 0.2 % (w/v) tryptone; 0.2 % (w/v) yeast extract; 10.8 % (w/v) saccharose; trace metals [10 mg/l ZnSO_4 ; 10 mg/l Fe(II)Cl_2 ; 10 mg/l MnCl_2]; 2 % (w/v) agar; pH 6.5) containing nourseothricin (50 $\mu\text{g/ml}$). To demonstrate the functionality of the DsRed-SKL protein in dual-color experiments, we combined the red peroxisomal label (pDsRed-SKL) with cytosolic GFP encoded by plasmid

pIG1783 (Pöggeler *et al.* 2003; Fig. 2C) or the green peroxisomal label GFP-SKL (pGPD::GFP-SKL; Ruprich-Robert *et al.* 2002; Fig. 2D). Co-transformants were selected either on CM-medium containing nourseothricin (50 $\mu\text{g/ml}$) and hygromycin (110 U/ml) (pDsRed-SKL and pIG1783) or on medium containing solely nourseothricin (50 $\mu\text{g/ml}$) (pDsRed-SKL and pGPD::GFP-SKL), since pGPD::GFP-SKL carries a phleomycin-resistance cassette, which is not selectable in *S. macrospora*.

Many yeast species and filamentous ascomycetes including *Acremonium chrysogenum*, *Neurospora crassa* and *A. nidulans* are sensitive to nourseothricin (Kück and Hoff 2006, Braus and Seiler, pers. communication). The antibiotic nourseothricin is commercially available (Werner BioAgents, Jena, Germany) and similarly expensive as hygromycin. Transformants and co-transformants were then inoculated on microscope slides overlaid with a thin layer of solid SWG minimal medium (0.4 % (w/v) arginine; 0.1 % (v/v) biotin; 2 % (w/v) glucose; 1x Westergaard (Westergaard and Mitchell 1947); 1.5 % (w/v) agar; pH 6.5) containing either nourseothricin or hygromycin and nourseothricin. After incubation for three days at 27 $^{\circ}\text{C}$, fluorescent proteins were visualized with appropriate filter combinations (Chroma filter set 49002, exciter ET470/40x, emitter ET525/50m and beamsplitter T495LP; Chroma filter set 49005 for green fluorescence, exciter ET545/30x, emitter ET620/60m and beamsplitter T570LP for red fluorescence) using an AxioImager M1 microscope (Zeiss, Jena, Germany) and an X-cite 120 PC lamp (EXFO, Ontario, Canada). Images were captured with a Photometrics CoolSNAP HQ^2 camera (Roper Scientific, Photometrics, Tucson, Arizona). Recorded images were edited with MetaMorph (VisitronSystems, Puchheim, Germany) and Adobe Photoshop CS2.

As expected, fluorescence appeared uniformly distributed throughout the cytoplasm of the hyphae in transformants carrying pRHN1 while transformation with pDsRed-SKL led to a punctate fluorescent pattern (Fig. 2A, B). An SKL-tagged DsRed used for peroxisomal studies has recently been described for the basidiomycete *Cryptococcus neoformans*. Similar to our results, a *C. neoformans* DsRED-SKL strain exhibited a punctate peroxisomal fluorescence (Idnurm *et al.* 2007).

Furthermore, we tested the fluorescence efficiency of DsRed-SKL in a cytoplasmic GFP-background by co-transformation of pDsRed-SKL and pIG1783 under nourseothricin/hygromycin double-selection (Fig. 2C). Transformants carrying both

plasmids displayed a well-defined punctate red fluorescent pattern clearly visible against the cytosolic background of GFP. To confirm that DsRed-SKL actually localizes to peroxisomes, we accomplished co-localization experiments with GFP-SKL. For this purpose, we analyzed 38 nourseothricin-resistant transformants from a pDsRed-SKL/pGPD::GFP-SKL co-transformation experiment. More than 55 % of the investigated transformants (21 transformants) exhibited DsRed as well as GFP fluorescence, while the remaining transformants either emitted only the red fluorescence of DsRed-SKL or no fluorescence. As shown in Fig. 2D, the DsRed-SKL fusion produced a fluorescence pattern, which coincides with the fluorescence pattern of the GFP-SKL. Thus the DsRed-SKL fusion construct is targeted to peroxisomes. However, because of rapid peroxisome movement in the cytosol, signals did not align perfectly in all cases (Fig. 2D).

In conclusion, expression of pDsRed-SKL leads to the import of DsRed into peroxisomes and results in the detection of fluorescence signals after excitation with green light. Vector pDsRed-SKL leads to high-level expression of the *DsRed-SKL* gene and the import of DsRed into peroxisomes.

Plasmid pDsRed-SKL is consequently a valuable tool for analyzing the sub-cellular localization, particularly the peroxisomal localization, of GFP-tagged proteins by means of dual-color experiments. It could be also used for the analysis of mutants affected in peroxisome biogenesis. The vector was primarily constructed for *S. macrospora*; however, it should prove very effective in a wide variety of filamentous ascomycetes in which transformation is based on nourseothricin resistance.

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