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Second-Order Nonlinear Optical Properties of Fluorescent Proteins for Second-Harmonic Imaging

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Addition to Materials and methods

Expression and purification of recombinant proteins

The fluorescent proteins were obtained by heterologous overexpression in 15 *Escherichia coli* and purification. Plasmids coding for EGFP and DsRed were used as described in Cotlet *et al.* (1). The gene coding for EYFP was cloned from pEYFP (Clontech) into pBAD/HisA in a similar way as described in Cotlet *et al.* The fluorescent proteins were expressed in *E. coli* Top-10 cells which were cultured in 4 1 LB-medium supplemented with ampicillin (100 mg/l). Expression of the proteins

- ²⁰ was induced by 0.2% arabinose at an optical density of 0.5 at 600 nm. The cells were harvested after 12 hours and a cell lyste was obtained. The fluorescent proteins, carrying an N-terminal His-tag, were purified under native conditions by Ni-affinity chromatography (1 ml HisTrap columns, GE Healthcare) according to the manufacturers recommended protocol and concentrated using a Vivaspin
- ²⁵ concentrator (cutoff 5000 Da). The final pH was 7.3. The proteins were checked for purity and mass by SDS-PAGE and mass spectrometry (Fig. S1). While the method used for MS suggests DsRed to be mono- and dimeric, a gel filtration experiment in the buffer used for HRS measurements demonstrates a tetrameric form of DsRed. The concentration of the samples used for HRS measurements was determined using
- ³⁰ the extinction coefficient of the chromophore at the wavelength of maximal absorption (listed in Table S1).

	EGFP	EYFP	DsRed
$\lambda_{max}(nm)$	488	513	558
$\varepsilon_{\rm max} ({\rm M}^{-1} {\rm cm}^{-1})$	56000	83400	35000
Abs _{max}	1,0458	2,1387	1,2199
Distance (cm)	0,2	0,2	0,2
Concentration (µM)	93,4	128	174

Table S1: Concentrations of proteins based on the extinction coefficient ε_{max} at the wavelength λ_{max} maximal absorption Abs_{max} of the chromophore (2).

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⁵ Figure S1: Mass spectrometry of (A) EGFP, (B) EYFP and (C) DsRed. The MALDI-TOF measurements were performed on a Bruker Ultraflex II mass spectrometer. Sinapinic acid was used as matrix and all measurements were carried out in positive linear ion mode, with a selected mass range of (A) and (B) 10 – 60 kDa, and (C) 20 – 140 kDa.

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References

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