

Supporting Information

Transient IR Spectroscopy Identifies Key Interactions and Unravels New Intermediates in the Photocycle of a Bacterial Phytochrome

Joachim Kübel,^{a†} Manoop Chenchiliyan,^{a†} Saik Ann Ooi,^a Emil Gustavsson,^a, Linnéa Isaksson,^a Valentyna Kuznetsova,^b
Janne A. Ihalainen,^b Sebastian Westenhoff^{*a} and Michał Maj^{*a}

^a Department of Chemistry and Molecular Biology, University of Gothenburg, Gothenburg 40530, Sweden; * E-mail: michal.maj@gu.se, westenho@chem.gu.se ^b Nanoscience Center, Department of Biological and Environmental Science, University of Jyväskylä, Jyväskylä 40014, Finland.

[†] These authors contributed equally to this work

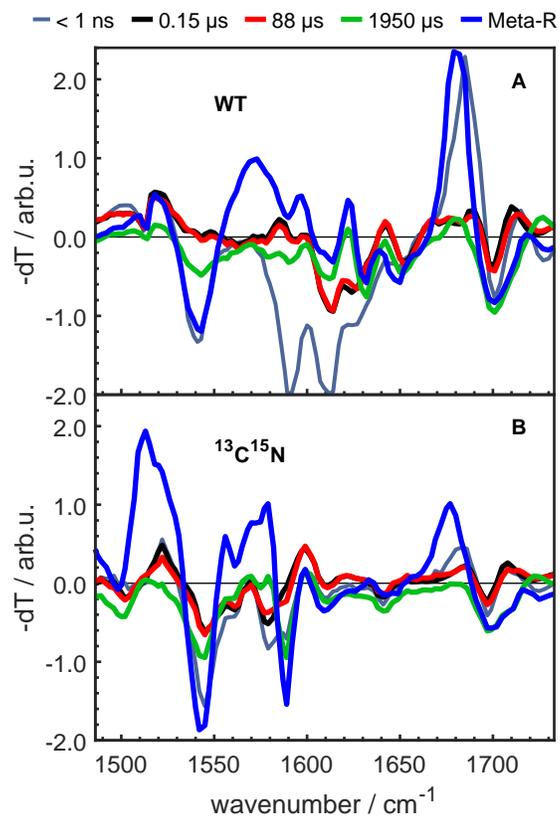


Fig. S 1 Species spectra obtained from a global fit of wild-type CBDPHY (A) and $^{13}\text{C}^{15}\text{N}$ -labeled CBDPHY (B), in which all protein carbon and nitrogen atoms are isotope labeled, except those of the chromophore.

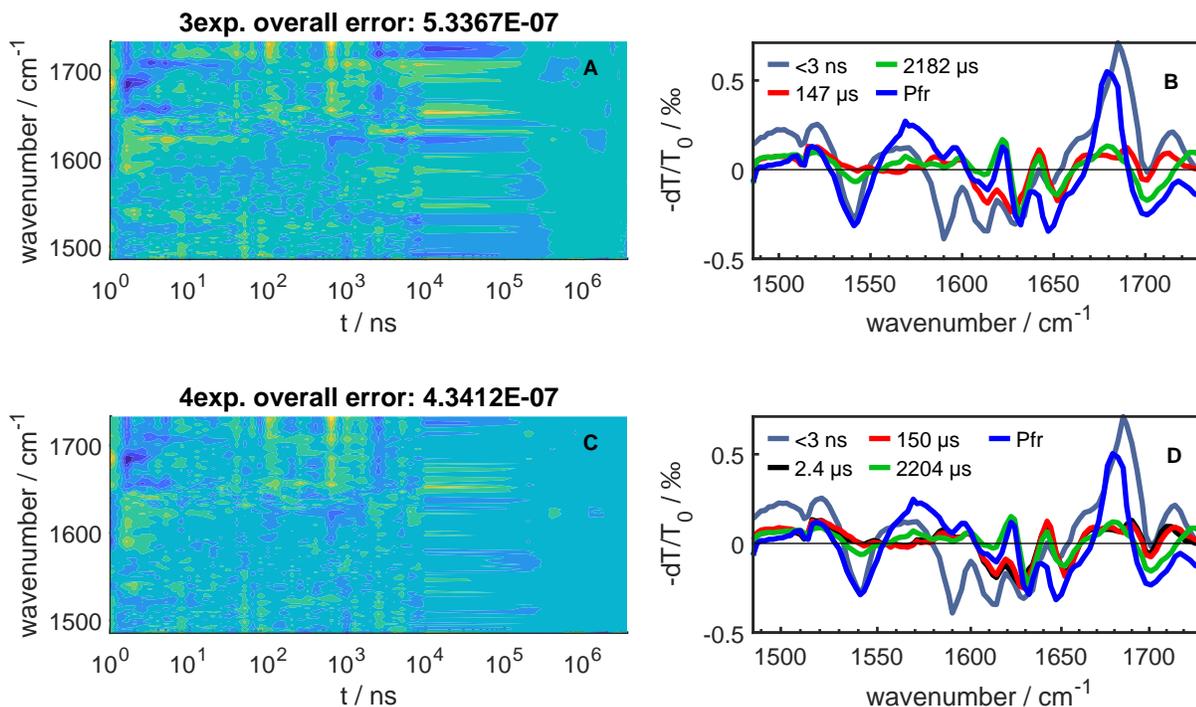


Fig. S 2 Comparison between 3- and 4-exponential fit for wild-type CDBPHY showing the presence of the second Lumi-R state. The contour plots show the residuals for a fit with 3 or 4 components. For the specific dataset shown here no timepoints were collected between 1E4 and 2E5 microseconds.

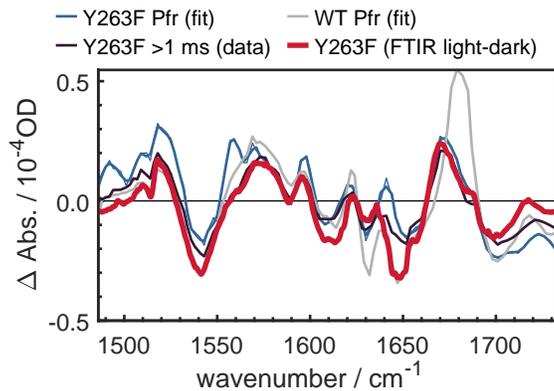


Fig. S 3 Pfr spectra determined from the global fit of nanosecond transient absorption data for wild-type CDBPHY and the Y263F mutant. The latter can be compared to the averaged spectra at delay times >1 ms, and the scaled FTIR difference spectrum. The discrepancy between the Pfr from the fit and the averaged spectra at long delays can be attributed to weak contributions from pre-0 signals. The transient transmission data has been converted to approximate transient absorption signals.