## **Supplementary Material**

# Biochemical properties of Paracoccus denitrificans FnrP: Reactions with

## molecular oxygen and nitric oxide

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#### Fusion protein

MSPILGYWKIKGLVQPTRLLLEYLEEKYEEHLYERDEGDKWRNKKFELGLEFPNLPYYIDGDVKLTQSMAIIRYI ADKHNMLGGCPKERAEISMLEGAVLDIRYGVSRIAYSKDFETLKVDFLSKLPEMLKMFEDRLCHKTYLNGDHVTH PDFMLYDALDVVLYMDPMCLDAFPKLVCFKKRIEAIPQIDKYLKSSKYIAWPLQGWQATFGGGDHPPKSDLVPRG SPGISGGGGGILDSMAHDAALCATQQPCEICPIRYRAVCAHCEGDELTELEEMKFYRRYEAGQVVVWAGDRMDFV ASVVAGMAGLTQQLEDGRTQMVGLLLPSDFLGRPGRDMAAYTVTATSDLVLCCFRRKPFEKLLIDNPRIASRLLE MTLDELDAARDWLLLLGRKSAREKIASLLVILARREAALIKRRPEGRITIELPLTREAMADYLGLTLETVSRQMS ALKREGVIELDGKRRVIVPSFVRLVTESGDDSDGGPLS

Expected mass: 55117.9

#### GST remnant

MSPILGYWKIKGLVQPTRLLLEYLEEKYEEHLYERDEGDKWRNKKFELGLEFPNLPYYIDGDVKLTQSMAIIRYI ADKHNMLGGCPKERAEISMLEGAVLDIRYGVSRIAYSKDFETLKVDFLSKLPEMLKMFEDRLCHKTYLNGDHVTH PDFMLYDALDVVLYMDPMCLDAFPKLVCFKKRIEAIPQIDKYLKSSKYIAWPLQGWQATFGGGDHPPKSDLVPR

Expected Mass: 26166.4

#### Linker FnrP

GSPGISGGGGGILDSMAHDAALCATQQPCEICPIRYRAVCAHCEGDELTELEEMKFYRRYEAGQVVVWAGDRMDF VASVVAGMAGLTQQLEDGRTQMVGLLLPSDFLGRPGRDMAAYTVTATSDLVLCCFRRKPFEKLLIDNPRIASRLL EMTLDELDAARDWLLLLGRKSAREKIASLLVILARREAALIKRRPEGRITIELPLTREAMADYLGLTLETVSRQM SALKREGVIELDGKRRVIVPSFVRLVTESGDDSDGGPLS

Expected Mass: 28969.5

**Fig. S1.** Sequence information and predicted molecular masses for GST-FnrP and thrombin derived cleavage products. The N-terminal Cys-rich region of FnrP occurs between the highlighted Met residues, and Cys residues are highlighted in yellow in the cleaved sequence. The released apo-FnrP species had a predicted mass of 28,969.5 Da.

		dimerization helix		
		Y Y		
EcFNR	:	-HHPSFAQALETSMVCEIPFETLDDLSGKMPNLRQQMMRLMSGEIKGDQDMILLLSKKNA	:	166
PdFnrP	:	AAYTVT <mark>A</mark> TSDLV <mark>L</mark> CCFRRKPFEKLLIDN <mark>P</mark> RIASRLLEMTLDELDAARDWLLLL <mark>L</mark> GRKSA	:	157
BjFixK2	:	HRLAAE <mark>A</mark> IIDTS <mark>V</mark> RLVKRSSLEKAAGIDVQVARKLWAMTAGELRHAEDHMLLLGRKTA	:	157
RrCooA	:	<mark>A</mark> TERTE <mark>V</mark> RFADIRTFEQKLQTC <mark>P</mark> SMAWGLIAILGRALTSCMRTIED <mark>L</mark> MFHDI	:	135
RsNnrR	:	-TYPATAV <mark>A</mark> AADCVTLSWPVRLWADFTANY <mark>P</mark> GFATESYRTLGARLGEMQNRITELATQQV	:	150
EcCRP	:	QERSAWVR <mark>A</mark> KTACEVAEISYKKFRQLIQVN <mark>P</mark> DILMRLSAQMARRLQVTSEKVGNLAFLDV	:	140
		and the second sec		
EcFNR	:	EERLAAFIYNLSRRFAQRGFSPREFRLTMTRGDIGNYLGLTVETISRLLGRFQKSG	:	222
PdFnrP	:	REKIASLLVILARREAALIKRRPEGRITIELPL <mark>TR</mark> EAMADYL <mark>G</mark> LTLE <mark>TVSRQM</mark> SALKREG	:	217
BjFixK2	:	MERVATFLLEMDRRLAVAGMMALPMCRRDIGDYLGLTLETVSRALSQLHTQG	:	209
RrCooA	:	KQRIAGFFIDHANTTGRQTQGGVIVSVDF <mark>TVEEIANLIG</mark> SSRQ <mark>TTST</mark> ALNSLIKEG	:	191
RsNnrR	:	EQRVAACLLRLVNQSGRKVEAGIEISFPITRQNISEMT <mark>GTTLHTVSR</mark> LLSAWEREG	:	206
EcCRP	:	TGRIAQTLLNLAKQPDAMTHP-DGMQIKITRQEIGQIVGCSRETVGRILKMLEDQN	:	195

**Fig. S2.** Partial sequence alignment of CRP/FNR family proteins. The dimerization helix is indicated. Two key residues within the helix that are important for *E. coli* FNR association state, Arg140 and Asp154, are indicated by orange arrowheads. Proteins are *E. coli* FNR (EcFNR), *Paracoccus denitrificans* FnrP (PdFnrP), *Bradyrhizobium japonicum* FixK2 (BjFixK2), *Rhodospirillum rubrum* CooA (RrCooA), *Rhodobacter sphaeroides* NnrR (RsNnrR) and *E. coli* CRP (EcCRP). The alignment was generated using Clustal Omega [1] and annotated using Genedoc [2].



**Fig. S3.** Temporal regulation of denitrification by FnrP and NnrR in *P. denitrificans*. Under semiaerobic conditions FnrP activates the expression of nitrate (*nar*) and nitrous oxide (*nos*) *reductases* [3]. Endogenously derived NO during nitrate/nitrate respiration causes a transient burst of NO production under anaerobic conditions, leading to nitrosylation of FnrP. Concomitant activation of NnrR by NO allows the timely expression of nitrite (*nir*), nitric oxide (*nor*) reductase, completing the denitrification metabolic pathway. The levels of O<sub>2</sub>, NO and N<sub>2</sub> are those found in the head space of the culture. Adapted from Bergaust *et al* [3].

### References

- Sievers F, Wilm A, Dineen D, Gibson TJ, Karplus K, Li W, Lopez R, McWilliam H, Remmert M, Soding J, Thompson JD, Higgins DG (2011) Fast, scalable generation of highquality protein multiple sequence alignments using Clustal Omega. Mol Sys Biol 7:539
- 2. Nicholas KB, Nicholas HBJ (1997) Genedoc: A tool for editing and annotating multiple sequence alignments, *Distributed by the authors*.
- Bergaust L, van Spanning RJ, Frostegard A, Bakken LR (2012) Expression of nitrous oxide reductase in Paracoccus denitrificans is regulated by oxygen and nitric oxide through FnrP and NNR. Microbiology 158:826-834.