Evidence for the Slow Reaction of Hypoxia-Inducible Factor Prolyl Hydroxylase 2 with Oxygen

Emily Flashman¹, Lee M. Hoffart², Refaat B. Hamed¹, J. Martin Bollinger, Jr.^{2‡}, Carsten Krebs^{2‡} and Christopher J. Schofield^{1‡}

¹ Department of Chemistry and Oxford Centre for Integrative Systems Biology, 12 Mansfield Road, Oxford, OX1 3TA, U.K.

² Department of Chemistry and Department of Biochemistry and Molecular Biology, The Pennsylvania State University, University Park, PA 16802, U.S.A.

[‡] Corresponding authors:	Email: christopher.schofield@chem.ox.ac.uk
	Tel. +44 1865 275625
	Fax: +44 1865 275674
	Email: cdk10@psu.edu
	Tel. +1 814 865 6089
	Fax: +1 814 865 2927
	Email: jmb21@psu.edu
	Tel. +1 814 863 5707
	Fax: +1 814 865 2927

Short title: Prolyl Hydroxylase 2 Reaction with Oxygen

Figure S1. Determination of rate constants for 2OG conversion to succinate and CODD hydroxylation

As seen in Figure 2 and described in the main text, LC-MS was used to monitor the timedependent conversion of 2OG to succinate (at 5°C) in the presence (black circles) and absence (white circles) of CODD (shown on a logarithmic scale); MALDI MS was used to quantify CODD hydroxylation with time (red circles). In order to determine rate constants, data were fitted by the equation, $f=y0+a^*(1-^{(-b^*x)})$, using SigmaPlotTM. Apparent first-order rate constants of (0.018 ± 0.0014) s⁻¹ and (0.0006 ± 0.0001) s⁻¹ were obtained for succinate production in the presence and absence of CODD, respectively. An apparent first-order rate constant of (0.013 ± 0.003) s⁻¹ was obtained for hydroxylation of CODD.

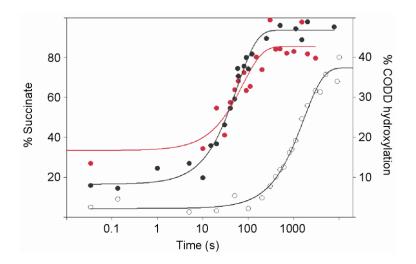
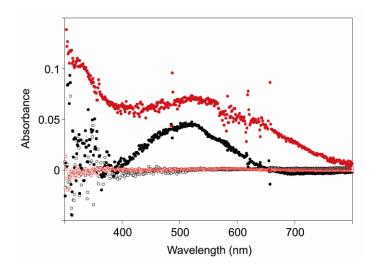


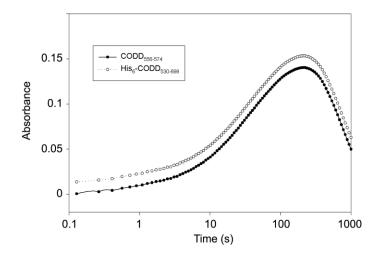
Figure S2. The PHD2.Fe.2OG and PHD2.Fe.2OG.CODD complexes demonstrate characteristic spectral features.



Addition of saturating 2OG to anoxic PHD2:Fe(II) (red open circles) and PHD2:Fe(II):CODD (black open circles) led to formation of the assigned PHD2:Fe(II):2OG (red filled circles) and PHD2:Fe(II):2OG:CODD complexes (black filled circles), with absorbance features at λ_{max} 527nm and 521nm, respectively. All spectra shown are 'blanked' against the spectrum of anoxic PHD2:Fe(II). Such features have been previously observed in

other 2OG-dependent oxygenases and shown to arise due to a metal-to-ligand charge-transfer band upon bidentate 2OG coordination to the active site Fe(II) [1]. A change in λ_{max} from ~530nm to ~520nm is often observed upon prime substrate binding.

Figure S3. UV-visible absorption spectra comparing rates of formation for the 320nm species in the presence of HIF-1 $\alpha_{556-574}$ CODD peptide and His₆-HIF₅₃₀₋₆₉₈ CODD protein substrates.



Oxygen-saturated buffer solution was rapidly mixed with a PHD2:Fe(II):2OG:CODD complex, where CODD was either a 19mer peptide (CODD₅₅₆₋₅₇₄, black circles) or a longer HIF-1 α protein substrate (His₆-CODD₅₃₀₋₆₉₈, white circles), which has a greater affinity for PHD2 [2, 3]. UV-visible absorption spectra monitored over time (shown on a logarithmic scale) demonstrated comparable rates of formation and degradation of the 320nm species (0.06s⁻¹ and 0.001s⁻¹, respectively) in the presence of both substrates. This indicates that the slow rate of formation of this species observed in the presence of CODD₅₅₆₋₅₇₄ is not due to inefficient substrate binding.

References

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