

## SUPPLEMENTARY INFORMATION

### **Converting cytochrome *c* into a DyP-like metalloenzyme**

Issei Omura<sup>a</sup>, Koichiro Ishimori,<sup>a,b</sup> and Takeshi Uchida<sup>\*a,b</sup>

<sup>a</sup>Graduate School of Chemical Sciences and Engineering, Hokkaido University, Sapporo 060-8628, Japan

<sup>b</sup>Department of Chemistry, Faculty of Science, Hokkaido University, Sapporo 060-0810, Japan

#### **Corresponding Author**

\*Department of Chemistry, Faculty of Science, Hokkaido University, Sapporo 060-0810, Japan. E-mail: uchida@sci.hokudai.ac.jp.

**Supplementary Table 1: Oligonucleotides used for construction of expression vectors for mutants. The underlined bases signify the introduced mutations.**

Mutants	Primers (top, sense; bottom, anti-sense)
M80V	5'— ACGAAAG <u>TG</u> ATCTTCGCGGGCATCAAA — 3' 5'— GAAGAT <u>CA</u> CTTTCGTGCCCCGGGATGTA — 3'
Y48H	5'— TTCACG <u>CA</u> TACGGACGCGAACAAAAAC — 3' 5'— GTCCGTAT <u>G</u> CGTGAAGCCCGGCCTG — 3'
Y67H	5'— ATGGA <u>AC</u> ATCTCGAGAACCCGAAAAAA — 3' 5'— CTCGAGAT <u>G</u> TTCATCAGCGTTTCTTC — 3'
Y97H	5'— ATCGCG <u>CA</u> TCTGAAAAAGGCGACGAAC — 3' 5'— CGCGAAG <u>AC</u> CTGATCGCGCATCTGAAA — 3'
P76W	5'— TACATC <u>T</u> GGGGCACGAAAATGATCTTC — 3' 5'— CGTGCCC <u>C</u> AGATGTATTTTTTCGGGTT — 3'
G41S	5'— AAAAC <u>G</u> TCTCAGGCGCCGGGCTTCACG — 3' 5'— CGCCTGAG <u>A</u> CGTTTTGCGGCCGAACAG — 3'
A51V	5'— ACGGAC <u>G</u> TGAACAAAAACAAAGGCATC — 3' 5'— TTTGTT <u>C</u> AGTCCGTGTACGTGAAGCC — 3'
G29D	5'— AAAAC <u>C</u> GACCCCAACCTGCACGGCCTG — 3' 5'— GTTGGG <u>G</u> TGTTTGTGTTTGCCGCC — 3'

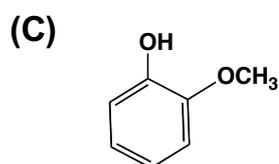
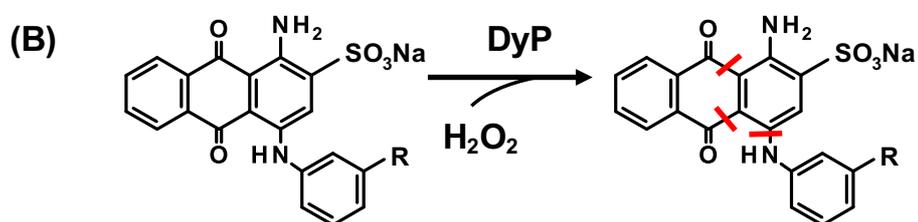
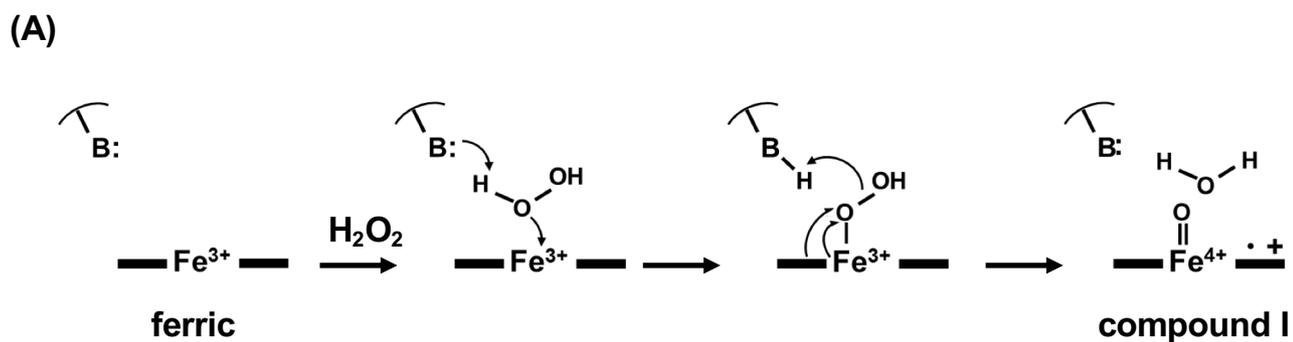


Fig. S1 (A) The mechanism of compound I formation by the reaction of DyP and  $\text{H}_2\text{O}_2$  (Sugano *et al.*, *J. Biol. Chem.*, 2007, 282, 36652–36658). (B) degradation of RB19. (C) molecular structure of guaiacol.

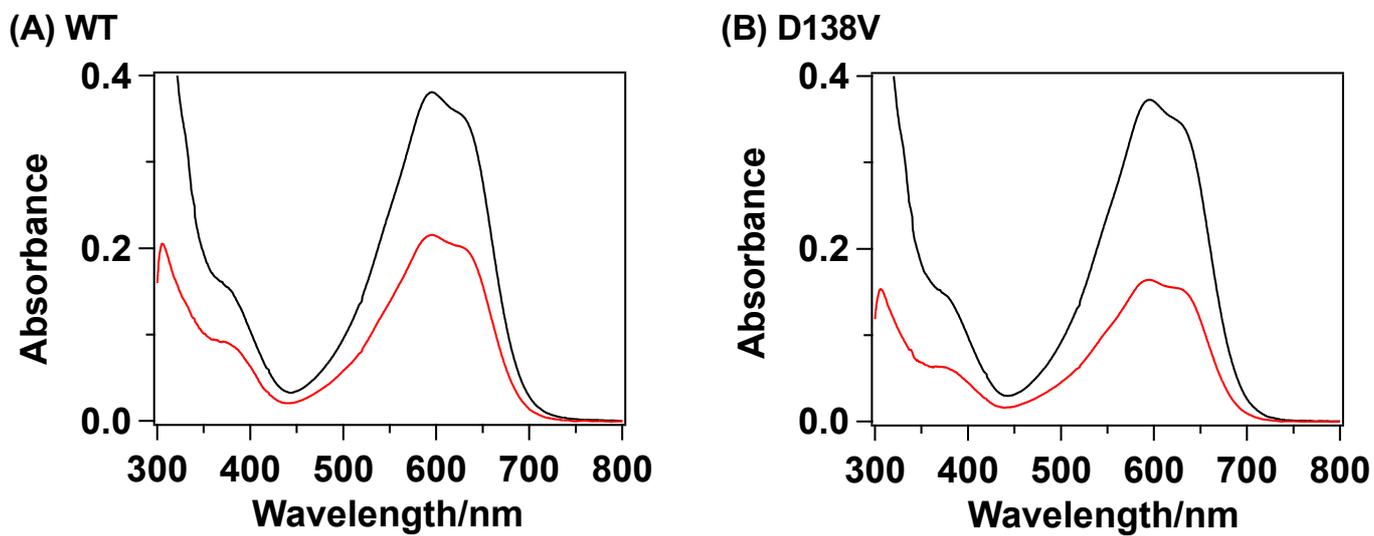


Fig. S2 Absorbance of the supernatant of culture medium with RB19 (40  $\mu$ M) of *E. coli* expressed (A) WT and (B) D138V *VcDyP*; 0h (black line) and 24h (red line).

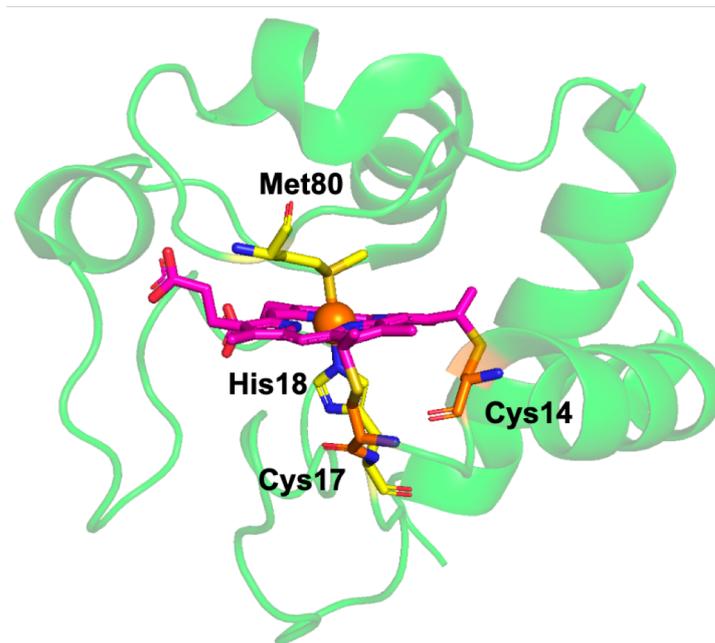
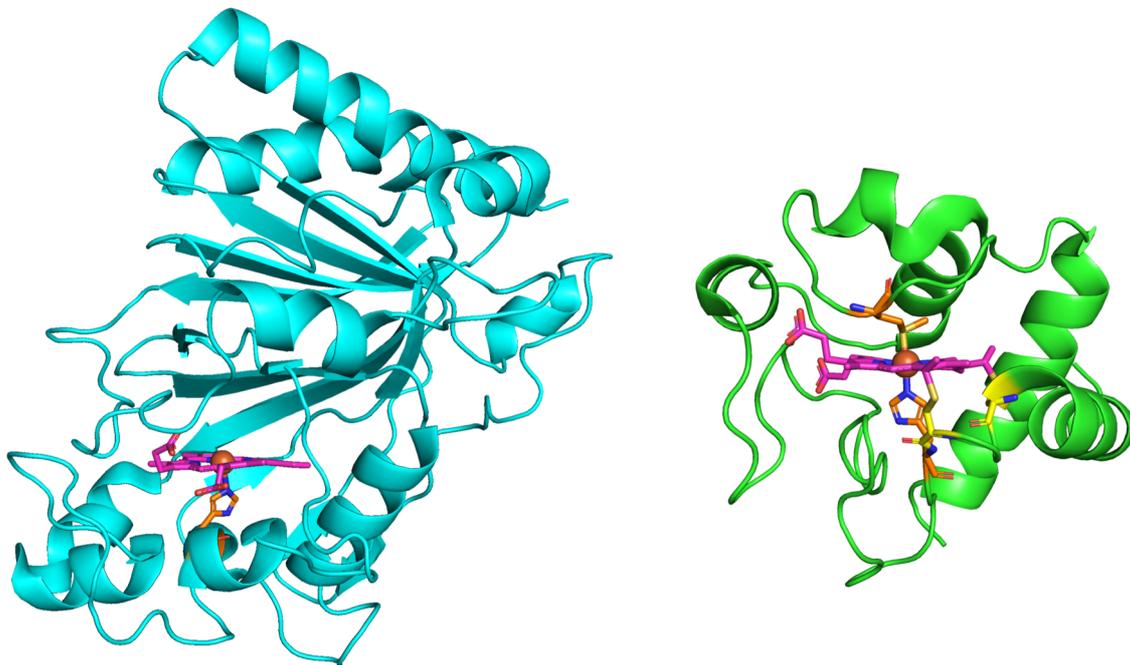


Fig. S3 X-ray crystal structure of cyt *c* (PDB ID: 6K9I)

(A)



(B)

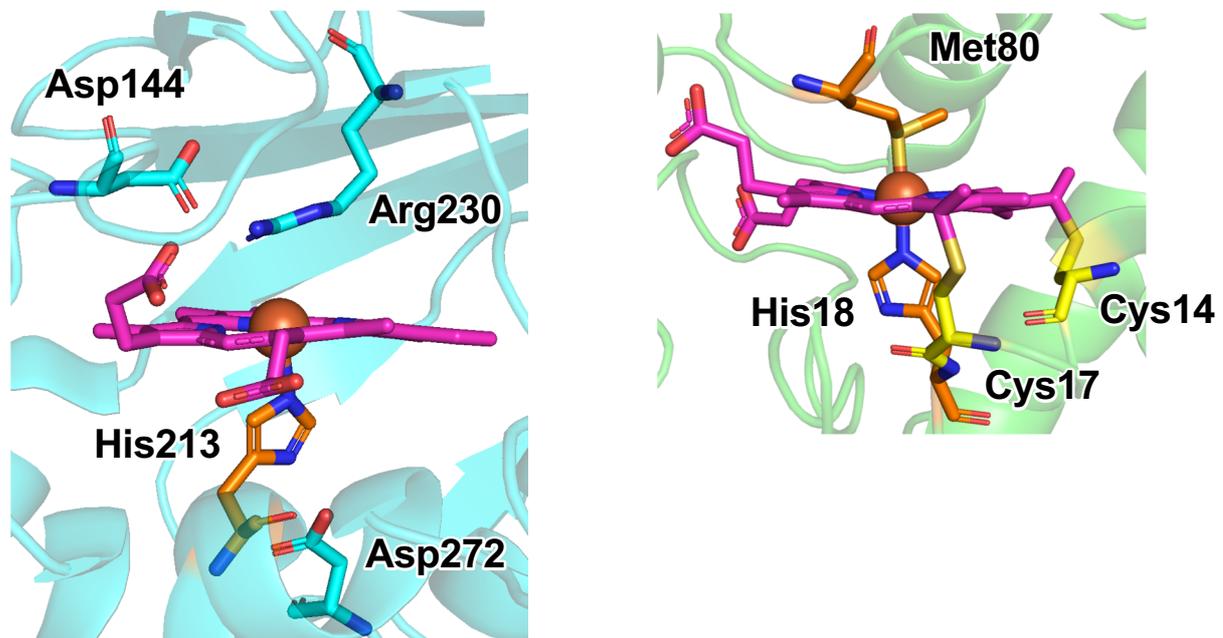


Fig. S4 Structural comparison between *VcDyP* and *cyt c*. Comparison of overall structures (A) and structures around heme (B).

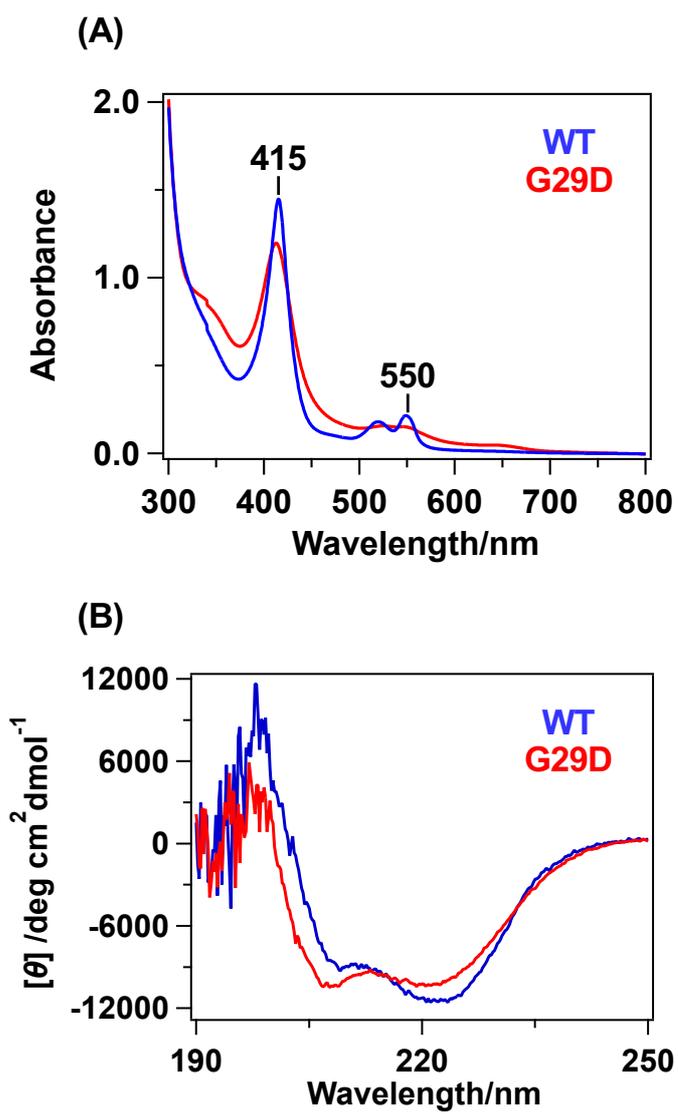


Fig. S5 (A) Absorbance of supernatant of lysate of *E. coli* expressed WT and G29D cyt *c*. (B) CD spectra of WT and G29D as purified.

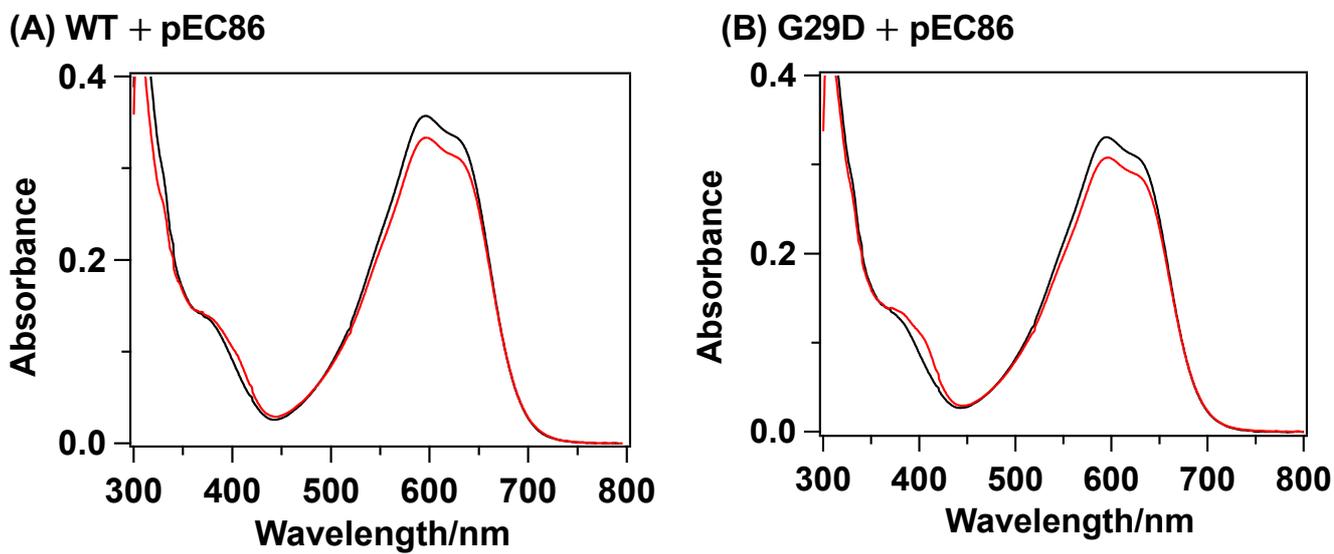


Fig. S6 Absorbance of the supernatant of culture medium with RB19 (40  $\mu$ M) of *E. coli* expressed (A) WT cyt *c* + pEC86, (B) WT cyt *c*, (C) G29D cyt *c* + pEC86, and (D) G29D cyt *c*; 0h (black line) and 24h (red line).