## Materials and methods

### 2.1 Materials

NiSO ${ }_{4}, \mathrm{ZnCl}_{2}, \mathrm{MnCl}_{2}, \mathrm{CoCl}_{2}, \mathrm{FeSO}_{4}$, L-ascorbic acid, and 2-oxopentanedioic acid (2-OG) were obtained from Sigma-Aldrich (St. Louis, MO, USA). Anti-PHD3 was obtained from Abcam (Cambridge, MA), and anti-PHD2 was from Cell Signaling Technology (Beverly, MA). The HIF-1 $\alpha$ peptide containing residues 556 to 575 denoted as F-P564 (fluorescein isothio-cyanate (FITC)-AminoCaproic Acid (ACA)-DLDLEALAPYIPADDDFQLR) was synthesized by AnyGen (KwangJu, Korea). All other chemicals were of the highest grade commercially available.

### 2.2 Site- directed mutagenesis of PHD3 <br> 2.2 Site-directed mutagenesis of PHD3

PCR primers were synthesized by Bioneer (Deajeon, Korea). Site-directed mutagenesis of Cys, Glu, Asp, Arg or His residues of PHD3 was conducted using the quikchange mutagenesis protocol provided by Stratagene, USA. All mutations were confirmed by DNA sequencing.

### 2.3 Preparation of recombinant proteins

Proteins were prepared as previously described (Cho et al, 2005). Briefly, plasmids for human VHL (amino acids 54-213) and human Elongin B (amino acids 1-118) in pGEX-4T-1 and for human Elongin C (amino acids 17-112) in pET29b were co-expressed in Escherichia coli (E. coli) BL21(DE3). Full-length human PHD2, truncated human PHD2 (amino acids 184-418) denoted as PHD2F and PHD2T, respectively, and full-length human PHD3 were subcloned into pET-28a (Novagen), pET-21b (Novagen), and pET-22b (Novagen), respectively and expressed in E. coli BL21(DE3). After induction with 0.5 mM isopropyl $\beta$ -D-thiogalactoside at $18^{\circ} \mathrm{C}$ for 15 hours, GST-VBC was purified using glutathione-Sepharose -
(Amersham Biosciences) and PHD2F, PHD2T, and PHD3 were lysed by sonication in the buffer containing 50 mM sodium phosphate $(\mathrm{pH}=8.0), 500 \mathrm{mM} \mathrm{NaCl}, 14 \mathrm{mM} \beta$ mercaptoethanol, 1 mM phenylmethanesulfonyl fluoride, and 10 mM imidazole. Triton X100 was added into the lysate to final concentration of $1 \%$ and the lysate was incubated for 10-30 minutes at $4{ }^{\circ} \mathrm{C}$. Proteins were purified using Ni-NTA agarose (Qiagen). The purified recombinant proteins were confirmed by SDS-PAGE and quantified by Bradford assay (BioRad).

### 2.4 Fluorescence polarization (FP)-based activity assays

For determination of inhibitory effects of divalent metal ions on PHDs, $1 \mu \mathrm{M}$ of F-P564 was incubated with $1 \mu \mathrm{~g}$ PHD2F, $0.25 \mu \mathrm{~g}$ PHD2T or $1 \mu \mathrm{~g}$ PHD3 in hydroxylation buffer containing $200 \mu \mathrm{M}$ ascorbic acid, $200 \mu \mathrm{M} 2-\mathrm{OG}, 5 \mathrm{mM} \mathrm{KCl}$, and $1.5 \mathrm{mM} \mathrm{MgCl}{ }_{2}$ in the absence and presence of increasing concentrations of $\mathrm{Zn}(\mathrm{II}), \mathrm{Ni}(\mathrm{II}), \mathrm{Mn}(\mathrm{II})$, and Co (II) at 30 ${ }^{\circ} \mathrm{C}$. To analyze effects of $\mathrm{Zn}(\mathrm{II})$ on PHDs’ activities, concentrations of $\mathrm{Zn}(\mathrm{II})$ were varied from 0 to $160 \mu \mathrm{M}$. For competition with Fe (II), varying concentrations of Zn (II) and $\mathrm{Fe}(\mathrm{II})$ were used. For $\mathrm{Zn}(\mathrm{II})$ pretreatment, PHD3 was incubated with 40 or $80 \mu \mathrm{M}$ of $\mathrm{Zn}(\mathrm{II})$ for 10 minutes at room temperature, followed by assaying in the presence of 0.5 mM DTT or 1 mM TCEP, respectively. After 2 hours of incubation at $30^{\circ} \mathrm{C}$, the hydroxylation reaction was stopped by heating for 1 minute at $95^{\circ} \mathrm{C}$. VBC $(200 \mathrm{nM})$ was then added to the reaction solution and diluted to a final peptide concentration of 100 nM in EBC buffer ( 50 mM Tris$\mathrm{HCl}, \mathrm{pH} 8.0,120 \mathrm{mM} \mathrm{NaCl}, 0.5 \%$ Nonidet P-40). The hydroxylation yields were analyzed by FP measurements using an Appliskan multimode microplate reader (Thermo Scientific, Waltham, MA, USA) with an excitation/emission filter set of 485/535 nm. Assays were performed in triplicate. $\mathrm{IC}_{50}$ values for compounds were determined by nonlinear regression with sigmoidal dose-response curves using Sigmaplot software.

### 2.5 Western blot analysis of the PHDs' oligomers

PHDs $(1 \mu \mathrm{~g})$ were incubated in the absence and presence of $80 \mu \mathrm{M} \mathrm{Zn}(\mathrm{II})$ and/or 1 mM TCEP at $30{ }^{\circ} \mathrm{C}$ for 10 minutes. For immunoblotting, reaction mixtures were prepared in the SDS sample buffer with or without 2-mercaptoethanol, boiled for 2 minutes, and run on 4-15\% linear gradient Tris- HCl ready gels (Bio-Rad). After electrophoresis, the protein was electroblotted onto poly(1,1-difluoroethylene) (PVDF) membranes (Millipore) by tankelectroblotting (Bio-Rad), and the membranes were blocked with $5 \%$ BSA for 1 hour. Then the membranes were incubated with anti-PHD2 (Cell signaling catalog \#: 3293S) or antiPHD3 (Abcam catalog \#: ab30782) as the primary antibody and anti-rabbit IgG as the secondary antibody. Blots were developed with the ECL Plus Western Blotting Detection System (Thermo Scientific).

Table S1 Primer list used in this study

| PHD3 Mutants | Primer name | Primer sequence ( $5^{\prime} \rightarrow 3^{\prime}$ ) |
| :---: | :---: | :---: |
| E37A | E37A F | GGA CAA CTT CCT GGG CGC CGT GGT GGG CGA CTG C |
|  | E37A R | GCA GTC GCC CAC CAC GGC GCC CAG GAA GTT GTC C |
| C42A | C42A F | CGA GGT GGT GGG CGA CGC CGT CCT GGA GCG CGT CA |
|  | C42A R | TGA CGC GCT CCA GGA CGG CGT CGC CCA CCA CCT CG |
| E45A | E45A F | GCG ACT GCG TCC TGG CCC GCG TCA AGC AGC TG |
|  | E45A R | CAG CTG CTT GAC GCG GGC CAG GAC GCA GTC GC |
| C52A | C52A F | GCG TCA AGC AGC TGC ACG CCA CCG GGG CCC TGC |
|  | C52A R | GCA GGG CCC CGG TGG CGT GCA GCT GCT TGA CGC |
| R57A | R57A F | CGG GGC CCT GGC GGA CGG CCA G |
|  | R57A R | CTG GCC GTC CGC CAG GGC CCC G |
| R71A | R71A F | CGC CGG CGT CTC CAA GGC ACA CCT GCG GGG CGA |
|  | R71A R | TCG CCC CGC AGG TGT GCC TTG GAG ACG CCG GCG |
| H72A | H72A F | CCG GCG TCT CCA AGC GAG CCC TGC GGG GCG ACC A |
|  | H72A R | TGG TCG CCC CGC AGG GCT CGC TTG GAG ACG CCG G |
| E85A | E85A F | GGG GGC AAC GCG GAG GGC TGC |
|  | E85A R | GCA GCC CTC CGC GTT GCC CCC |
| E86A | E86A F | GGC AAC GAG GCG GGC TGC GAG |
|  | E86A R | CTC GCA GCC CGC CTC GTT GCC |
| C88A | C88A F | GGC AAC GAG GAG GGC GCC GAG GCC ATC AGC TTC C |
|  | C88A R | GGA AGC TGA TGG CCT CGG CGC CCT CCT CGT TGC C |
| E89A | E89A F | ACG AGG AGG GCT GCG CCG CCA TCA GCT TCC TC |
|  | E89A R | GAG GAA GCT GAT GGC GGC GCA GCC CTC CTC GT |
| C124A | C124A F | AAG GCA ATG GTG GCT GCC TAT CCG GGA AAT GG |
|  | C124A R | CCA TTT CCC GGA TAG GCA GCC ACC ATT GCC TT |
| D183A | D183A F | GTG GAG CCC ATT TTT GCC AGA CTC CTG TTC TTC |
|  | D183A R | GAA GAA CAG GAG TCT GGC AAA AAT GGG CTC CAC |
| R71AH72A | R71AH72A F | CGC CGG CGT CTC CAA GGC AgC CCT GCG GGG CGA CCA |
|  | R71AH72A R | TGG TCG CCC CGC AGG GCT GCC TTG GAG ACG CCG GCG |
| E85AE86A | E85AE86A F | CGG GGG CAA CGC GGC GGG CTG CGA G |
|  | E85AE86A R | CTC GCA GCC CGC CGC GTT GCC CCC G |

