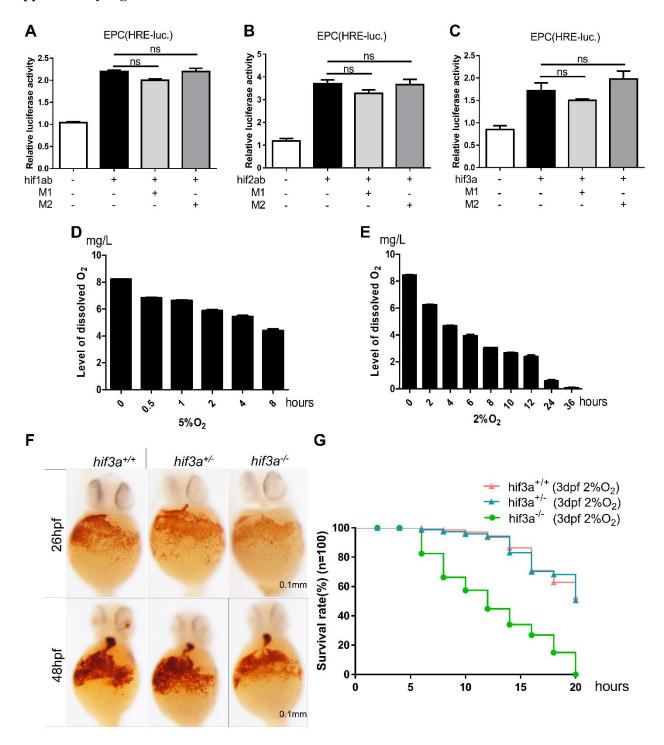
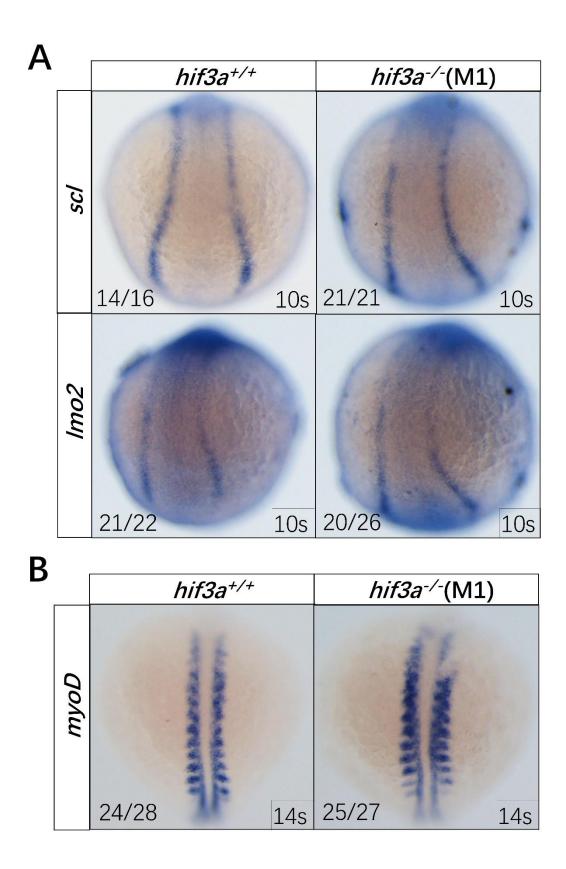


**Figure S1**. Generation of *hif-3a*<sup>-/-</sup> zebrafish using CRISPR/Cas9 technology. (A) Schematic of the targeting site in *hif-3a* and the resulting nucleotide sequence in mutant 1 (M1:

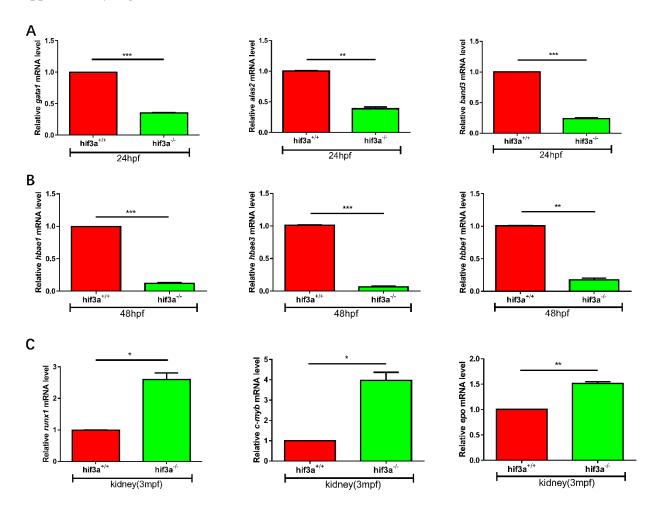
Hif- $3a^{ihb20180620/ihb20180620}$ ) and mutant 2 (M2: hif- $3a^{ihb20180621/ihb20180621}$ ). (B) The predicted protein products of hif-3a in the mutants and their wild-type siblings. (C) Verification of the efficiency of CRISPR/Cas9-mediated disruption of zebrafish hif-3a disruption by heteroduplex mobility assay (HMA). (D) The relative mRNA expression levels of hif-3a in the wild-type zebrafish and the homozygous mutants (10 embryos for each, 3 replicates; 36 hpf) were quantified by qRT-PCR. (E) hif-3a protein level in the wild-type and homozygous mutant embryos (200 embryos) under normoxic conditions detected by anti-hif-3a antibody. \*\*\* p < 0.001.



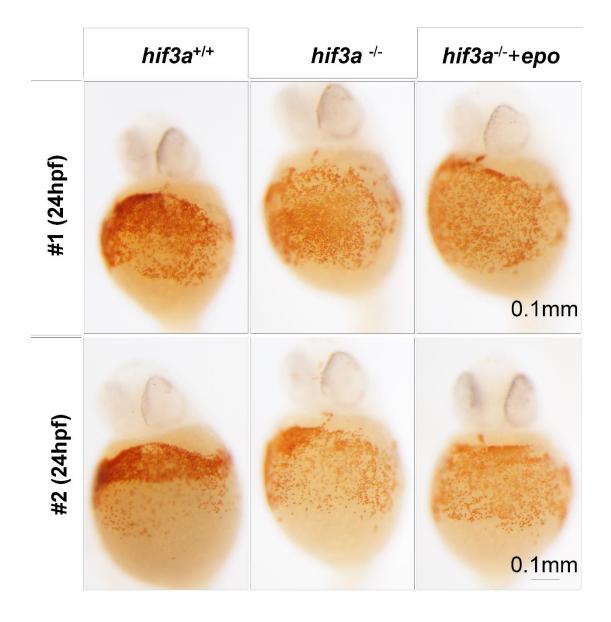
**Figure S2.** (A, B, C) The predicted truncated proteins in M1 and M2 mutants had no effect on the transcriptional activity of *hif-1ab*, *hif-2ab* and *hif-3a*. (D, E) The actual levels of dissolved O<sub>2</sub> in water were measured with an LDO101 probe at different time points when the flasks were put into Invivo2 Hypoxia workstation set at 5% O<sub>2</sub> and 2% O<sub>2</sub> respectively (3 replicates). (F) O-dianisidine staining of functional hemoglobin in the mature primitive erythrocytes in *hif-3a*<sup>+/+</sup> (left), *hif-3a*<sup>+/-</sup> (middle) and *hif-3a*<sup>-/-</sup> zebrafish (right) at 26 hpf and 48hpf. (G) The survival rate curve of *hif-3a*<sup>-/-</sup> zebrafish larvae, *hif-3a*<sup>+/-</sup> zebrafish larvae and their WT siblings (100 larvae). The oxygen concentration of the hypoxia workstation (Ruskinn INVIVO2 400) was adjusted to 2% prior to experimentation. The dead larvae were counted once every two hours. M1, mutant1; M2, mutant 2; hpf, hours post-fertilization; dpf, days post-fertilization.



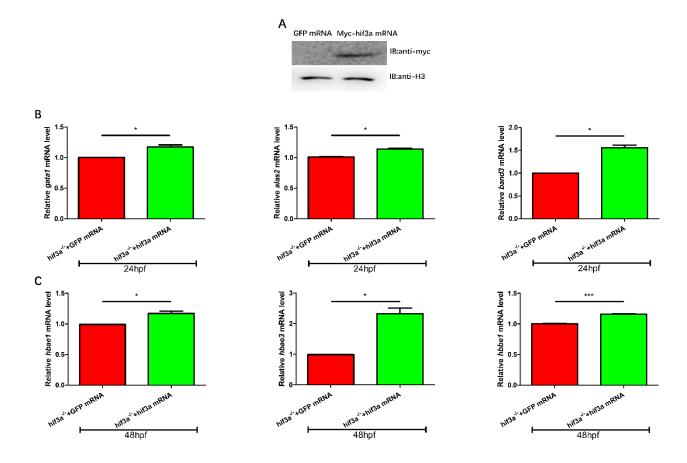
**Figure S3.** (A) Disruption of *hif-3a* did not alter the expression of *scl* and *lmo2* at the 10 s stage in the posterior lateral mesoderm. (B) Disruption of *hif-3a* did not alter the expression of *myoD* (the somatic mesoderm marker) at 14 s. The number of stained embryos was indicated in the left lower corner of each representative picture. M1, mutant 1; s, somite.



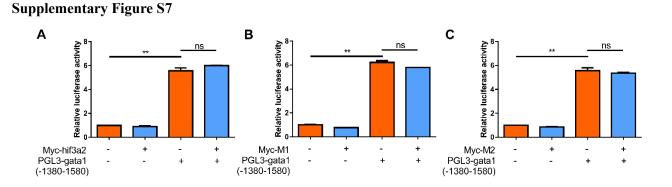
**Figure S4.** (A) qRT-PCR assays confirmed that the expression levels of gata1, alas2, and band3 were reduced in hif-3a-null larvae at 24 hpf (30 embryos for each, 3 replicates). (B) Quantitative RT-PCR assays confirmed that the expression levels of hbae1, hbae3 and hbbe1 were reduced in hif-3a-null larvae at 48 hpf (30 embryos for each, 3 replicates). (C) qRT-PCR assays showing that the expression levels of runx1, c-myb and epo were increased in hif-3a-null kidneys at 3 mpf (3 replicates). hpf, hours post-fertilization; mpf, months post-fertilization. \* p < 0.05; \*\* p < 0.01; \*\*\* p < 0.001.



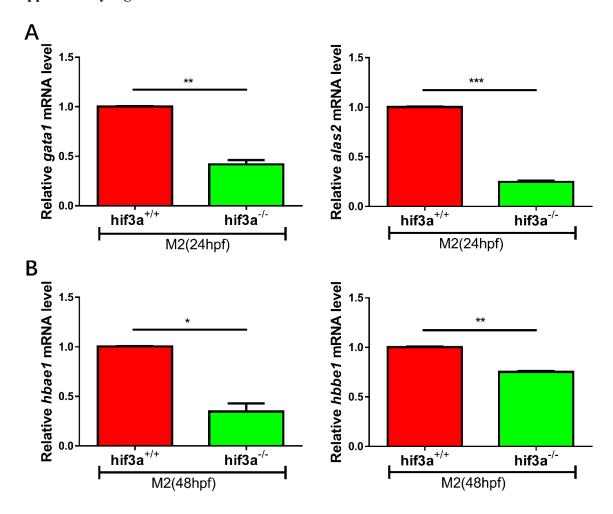
**Figure S5.** O-dianisidine staining indicated that co-injection of *epo* mRNA (500 pg/embryo) could not restore hemoglobin levels in hif- $3a^{-/-}$  larvae at 24 hpf. hpf, hours post-fertilization.



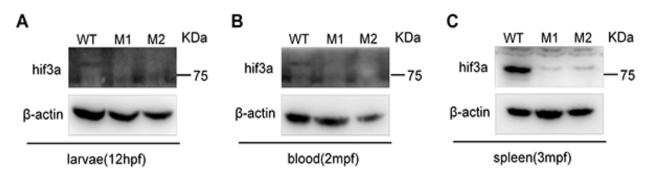
**Figure S6.** (A) Expression of myc-hif-3a in the injected embryos was confirmed by western blot assays (200 embryos). (B) qRT-PCR confirmed the restoration of gata1, alas2 and band3 by injection of hif-3a mRNA in hif-3a-null embryos at 24 hpf as compared with by the injection of the GFP mRNA control. (C) qRT-PCR confirmed the restoration of hbae1, hbea3 and hbbe1 by the injection of hif-3a mRNA in hif-3a-null embryos at 48 hpf as compared to the injection of the GFP mRNA control. IB, immunoblotting; hpf, hours post-fertilization. 30 embryos for each, 3 replicates; \* p < 0.05;\*\*\*\* p < 0.001.



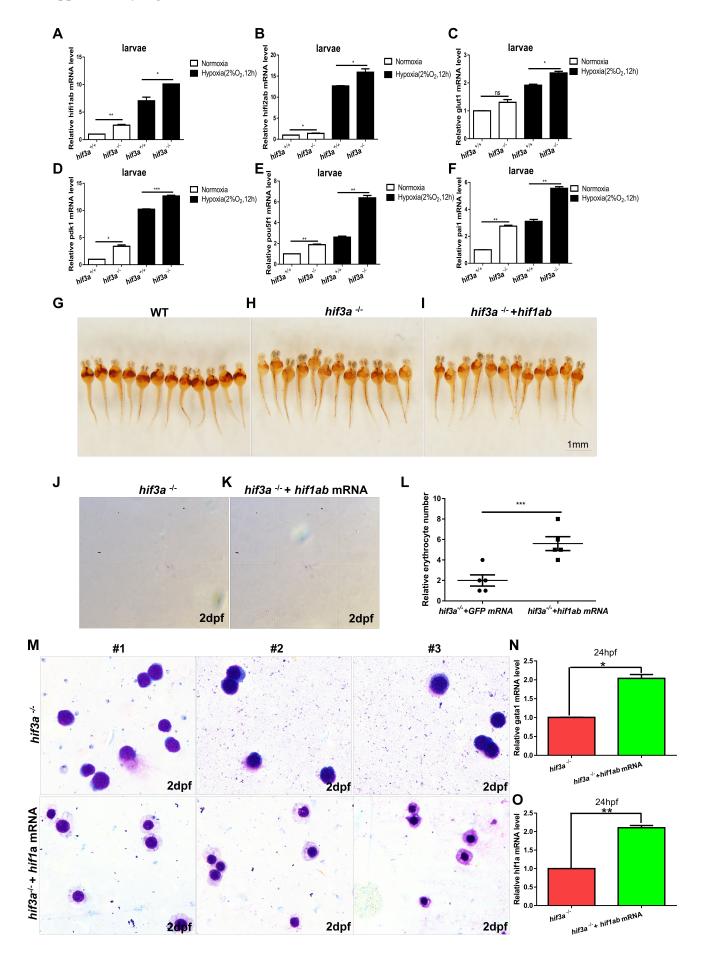
**Figure S7.** Luciferase reporter assays indicate that hif-3a2(A), hif-3a mutant M1(B) and M2(C) could not active gata1 promoter in EPC cells. \*\* p < 0.01; ns, no significance.



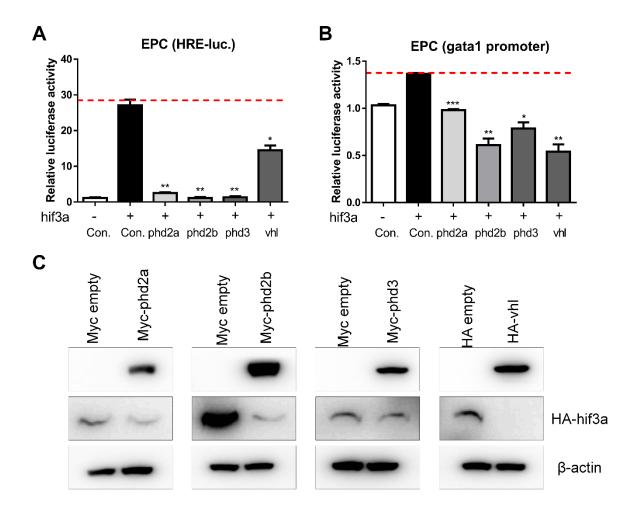
**Figure S8.** (A) Expression levels of erythrocytic markers *gata1* and *alas2* in wild-type and *hif3a*<sup>-/-</sup> (M2:  $hif1al^{ihb20180621/ihb20180621}$ ) zebrafish larvae at 24 hpf. (B) Expression levels of *hbae1* and *hbbe1* were quantified by qRT-PCR at 48hpf. Values graphed are the means of three independent experiments performed in triplicates; error bars indicate the standard error of the mean (S.E.M.). hpf, hours post-fertilization. n=30; \* p < 0.05; \*\* p < 0.01; \*\*\* p < 0.001.



**Figure S9.** (A, B, C) Western blot analysis of hif-3a protein level in larvae (12hpf; 200 embryos), blood (2mpf; 3 zebrafish for each, 3 replicates) and spleens (3mpf; 3 zebrafish for each; 3 replicates) from wild-type and M1, M2 mutants. WT, wildtype; M1, mutant 1; M2, mutant 2; mpf, months post-fertilization.



**Figure S10.** (A, B) qRT-PCR analysis of hifla(A) and hif2a(B) expression in WT(hif- $3a^{+/+}$ ) and hif-3a-null(hif- $3a^{-/-}$ ) zebrafish embryo (10 embryos for each, 3 replicates; 3dpf) under normoxia and hypoxia (2% O<sub>2</sub> for 12 hours). (C-F) Expression levels of the hif-1a down-stream targets glutl(C), pdkl(D) and hif-2a down-stream targets pou5fl(E), pail(F) were increased in hif- $3a^{-/-}$  larvae (10 embryos for each, 3 replicates; 3 dpf) under normoxia and hypoxia (2% O<sub>2</sub> for 12 hours). (G-I) O-dianisidine staining indicated that co-injection of hif-1ab mRNA (500 pg/embryo) partially restored hemoglobin levels in hif- $3a^{-/-}$  larvae at 2 dpf. (J-K) Erythrocyte number counting indicated that co-injection of hif-1ab mRNA (500 pg/embryo) increased erythrocyte in hif- $3a^{-/-}$  larvae at 2 dpf. (M) May-Grunwald-Giemsa staining indicated that co-injection of hif-1ab mRNA (500 pg/embryo) restored erythrocytic maturation in hif- $3a^{-/-}$  larvae at 2 dpf. (N) Co-injection of hif-1ab mRNA (500 pg/embryo) restored gata1 expression in hif- $3a^{-/-}$  embryos at 24 hpf. (O) Expression of micro-injected hif-1ab mRNA in hif- $3a^{-/-}$  embryos at 24 hpf was confirmed by qRT-PCR. Hpf, hours post fertilization; dpf, days post-fertilization. \*p < 0.05; \*\*p < 0.01; \*\*\*p < 0.001.



**Figure S11.** (A) Co-expression of *phd2a*, *phd2b*, *phd3* and *vhl* together with *hif-3a* suppressed the activity of HRE luciferase reporter induced by hif-3a in EPC cells. (B) Co-expression of *phd2a*, *phd2b*, *phd3* and *vhl* together with *hif-3a* suppressed the activity of gata1 promoter luciferase reporter induced by hif-3a in EPC cells. \*p < 0.05; \*\*p < 0.01; \*\*\*p < 0.001. (C) Western blot analysis indicated that hif-3a protein level was decreased when *phd2a*, *phd2b*, *phd3* or *vhl* were co-expressed in EPC cells. Con, control. Myc empty, pCMV-Myc empty vector.



**Movie 1.** Wildtype (left, WT) and *hif3-a* null (right, KO) zebrafish (3 mpf, body weight =  $0.32 \pm 0.02$  g) placed in a hypoxia workstation at the beginning (5% O<sub>2</sub>) (before 30 min).



**Movie 2.** Wildtype (left, WT) and *hif3-a* null (right, KO) zebrafish (3 mpf, body weight =  $0.32 \pm 0.02$  g) placed in a hypoxia workstation for a while (5%  $O_2$ ).