

DNA demethylase ALKBH1 promotes adipogenic differentiation via regulation of HIF-1 signaling

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Supplementary Information

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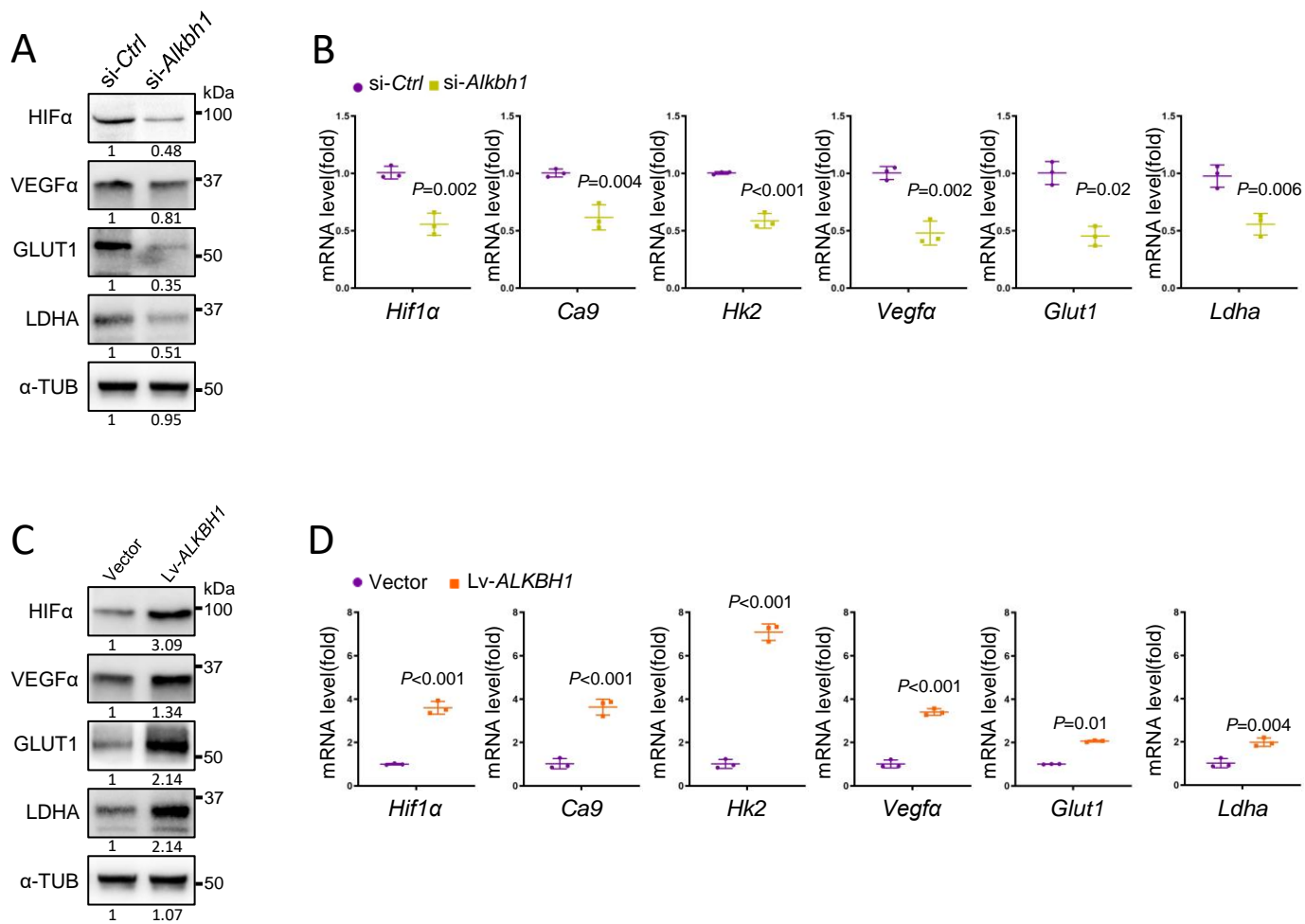


Figure S1. ALKBH1 activates HIF-1 pathway in 5% oxygen conditions. (A) Western blots of hypoxia-related proteins in *Alkbh1*-knocked 3T3-L1 cells in 5% oxygen conditions. **(B)** qRT-PCR analysis of *Hif-1α*, *Ca9*, *Hk2*, *Vegfa*, *Glut1* and *Ldha* in *Alkbh1*-deficient 3T3-L1 lines in 5% oxygen conditions (n=3). **(C)** Western blots and **(D)** qRT-PCR analysis confirm the activation of hypoxia axis in *ALKBH1*-overexpressed cells under 5% concentration of oxygen (n=3). The *P* values were calculated by two-tailed Student's *t* test. Scatter plots show individual data points \pm SD.

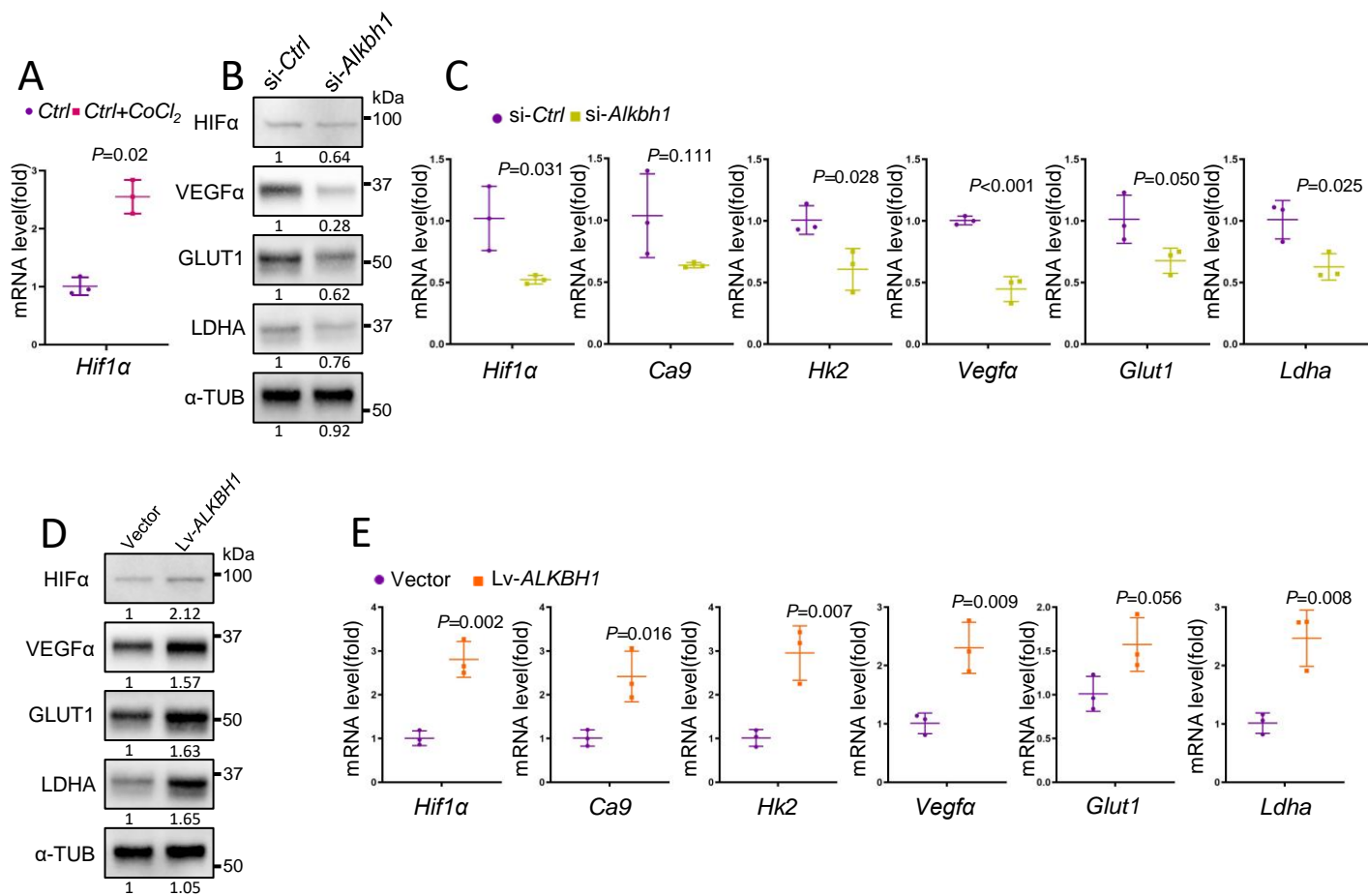


Figure S2. ALKBH1 activates HIF-1 pathway in hypoxia induced by CoCl₂. (A) qRT-PCR of *Hif-1 α* at 24-hour treatment of CoCl₂. CoCl₂ activates HIF-1 pathway in 3T3-L1 cells (n=3). (B) Western blots of hypoxia-related proteins in *Alkbh1*-knocked 3T3-L1 cells in hypoxia induced by CoCl₂. (C) qRT-PCR analysis of *Hif-1 α* , *Ca9*, *Hk2*, *Vegfa*, *Glut1* and *Ldha* in *Alkbh1*-deficient 3T3-L1 lines in hypoxia (n=3). (D) Western blots and (E) qRT-PCR analysis confirm the overactivation of hypoxia axis in *ALKBH1*-overexpressed cells (n=3). The *P* values were calculated by two-tailed Student's *t* test. Scatter plots show individual data points \pm SD.

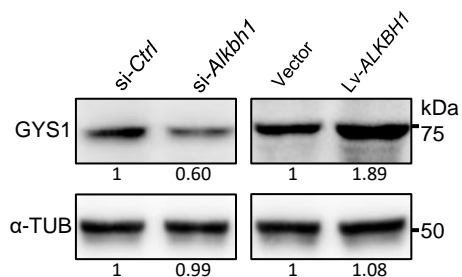


Figure S3. Positive relationship between ALKBH1 and GYS1 in hypoxia. Western blots confirms the relationship of ALKBH1 and GYS1 in 5% oxygen. GYS1 work as a target of ALKBH1.

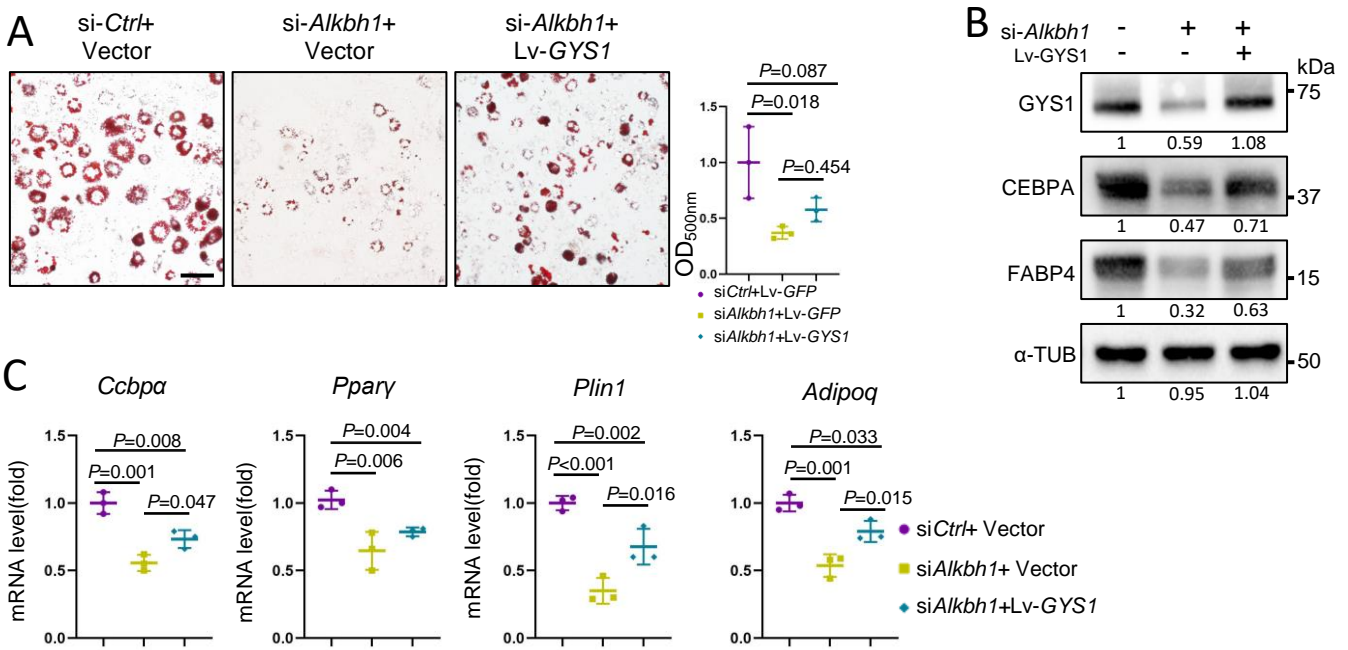


Figure S4. ALKBH1 demethylates DNA 6mA of GYS1. (A) Representative images of Oil red O staining in 3T3-L1 cells and quantitation analysis (n=3). Scale bar: 25µm. (B) Western blot analysis of HIF-1α, CEBPA and FABP4. Overexpression of GYS1 failed to fully rescue the decrease of adipogenic markers. (C) qRT-PCR shows the mRNA level of *Cebpa*, *Pparγ*, *Plin1* and *Adipoq* in si-Ctrl+Vector, si-*Alkbh1*+Vector and si-*Alkbh1*+Lv-GYS1 group at 5 days of differentiation (n=3). The P values were calculated one-way ANOVA with Tukey's post hoc test. Scatter plots show individual data points±SD.

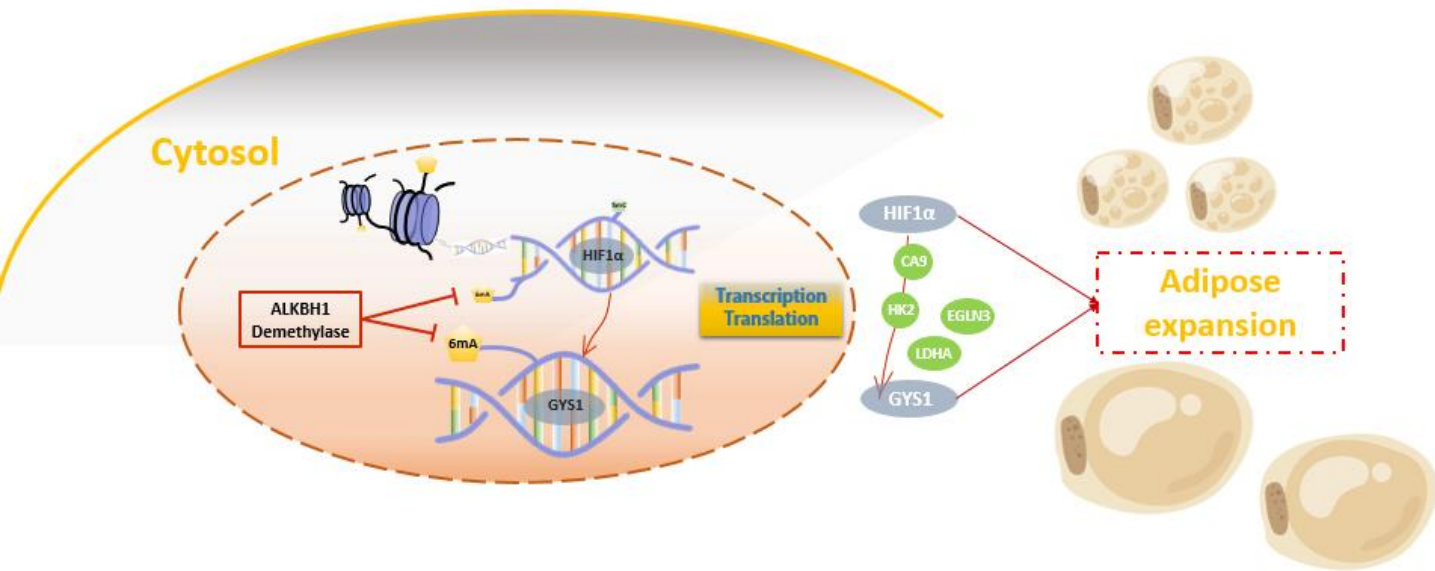


Figure S5. ALKBH1 enhances adipogenesis through activating HIF-1 axis. Intranuclear ALKBH1 demethylates the 6mA of *HIF1α* and *GYS1*, leading to the active transcription of these two genes. Numerous transcriptions result in the excessive accumulation of HIF1α and GYS1 proteins. The activation of HIF-1 pathway promotes the adipogenic differentiation of mesenchymal stem cells.

siRNA sequences	
siRNA-sense	GGUCUAAAGAAGUGACUAATT
siRNA-antisense	UUAGUCACUUCUUUAGACCTT

Table S1. mouse siRNA sequences used in this study

Figure 1B

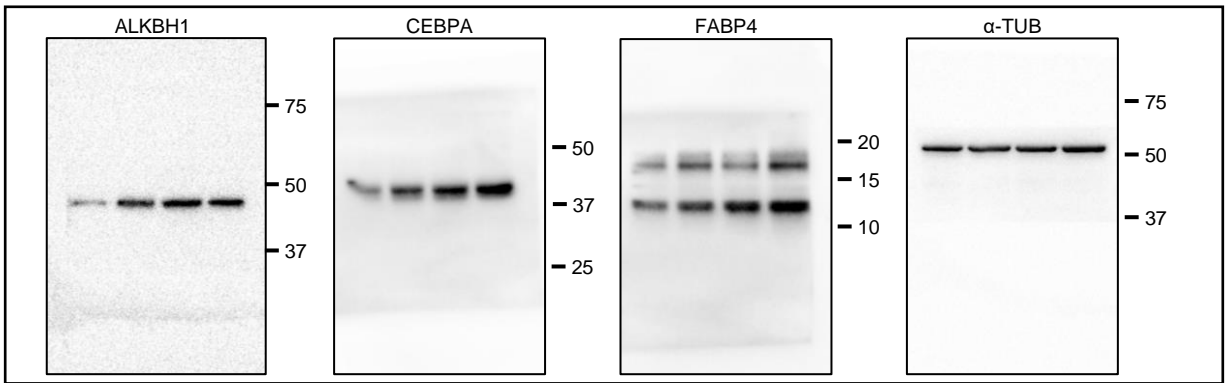


Figure 1D

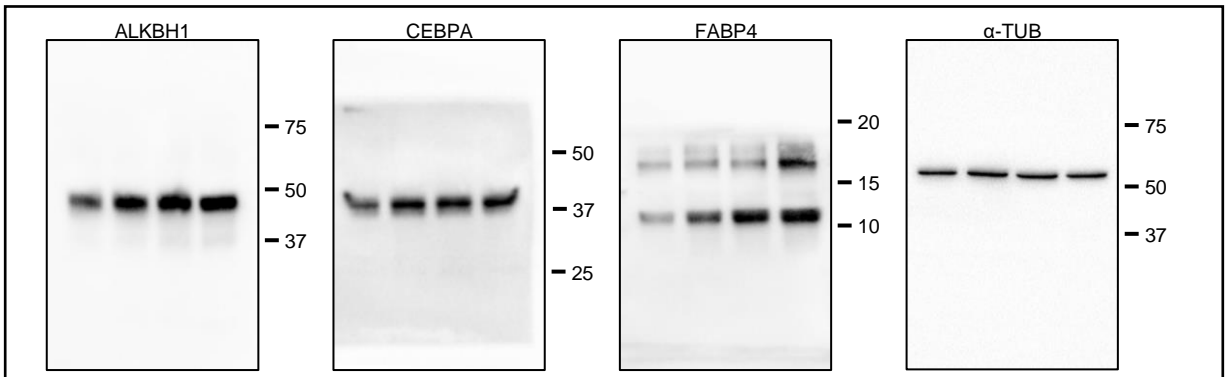


Figure 2A

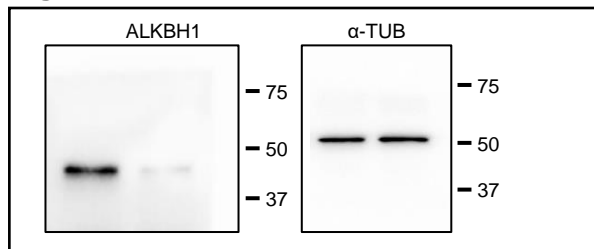


Figure 2E

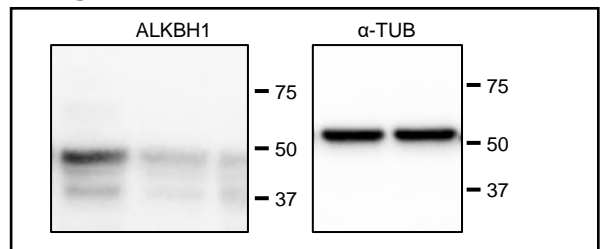


Figure 3A

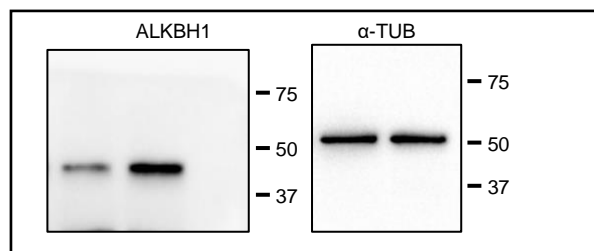


Figure 3D

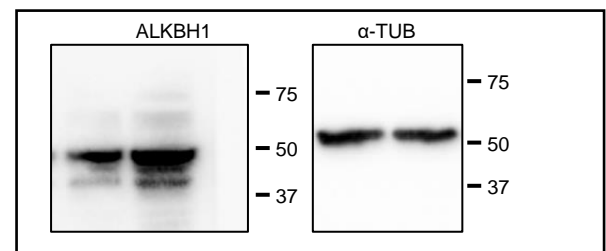


Figure 4E

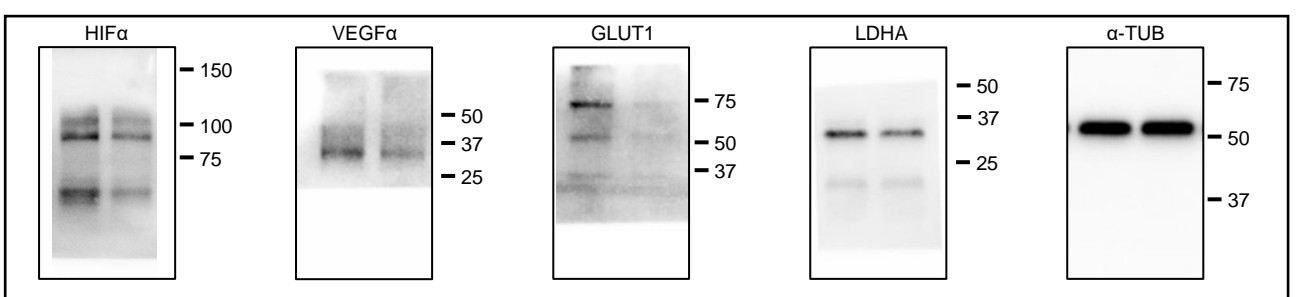


Figure 4G

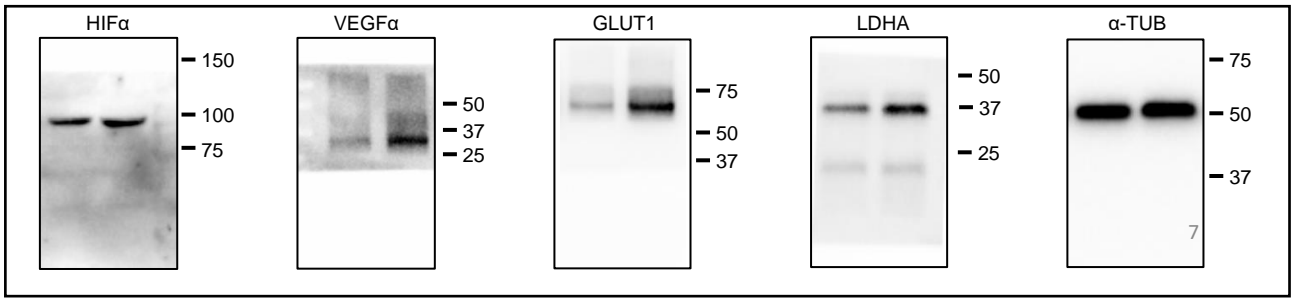


Figure 5A

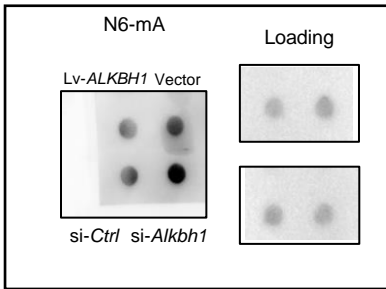


Figure 5E

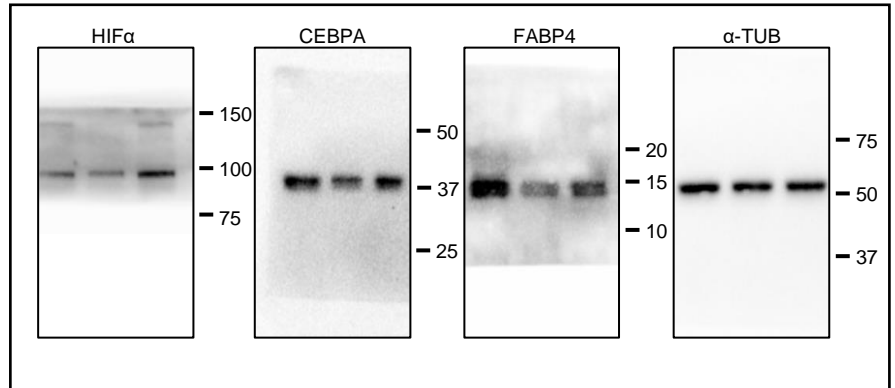


Figure 6C

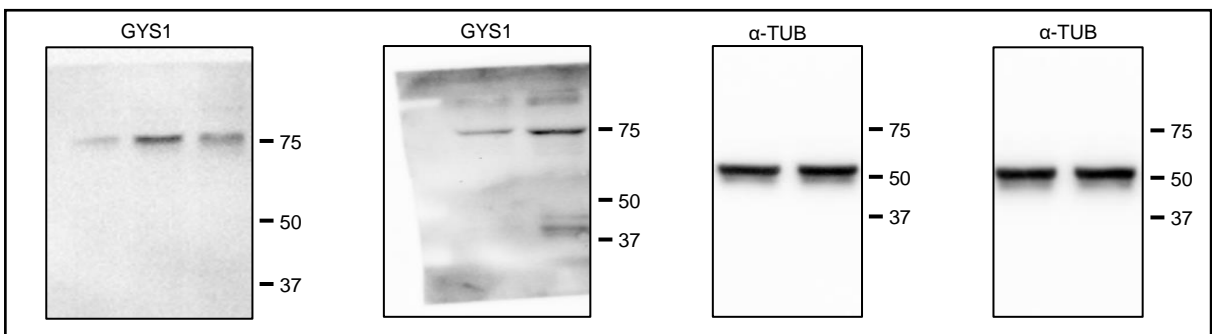


Figure 6E

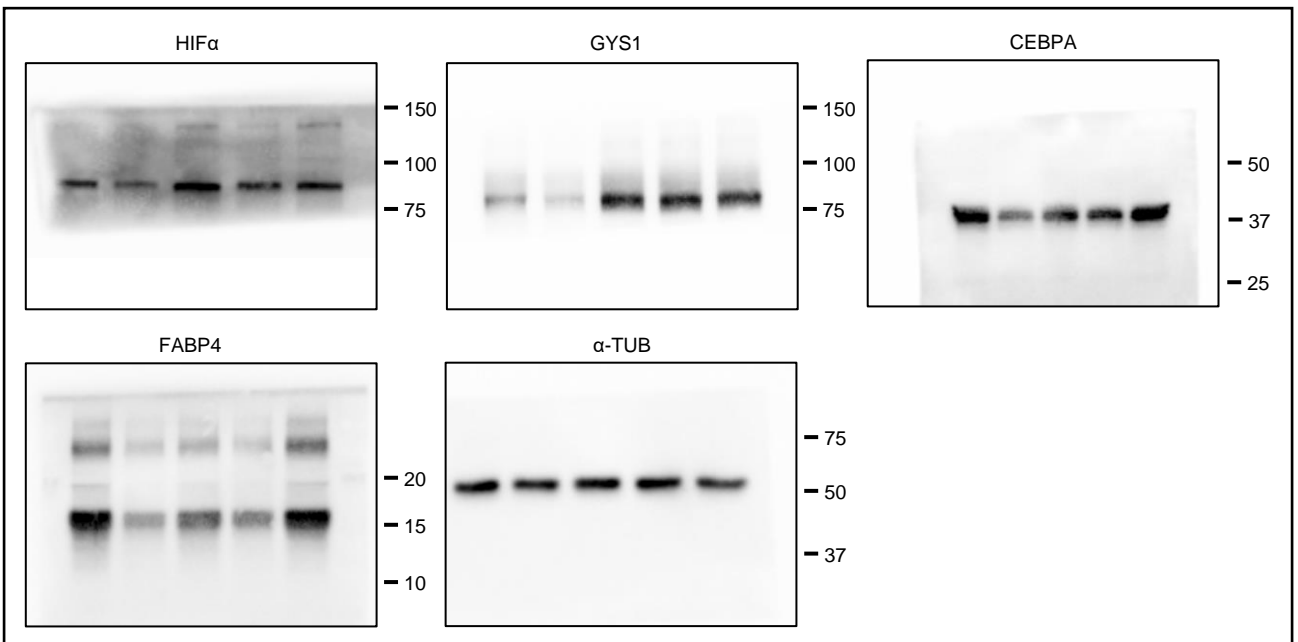


Figure S1A

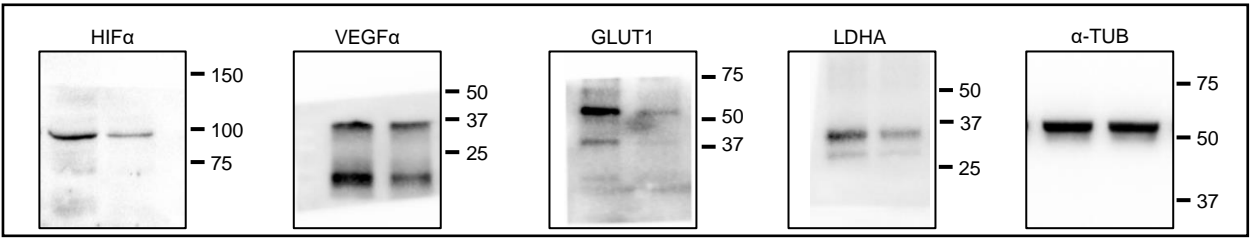


Figure S1C

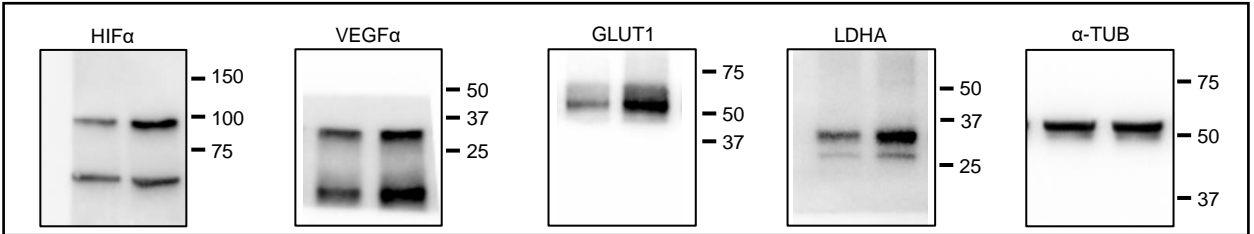


Figure S2B

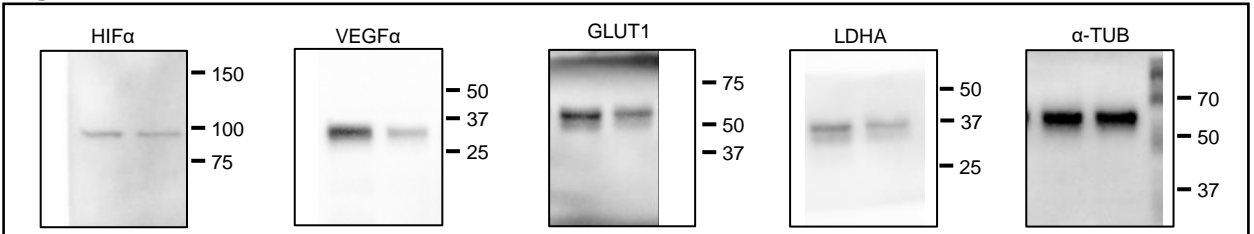


Figure S2D

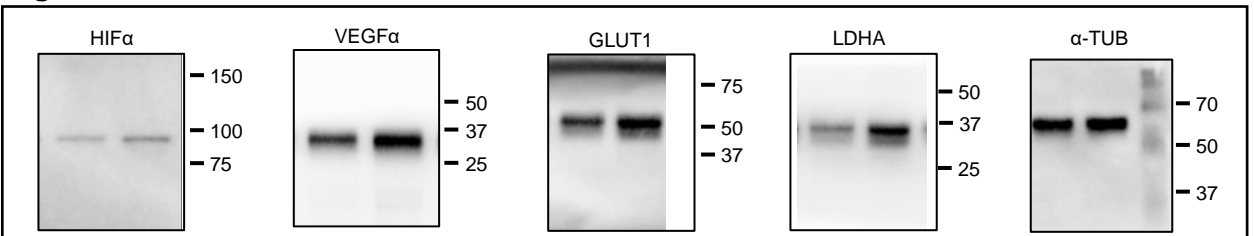


Figure S3A

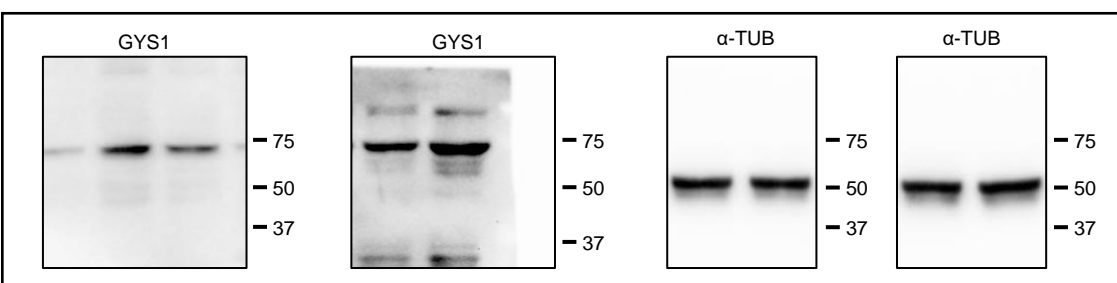


Figure S4B

