Supplementary Information 1 2 Proximity labeling expansion microscopy (PL-ExM) evaluates interactome labeling 3 4 techniques 5 6 Authors: Sohyeon Park¹, Xiaorong Wang², Yajin Mo¹, Sicheng Zhang³, Xiangpeng Li⁴, Katie C. Fong^{3,5}, Clinton Yu², Arthur A. Tran⁶, Lorenzo Scipioni^{7,8}, Zhipeng Dai⁴, Xiao 7 8 Huang⁹, Lan Huang², Xiaoyu Shi^{1,3,8,10,*} 9 10 Affiliations: 11 12¹ Center for Complex Biological Systems, University of California, Irvine; Irvine, 92697, 13 United States. 14² Physiology and Biophysics, University of California, Irvine; Irvine, 92697, United States. 15³ Department of Developmental and Cell Biology, University of California, Irvine; Irvine, 16 92697, United States. 17⁴ Department of Bioengineering and Therapeutic Sciences, University of California, San 18 Francisco; San Francisco, 94143, United States. 19 ⁵ Current Address: School of Criminal Justice and Criminalistics, California State 20 University, Los Angeles; Los Angeles, 90042, United States. 21 ⁶ Cardiovascular Research Institute, School of Medicine, University of California, San 22 Francisco; San Francisco, 94143, United States. ²³ ⁷ Laboratory for Fluorescence Dynamics, University of California, Irvine; Irvine, 92697, 24 United States. ²⁵ ⁸ Department of Biomedical Engineering, University of California, Irvine; Irvine, 92697, 26 United States. ²⁷ ⁹ School of Biomedical Engineering, Science and Health Systems, Drexel University; 28 Philadelphia, PA19104. ²⁹ ¹⁰ Department of Chemistry, University of California, Irvine; Irvine, 92697, United States. 30 31 *Corresponding author: Xiaoyu Shi. Email: xiaoyu.shi@uci.edu

32 Image resolution measurement

33 0.1µm size fluorescent beads (TetraSpeck Microspheres, Invitrogen; T7279) were used

- 34 to measure the resolution of the Airyscan LSM980 resolution with 63x water immersion
- 35 objective (NA1.15). 30 different beads were sampled to obtain the average full width half
- 36 maximum (FWHM) with standard error. The effective resolution of PL-ExM was measured
- 37 by calculating FWHM divided by the physical length expansion factor of the hydrogel.



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Figure S1. Resolution measurement. (A) A 0.1µm size fluorescent bead slide is used to measure the resolution of the Airyscan LSM980 with 63x water immersion objective (NA1.15), which is the system used for this paper. 30 different beads are sampled to obtain average full width half maximum (FWHM) value with standard error. Measured value is 139 ± 3 nm (mean \pm standard error). (B) The effective resolution of PL-ExM after different times of expansion. The measured effective resolution of PL-ExM in this paper was in the range of 14 - 70 nm depending on the length expansion factors and the type of microscope.

47 Chemical reactions in PL-ExM



49

- 50 Figure S2. Chemical reactions involved in the PL-ExM workflow, including
- 51 proximity labeling, protein anchoring, gelation, and fluorescent staining steps.

52 PL-ExM imaging using confocal microscopy



54 Figure S3. Confocal PL-ExM images of mitochondria in U2OS cells. Proteins in the

55 proximity of TOMM20 were labeled with antibodies conjugated with HRP (green). DNA 56 was stained with DAPI (blue). The images were taken on a laser scanning confocal

57 microscopy LSM780 with 63x water immersion objective (NA1.15). Length expansion

58 factor: 4.1. Scale bars: 5 μm (physical size post-expansion, 20.5 μm).

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60 APEX and HRP approaches of PL-ExM



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62 Figure S4. Airyscan PL-ExM images of mitochondria labeled with APEX2 or HRP as

63 the peroxidase. (A) Proteins in the proximity of APEX2 expressed with the outer

64 mitochondrial membrane were labeled (magenta). DNA was stained with DAPI (blue). (B)

65 Proteins in the proximity of TOMM20 were labeled with antibodies conjugated with HRP 66 (green). DNA was stained with DAPI (blue). The images were taken on an Airyscan LSM 67 980 with 63x water immersion objective (NA1.15). Length expansion factor: 4.0. Scale

 $\,68\,$ bars: 5 μm (physical size post-expansion, 20 μm).

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70 TurbolD approach of PL-ExM



Figure S5. Airyscan PL-ExM images of mitochondria labeled with TurbolD approach. Live cell PL was initiated for 30 min in cells stably expressing the TurbolD-OMM fusion construct. (A) Biotinylated proteins were visualized by post-gelation staining with streptavidin- Alexa Fluor 488 and (B) Immunostaining of TOMM20 localized mitochondria. DAPI is a nuclear marker. Length expansion factor: 4.4. Scale bars, 5 µm

in pre-expansion unit. All images in this figure were taken on an Airyscan microscope.

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79 Validation of APEX2-OMM cells shows negligible overexpression artifact

We are careful about the potential artifact that overexpressing APEX2-OMM could cause. As stated in the Methods, we made an APEX2-OMM plasmid with a moderate promoter SFFV rather than CMV to transfect U2OS cells. The strong CMV promoter can easily introduce overexpression. However, we didn't observe noticeable artifacts in our cell lines expressing APEX2-OMM with the SFFV promotor. In our SFFV APX2-OMM cell line, a FLAG- tag is fused on the N terminus of APEX2. We immunostained TOMM20 and FLAG-tag and performed APEX2catalyzed PL. The colocalization of the signals from TOMM20, FLAG-tag, and APEX PL (Figures S6A-C) confirms that the APX2 was expressed specifically in mitochondria. The stringy and tubular morphology of mitochondria in our APEX2-OMM cells (Figure S6A) is similar to the typical mitochondrial morphology in U2OS cells (Figures 2A&G). This similarity indicates negligible overexpression artifact. More structural details are visualized in the multicolor PL-ExM images of APEX PL and immunostained TOM20 (Figures S6F-I).



- 93 Figure S6. Validation of APEX2-OMM cells with Airyscan and PL-ExM. (A-E)
- 94 Fluorescence imaging of APEX2-mitochondria localization and biotinylation activity in a
- 95 cell line stably expressing APEX2-OMM with SFFV promotor. A FLAG-tag is fused on the
- 96 N terminus of APEX2 Live cell PL was performed for 1 min. Mitochondria (A) and FLAG-
- 97 APEX2 (B) expression were visualized with antibody staining, while biotinylation was 98 visualized with streptavidin-Alexa Fluor 488 (C). DAPI is a nuclear marker (D). Scale bars, 99 5 um.
- 100 (F-I) Multicolor PL-ExM images of APEX2-OMM cells. Live cell PL was performed for 1
- 101 min in activity in the APEX2-OMM, followed by expansion microscopy. Mitochondria were
- 102 marked with anti-TOMM20 antibodies (F), and biotinylated proteins were visualized with
- 103 streptavidin- Alexa Fluor 488 (G). DAPI is a nuclear marker (H). Length expansion factor:
- 104 4.4. Scale bars, 5 µm in pre-expansion unit.
- 105 All images in this figure were taken on an Airyscan microscope.

106 **3D PL-ExM imaging of brain tissues**



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108 **Figure S7. Orthogonal views of PL-ExM imaging of mouse brain section.** The images, 109 captured using Airyscan PL-ExM, depict a 20-µm mouse brain section expressing Thy1-YFP.

10 Proximity labeling was applied to Thy1-YFP (green) and GFAP was immunostained (magenta).

111 Antibodies permeated tissue thoroughly for proximity labeling to occur robustly along the z-

112 axis of the tissue. The tissue was expanded by a factor of 4.0. Scale bar: 20 µm in pre-expansion

113 units. All images in this figure were taken on an Airyscan microscope.

114 Negative Controls for HRP-, APEX2-, and TurbolD-catalyzed PL.



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- 116 Figure S8. Three approaches of PL-ExM with negative controls in cells, and tissues.
- 117 (A-B) HRP-catalyzed PL-ExM. PL was initiated for 30s. Biotinylation was visualized with
- 118 streptavidin- Alexa Fluor 488 (A), while no primary antibody was used in the negative
- 119 control (B). Length expansion factors: 4.0(A) and 4.4(B). Scale bars, 5 µm in pre-
- 120 expansion unit.

121 (C-D) APEX2-catalyzed PL-ExM. Live cell PL was initiated for 1 min, and biotinylation 122 was visualized with streptavidin- Alexa Fluor 488 (C), while wild-type U2OS cells without 123 APEX2 were used in the negative control (D). Length expansion factors: 4.3(C) and 124 4.4(D). Scale bars, 5 µm in pre-expansion unit.

125 (E-F) TurboID-catalyzed PL-ExM. Live cell PL was initiated for 30 min, and biotinylation 126 was visualized with streptavidin- Alexa Fluor 488 (E), while wild-type U2OS cells without 127 TurboID were used in the negative control (F). Length expansion factors: 4.4(E) and 128 4.4(F). Scale bars, 5 µm in pre-expansion unit.

129 (G-H) HRP-catalyzed PL-ExM in a 20 μ m tissue section of a mouse brain expressing 130 Thy1-YFP. The samples were immunostained with Rabbit anti-GFP antibodies, followed 131 by Donkey x Rabbit and HRP Horse x Goat antibodies. PL duration was 1 min. 132 Biotinylation was visualized with streptavidin- Alexa Fluor 488 (G), while no primary 133 antibody was used in the negative control (H). Length expansion factors: 4.0 (G) and 134 4.1(H). Scale bars, 2 mm in pre-expansion unit.

135 All images in this figure were taken on an Airyscan microscope.