# scientific reports



# **OPEN** Activation of L-lactate oxidase by the formation of enzyme assemblies through liquid-liquid phase separation

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The assembly state of enzymes is gaining interest as a mechanism for regulating the function of enzymes in living cells. One of the current topics in enzymology is the relationship between enzyme activity and the assembly state due to liquid-liquid phase separation. In this study, we demonstrated enzyme activation via the formation of enzyme assemblies using L-lactate oxidase (LOX). LOX formed hundreds of nanometer-scale assemblies with poly-L-lysine (PLL). In the presence of ammonium sulfate, the LOX-PLL clusters formed micrometer-scale liquid droplets. The enzyme activities of LOX in clusters and droplets were one order of magnitude higher than those in the dispersed state, owing to a decrease in  $K_{\rm M}$  and an increase in  $k_{\rm cat}$ . Moreover, the clusters exhibited a higher activation effect than the droplets. In addition, the conformation of LOX changed in the clusters, resulting in increased enzyme activation. Understanding enzyme activation and assembly states provides important information regarding enzyme function in living cells, in addition to biotechnology applications.

Enzymes are observed in various assembly states in living systems. For example, enzymes exist in aggregate<sup>1,2</sup>, fibril<sup>3,4</sup>, and condensate<sup>5,6</sup> states in response to stress. Hence, it is assumed that the function of enzymes is precisely controlled by their assembly states. However, there is no unified view of the driving force of these assemblies and their effect on enzyme activity<sup>7</sup>. This may be because historically, several in vitro enzymatic studies have been performed under conditions that do not promote enzyme assembly, such as using diluted and purified enzymes.

In the last decade, liquid–liquid phase separation (LLPS) is one of the subjects of debate in cell biology<sup>8,9</sup>. LLPS is a phenomenon in which a one-phase solution changes to a two-phase solution according to thermodynamic equilibrium<sup>8,9</sup>. Liquid droplets formed by LLPS can regulate enzyme activities<sup>10-12</sup>. Extensive studies on intercellular droplets reveals the importance of intrinsically disordered proteins (IDPs) or nucleic acids in droplet formation<sup>13-15</sup>. IDPs comprise intrinsically disordered regions (IDRs) that lack a fixed three-dimensional structure<sup>16</sup>. Some enzymes interact favorably with IDPs and nucleic acids, known as droplet scaffold molecules. For example, RubisCO, a well-known enzyme of more than 500 kDa for carbon dioxide fixation, functions by forming droplets with a small IDP<sup>17</sup>. Multienzyme assemblies that activate multistep reactions also exhibit liquid-like properties and require IDP domains<sup>18</sup>. Furthermore, several metabolic enzymes have RNA-binding ability<sup>19,20</sup>, which are prone to forming droplets. Numerous in vitro studies have revealed that enzymes such as ribozyme<sup>21</sup>, kinase<sup>12,22</sup>, multienzyme complexes<sup>23</sup>, RNA polymerase, and ribosomes<sup>24</sup> are activated in the formation of droplets. The droplets in these studies range in size from a few to tens of µm, making them easy to detect using an optical microscope. Moreover, several in vivo and in vitro studies report the presence of submicron-sized clusters as a precursor to micron-sized droplets<sup>25-27</sup>. Thus, the relationship between the assembly state and biological functions, such as enzymatic activation, is therefore of interest, however, this has not been demonstrated experimentally.

In this study, we investigated the relationship between assembly state and enzyme activity using L-lactate oxidase (LOX) as a model. Poly-L-lysine (PLL), which mimics the intrinsically disordered region of IDP<sup>28</sup>, was used as a scaffold molecule that electrostatically interacts with LOX. The addition of small amounts of salt (<100 mM) has been reported to promote droplet formation, presumably by modulating electrostatic interactions<sup>29</sup>. Furthermore, kosmotropic salts tend to promote droplet formation more efficiently<sup>29</sup>. When we investigated effective

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salts for LOX droplet formation, ammonium sulfate  $(NH_4)_2SO_4$ , a kosmotropic salt, caused LOX droplets to form most efficiently as expected (Supplementary Fig. S1). Thus, in this study, we employed  $(NH_4)_2SO_4$  salt to stabilize droplets. In the presence of PLL and absence of  $(NH_4)_2SO_4$ , LOX formed a soluble hundreds-nm-scale assembly, called "cluster<sub>lox</sub>" The addition of a small amount of  $(NH_4)_2SO_4$  formed several visible droplets in the scale of micrometers. The activities of both the assembly states were one order of magnitude higher than that of the enzyme in the dispersed state. The clusters<sub>lox</sub> and droplets have the common features of increasing  $k_{cat}$  and decreasing  $K_M$ , however, the activation effect was higher for clusters<sub>lox</sub> than for droplets.

# Results

LOX assembly states in the presence of PLL and ammonium sulfate. We investigated the droplet formation conditions of LOX by adding scaffold molecules and salts (Fig. 1a). LOX has an isoelectric point at approximately pH 6 (calculated); hence, it is negatively charged at the physiological pH. PLL was used as a scaffold molecule for the formation of droplets with LOX because it has an isoelectric point at approximately pH 10 (calculated), and is a disordered structure at physiological pH<sup>30</sup>. It has been shown that salts play an important role in the formation of droplets via electrostatic interactions using synthetic polymers<sup>29</sup>. We investigated the effects of (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> on the formation of LOX-PLL droplets because kosmotropic salts promote droplet formation<sup>30</sup>. Microscopic images 1 h after solution preparation showed no assembly in the presence of 0.1 µM LOX and 0.02 mM PLL, whereas droplets were observed in the presence of 0.1 µM LOX, 0.02 mM PLL, and 10 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> (Fig. 1a). We further investigated the effects of the salts on LOX-PLL droplets (Supplementary Fig. S1). In the presence of 25 mM sodium chloride (NaCl) and sodium thiocyanate (NaSCN), LOX-PLL did not form droplets. With increasing concentrations of NaCl and NaSCN, aggregate-like assemblies were observed above 100 mM. In contrast, in the presence of 25 mM sodium sulfate  $(Na_2SO_4)$  and  $(NH_4)_2SO_4$ , LOX-PLL droplets were observed (Supplementary Fig. S1). These data indicate that a small amount of sulfate ions stabilizes the LOX-PLL droplets via the kosmotropic effect, which is consistent with a previous study<sup>29</sup>. We further investigated the mixing ratios of LOX and PLL (Supplementary Fig. S2 and S3) in the presence of  $(NH_4)$ , SO<sub>4</sub>, indicating that 5  $\mu$ M LOX and 1 mM PLL formed well-shaped and stable droplets in the presence of 5-20 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>. Under these conditions, the droplets contained approximately 90% of the LOX in the solution (Supplementary Fig. S4).

Additionally, the presence of submicron-sized assemblies was investigated using dynamic light scattering (DLS) (Fig. 1b), because the size of these assemblies was below the detection limit of the optical microscope (~400 nm). In the absence of  $(NH_4)_2SO_4$ , 0.1  $\mu$ M LOX and 0.02 mM PLL formed two types of assemblies with diameters of approximately 50–60 and 200–300 nm, respectively, which could not be observed under the microscope (Fig. 1a,b). Here, we define these submicron-sized assemblies formed in the absence of  $(NH_4)_2SO_4$  as "clusters<sub>lox</sub>" (Fig. 1c). However, 0.1  $\mu$ M LOX and 0.02 mM PLL with 10 mM  $(NH_4)_2SO_4$  formed broad size of droplets with diameters of approximately 200–500 nm (Fig. 1a,b), which could be observed under the microscope. The larger peak of cluster<sub>lox</sub> overlapped with the peak of the droplet, suggesting that cluster<sub>lox</sub> formed a part of the droplet-like assembly.

**Features of LOX-PLL droplets.** We observed the LOX-PLL droplets using bright-field and fluorescence microscopy. Figure 2 shows microscopic images of the sample containing 5  $\mu$ M LOX, 1 mM PLL, and 6 mM ammonium sulfate at pH 8. PLL was monitored using chemically modified Rhodamine B isothiocyanate (RBITC) (red), and LOX was monitored using an intrinsic flavin mononucleotide (green). Bright-field microscopic images showed droplets with spherical structures and diameters of 10  $\mu$ m or less (Fig. 2a), indicating the



**Figure 1.** LOX assembly states in the presence of PLL and  $(NH_4)_2SO_4$ . (a) Microscopy images of 0.1  $\mu$ M LOX, 20 mM Tris–HCl (pH 8), 0 or 0.02 mM PLL (concentrations refer to lysine monomer units), 0 or 10 mM  $(NH_4)_2SO_4$ . Scale bar, 10  $\mu$ m (b) DLS data of sample solutions containing 0.1  $\mu$ M LOX, 0.02 mM PLL, 20 mM Tris–HCl (pH 8), with 0 or 10 mM  $(NH_4)_2SO_4$ . (c) Schematic image of assembly states of LOX with PLL and  $(NH_4)_2SO_4$ .





Figure 2. Features of LOX-PLL droplets. (a) Bright-field microscopic images of droplets (left) and fluorescent microscopic images of LOX (middle) and PLL-RBITC (right). The solution contained 5 µM LOX, 1 mM PLL, 6 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>, 20 mM Tris-HCl, and 20 mM MES (pH 8). Scale bar, 20 µm. (b) Localization of LOX and PLL-RBITC in the droplet. Fluorescence intensity along the dashed white lines was quantified from the brightness of each pixel. (c) Bright-field microscopic images of coalescing LOX-PLL droplets. The solution contained 5 µM LOX, 1 mM PLL, 6 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub> 20 mM Tris-HCl, and 20 mM MES (pH 8). Scale bar, 10 µm.

typical appearance of liquid droplets. Fluorescence microscopy revealed that the droplets contained both LOX and PLL molecules uniformly, as indicated by green and red fluorescence (Fig. 2b). Furthermore, the LOX-PLL droplets coalesced in approximately 10 s (Fig. 2c), indicating that the LOX-PLL droplets have liquid-like fluidity. This coalescence also seems to occur in small droplets formed with 0.1 µM LOX because the particle size increased in solution with time (Supplementary Fig. S5). Moreover, because clusters<sub>lox</sub> also tended to increase in particle size, it can be assumed that they have the same properties as droplets (Supplementary Fig. S5). To investigate the contribution of the electrostatic interaction between LOX and PLL in the formation of LOX-PLL droplets, we determined the effect of pH on the formation of LOX-PLL droplets (Supplementary Fig. S6). At pH 5, only amorphous aggregates were observed. At pH 6 and 7, both amorphous aggregates and liquid droplets were seen. At pH 8 and 9, however, only liquid droplets were observed. These results demonstrate that the electrostatic interaction between LOX and PLL plays a vital role in forming the liquid droplets because LOX and PLL have isoelectric points around pH 6 and 10, respectively.

LOX activation in clusters<sub>lox</sub> and droplets. We investigated changes in the assembly state and LOX activity depending on the concentration of ammonium sulfate. Microscopic images showed the formation of several micrometer-scale droplets above approximately 5 mM ammonium sulfate (Fig. 3a). Next, we investigated the presence of the LOX assembly using DLS measurements (Fig. 3b). Two peaks appeared in the absence of ammonium sulfate, as shown in Fig. 1c; the addition of  $1 \text{ mM} (\text{NH}_4)_3 \text{SO}_4$  changed these two peaks into one large peak. With an increase in the concentration of  $(NH_4)_2SO_4$ , this large peak shifted to the right and broadened (Fig. 3b). This result indicates that the size of the assembly increased with increasing ammonium sulfate concentration, which is consistent with the microscopic results (Fig. 3a). Under these conditions, we investigated how the enzyme activity of LOX changed depending on ammonium sulfate concentration (Fig. 3c). In the absence of PLL, the activity slightly increased depending on the ammonium sulfate concentration. In the presence of PLL and absence of ammonium sulfate, that is, in cluster<sub>lox</sub>, the enzyme activity of LOX was approximately 15 times higher (Fig. 3c). Although the activation rate decreased as the concentration of ammonium sulfate increased, it remained approximately five times more active in the presence of 10 mM ammonium sulfate (Fig. 3c). Furthermore, we confirmed that removing clusters<sub>lox</sub> and droplets from solutions by centrifugation impaired LOX activity (Supplementary Fig. S7). The findings revealed that clusters<sub>lox</sub> and droplets both increased LOX enzyme activity, with clusters<sub>lox</sub> being more favorable for activation than droplets. The activating effects of LOX in the droplets and clusterlox remained almost unchanged after 2 h and were retained at approximately 1 and sixfold for droplets and clusters<sub>lox</sub>, respectively, even after 24 h (Supplementary Fig. S8).

Kinetic analysis of LOX in clusters<sub>lox</sub> and droplets. To elucidate the detailed mechanism of enzyme activation by the formation of clusters<sub>lox</sub> and droplets, we determined the kinetic parameters of LOX in 20 mM Tris-HCl (pH 8.0) at 25 °C (Fig. 4 and Table 1). The K<sub>M</sub> of LOX in clusters<sub>lox</sub> was 0.05 mM, approximately 64-fold smaller than that of LOX alone, indicating that cluster<sub>lox</sub> is favorable for binding between LOX and its substrate (Fig. 4a). Additionally, the  $k_{cat}$  of LOX in clusters<sub>lox</sub> was approximately sixfold higher than that in the dispersed state, indicating that the turnover number of LOX increased in clusters<sub>lox</sub> (Fig. 4a). Next, the  $K_M$  of LOX-PLL in the droplet was 0.36 mM, approximately threefold smaller than that without PLL, indicating that the droplet is also favorable for binding between LOX and its substrate (Fig. 4b). The  $k_{cat}$  value of LOX in the droplets was approximately fourfold higher than that in the dispersed state (Fig. 4b). These results showed that the mecha-



**Figure 3.** Assembly state and activity of LOX depend on the concentration of ammonium sulfate. (a) Bright-field microscopic images of enzyme assemblies. The solution contained 0.1  $\mu$ M LOX, 0.02 mM PLL, 0–10 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>, and 20 mM Tris–HCl (pH 8). Scale bar, 20  $\mu$ m. (b) DLS data of sample solution containing 0.1  $\mu$ M LOX, 0.02 mM PLL, and 20 mM Tris–HCl (pH 8), with 0–10 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>. (c) Enzymatic activity of LOX in the presence of PLL and ammonium sulfate. Relative enzyme activity was defined as the initial reaction velocity in each condition divided by that in the absence of PLL and (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>.



**Figure 4.** Enzyme kinetics of LOX in clusters<sub>lox</sub>. (a) and droplets (b). The sample solution contained 0.1  $\mu$ M LOX, 0 or 0.02 mM PLL, 0–8 mM L-lactic acid, 20 mM Tris–HCl (pH 8), and 0 or 10 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>.

Assembly state	0.02 mM PLL	10 mM (NH <sub>4</sub> ) <sub>2</sub> SO <sub>4</sub>	$k_{\text{cat}}$ (s <sup>-1</sup> )	$K_{\rm M}$ (mM)	$k_{\rm cat}/K_{\rm M}$
Disperse	-	-	$3.58 \pm 0.34$	3.20	1.13
Disperse	-	+	$3.29 \pm 0.09$	0.97	3.40
Clusters <sub>lox</sub>	+	-	$20.92 \pm 0.75$	0.05	418.36
Droplets	+	+	$14.06 \pm 0.54$	0.36	39.06

Table 1. Michaelis-Menten parameters for LOX in different assembly states.

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nism of enzyme activation in clusters<sub>lox</sub> and droplets is common; the synergistic effect of an increase in substrate affinity and catalytic turnover; however, clusters<sub>lox</sub> are more effective than droplets for LOX activation.

**Secondary structure of LOX in clusters**<sub>lox</sub>. The increase in  $k_{cat}$  may have resulted from the conformational change in LOX within clusters<sub>lox</sub> and droplets. Thus, we investigated the secondary structure of LOX via far-ultraviolet (UV) circular dichroism (CD) spectroscopy. The far-UV CD spectra of LOX showed negative peaks at 208 and 218 nm, whereas the PLL spectra showed positive peaks between 210 and 230 nm (Fig. 5a). For samples forming clusters<sub>lox</sub>, the CD spectra of the mixture of LOX and PLL did not match those calculated from the individual spectra of LOX and PLL alone (Fig. 5b). These findings indicated that some structural changes in LOX and PLL were induced by their interactions.

PLL and LOX alone showed large far-UV CD ellipticity (Fig. 5a). Thus, we employed poly-(D,L)-lysine (PDLL), which also induces LOX droplets and activation (Supplementary Fig. S9) but is achiral in CD measurements to detect subtle changes in the secondary structure of LOX (Fig. 5c). As expected, PDLL showed almost no far-UV CD signal owing to its lack of optical activity (Fig. 5c). The spectrum of LOX drastically changed in the presence of 0.2 mM PDLL. Notably, the intensity of LOX at 210 nm was higher than that in the absence of PDLL. These results showed that a conformational change in LOX was induced in clusters<sub>lox</sub>, which may have caused an increase in  $k_{cat}$ . For samples containing droplets, the CD spectra could not be obtained stably within the measurement time, probably due to droplet coalescence and sedimentation (data not shown).

**Interaction between substrate and PLL.** Polymers with opposite charges to the substrate reduce  $K_{\rm M}^{31}$ . The substrate l-lactic acid is negatively charged, hence, positively charged PLL may contribute to the decrease in  $K_{\rm M}$ . Therefore, we investigated the interaction of PLL and L-lactate by isothermal titration calorimetry (ITC) and nuclear magnetic resonance (NMR). The titration of L-lactate into the PLL solution showed no significant changes during titration (Supplementary Fig. S10a), indicating that L-lactate did not bind to PLL. To further confirm the binding between L-lactate and PLL, <sup>1</sup>H-NMR spectrum of L-lactate was analyzed in the presence and absence of PLL (Supplementary Fig. S10b). L-lactate alone was assigned by three peaks ranging from 1.22–1.24, 1.34–1.38, and 1.48–1.50 ppm. The addition of PLL to the L-lactate does not interact with PLL. Thus, cluster<sub>lox</sub> and droplet play an essential role in increasing the affinity between the substrate and enzyme, rather than between the substrate and PLL.

#### **Discussion and conclusions**

In this study, we reported that the enzyme activity of LOX is increased by the formation of clusters<sub>lox</sub> and droplets induced by the addition of PLL. The mechanism of LOX activation may be derived from kinetic parameters (Fig. 4 and Table 1) and conformational properties (Fig. 5). The enzyme activity of LOX increased in both clusters<sub>lox</sub> and droplets, resulting from decreased  $K_{\rm M}$  and increased  $k_{\rm cat}$ . The  $k_{\rm cat}$  value of clusters<sub>lox</sub> was higher than that of the droplets. In addition, the degree of activation decreased as the droplet size increased (see Fig. 3). This suggests that the degree of activation may not be uniform throughout the droplet, such that the LOX activity is higher near the droplet surface. In contrast, the  $K_{\rm M}$  of the droplets decreased compared to the dispersed state, and it is likely that the compartmentalization effect of the substrate inside the droplets<sup>10</sup>. In addition to the compartmentalization effect, hundreds of nanometer-sized clusters<sub>lox</sub> are more dispersible in solution than several micrometer-sized droplets; therefore, they probably have a higher probability of collision with the substrate. This may be the reason the  $K_{\rm M}$  value of the clusters<sub>lox</sub> was smaller than that of the droplets.

The structural change of LOX in the clusters<sub>lox</sub> detected in CD is likely to be related to LOX activation, as LOX has been well investigated for its activity-structure relationship<sup>32–35</sup>. The LOX structure contains a core TIM barrel



**Figure 5.** Far-UV CD spectra of LOX and PLL in clusters<sub>lox</sub>. (a) CD spectra of 1  $\mu$ M LOX (solid line) and 0.2 mM PLL (broken line). (b) CD spectrum of LOX and PLL mixture (solid line), and CD spectrum calculated from that of LOX and PLL in A (broken line). (c) CD spectra of PDLL, LOX and LOX with PDLL.

of eight  $\alpha$ -helices and eight  $\beta$ -strands, binds FMN at the C-terminus of the  $\beta$ -barrel, and, importantly, has a very flexible loop and a short helix covering the active site<sup>32</sup>. Changes in this flexible loop structure and dynamics alter LOX activity by controlling substrate uptake and product release as shown in previous studies using mutations to this flexible loop<sup>33-35</sup>. The active site is open when this flexible loop adopts a disordered structure, and the active site is closed when it adopts a folded structure<sup>33-35</sup>. Specifically, in the crystal structure,  $\alpha$ -helix decreases when active site is open compared to when active site is closed<sup>33-35</sup>. Thus, the decrease in  $\alpha$ -helical content upon clusters<sub>lox</sub> formation may reflect structural changes and/or dynamics in the flexible loop covering the active site (Fig. 5). As no method has been established to investigate the conformational changes of folded enzymes in the enzyme assemblies, investigation of detailed conformational changes of LOX will be a future challenge.

Changes in enzyme structure and/or dynamics that accompany cluster<sub>lox</sub> and droplet formation can be attributed to the following possibilities. First, enzyme interaction with the polymer in cluster<sub>lox</sub> and droplets may change the structural stability of the native enzyme, leading to an increased  $k_{cat}$ <sup>36,37</sup>. Second, the enzyme assemblies like droplets are highly crowded with macromolecules, resulting in the exclusion of water molecules<sup>38,39</sup>. This crowding environment may stabilize non-native structures that differ from those in dilute conditions<sup>40</sup>. Therefore, crowding may promote the transition state of the enzyme, leading to enhanced enzyme activity. Finally, droplets represent a nonpolar environment compared to a dilute solution<sup>39,41</sup>. A nonpolar solution increases the stability of hydrophobic interaction<sup>38,39,42</sup>, hydrogen bonds, and electrostatic interactions<sup>43</sup>, which change enzyme activity. Further structural analysis of the enzyme in clusters<sub>lox</sub> and droplets will be an interesting subject for future research.

The activation of LOX within the enzyme assemblies can provide valuable information for industrial applications. Currently, the primary methods of enzyme activation are protein engineering<sup>44</sup> and directed evolution<sup>45</sup>. These methods generate favorable mutants with high activity through repeated mutagenesis and screening. However, it is time-consuming and costly to produce desirable mutants. In contrast, it is very simple to form an enzyme assembly using a polyelectrolyte to improve activity. LOX is an oxidoreductase applied to biofuel cells<sup>46,47</sup> and biosensors<sup>48</sup>. Thus, the formation of enzyme assemblies represents a versatile approach for improving enzyme activity in practical applications using LOX or other enzymes.

# Methods

**Materials.** Poly-L-lysine hydrobromide (MW, 70,000–150,000 Da), poly-(D,L)-lysine hydrobromide (MW, 25,000–40,000 Da), and 2,6-dichloroindophenol sodium salt hydrate (DCIP) were obtained from Sigma-Aldrich Co. (St Louis, MO, USA). NaCl,  $(NH_4)_2SO_4$ , Na<sub>2</sub>SO<sub>4</sub>, NaSCN, and dimethyl sulfoxide (DMSO) were obtained from Kanto Chemical Co., Inc. (Tokyo, Japan). Tris(hydroxymethyl)aminomethane was obtained from Nacalai Tesque (Kyoto, Japan). 2-Morpholinoethanesulfonic acid monohydrate and 3-[4-(2-Hydroxyethyl)-1-piperazi-nyl] propanesulfonic acid (EPPS) were obtained from Dojindo Laboratories (Kumamoto, Japan). l-lactic acid was obtained from Tokyo Chemical Industry Co., Ltd. (Tokyo, Japan). Rhodamine B isothiocyanate (RBITC) was obtained from Santa Cruz Biotechnology (Dallas, TX). LOX was prepared as previously described<sup>46</sup>.

**Fluorescent labeling of PLL.** PLL labeled with the amine-reactive dye RBITC (excitation/emission:555/580 nm) was prepared following the manufacturer's instructions. Briefly, a solution of RBITC (1.77 mM) in DMSO (50  $\mu$ L) was quickly added to a stirred solution of 20 mM PLL and 20 mM EPPS (950  $\mu$ L; pH 8.5) at 25 °C. After the reaction mixture was gently stirred for 1.5 h, 200 mM Tris–HCl (100  $\mu$ L, pH 8.5) was added. The PLL-dye conjugates were purified by filtration through Amicon Ultra-0.5 mL centrifugal filters with a molecular weight cutoff (MWCO) of 3 kDa (Millipore Sigma). The final concentration of PLL was determined using bicinchoninic acid (BCA) assay. The number of RBITC molecules conjugated to PLL in 10 mM Tris–HCl (pH 8.0) was determined from the absorbance at 556 nm, using the molar absorption coefficient  $\varepsilon_{556}$ =87,000 M<sup>-1</sup> cm<sup>-1</sup>. The number of dye molecules per PLL molecule was 0.2.

**Enzyme assays.** Enzyme solutions containing 0.1  $\mu$ M LOX, 0–0.2 mM PLL, and 0–10 mM ammonium sulfate in 20 mM Tris–HCl (pH 8) were prepared and left standing for 20 min. A 90  $\mu$ L aliquot of enzyme solution was mixed with a 10  $\mu$ L aliquot of substrate solution containing 0–80 mM L-lactic acid and 1 mM DCIP solution. The initial reaction velocities ( $v_0$ ) were determined from the slope of the initial decrease in absorbance at 555 nm using an Ultrospec 2100 pro UV/visible spectrophotometer (Amersham Biosciences Corp, Amersham, UK). Relative enzyme activity was defined as the initial reaction velocity under each condition divided by that in the absence of PLL and (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>. The  $K_M$  and  $k_{cat}$  values were determined by the initial reaction velocity on a theoretical Michaelis–Menten curve by nonlinear regression.

**Dynamic light scattering.** Dynamic light scattering (DLS) experiments were performed using a Zetasizer Nano ZS light scattering photometer (Malvern Instruments, Worcestershire, UK) equipped with a 4 mW He–Ne ion laser ( $\lambda = 633$  nm). To determine the size of LOX assemblies, solutions containing 20 nM LOX, 0–1 mM PLL, 0–10 mM (NH<sub>4</sub>)<sub>2</sub>SO<sub>4</sub>, and 20 mM Tris–HCl were placed in a 1-cm path length disposable cuvette, and DLS measurements were performed at 25 °C at a detection angle of 173°. The viscosities of the solutions were approximated using water ( $\eta = 0.87$  cP). All measurements were performed in 15 min after the solution preparation to prevent a decrease in scattering intensity by the sedimentation of enzyme assemblies. All the results are presented as the mean values of three independent experiments.

**Optical microscopy.** Images were recorded using an all-in-one fluorescence microscope BZ-X710 (KEY-ENCE, Osaka, Japan) 1 h after sample preparations. Aliquots ( $100 \mu$ L) of the samples were placed in an ultra-low-attachment 96-well plate (Corning, NY, USA). All images were prepared using BZ-X Analyzer (KEYENCE).

**Isothermal titration calorimetry.** Isothermal Titration Calorimetry (ITC) was performed using a Microcal Auto-iTC200 calorimeter (Malvern Instruments). The experiments consisted of a series of 0.2  $\mu$ L injections of 4 mM PLL into 200  $\mu$ L of 200  $\mu$ M LOX solution or 1 mM l-lactic acid in the thermostatic cell with an initial delay of 60 s, a 0.4 s duration of injection, and a spacing of 120 s between injections. In all cases, the samples were dialyzed in the same buffer containing 20 mM Tris–HCl and 20 mM MES (pH 8) to minimize interference from mixing and dilution heat signals.

**Hydrogen-1 nuclear magnetic resonance spectroscopic analysis.** Hydrogen-1 (<sup>1</sup>H) nuclear magnetic resonance (NMR) spectra were recorded in 20 mM Tris–HCl buffer at pH 8. The experiments were performed using a Bruker BioSpin Avance III 700 MHz NMR spectrometer at 25 °C.

**Circular dichroism.** Circular dichroism (CD) experiments were performed in a 1-cm path-length quartz cuvette using a spectropolarimeter (J-720 W; JASCO Co., Ltd). For clusters<sub>lox</sub> measurements, the enzyme solution containing 1  $\mu$ M LOX and 20 mM Tris–HCl buffer (pH 8.0) was incubated with 0.2 mM PLL or 0.2 mM Poly-(D,L)-lysine at 25 °C for 20 min before measurement. The CD spectra of the samples were corrected by subtracting the corresponding spectra of buffers.

**Precipitation rate of LOX.** The formation of liquid droplets of LOX with PLL was investigated based on the precipitation rate of the LOX. The concentration dependence of the PLL on liquid droplet formation was measured as follows: The PLL stock solution was prepared using 0–2 mM PLL in 20 mM Tris–HCl (pH 8). Aliquots (100  $\mu$ L) of various solutions containing 1 mM LOX in 20 mM Tris–HCl (pH 8) were mixed with 100  $\mu$ L of the PLL stock solution. Furthermore, the samples were centrifuged at 18,000×g for 20 min at 25 °C. The concentration of LOX in the supernatant was determined from the absorbance at 280 nm using an ND-1000 spectrophotometer (NanoDrop Technologies, Wilmington, DE, USA). Precipitation rates were calculated as follows:

Precipitation rate(%) =  $[1 - C_n/C_0] \times 100(\%)$ 

where  $C_0$  (mM) is the concentration in the supernatant without PLL and  $C_n$  (mM) is the LOX concentration in the supernatant with PLL.

#### Data availability

The data supporting the findings of this study are available in the paper and Supplementary file. All other data are available from the corresponding authors upon request.

Received: 22 November 2022; Accepted: 11 January 2023 Published online: 25 January 2023

#### References

- 1. Saad, S. *et al.* Reversible protein aggregation is a protective mechanism to ensure cell cycle restart after stress. *Nat. Cell Biol.* **19**, 1202–1213 (2017).
- Narayanaswamy, R. et al. Widespread reorganization of metabolic enzymes into reversible assemblies upon nutrient starvation. Proc. Natl. Acad. Sci. U. S. A. 106, 10147–10152 (2009).
- Cereghetti, G. *et al.* Reversible amyloids of pyruvate kinase couple cell metabolism and stress granule disassembly. *Nat. Cell Biol.* 23, 1085–1094 (2021).
- Simonet, J. C., Burrell, A. L., Kollman, J. M. & Peterson, J. R. Freedom of assembly: Metabolic enzymes come together. *Mol. Biol. Cell* 31, 1201–1205 (2020).
- Miura, N. et al. Spatial reorganization of Saccharomyces cerevisiae enolase to alter carbon metabolism under hypoxia. Eukaryot. Cell 12, 1106–1119 (2013).
- An, S., Kumar, R., Sheets, E. D. & Benkovic, S. J. Reversible compartmentalization of de novo purine biosynthetic complexes in living cells. Science 320, 103–106 (2008).
- Schmitt, D. L. & An, S. Spatial organization of metabolic enzyme complexes in cells. *Biochemistry* 56, 3184–3196. https://doi.org/ 10.1021/acs.biochem.7b00249 (2017).
- 8. Shin, Y. & Brangwynne, C. P. Liquid phase condensation in cell physiology and disease. Science 357, 1253 (2017).
- Banani, S. F., Lee, H. O., Hyman, A. A. & Rosen, M. K. Biomolecular condensates: organizers of cellular biochemistry. Nat. Rev. Mol. Cell Biol. 18, 285–298 (2017).
- 10. Peeples, W. & Rosen, M. K. Mechanistic dissection of increased enzymatic rate in a phase-separated compartment. *Nat. Chem. Biol.* 17, 693–702 (2021).
- 11. Nakashima, K. K., Vibhute, M. A. & Spruijt, E. Biomolecular chemistry in liquid phase separated compartments. *Front. Mol. Biosci.* 6, 21 (2019).
- 12. Ura, T., Tomita, S. & Shiraki, K. Dynamic behavior of liquid droplets with enzyme compartmentalization triggered by sequential glycolytic enzyme reactions. *Chem. Commun. (Camb)* 57, 12544–12547 (2021).
- Oldfield, C. J. & Dunker, A. K. Intrinsically disordered proteins and intrinsically disordered protein regions. Annu. Rev. Biochem. 83, 553–584 (2014).
- Shapiro, D. M., Ney, M., Eghtesadi, S. A. & Chilkoti, A. Protein phase separation arising from intrinsic disorder: first-principles to bespoke applications. J. Phys. Chem. B. 25, 6740–6759 (2021).
- Martin, E. W. & Holehouse, A. S. Intrinsically disordered protein regions and phase separation: Sequence determinants of assembly or lack thereof. *Emerg. Top Life Sci.* 4, 307–329 (2020).
- Uversky, V. N. Intrinsically disordered proteins in overcrowded milieu: Membrane-less organelles, phase separation, and intrinsic disorder. Curr. Opin. Struct. Biol. 44, 18–30 (2017).
- 17. Wang, H. *et al.* RuBisCO condensate formation by CcmM in  $\beta$ -carboxysome biogenesis. *Nature* 566, 131–135 (2019).
- 18. Jin, M. et al. Glycolytic enzymes coalesce in G bodies under hypoxic stress. Cell Rep. 20, 895–908 (2017).
- Castello, A., Hentze, M. W. & Preiss, T. Metabolic enzymes enjoying new partnerships as RNA-binding proteins. Trends Endocrinol. Metab. 26, 746–757 (2015).

- Beckmann, B. M. et al. The RNA-binding proteomes from yeast to man harbour conserved enigmRBPs. Nat. Commun. 6, 10127 (2015).
- Strulson, C. A., Molden, R. C., Keating, C. D. & Bevilacqua, P. C. RNA catalysis through compartmentalization. Nat. Chem. 4, 941–946 (2012).
- Koga, S., Williams, D. S., Perriman, A. W. & Mann, S. Peptide-nucleotide microdroplets as a step towards a membrane-free protocell model. *Nat. Chem.* 3, 720–724 (2011).
- 23. Crosby, J. et al. Stabilization and enhanced reactivity of actinorhodin polyketide synthase minimal complex in polymer-nucleotide coacervate droplets. *Chem. Commun. (Camb)* 48, 11832–11834 (2012).
- Sokolova, E. et al. Enhanced transcription rates in membrane-free protocells formed by coacervation of cell lysate. Proc. Natl Acad. Sci. U. S. A. 110, 11692–11697 (2013).
- Forman-Kay, J. D., Ditlev, J. A., Nosella, M. L. & Lee, H. O. What are the distinguishing features and size requirements of biomolecular condensates and their implications for RNA-containing condensates?. RNA 28, 36–47 (2022).
- Kar, M. et al. Phase-separating RNA-binding proteins form heterogeneous distributions of clusters in subsaturated solutions. Proc. Natl. Acad. Sci. U. S. A. 119, e2202222119 (2022).
- Penga, S. et al. Phase separation at the nanoscale quantified by dcFCCS. Proc. Natl. Acad. Sci. (PNAS) 117(44), 27124–27131. https://doi.org/10.1073/pnas.2008447117 (2020).
- 28. Fisher, R. S. & Elbaum-Garfinkle, S. Tunable multiphase dynamics of arginine and lysine liquid condensates. *Nat. Commun.* 11, 4628 (2020).
- Perry, S. L., Li, Y., Priftis, D., Leon, L. & Tirrell, M. The effect of salt on the complex coacervation of vinyl polyelectrolytes. *Polymers* 6, 1756–1772 (2014).
- 30. Perry, S. L. et al. Chirality-selected phase behaviour in ionic polypeptide complexes. Nat. Commun. 6, 6052 (2015).
- Kurinomaru, T., Tomita, S., Hagihara, Y. & Shiraki, K. Enzyme hyperactivation system based on a complementary charged pair of polyelectrolytes and substrates. *Langmuir* 30, 3826–3831 (2014).
- Leiros, I. et al. The 2.1 Å structure of Aerococcus viridans L-lactate oxidase (LOX). Acta Crystallogr. Sect. F Struct. Biol. Cryst. Commun. F62, 1185–1190 (2006).
- Hiraka, K. et al. Structure of lactate oxidase from Enterococcus hirae revealed new aspects of active site loop function: Productinhibition mechanism and oxygen gatekeeper. Protein Sci. 10, c4434 (2022).
- Stoisser, T., Brunsteiner, M., Wilson, D. K. & Nidetzky, B. Conformational flexibility related to enzyme activity: evidence for a dynamic active-site gatekeeper function of Tyr(215) in Aerococcus viridans lactate oxidase. Sci. Rep. 6, 27892 (2016).
- Stoisser, T., Klimacek, M., Wilson, D. K. & Nidetzky, B. Speeding up the product release: A second-sphere contribution from Tyr191 to the reactivity of L-lactate oxidase revealed in crystallographic and kinetic studies of site-directed variants. FEBS J. 282, 4130–4140 (2015).
- Thiele, J. M. et al. Enzyme-polyelectrolyte complexes boost the catalytic performance of enzymes. ACS Catal. 8, 10876–10887 (2018).
- 37. Lee, E.-H. et al. Enhancement of enzyme activity and stability by poly(γ-glutamic acid). Polym. J. 42, 818-822 (2010).
- Brady, J. P. et al. Structural and hydrodynamic properties of an intrinsically disordered region of a germ cell-specific protein on phase separation. Proc. Natl Acad. Sci. U. S. A. 114, E8194–E8203 (2017).
- Nott, T. J. et al. Phase transition of a disordered nuage protein generates environmentally responsive membraneless organelles. Mol. Cell 57, 936–947 (2015).
- Harada, R., Tochio, N., Kigawa, T., Sugita, Y. & Feig, M. Reduced native state stability in crowded cellular environment due to protein-protein interactions. J. Am. Chem. Soc. 135, 3696–3701 (2013).
- Tibble, R. W., Depaix, A., Kowalska, J., Jemielity, J. & Gross, J. D. Biomolecular condensates amplify mRNA decapping by biasing enzyme conformation. *Nat. Chem. Biol.* 17, 615–623 (2021).
- 42. Privalov, P. L. & Gill, S. J. Stability of protein structure and hydrophobic interaction. Adv. Protein Chem. 39, 191-234 (1988).
- Pace, C. N., Treviño, S., Prabhakaran, E. & Scholtz, J. M. Protein structure, stability and solubility in water and other solvents. *Philos. Trans. R. Soc. Lond. B Biol. Sci.* 359, 1225–1234 (2004) (discussion 1234).
- 44. Goldsmith, M. & Tawfik, D. S. Enzyme engineering: Reaching the maximal catalytic efficiency peak. *Curr. Opin. Struct. Biol.* 47, 140–150 (2017).
- Bornscheuer, U. T. & Pohl, M. Improved biocatalysts by directed evolution and rational protein design. Curr. Opin. Chem. Biol. 5, 137–143 (2001).
- 46. Shitanda, I. *et al.* High-power lactate/O<sub>2</sub> enzymatic biofuel cell based on carbon cloth electrodes modified with MgO-templated carbon. *J. Power Sources* **436**, 226844 (2019).
- Shitanda, I. et al. High-performance, two-step/Bi-enzyme lactate biofuel cell with lactate oxidase and pyruvate oxidase. J. Power Sources 498, 229935 (2021).
- Bollella, P., Sharma, S., Cass, A. E. G. & Antiochia, R. Microneedle-based biosensor for minimally-invasive lactate detection. Biosens. Bioelectron. 123, 152–159 (2019).

## Acknowledgements

This study is based on results obtained from a project, JPNP14004, commissioned by the New Energy and Industrial Technology Development Organization (NEDO). This work was also partially supported by JSPS KAKENHI (Grant No. 22H05437) to T. M., JSPS KAKENHI (Grant No. 18H02383) to K. S., JSPS Fellows (Grant No. 20J21734) and Leave a Nest grant Fundamental Biotechnology award to T.U. We would like to thank Editage (www.editage.com) for English language editing.

## Author contributions

T.U., K.S., and T.M. conceived and designed the experiments. A.K. purified LOX. Y.H. performed NMR experiments and analyzed the data. T.U. performed most of the experiments and analyzed the data together with S.N. N.T., and T.K. contributed to the interpretation of results. T.U., K.S., and T.M. wrote the manuscript with input from all co-authors.

### **Competing interests**

The authors declare no competing interests.

#### Additional information

**Supplementary Information** The online version contains supplementary material available at https://doi.org/ 10.1038/s41598-023-28040-1.

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