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A fluorescent peptidyl substrate for visualizing peptidyl-prolyl *cis/trans* isomerase activity in live cells

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This paper reports on a fluorescent probe (PPI-P) for imaging active peptidyl-prolyl *cis/trans* isomerases in live cells. PPI-P is capable of responding to both recombinant and cellular PPIases fluorogenically, and has been shown to specifically image active PPIases in live cells.

Due to the partial double bond character of peptidyl-prolyl bonds and the high activation enthalpies for *cis/trans* isomerization, prolyl isomerization is intrinsically slow but can be accelerated by peptidylprolyl cis/trans isomerases (PPlases).¹ PPlases are therefore considered as important chaperones in modulating the folding, trafficking and function of target proteins.²⁻⁴ PPIases are divided into three groups, the FK506-binding protein (FKBP) family, Cyclophilin and the Parvulin family (Pin1 and Par14).5,6 PPlases directly or indirectly regulate pathogenic protein multimerization in human diseases and represent a family rich in targets for modulating mitochondrial function, chaperone activity, stress response, transcription gene regulation, chromatin dynamic, and kinase activity.7-11 Many individual genes within the PPIase family are associated with age-related diseases,12 including cardiovascular diseases, cancer and age-related macular degeneration, e. g., Pin1 has been implicated in the pathogenesis of cancers and Alzheimer's disease,13, 14 while FKBP25 participates in epigenetic regulation of gene expression and ribonucleoprotein complexes.¹⁵ Despite the biological importance, the relationship between PPIase activity and their precise function, and the underlying mechanism responsible for physiological regulation and pathological dysregulation of prolyl isomerization remain largely unknown, which emphasizes the necessity of reliable assays for monitoring PPIase activity.

To date, the catalytic activity of PPIases is usually monitored spectrophotometrically by the chymotrypsin-coupled assay using *N*-succinyl-Ala-Ala-Pro-Phe-*p*-nitroanilide peptide as a probe. This

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assay exploits the finding that α -chymotrypsin can proteolyze the *p*nitroanilide amide bond to enhance the absorbance at 390 nm when the Ala-Pro bond is in the *trans* conformation.^{16,17} While this assay has facilitated PPIase evaluation and yielded important results, it can only be used with lysates or recombinant proteins thus precluding the application to live cells. Since cell lysis disrupts the carefully controlled cellular environment and may affect the stability and function of target proteins, it is very important to study proteins in their native environments in order to achieve more biologically relevant results.¹⁸

In recent years, chemical tools have become the method of choice for detecting protein activity in intact live samples.^{19,20} Though there have been a number of probes reported for imaging various proteins, probes that could significantly advance PPIase research remain undeveloped. With this research we introduce an exciting new chemical tool to the currently available toolbox of probes dedicated to understanding protein function: fluorogenic probe (PPI-P) for monitoring PPIase activity in live cells. The probe is a peptide with the sequence of Ala-Ala-Pro-Phe labeled with a 6-(dimethylamino)-2-naphthoyl fluorophore at the N terminus and a p-nitroanilide group at the C terminus. We reasoned that the cis geometry of PPI-P would bring the naphthalene fluorophore and the nitroanilide quencher into close proximity and therefore cause fluorescence quenching.²¹ Prolyl isomerases can catalyze the transformation of the *cis* prolyl bond to *trans*, separating the fluorophore from the quencher and therefore restoring the fluorescence. We have shown that **PPI-P** is sensitive to PPIases in vitro. Notably, it facilitates the visualization of the dynamic changes involved during PPIase activity in live cells.

Initially, to facilitate the imaging of functional PPIases in live cells, we took advantage of their ability to catalyze prolyl-containing peptide isomerization. Since the distance between the *N* and *C* terminals of a peptide may change significantly before and after the peptidyl-prolyl isomerization, a fluorophore and a quencher were incorporated into a PPIase substrate, resulting in a probe where fluorescence intensity changes can be used to monitor the isomerization. For this purpose, Ala-Ala-Pro-Phe was selected as a representative PPIase substrate due to its broad-spectrum activity

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among various PPIases,²² and probe **PPI-P** was developed by tagging the tetrapeptide with the 6-(dimethylamino)-2-naphthoyl fluorophore at the N terminus and p-nitroanilide quencher at the C terminus. We reasoned that as the *cis* isomer, the probe would be weakly fluorescent due to the close proximity of the fluorophore and the quencher. Then, PPIases would catalyze the isomerization from cis to trans, separating the fluorophore from the quencher, resulting in a concomitant fluorescence increase (Fig. 1). The tagged PPI-P peptide was synthesized as detailed in the supporting information. As shown by the attached NMR traces in the supporting information, both the ¹H NMR signals and ¹³C NMR signals of **PPI-P** appear in pairs, suggesting the presence of both the *cis* and *trans* isomers. However, high performance liquid chromatography (HPLC) analysis gave only one peak under several elution conditions. After failing to differentiate the cis isomer from the trans one, we decide to use PPI-**P** as such for the following experiments.



Fig. 1 Structure of PPI-P and the proposed mode of action.

With PPI-P in hand, we first investigated its fluorescence response towards PPIases in lysates. Given that PPIases exist in most cell types, we then measured the fluorescence response of PPI-P to various lysates. Having confirmed that the lysis buffer caused no change to the PPI-P fluorescence (Fig. S1). We then recorded the spectra of PPI-P in the presence of various lysates. As shown in Fig. 2A, PPI-P itself in PBS displayed moderate fluorescence centered at 460 nm (Φ 0.101), probably due to presence of residual trans isomer. Interestingly, all the cell lysates tested were able to induce fluorescence enhancement of PPI-P. And an increase of fluorescence quantum yield to 0.132 was observed when PPI-P fluorescence plateaued after the treatment of lysate from HEK-293 cells. Furthermore, using lysate from N2a cells, we were able to demonstrate that the fluorescence increase of PPI-P was dose dependent, with increasing lysate resulting in a stronger fluorescence response (Fig. 2B, Fig. S2), similar results were obtained when PPI-P was treated with lysate from HEK-293 cells (Fig. S3). To make sure that this observed fluorogenic response towards cell lysates was not due to probe aggregation, we determined the solubility of PPI-P in PBS (pH 7.4, 10 mM) by UV absorption analysis, and a linear correlation between absorption intensity and PPI-P concentration was observed in the range of 0-50 μ M (Fig. S4), suggesting the good solubility of the probe under the working concentration. We also confirmed that PPI-P was stable in cell lysate by monitoring the probe-lysate mixture with liquid chromatography - mass spectrometry (LC-MS) analysis. As shown in Fig. S5, when PPI-P was treated with HEK-293 lysate, a time-dependent fluorescent intensification was observed. When aliquots of the mixture at indicated time was analysed by LC-MS, no new peak other than PPI-P could be found, indicating that PPI-P is stable enough to resist decomposition at least during the tested period. These results also

indicate that the fluorogenic response of **PPI-P** towards cell lysates was due to its isomerization from *cis* to *trans*.



Fig. 2 (A) Fluorescence spectra of PPI-P (5 μ M) before and after the treatment (10 min) of lysates (20 μ L) from various cell lines. (B) Fluorescence spectra of PPI-P (5 μ M) after the treatment (10 min) of various amounts of lysate from N2a cells. All cell lysates were adjusted to a total protein concentration of 1 μ g/ μ L before addition. Spectra were taken in PBS (10 mM, pH 7.4) at ambient temperature.

To further confirm that the fluorescence response of **PPI-P** to cell lysates was due to PPlases, two additional experiments were performed. First, the response of PPI-P towards recombinant PPIases was investigated. For this purpose, recombinant FKBP25 and Pin1 were chosen due to their relevance in the pathological processes of cardiovascular and cerebrovascular disorders.23,24 The catalytic activity of FKBP25 was verified using the traditional chymotrypsincoupled assay which displayed a dramatically accelerated enhancement for the absorption of N-succinyl-Ala-Ala-Pro-Phe-pnitroanilide at 405 nm in the co-presence of α -chymotrypsin (Fig. S6), indicating the efficient catalytic activity of recombinant FKBP25. Interestingly, when the **PPI-P** probe was treated with the same recombinant FKBP25, significant fluorescence enhancement was observed (Fig. 3A). Consistently, similar results were obtained for recombinant Pin1 treatment (Fig. 3B), demonstrating the potential of **PPI-P** as a reliable and facile probe for PPIase activity.

Second, we investigated the effect of FKBP inhibitors on the lysateinduced fluorescence enhancement. When **PPI-P** was treated with whole lysate derived from HEK293 cells, a significant fluorescence enhancement was observed (Fig. 3C). However, the fluorescence increase was significantly reduced when the system was coincubated with rapamycin or FK506, two potent inhibitors of FKBPs (Fig. 3D), clearly suggestive of **PPI-P** being the molecular target of PPIases. All these results taken together, demonstrate the specificity of **PPI-P** towards PPIases among the myriad of other biological components found in lysates.



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Fig. 3 Characterization of the kinetic response of **PPI-P** to PPIases. **PPI-P** fluorescence was recorded at 460 nm after recombinant FKBP25 (1 µg/µl, 10 µl) (A), or recombinant Pin1 (1 µg/µl, 10 µl) (B) treatment. (C) Lysate from HEK293 cells (1 µg/µl, 10 µl) could also induce a **PPI-P** fluorescence increase and (D) this increase by cell lysate (1 µg/µl, 5 µl) could be blocked in the presence of FKBP inhibitor rapamycin (final concentration, 10 µM) and FK506 (final concentration, 1 mM).

Having confirmed the capability of **PPI-P** to evaluate PPIase activity from lysates, we moved on to evaluate the feasibility to image endogenous PPIase activity in live cells. We first confirmed that **PPI-P** was of little cytotoxicity by CCK-8 assay (Fig. S7). Then we stained EA.hy926 cells with **PPI-P** after cells were transfected with lentiviral vector encoding EGFP for dual-colour confocal imaging, and the dynamic change of **PPI-P** fluorescence in the transfected cells was monitored (video S1).



Fig. 4 Detecting endogenous PPIase activity in live EA.hy926 cells with PPI-P. Data shown are the time-lapse confocal images of EA.hy926 cells treated with PPI-P (final

concentration, 2.5 μ M) in the absence (A) or presence (B) of rapamycin (final concentration, 1 μ M). Lentiviral vector encoding EGFP were transfected into EA.hy926 cells for confocal imaging. The merge panels indicate an overlap of the EGFP (Green) signal with **PPI-P** (Blue) signal. Probe fluorescence was collected at 420–480 nm with λ_{ex} 405 nm. EGFP fluorescence was collected at 500–550 nm with λ_{ex} 488 nm.

The series of images in Fig. 4A are individual frames from the continuous time-lapse movie, from which we observed the gradual elevation of intracellular **PPI-P** fluorescence in a time dependent manner, which was confirmed by quantification of the fluorescence data (Fig. S8), indicating the presence of native PPlases and their catalytic effect on **PPI-P** to catalyze the isomerization of the probe from *cis* to *trans*. In order to assess the specificity of **PPI-P** in live cells, its fluorescence response in EA.hy926 cells pretreated with rapamycin was monitored (video S2). As shown in Fig. 4B and S8, PPlase inhibition by rapamycin resulted in a reduced and delayed fluorescent response for **PPI-P** to endogenous PPlases, further demonstrating the specificity of **PPI-P** for PPlases in live cells.

To further evaluate the spatiotemporal resolution of **PPI-P** for determining PPlase activity in live cells, the response towards overexpressed FKBP25 fused with EGFP was investigated. The lentivirus vector encoding EGFP-FKBP25 was transfected into live EA.hy926 cells, and FKBP25-overexpression-induced **PPI-P** fluorescence changes were monitored in real-time with EGFP as an overexpression indicator (video S3). Representative images from video S3 are given in Fig. 5. Notably, intracellular **PPI-P** fluorescence was positively correlated with the FKBP25 expression level with cells overexpressing EGFP-FKBP25 demonstrating stronger **PPI-P** fluorescence than its non-overexpressing counterparts (Fig. 5 and S9), which agrees well with the data in Fig. 2B, further suggesting the specificity of **PPI-P** towards PPlases. Taken together, these experiments provide convincing evidence that **PPI-P** is capable of detecting endogenous PPlase activity in live cells.



Fig. 5 Confocal fluorescence images of FKBP25-induced PPIase activity detected by PPI-P in live EA.hy926 cells. Lentiviral vector encoding FKBP25-EGFP were transfected into EA.hy926 cells. Cells were then treated with PPI-P (2.5 μ M). Data shown are the timelapse series for confocal plane images recorded from live EA.hy926 cells over a 20-min

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time frame. The dotted line in white or yellow respectively indicated the cells overexpression the FKBP25-EGFP or not. Probe fluorescence was collected at 420–480 nm with λ_{ex} 405 nm. EGFP fluorescence was collected at 500–550 nm with λ_{ex} 488 nm.

In summary, by taking advantage of the mechanism of PPIase activity, we have developed a fluorogenic probe to report on the catalytic activity of PPIases in both lysates and live cells. The specificity and sensitivity of the probe has been demonstrated using the imaging of exogenous and endogenous PPIases in live cells. It should be noted that although the sensitivity of the probe is not good enough due to the apparent background fluorescence, it represents the first example where PPIase activity has been directly imaged in live cells, and should be inspiring for future work. Given the importance of PPIases in regulating protein conformation and function, we envision that further work may be conducted with the construction of new probes whose spectra properties at *cis* and *trans* geometry could be well resolved and those that have brighter and red-shifted excitation/emission spectra.

Conflicts of interest

There are no conflicts to declare.

Notes and references

All methodology procedures are detailed in the Supporting Information. This work was supported by the National Key Research and Development Program of China (2016YFE0125400), National Natural Science Foundation of China (81573411, 21778048), Natural Science Foundation of Zhejiang Province (LZ16H310001, LR18H300001), Zhejiang Province Program for Cultivation of Highlevel Health talents.

- 1 G. Fischer, Chem. Soc. Rev., 2000, 29, 119-127.
- 2 K. Lang and F. X. Schmid, *Nature*, 1988, **331**, 453-455.
- 3 K. Lang, F. X. Schmid and G. Fischer, *Nature*, 1987, **329**, 268-270.

- 4 G. Fischer and F. X. Schmid, *Biochemistry-Us*, 1990, **29**, 2205-2212.
- 5 C. Scholz, J. Rahfeld, G. Fischer and F. X. Schmid, *J. Mol. Biol.*, 1997, **273**, 752-762.
- 6 J. Fanghanel and G. Fischer, Front Biosci, 2004, 9, 3453-3478.
- 7 A. Bell, P. Monaghan and A. P. Page, *Int J Parasitol*, 2006, **36**, 261-276.
- 8 L. J. Blair, J. D. Baker, J. J. Sabbagh and C. A. Dickey, J Neurochem, 2015, 133, 1-13.
- 9 S. D. Hanes, Biochim Biophys Acta, 2015, 1850, 2017-2034.
- 10 P. E. Shaw, EMBO Rep, 2007, 8, 40-45.
- 11 Q. Yao, M. Li, H. Yang, H. Chai, W. Fisher and C. Chen, World J Surg, 2005, 29, 276-280.
- 12 L. McClements, S. Annett, A. Yakkundi and T. Robson, *Curr Mol Pharmacol*, 2015, **9**, 165-179.
- 13 L. Pastorino, A. Sun, P. J. Lu, X. Z. Zhou, M. Balastik, G. Finn, G. Wulf, J. Lim, S. H. Li, X. Li, W. Xia, L. K. Nicholson and K. P. Lu, *Nature*, 2006, **440**, 528-534.
- 14 T. H. Lee, L. Pastorino and K. P. Lu, *Expert Rev Mol Med*, 2011, **13**, e21.
- 15 R. Thapar, Biomolecules, 2015, 5, 974-999.
- 16 B. Janowski, S. Wollner, M. Schutkowski and G. Fischer, Anal Biochem, 1997, 252, 299-307.
- 17 Y. J. Jin, S. J. Burakoff and B. E. Bierer, *J Biol Chem*, 1992, **267**, 10942-10945.
- 18 J. A. Prescher and C. R. Bertozzi, Nat Chem Biol, 2005, 1, 13-21.
- 19 J. A. Gonzalez-Vera and M. C. Morris, *Proteomes*, 2015, **3**, 369-410.
- 20 K. M. Dean and A. E. Palmer, *Nat Chem Biol*, 2014, **10**, 512-523.
- 21 Y. Hori, K. Nakaki, M. Sato, S. Mizukami and K. Kikuchi, Angew Chem Int Ed Engl, 2012, **51**, 5611-5614.
- 22 G. Zoldak, T. Aumuller, C. Lucke, J. Hritz, C. Oostenbrink, G. Fischer and F. X. Schmid, *Biochemistry-Us*, 2009, 48, 10423-10436.
- 23 C. Mas, I. Guimiot-Maloum, F. Guimiot, M. Khelfaoui, V. Nepote, F. Bourgeois, B. Boda, B. Levacher, A. Galat, J. M. Moalic and M. Simonneau, *Gene Expr Patterns*, 2005, 5, 577-585.
- 24 G. L. Perrucci, A. Gowran, M. Zanobini, M. C. Capogrossi, G. Pompilio and P. Nigro, *Cardiovasc Res*, 2015, **106**, 353-364.