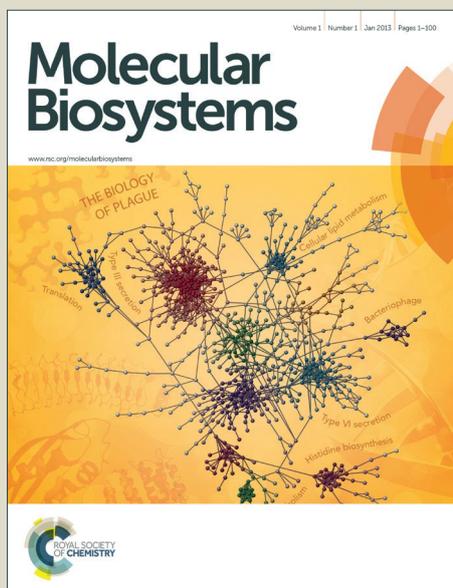


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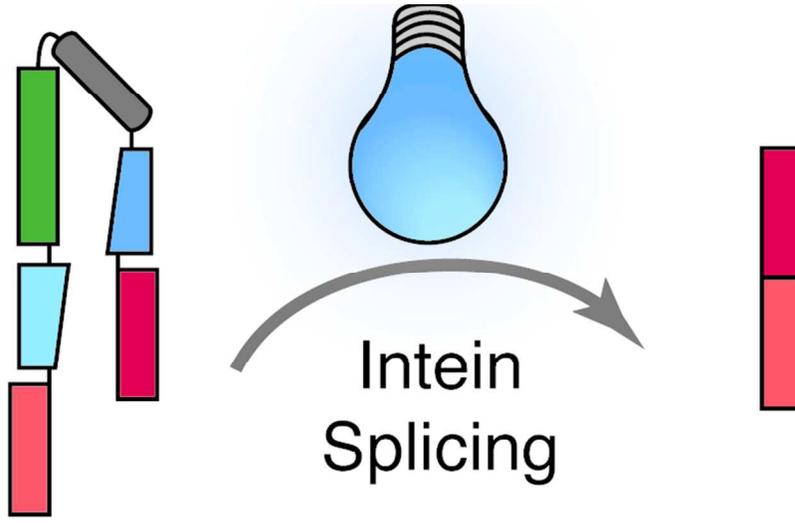
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ARTICLE

Post-translational control of protein function with light using a LOV-intein fusion protein.

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Methods for the post-translational control of protein function with light hold much value as tools in cell biology. To this end, we report a fusion protein that consists of DnaE split-inteins, flanking the light sensitive LOV2 domain of *Avena Sativa*. The resulting chimeria combines the activities of these two unrelated proteins to enable controlled formation of a functional protein via upregulation of intein splicing with blue light in bacterial and human cells.

Introduction

Inteins are naturally occurring protein domains that self-excise from a host protein by an autocatalytic process. Intein-mediated protein splicing results in the adjacent N- and C-terminal peptide chains (known as exteins) being joined by a peptide bond, to give a functional, mature protein. The ability of inteins to join two protein fragments with high specificity and selectivity has resulted in their extensive use for a variety of biotechnological applications. There is also significant potential for methods that enable intein splicing to be selectively triggered by an external stimulus; this would allow the activity of a given protein to be regulated post-translation, enabling initiation of a given protein's function with high special and temporal control in living cells (*via* activation of intein splicing and excision of the intein domain). The majority of current approaches involve fusing an additional domain to an intein so that splicing is allosterically triggered by either a small molecule,¹⁻⁴ changes in pH,⁵ or temperature.^{6,7} Although these approaches allow protein splicing to be triggered throughout a given sample, they do not allow special or temporal control of activity within a subset of the sample. This limitation has been overcome by replacing key residues required for splicing with photocaged non-natural amino acids, which enables intein-splicing to be initiated with UV light.⁸⁻¹² While potentially very powerful, these approaches necessitate amber codon reassignment and so require the incorporation of an orthogonal tRNA/tRNA synthetase pair into the targeted living system.

We envisaged an alternative approach to engineering photo-regulation into inteins, whereby the coupling of a light sensitive protein domain to an intein results in functional

combination of the activities of these unrelated proteins. The *Avena Sativa* Light Oxygen Voltage 2 (LOV2) domain, which undergoes a marked change in protein structure upon exposure to blue light, has been previously used to directly engineer photocontrol into several proteins.¹³ The photochemistry of the LOV2 domain is determined by the properties of a flavin mononucleotide ligand whose excitation by a photon leads to the formations of a cysteinyl adduct, resulting in a number of conformational changes that lead to the dissociation of an α -helix (called the $J\alpha$ -helix) from the main body of the protein.¹⁴ We hypothesized that the change in conformation and spatial orientation of the $J\alpha$ -helix upon exposure of LOV2 to blue light may be exploited to regulate the splicing a split intein (Figure 1A). We chose a mutant form of the *Nostoc punctiforme* (*Npu*) DnaE trans intein for this purpose for two reasons;¹⁵ first, its high tolerance to amino acid variations at the splicing junction would allow the function of a variety of exteins to be controlled with the resulting chimeric protein.¹⁶ Second, we reasoned that as the splice time of *Npu* inteins^{17,18} is less than the recovery half life of the LOV2 photo-adduct,¹⁹ intein splicing would occur prior to recovery of LOV2 back to its dark state.

Whilst this manuscript was in preparation, Truong and co-workers reported photoactivated trans splicing by fusing LOV2 to a truncated C-terminal *Npu* intein.²⁰ Here, we detail the a generally applicable method that utilizes LOV2 for the photocontrol of cis splicing of *Npu* inteins.

Results

Light-activated intein splicing in *E. coli*

We initially constructed a cassette encoding a fusion protein consisting of the N-terminal *Npu* intein, followed by LOV2, followed by the C-terminal *Npu* intein (referred to as InN-LOV-

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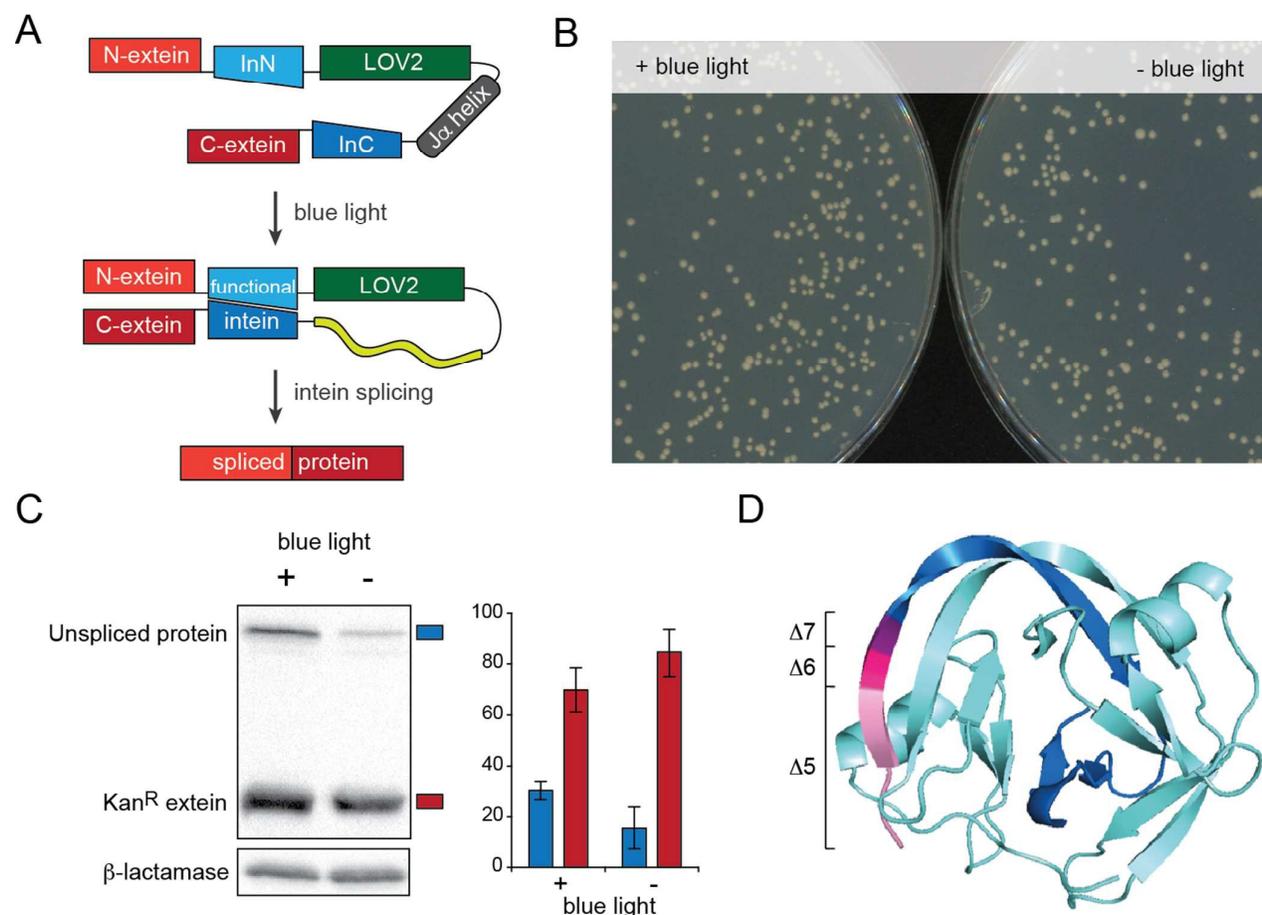


Figure 1. Light activated protein splicing with a LOV-intein chimeric protein. **A**) A fusion protein consisting of the LOV2 domain, flanked by N- and C-terminal inteins (InN and InC) and N- and C-terminal extein halves would be expected to undergo conformational change upon exposure to blue light, initiating protein splicing to give the mature extein. **B**) Blue-light dependent growth advantage is not observed in *E. coli* transformed with the plasmid encoding Kan^{R188}-InN-LOV-InC-Kan^{R189-272} and incubated on media containing kanamycin (25 μ g/mL). **C**) Western blot analysis of intein splicing in the above cells, shows a similar rate of intein splicing for samples grown in the dark, or under blue light. **D**) Structure of the *Npu* inteins, showing the amino acids deleted in the InN-LOV- Δ 5InC, InN-LOV- Δ 6InC, and InN-LOV- Δ 7InC. PDB ID = 4LX3.

InC from here). Neomycin phosphotransferase, which imparts kanamycin resistance (referred to as kan^R from here), was split between F188 and S189 and the two resulting fragments used as exteins that flanked either side of InN-LOV-InC to give a plasmid encoding Kan^{R188}-InN-LOV-InC-Kan^{R189-272} (referred to as Kan^R-LOV-intein). It should be noted when choosing the site to split a potential extein, that the mechanism of intein splicing requires a cysteine or serine as the first amino acid of the second extein fragment (joined to the C-intein). *E. coli* were transformed with the above plasmid and incubated overnight on media supplemented with kanamycin at 37 °C, either in the dark or under blue light from a LED array (Supplementary Figure 1).

If splicing of Kan^R-LOV-intein is regulated by blue light, more colonies would be expected to be present on plates incubated under blue light than those grown in the dark. There was however no change in the number or size of surviving colonies between the samples (Figure 1B), indicating that the InN-LOV-InC fusion protein splices even in the absence of blue light. We further assessed this by western blot (Figure 1C), and observed a high rate of splicing, with no difference between samples

incubated under blue light and those grown in the dark. Given this observation, we reasoned that the high affinity of the N- and C-terminal *Npu* intein fragments for each other is either overriding, or affecting LOV2 photoswitching. To probe this, we measured the recovery of photoactivated LOV2 via the half-life of the flavin mononucleotide-cysteiny adduct.²¹ We observed a significant reduction in the half-life of photoadduct decay in the InN-LOV-InC chimera, from 58 ± 0.3 seconds for LOV2 (Supplementary Figure 2) to 24 ± 0.1 seconds for InN-LOV-InC (Supplementary Figure 3). This suggested that the *Npu* inteins destabilize the photoactivated form of LOV2. We reasoned that this was due to the high affinity of *Npu* intein fragments for each other, and that reducing this affinity should also reduce the effect of the inteins on LOV2 photoswitching, potentially enabling light-activated protein splicing. Previous work has shown that deleting the first 6 amino acids from the N-terminus of the *Synechocystis* sp. PCC6803 DnaE C-intein is sufficient to eliminate the binding and splicing of N- and C-inteins.²² We therefore constructed plasmids encoding InN-LOV-InC with deletions of 5, 6 or 7 amino acids from the N-terminus of the *Npu* C-intein (Figure 1D). As before, Split Kan^R

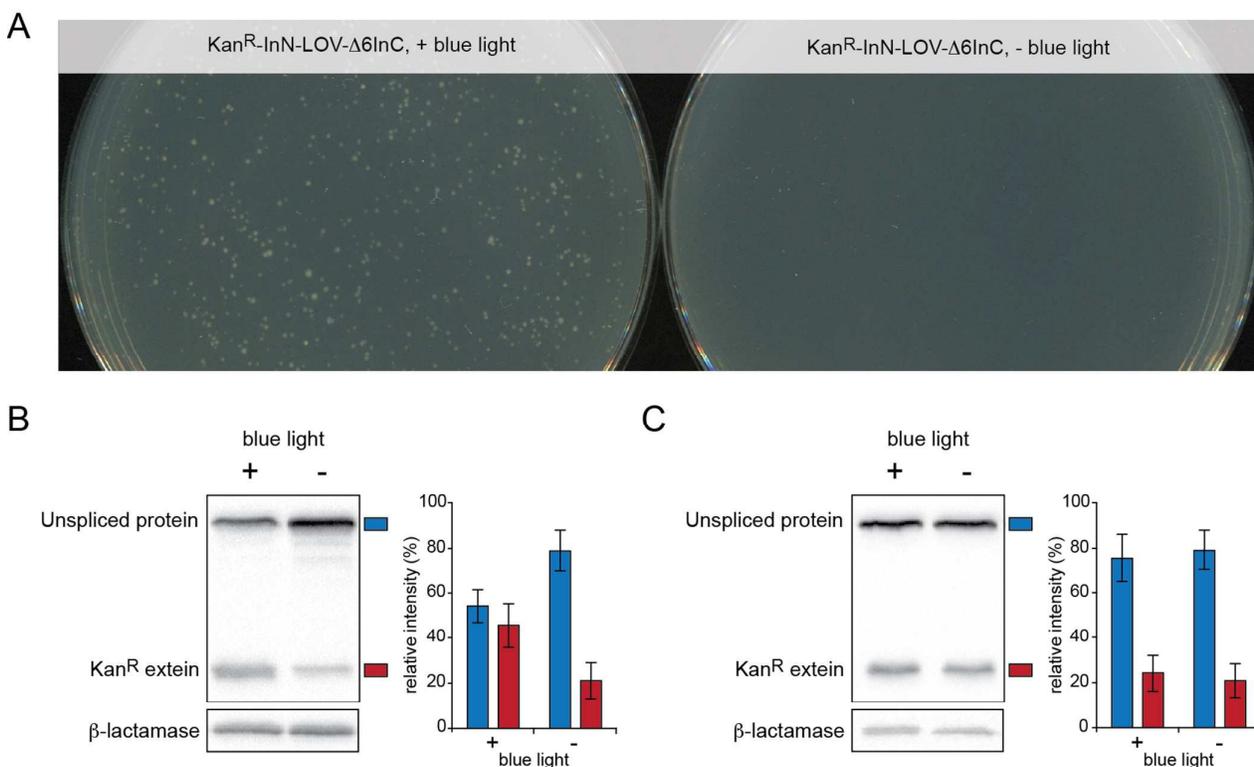


Figure 3. Light-activated intein splicing in *E. coli*. **A)** Bacteria transformed with the Kan^{R1-188}-InN-LOV-Δ6InC-Kan^{R189-272} plasmid were incubated overnight on LB-agar containing 30 µg/mL of Kanamycin, in the presence or absence of blue light. Colonies incubated under blue light show a growth advantage over those incubated in the dark. **B)** Western blot analysis of intein splicing shows a ~2-fold increase in spliced extein (Kan^R) in bacterial incubated under blue light, compared to those incubated in the dark. **C)** The effect from blue light is eliminated when using dark-locked LOV^{C450S} protein, showing that the above effect is dependent on a functional LOV2, and not a result of blue light directly affecting intein splicing.

was used as the extein. *E. coli* were transformed with these plasmids and incubated overnight on LB agar containing kanamycin, either in the dark, or under blue light. Bacteria transformed with the plasmid encoding Kan^{R1-188}-InN-LOV-Δ5InC-Kan^{R189-272} did not show growth advantage under blue light (Supplementary Figure 4), while those transformed with the plasmid encoding Kan^{R1-188}-InN-LOV-Δ6InC-Kan^{R189-272} grew significantly better under blue light than in the dark (Figure 2A). Bacteria transformed with the plasmid encoding Kan^{R1-188}-InN-LOV-Δ7InC-Kan^{R189-272} did not grow in the dark, or under blue light (Supplementary Figure 5). These results suggest that deletion of 5 amino acids from the N-terminus of the C-terminal intein does not sufficiently reduce the affinity of the *Npu* intein fragments to allow control of splicing by LOV, whereas deletion of 7 amino acids reduces the affinity of the intein fragments to the extent that they are no longer able to splice. Deletion of 6 amino acids from the C-intein however, reduces the affinity of the intein fragments to the extent that the conformation change of LOV upon photo-activation regulates splicing. We further examined the effect of blue light on InN-LOV-Δ6InC splicing in these cells by western blot. Cells transformed with this plasmid were grown without selective pressure for the spliced extein (i.e. in the absence of kanamycin), and protein levels between samples was normalized using β-lactamase levels, which is expressed from the same plasmid as the InN-LOV-Δ6InC. We observed around ~2.5-fold higher levels of spliced kan^R protein in bacteria

incubated under blue light than those grown in the dark (Figure 2B). Comparing the ratio of the unspliced intein to the spliced extein between InN-LOV-InC (Figure 1C) and InN-LOV-Δ6InC (Figure 2B) suggests that the splicing of the InN-LOV-Δ6InC is less efficient in comparison, probably due to the reduced affinity of the *Npu* intein fragments, but is nonetheless regulated by LOV2 photoactivation. Analysis of the kinetics of LOV switching in InN-LOV-Δ6InC sheds further light on the mode of action; the half life of the LOV2 flavin mononucleotide-cysteinylyl adduct in the InN-LOV-Δ6InC protein was measured as 38 ± 0.7 seconds (Supplementary Figure 6), which is around 2-fold slower than the original InN-LOV-InC protein. This supports our hypothesis that the affinity of the *Npu* intein fragments was affecting the photoswitching of LOV2. The kinetics of photoswitching of similar inteins has recently been reported to be 2-3 hours (an instant response is observed, which plateaus to a maximum after ~150 minutes).²⁰ The level of LOV-independent splicing, and the effect of blue light on the splicing of *Npu* inteins was measured using a C450S mutant of LOV2 that has previously been shown to be locked in its dark state even in the presence of blue light.²³ We observed no difference in spliced extein levels in samples incubated in the dark or in the presence of blue light (Figure 2C), indicating that a functional LOV2 domain is required for the regulation InN-LOV-Δ6InC splicing with blue light. Interestingly, incorporation of LOV^{C450S} into the original Kan^R-InN-LOV-InC complex had no effect on the relative levels of

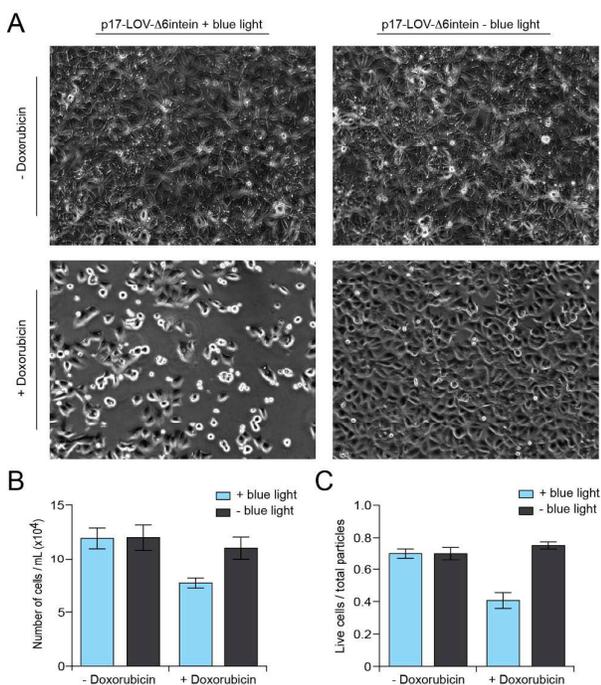


Figure 4. Light-activated intein splicing in human MCF7 cells. **A)** Micrographs of MCF7 cells transfected with the p17-InN-LOV-Δ6InC-p12 plasmid, cultured with or without 10 μM doxorubicin, in the dark or under blue light. A reduction in viability and the presence of dead cells is clearly visible in the panel containing cells cultured in blue light and with doxorubicin. **B)** Effect of doxorubicin on cell number of MCF7 cells transfected with the p17-InN-LOV-Δ6InC-p12 plasmid, incubated in the dark or under blue light. **C)** Effect of doxorubicin on the ratio of living cells versus total particles in MCF7 cells transfected with p17-InN-LOV-Δ6InC-p12 plasmid, incubated in the dark or under blue light.

spliced extein (Supplementary Figure 7), further indicating that the high affinity of the *Npu* split-intein fragments overrides any conformational inhibition of splicing when LOV is in its dark state.

Light-activated intein splicing in human cells.

We next sought to assess the possibility of using the InN-LOV-Δ6InC for photo-activated splicing in human cells. MCF7 breast cancer cells are deficient in caspase 3 (due to a deletion mutation in exon 3 of its gene),²⁴ and the absence of caspase 3 is required for the resistance of MCF7 cells to the chemotherapeutic agent Doxorubicin; the introduction of a plasmid encoding caspase 3 into MCF7 cells is sufficient to re-sensitize them to doxorubicin and trigger apoptosis.²⁵ We sought to use this property to demonstrate photocontrolled unmasking of caspase 3 function with our InN-LOV-Δ6InC in MCF7 cells. Caspase 3 has been previously demonstrated to be functionally reconstitutable when split between its p17 and p12 domains (amino acids D175 and S176).²⁶ We therefore generated a plasmid that encoded these two halves of caspase 3 as exteins that flank the sequence encoding the InN-LOV-Δ6InC protein. The encoded fusion protein would be expected to splice in the presence of blue light to reconstitute functional caspase 3, causing a reduction in the viability of MCF7 cells in the presence of doxorubicin.

MCF7 cells were transfected with the plasmid encoding a p17-InN-LOV-Δ6InC-p12 fusion protein, and incubated in the dark

for 24 hours. These cells were then incubated either in the dark or under blue light for 3 hours, in the presence or absence of 10 μM doxorubicin. Significant loss of cell viability was observed in the micrographs of cells incubated in the presence of blue light and doxorubicin compared to those incubated in the dark; there was no difference between samples incubated with blue light or in the dark in the absence of doxorubicin (Figure 3A). This observation was confirmed when the number of cells was quantified under each condition; $1.20 \times 10^5 \pm 2.0 \times 10^4$ cells were present when transfected MCF7 cells were cultured in blue light, but in absence of doxorubicin, and $1.10 \times 10^5 \pm 1.8 \times 10^4$ cells were present for equivalent samples incubated in the dark, showing no effect on cell viability from blue light alone. In the presence of 10 μM doxorubicin however, a 35% reduction in cell number was observed in the presence of blue light; there were $1.18 \times 10^5 \pm 1.6 \times 10^4$ cells present when transfected MCF7 cells were cultured in the dark, with $7.69 \times 10^4 \pm 7.9 \times 10^3$ cells present in samples cultured under blue light (Figure 3B). The ratio of live cells versus total particles (arising from cell death and apoptosis) was quantified with a microfluidic-based cell counter. In the absence of doxorubicin, no difference was observed in this ratio for cells incubated in the dark (0.70 ± 0.06) or under blue light (0.75 ± 0.04) indicating that blue light in combination with the p17-InN-LOV-Δ6InC-p12 protein was not sufficient to trigger cell death (Figure 3C). In the presence of 10 μM doxorubicin however, a 42% reduction in this ratio was observed for cells under blue light (0.41 ± 0.08) versus those incubated in the dark (0.70 ± 0.05). This data further indicates that blue light triggers splicing of the p17-InN-LOV-Δ6InC-p12 protein to give active caspase 3, which sensitizes MCF7 cells to doxorubicin and results in cell death (Figure 3C).

Conclusions

Taken together, the above data demonstrates that the InN-LOV-Δ6InC protein combines the function of the light sensitive LOV2 domain with that of the *Npu* split-inteins, to enable allosteric control of protein splicing with blue light in *E. coli* and in human cells. It should be noted that although the magnitude of light-activation of splicing is relatively modest (~2.5-fold), this is achieved using the light induced conformational change of the InN-LOV-Δ6InC fusion protein, which consists of only the 20 proteinogenic amino acids. Furthermore, the level of photocontrol observed here is in line with a previous report that used LOV2 to control the activity of dihydrofolate reductase with light,²⁷ and a recent report that uses LOV-fused *Npu* inteins for photocontrol of trans splicing.²⁰ This level of photocontrol by InN-LOV-Δ6InC is therefore likely to be an inherent property of the LOV-intein fusion protein, rather than a result of the chosen extein. It should be noted that our data suggests this level of photoswitching is sufficient for the induction of cellular phenotypes with light. InN-LOV-Δ6InC may therefore be utilized for a variety of biological studies, especially given the relative ease with it can be deployed in a variety of systems.

Materials and Methods

All reagents were purchased from ThermoFisher unless otherwise stated. UV-Vis spectra were recorded on a Vairan Cary 100. Cell culture reagents were purchased from Life Technologies unless otherwise stated. MCF7 cells were sourced from the American Type Culture Collection and not grown continually for more than 3 months. Cells were cultured at 37°C in a humidified 5% CO₂ atmosphere in DMEM containing 10% fetal bovine serum (FBS) and 1% penicillin/streptomycin.

Plasmid construction

All genes were synthesized de novo by Integrated DNA Technologies. The amino acid sequence of the light switchable InN-LOV-Δ6InC protein is given in supplementary information. The KanR protein was fragmented in two between F188 and S189, and Caspase 3 was split between D175 and S176. The genes used for *E. coli* expression were inserted into pRSET, the T7 promoter of this plasmid was replaced with a constitutive promoter. The genes used in human cell studies were cloned into the backbone of pEGPF-N1 (the eGFP gene was removed).

Light/dark growth of samples

Plasmids were transformed in to *E. coli*, and after recovery, incubated overnight at 37°C in darkness with shaking, in media containing 100 µg/mL of carbenecillin. 1% of the overnight culture was plated onto LB agar plates containing 25 µg/mL of Kanamycin and 100 µg/mL of carbenecillin. These plates were divided into two sets and incubated at 37°C for 16 hours, either in the dark, or under blue light. Plates grown under light were illuminated with an LED array composed of high-emission low power InGaN bulbs (Supplementary Figure 1C). The light intensity was ~5 mW/cm².

Measuring intein splicing by western blot.

An overnight culture of *E. coli* containing the appropriate plasmid was used to inoculate 2 x 500mL of LB (1:200 dilution). Once the OD₆₀₀ of the cultures had reached 0.6 they were separated in two. The first flask was incubated with shaking at 18°C overnight in a darkroom (to exclude light). The second was incubated with shaking at 18°C overnight, under blue light (Supplementary Figure 1D). Each culture was harvested by centrifugation at 4000rpm and the pellets were frozen at -80°C. Pellets were lysed using lysis buffer (Imidazole 400 mM, 0.5 M NaCl, 50 mM Tris-HCl, pH 8.0) in the dark. The resulting lysate was separated from cell debris by centrifugation and filtered using a sterile syringe filter (Millipore UK). Proteins were analyzed by SDS-PAGE and western blot as detailed below.

Protein Immunoblotting

Cell lysates were prepared as above, with the protein content of each sample quantified using the Bradford assay. Proteins were separated on 12% SDS-polyacrylamide Bis-Tris gels under denaturing conditions, transferred to unsupported pure nitrocellulose membrane (Amersham) and subjected to

immunoblot analysis. Mouse monoclonal antibody raised against the 6-histidine epitope tag present on the N-terminus of the extein (Novus, 1:2,500) were used for blotting. A mouse antibody raised against β-lactamase (Novus, 1:1,000), which is expressed from the same plasmid as the light-activated intein was used as a loading control. Antibodies were incubated with the membrane overnight at 4 °C. Horseradish peroxidase-conjugated anti-mouse immunoglobulin G antibody (Novus, 1:10,000) was used as the secondary antibody. Bound immunocomplexes were detected using ECL prime western blot detection reagent (GE Healthcare) and analysed using Image Lab 4.0 (Bio-Rad). Molecular mass of visualized proteins was confirmed using molecular markers (for example image see Supplementary Figure 8).

Kinetics of photo-adduct decay

Recombinant LOV-intein proteins were expressed as above, and purified in a darkroom using FPLC and a Nickel affinity column (GE). Proteins were removed from the column with elution buffer (1 M NaCl, 400 mM imidazole, 100 mM Tris-HCl, pH 8.0). The purified proteins were dialyzed into analysis buffer (Sodium Phosphate 10 mM, NaCl 10 mM, pH 8.0), followed by filtration (Millipore UK). Samples were kept in darkness at all times. Photo-adduct recovery was measured by UV-Vis spectroscopy at 25°C. For a dark state spectrum, the sample was left in the UV-Vis spectrometer in total darkness for 10 minutes to ensure depletion of any photo-activated adduct. The absorption spectrum (360-510nm) of the proteins was then measured. For light state spectra, samples were exposed to the LED array for 5 minutes and scanning UV-Vis absorption spectra (360-510nm) were continuously recorded for 600 seconds immediately afterwards. Photo-adduct levels were calculated as $1-(A_t-A_0)/(A_\infty-A_0)$, where A_t , A_∞ , and A_0 are absorptions at 450 nm at time t in seconds, in darkness, and under blue light, respectively. The logarithm of the remaining photo-adduct level was plotted against to indicate the fast and slow components of reversion kinetics. The the decay rate of the C(4a)-cysteiny adduct was determined using the absorption spectrum of the samples after exposure to the LED array. The absorption at 447nm was used to determine the half-life of the adduct using $t_{1/2} = t \times \ln(2) / \ln(A_t/A_0)$, where t , A_t and A_0 are time, absorption at time t in seconds and at time 0 respectively. The rate of photo-adduct decay was determined to be first order or pseudo first order for all samples half lives were quantified from the photo-adduct decay plots using Prism 6 (Graphpad).

Transfection, cell treatments and cell counting

Cells were transiently transfected with the p17-LOV-Δ6intein-p12 plasmids using FuGENE HD (Promega) according to the manufacturers' instructions. Transfected cells were incubated in the dark for 24 h, then treated with 10 µM doxorubicin then either kept in the dark or exposed to blue light for 3 h. Visualization of treated cells under phase contrast was carried out with a Zeiss Axio Vert.A1 microscope at 20 X and 40 X magnification. Cell counts and live/dead cell ratios were obtained using a Moxi Z Mini automated cell counter (Orflo).

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References

- (1) Buskirk, A. R.; Ong, Y. C.; Gartner, Z. J.; Liu, D. R. *Proceedings of the National Academy of Sciences of the United States of America* **2004**, *101*, 10505.
- (2) Mootz, H. D.; Blum, E. S.; Tyszkiewicz, A. B.; Muir, T. W. *Journal of the American Chemical Society* **2003**, *125*, 10561.
- (3) Mootz, H. D.; Muir, T. W. *Journal of the American Chemical Society* **2002**, *124*, 9044.
- (4) Skretas, G.; Wood, D. W. *Protein Sci* **2005**, *14*, 523.
- (5) Wood, D. W.; Wu, W.; Belfort, G.; Derbyshire, V.; Belfort, M. *Nat Biotechnol* **1999**, *17*, 889.
- (6) Zeidler, M. P.; Tan, C.; Bellaiche, Y.; Cherry, S.; Hader, S.; Gayko, U.; Perrimon, N. *Nat Biotechnol* **2004**, *22*, 871.
- (7) Adam, E.; Perler, F. B. *J Mol Microb Biotech* **2002**, *4*, 479.
- (8) Berrade, L.; Kwon, Y.; Camarero, J. A. *Chembiochem : a European journal of chemical biology* **2010**, *11*, 1368.
- (9) Bocker, J. K.; Friedel, K.; Matern, J. C.; Bachmann, A. L.; Mootz, H. D. *Angewandte Chemie* **2015**, *54*, 2116.
- (10) Cook, S. N.; Jack, W. E.; Xiong, X. F.; Danley, L. E.; Ellman, J. A.; Schultz, P. G.; Noren, C. J. *Angew Chem Int Edit* **1995**, *34*, 1629.
- (11) Ren, W.; Ji, A.; Ai, H. W. *Journal of the American Chemical Society* **2015**, *137*, 2155.
- (12) Vila-Perello, M.; Hori, Y.; Ribo, M.; Muir, T. W. *Angewandte Chemie* **2008**, *47*, 7764.
- (13) Gautier, A.; Gauron, C.; Volovitch, M.; Bensimon, D.; Jullien, L.; Vriz, S. *Nature chemical biology* **2014**, *10*, 533.
- (14) Halavaty, A. S.; Moffat, K. *Biochemistry* **2007**, *46*, 14001.
- (15) Lockless, S. W.; Muir, T. W. *Proceedings of the National Academy of Sciences of the United States of America* **2009**, *106*, 10999.
- (16) Iwai, H.; Zuger, S.; Jin, J.; Tam, P. H. *FEBS letters* **2006**, *580*, 1853.
- (17) Zettler, J.; Schutz, V.; Mootz, H. D. *FEBS letters* **2009**, *583*, 909.
- (18) Cheriyan, M.; Pedomallu, C. S.; Tori, K.; Perler, F. *J Biol Chem* **2013**, *288*, 6202.
- (19) Schwartz, E. C.; Saez, L.; Young, M. W.; Muir, T. W. *Nature chemical biology* **2007**, *3*, 50.
- (20) Wong, S.; Mosabbir, A. A.; Truong, K. *Plos One* **2015**, *10*, e0135965.
- (21) Swartz, T. E.; Corchnoy, S. B.; Christie, J. M.; Lewis, J. W.; Szundi, I.; Briggs, W. R.; Bogomolni, R. A. *J Biol Chem* **2001**, *276*, 36493.
- (22) Ghosh, I.; Sun, L.; Xu, M. Q. *J Biol Chem* **2001**, *276*, 24051.

(23) Alexandre, M. T.; van Grondelle, R.; Hellingwerf, K. J.; Robert, B.; Kennis, J. T. *Physical chemistry chemical physics : PCCP* **2008**, *10*, 6693.

(24) Janicke, R. U.; Ng, P.; Sprengart, M. L.; Porter, A. G. *J Biol Chem* **1998**, *273*, 15540.

(25) Yang, X. H.; Sladek, T. L.; Liu, X.; Butler, B. R.; Froelich, C. J.; Thor, A. D. *Cancer research* **2001**, *61*, 348.

(26) Chelur, D. S.; Chalfie, M. *Proceedings of the National Academy of Sciences of the United States of America* **2007**, *104*, 2283.

(27) Lee, J.; Natarajan, M.; Nashine, V. C.; Socolich, M.; Vo, T.; Russ, W. P.; Benkovic, S. J.; Ranganathan, R. *Science* **2008**, *322*, 438.