

**Azobenzene photocontrol of peptides and proteins**

Journal:	<i>ChemComm</i>
Manuscript ID	CC-FEA-05-2016-004004.R1
Article Type:	Feature Article
Date Submitted by the Author:	06-Jul-2016
Complete List of Authors:	Allemann, Rudolf; Cardiff University, School of Chemistry Mart, Robert; Cardiff University, Chemistry

Azobenzene photocontrol of peptides and proteins

 Robert J. Mart,^a and Rudolf K. Allemann^{a*}

 Received 00th January 20xx,
Accepted 00th January 20xx

DOI: 10.1039/x0xx00000x

www.rsc.org/

The last few years have witnessed significant advances in the use of light as a stimulus to control biomolecular interactions. Great efforts have been devoted to the development of genetically encoded optobiological and small photochromic switches. Newly discovered small molecules now allow researchers to build molecular systems that are sensitive to a wider range of wavelengths of light than ever before with improved switching fidelities and increased lifetimes of the photoactivated states. Because these molecules are relatively small and adopt predictable conformations they are well suited as tools to interrogate cellular function in a spatially and temporally controlled fashion and for applications in photopharmacology.

Introduction

The development of tools to control biomacromolecular function with light offers the potential to change the properties of defined molecules in biological systems with minimal perturbation to the rest of the system. This article presents highlights of current developments alongside their historical context to complement recent reviews of this field that have covered molecular switches,^{1,2} light-responsive materials,^{3,4} optogenetic tools⁵⁻⁷ and biological control.⁸

Principles of Photocontrol

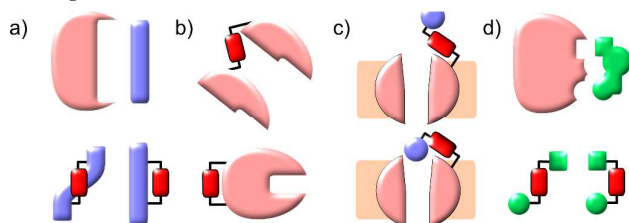


Figure 1. Photochromic molecules can influence biomolecules by stabilising or destabilising their interacting conformations. This approach can be applied to a) peptides corresponding to elements of protein-protein interaction partners, b) loops or domain interfaces of proteins themselves, c) tethered ligands to control ion channels or d) incorporated into existing non-photochromic drugs.

Protein-protein interaction networks allow cells to integrate chemical signals about their state and environment and respond by regulating protein production, enzyme activities and metabolism to suit their circumstances. Photocontrol of peptides, proteins or drugs provides a non-invasive way to perturb these networks to investigate their effect or cause a defined outcome. Achieving photocontrol requires the addition to a photoresponsive element to a protein, peptide or

drug in such a way that its photochromism alters the properties of the conjugated molecule (Figure 1). Some photochromic molecules respond to light by making or breaking bonds² leading to changes in their geometry and polarity, but the increased degrees of freedom of one of the states can lessen conformational control. In contrast, molecules capable of acting as photoswitches that interconvert between *E* and *Z* double bonds offer highly predictable light-state structural properties. In such molecules isomerisation is induced by absorption of a photon to generate an electronically excited state, in which the barrier to rotation between the *E* and *Z* forms is reduced. Decay to the electronic ground state re-establishes the barrier, kinetically trapping the thermally less stable isomer. The barrier height relative to the available thermal energy then determines the rate of relaxation to the thermodynamically more stable isomer. Although stilbenes,⁹⁻¹³ hemithioindigos,^{10,12,14-23} thioxopeptides,²⁴⁻²⁸ selenoxopeptides,²⁹ diarylethenes³⁰⁻³² and acylhydrazones³³ and rhodopsin-like molecules^{30,34} have been used as switches, azobenzenes^{35,36} currently offer the largest range of activation wavelengths and photoactivated state stabilities. In azobenzene-derived photoswitches rotation around the central double bond is transmitted with high fidelity through the rigid rings resulting in a predictable distance changes, particularly between substituents on the *para* positions. Azobenzenes are readily synthesised either non-symmetrically by the condensation of a nitroso-aryl compound and anilines or symmetrically by *in situ* nitroso formation under oxidising conditions. Classical aromatic chemistry allows access to various photo-switching wavelengths and thermal reversion characteristics tailored to suit a particular application (Figure 2). The choice of azobenzene derivative is dependent on its spectral properties, the ratio of isomers at its photostationary state, its solubility, its resistance to reduction and the distance between the intended points of control.

^a School of Chemistry & Cardiff Catalysis Institute, Cardiff University, Park Place, Cardiff, UK.

Azobenzene photoswitches

Improving photophysical properties

Photoswitches can be conveniently linked to proteins by selective reactions with the thiol side chains of cysteine residues. Surface accessible cysteine side chains react readily with eg. *para*-haloacetamide substituted azobenzenes such as iodoacetamidoazobenzene (**1**) (Figure 3).³⁷ Early work demonstrated that with pairs of cysteines in *i,i+4* or *i,i+7* spacings along a peptide chain, an α -helical conformation is stabilised by the *Z*-isomer of **1** while the *E*-isomer is α -helix stabilising for *i,i+11* spacings.³⁸ The water-soluble azobenzene **2** expanded the scope of this technique and allowed protein-protein, protein-DNA and protein-RNA interactions to be controlled through α -helical motifs.^{39, 40} While the excitation wavelengths in the UV-range limited the biological applications of these early azobenzene photoswitches, new photochromic molecules that absorb at higher wavelengths have recently been developed and can be actuated with non-toxic visible light.⁴¹

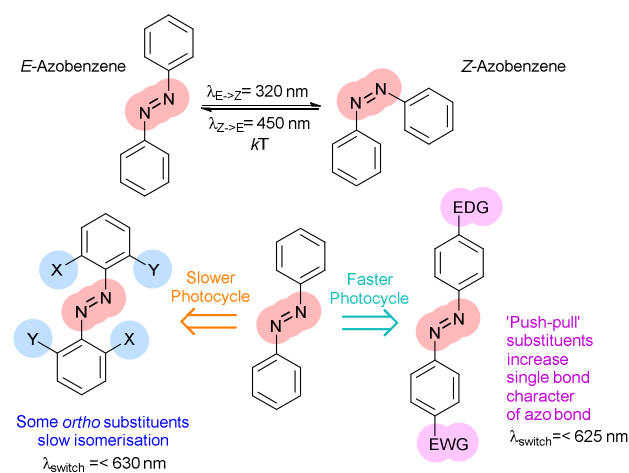


Figure 2. Substituents can alter the photophysical properties of azobenzenes with wavelength ranges between 300 and 630 nm and drive isomerisation and provide the means to tune the half-life of the thermodynamically less stable *Z*-isomer.

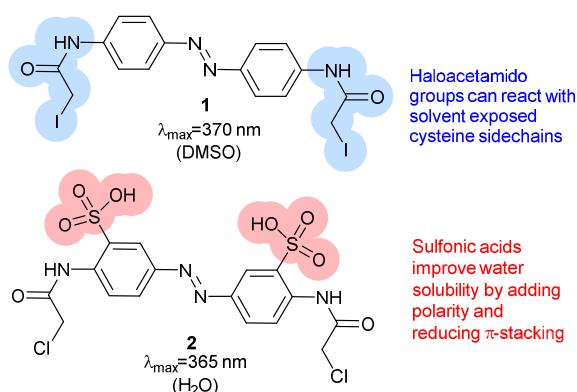


Figure 3. Bis-*para*-(iodoacetamido)azobenzene (**1**) reacts with the sidechains of a pair of cysteines of a peptide to form a photo-responsive 'staple'. The addition of sulfonic acid groups render **2** water-soluble.

Introduction of *ortho*-amino groups yielded azobenzenes (**3-8**) with a wide range of photochemical properties.⁴² Most have significantly red-shifted absorbance spectra (445-470 nm), but protonation of the amines produces forms that diminish the double bond character of the azo bond and lead to shorter relaxation half-lives of between 7 seconds and 5 minutes compared to ~30 minutes for **2**. A red-shifted π - π^* absorbance maximum was also observed with a pair of *ortho*-methoxy substituents (**9**), but the introduction of four methoxy-groups *ortho* to the azo-bond (**10**) produced different properties.⁴³ The absorbance spectrum of *Z*-**10** reveals a 36 nm shift of its n - π^* band compared to *E*-**10** (Figure 6). This separation of the n - π^* bands permits green light irradiation (530 to 560 nm) to convert *E*-**10** to ~80% *Z*-**10**, eliminating the need for damaging UV light altogether. Furthermore, the relaxation half-life of *Z*-**10** is 12 hours in buffer at 40 °C, making it ideal for biological applications. However, the electron donating effect makes **10** vulnerable to reduction with a half-life of 90 min in glutathione-containing buffers that reproduce the cytosolic reduction potential.⁴³ Replacing the methoxy groups with methylthioethers in **11** retained the separation between the n - π^* transitions of the isomers, which were red shifted and strengthened compared to **10**, but model bioconjugates of **11** had short half lives of the light activated state (3.3 minutes at 37 °C).⁴⁴ In contrast, the extended analogue *Z*-**12** had a thermal reversion half-life of several hours in water at 37 °C.⁴⁵

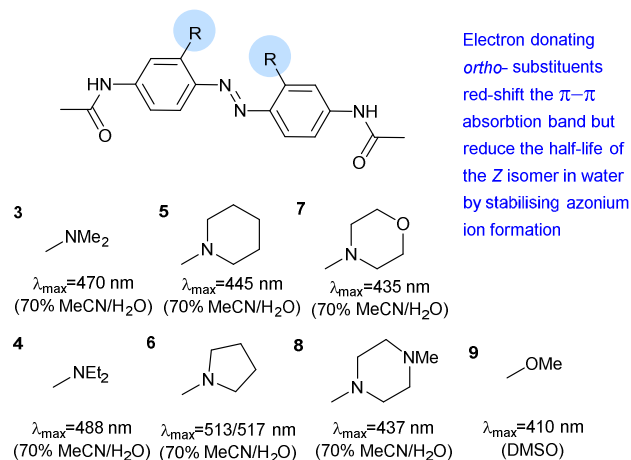


Figure 4. Amino-substituted azobenzenes with red-shifted π - π^* transitions and rapid relaxation.

Given the destabilising effect of amino (rather than amide) substituents of **3-8** on their *Z* isomers, it is understandable that installing piperidine as a more flexible spacer in the *para* position resulted in *Z*-**13** having a half life of less than 4 seconds in buffer at 25 °C.⁴⁶

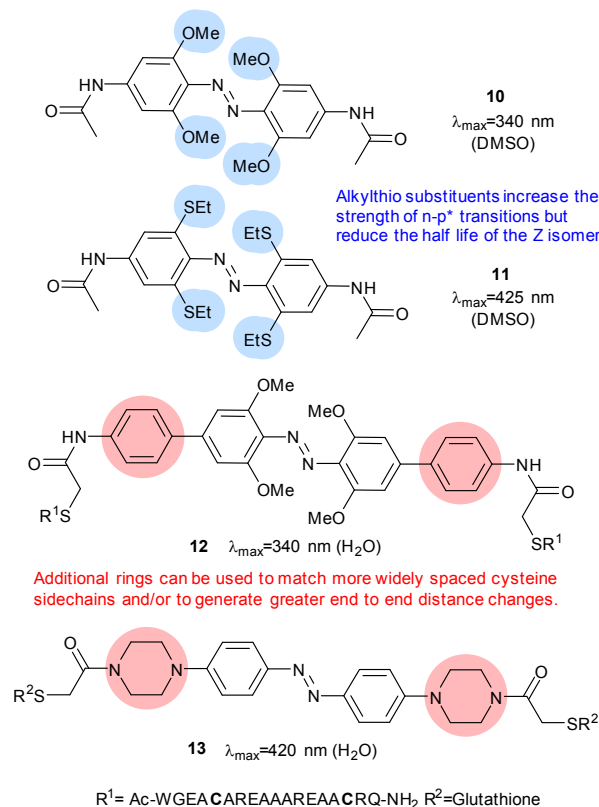


Figure 5. Tetra-*ortho* ether or thioether substitution has the beneficial effect of shifting the n- π^* transition bands so that at certain wavelengths the absorption spectra of the *E*- and *Z*-isomers do not overlap, allowing visible light switching. Extended azobenzenes may either be used with more widely spaced cysteine sidechains or to increase the change in end-to-end distance to provide greater photocontrol.

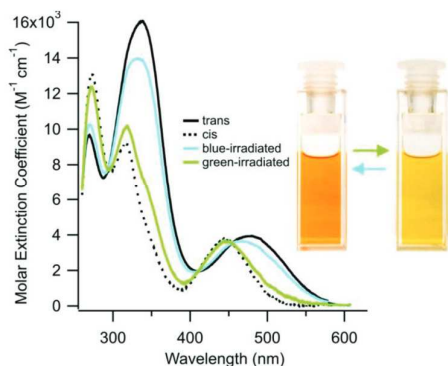


Figure 6. UV/visible spectra of **10** in dimethylsulfoxide show the separation of the n- π^* bands at 450 and 500 nm. The separation is less pronounced in water, but bidirectional visible light switching is still possible. Reprinted with permission from Beharry *et al.* *J. Am. Chem. Soc.* 2011, **133**, 19684. Copyright 2011 American Chemical Society.

However, when this motif was combined with tetra-*ortho*-methoxy substitution, the resulting molecule (**14**) displayed unexpected properties.⁴⁷ With methoxy and amino substituents in place the pK_a of the protonated form of **14** rises to the physiologically accessible value of 7.2 and the absorption spectrum of the azonium species is broad enough for irradiation with 630 nm light to cause isomerisation. This

effect was ascribed to a difference in pK_a values between the protonated states of *E*-**14** and *Z*-**14**, with the azonium species stabilised by a hydrogen bond in *E*-**14** but not in *Z*-**14**.

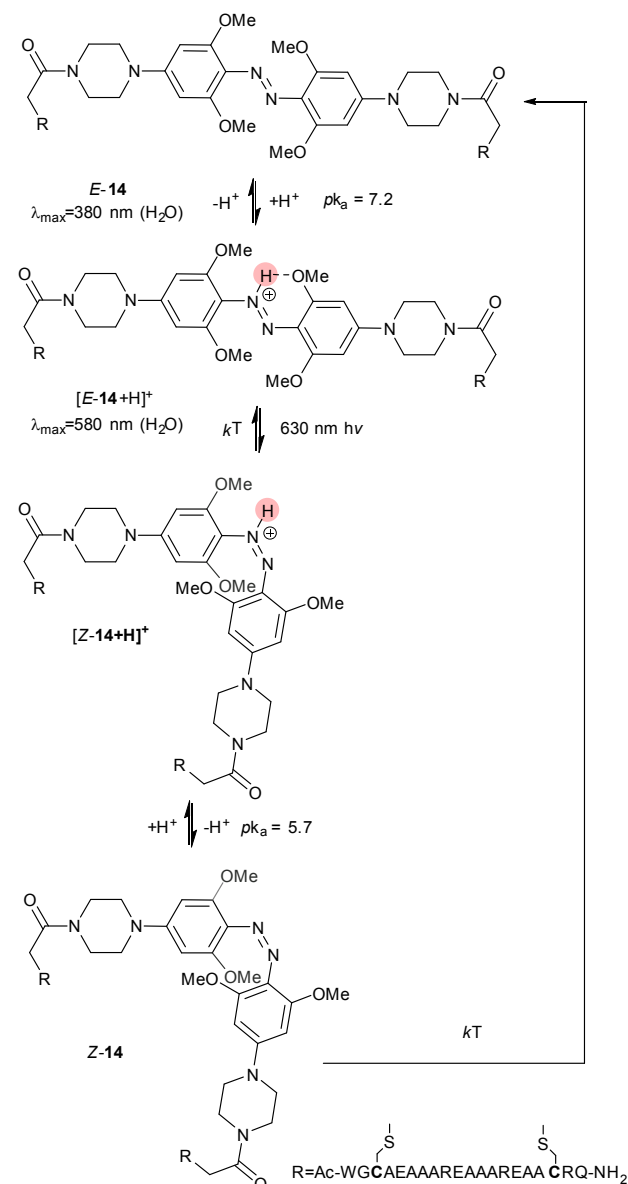


Figure 7. The combination of tetra-*ortho* methoxy ether and *para* amino substitution increases the pK_a of the diazo bond, so that azonium species are formed at neutral pH. However, the pK_a of *E*-**14** is approximately 1.5 units lower, resulting in deprotonation and slower relaxation to *Z*-**14** than would be expected from the azonium ion.

Therefore, at pH 7, **14** is protonated, photoswitches and then deprotonates. Whilst the half-life of *Z*-**14** is only 13 seconds, it can reliably and repeatedly be photoswitched by red light in the presence of whole blood, although it is sensitive to reduction by glutathione.⁴⁷ A second generation of azonium ion precursors (Fig. 6) were designed to determine the optimal methoxy substitution pattern to donate electrons to the ring without sterically hindering azonium ion formation.⁴⁸ These efforts resulted in **15**, which absorbs significantly at 650 nm.

However, the *meta* methoxy group increases steric hindrance and forces the piperazine ring to rotate into a conformation that is less able to stabilise azonium ion formation, reducing the apparent *pK*_a to 2.6. This effect is alleviated in **16** by the less bulky bicyclic pyrrolidine substituent and the apparent *pK*_a is increased to 5.4. The absorption spectrum of **16** stretches into the near IR region with significant absorbance up to 750 nm.

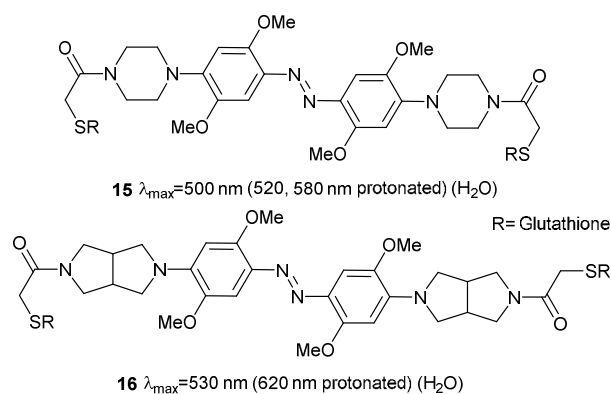


Figure 8. Moving one methoxy group to the *meta* position maintains the electron rich nature of the azobenzene ring system, but removes steric congestion from the azonium ion formed on protonation.

Electron rich azobenzenes can support protonation prior to or concomitant with nucleophilic attack by thiols and are vulnerable to reduction by glutathione. In contrast, tetra-*ortho*-halogen substituents (**17**) including tetra-*ortho*-fluoro,⁴⁹ chloro and bromo variants⁴⁵ are all resistant to glutathione reduction. Tetra-*ortho*-fluorodi-*para*-bromo azobenzene **18** has a clear (42 nm) separation between the *n*- π^* transitions of its *E* and *Z* isomers, which allows visible light switching.⁴⁹ A mildly electron releasing group in the *para* position (**17a**) reduced the *n*- π^* band separation to 22 nm compared to **18**, whilst a more electron withdrawing ester substituent (**19**) increased it to 50 nm. Detailed studies of fluorinated azobenzene derivatives with symmetrical or non-symmetrical rings and various electron withdrawing groups in the *para*-position including fluorine, esters, nitriles and amides were carried out. Symmetrical (**20**) or non-symmetrical (**21**) difluoro azobenzenes exhibited a smaller (28-32 nm) *n*- π^* band separation than **18** (42 nm), but the presence of further additional fluorine atoms did not result in increased separation in hexa-, octa- or deca-fluorinated azobenzenes (36-43 nm).⁵⁰ Introduction of a symmetric pair of more electron withdrawing trifluoromethyl groups in the *ortho* positions decreased the *n*- π^* band separation, an effect attributed to the increased steric demand of this substituent resulting in the *E*-isomers phenyl rings not being coplanar. This reduction in conjugation in the ground state is likely to reduce the difference in energy and hence the *n*- π^* band separation between the *E*- and *Z*-isomers. Time dependant density functional theory calculations qualitatively reproduced the spectral properties for this series of compounds and further modelling suggested that the extended half-lives of their *Z* isomers are a consequence of the

dipole moments in the transition states.⁵⁰ Tetra-*ortho*-chloro azobenzene **17b** had a similar *n*- π^* band separation (20 nm) to **17a** (22 nm) and was shown to photoswitch under irradiation by a 635 nm LED.⁴⁵ Rapid reversion to *E*-**17b** was conveniently induced by irradiation with 450 nm light and neither isomer of **17b** was degraded by exposure to 10 mM reduced glutathione. When the chloroacetamido analogue of **17b** was attached to a test peptide, a thermal reversion half-life of 3.5 hours was observed at 37 °C. Subsequently, **17b** was conjugated to a *D*-peptide labelled with fluorescein and microinjected into zebrafish embryos.⁴⁵ Photoswitch-induced quenching of the fluorophore⁵¹ was possible for 30 hours with no obvious ill-effects (Figure 11).⁴⁵

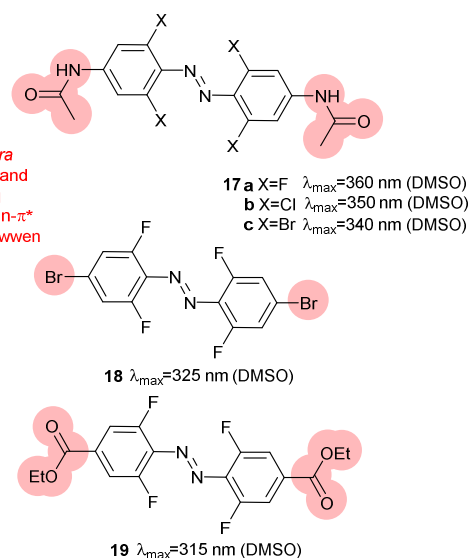


Figure 9. Tetra-*ortho* halogen substituents separate the *n*- π^* transition energies of *E* and *Z* isomers, impart resistance to glutathione reduction and lead to long lived *Z*-isomers.

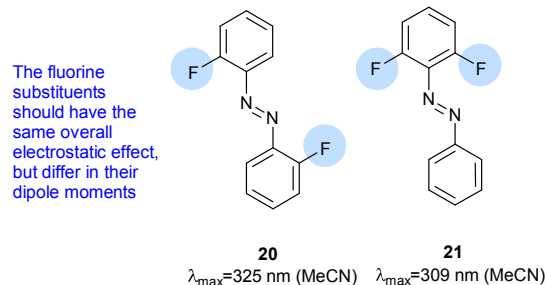


Figure 10. Symmetrical and non-symmetrical difluoroazobenzenes used to examine the origins of the *n*- π^* band shifts and extended *Z*-isomer half-lives.

This system represents the current state of the art for a slow cycling switch in that it is stable against reductive and metabolic reduction and offers the capability of bi-directional, visible-light switching *in vivo*. Further developments in slow-cycling photo-switches have been driven by potential uses for molecules that can respond to infrared activation to which the human body is mostly transparent. Such switches could be

activated at defined points in 3D space using multiphoton techniques.

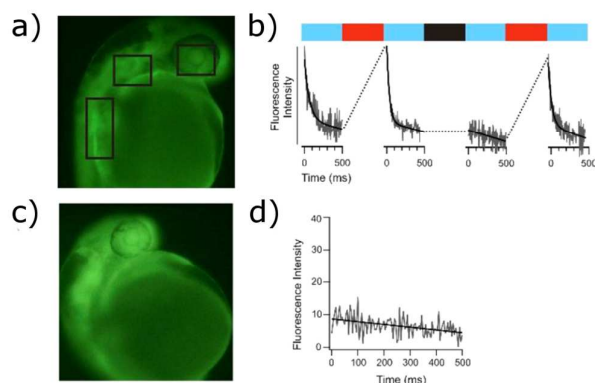


Figure 11. a) Fluorescence image of a zebrafish embryo (early pharyngula stage/ \sim 30 hours post fertilisation) containing the fluorescent reporter peptide cross-linked with **4**. b) Measurements of fluorescence intensity at the zones indicated by black boxes in a), as a function of time during blue light irradiation. A change in fluorescence emission was seen only if the peptide was first switched with red light (indicated by red bars). No pre-irradiation with red light (indicated by black bar) before blue light produces no exponential (switching) transient, only a background slope due to fluorescein reporter photobleaching. c) Fluorescence image of a zebrafish embryo (\sim 30 hours post fertilisation) containing the fluorescent reporter peptide without a photoswitch. d) Measurement of fluorescence intensity of the nonswitchable peptide as a function of time during blue-light irradiation after preirradiation with red light as in (b). Adapted with permission from Samanta *et al. J. Am. Chem. Soc.* 2013, **135**, 9777. Copyright 2013 American Chemical Society.

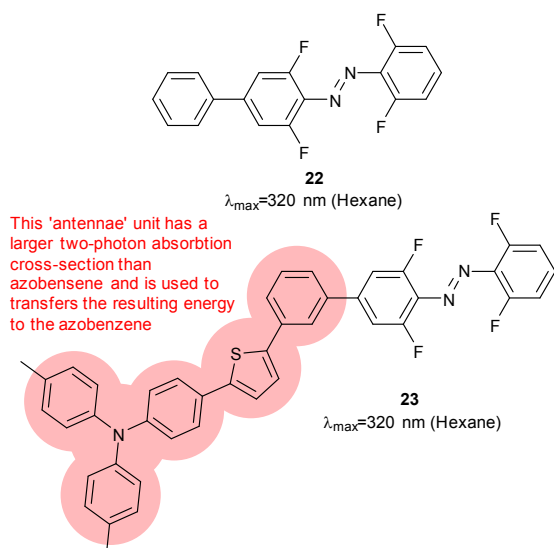


Figure 12. Tetra-*ortho*-fluoro substitution increases the two photon absorption cross section compared to azobenzene itself. A two photon antennae complex is excited, then transfers energy to the azobenzene.

Two-photon photoswitching of **2** had previously been reported to be effective in blood lysates⁵² but not for tetra-*ortho*-methoxy dyes.⁴⁵ However, a slightly modified tetra-*ortho*-fluoro azobenzene (**22**) was found to have a two-photon cross-section seven times greater than azobenzene alone; two-photon excitation of the π - π^* band was possible at 640 or 720

nm.⁵³ Installation of a larger aromatic system with extended conjugation to act as an antennae complex yielded **23**, which can be converted to *Z*-**23** by 510 nm light and back to *E*-**23** by two-photon absorption at 750 nm.⁵⁴

Another series of photoswitches employs different types of bridged azobenzenes (Fig. 10).⁵⁵⁻⁵⁸ The bridge has the effect of reversing the normal thermodynamic preference for the *E*-isomer so that **24** adopts almost exclusively the *Z*-form in its thermal equilibrium state. **24** is converted to $>90\%$ of the *E*-form on irradiation with 370-400 nm light. Reversion to *Z*-**24** can be achieved with irradiation with 480-550 nm light. When a chloroacetimido analogue of **25** was attached to a test peptide, a photostationary state of 70% *E*-**25** was formed upon irradiation with 407 nm light.⁵⁹ Thermal relaxation from the *E* to the *Z* form occurred with a half-life of 8.3 hours at 20 °C, but reversion could also be driven with 518 nm light irradiation. Whilst *Z*-**25** was completely resistant to glutathione reduction, *E*-**25** degraded with a half-life of approximately 3 hours. A different ring substitution position resulted in **26**, whose n - π^* transitions were separated by almost 90 nm (*Z*-**26**: 404 nm, *E*-**26**: 490 nm) although only 53% *E*-**26** was present in the photostationary state.⁶⁰ Other examples of constrained azobenzenes are found in structures containing a borodifluoride bridge (**27**). These molecules are strongly red shifted (570 nm) compared to conventional azobenzenes.⁶¹ Electron donating substituents can be used to push red absorbing **28** (630 nm) further towards the IR range in **29** (710 nm). An additional methoxy substituent *ortho* to the azo substituent in **30** caused a blue-shift (640 nm) because its steric demands twisted the phenyl ring out of the plane of the azo bond, thereby reducing its electronic influence.⁶²

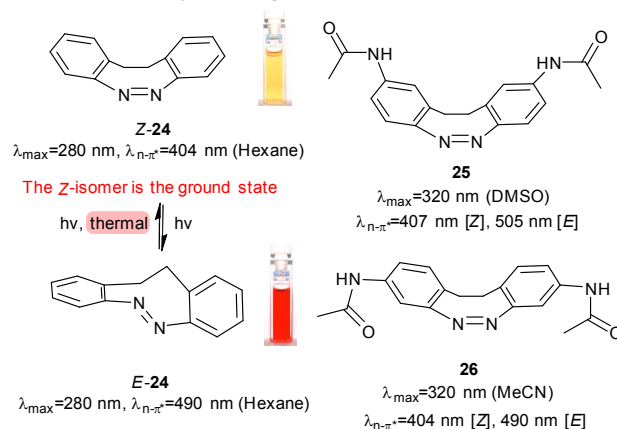


Figure 13. The conformational constraints applied to bridged azobenzenes make their *Z*-isomers their thermodynamic ground states. Many bridged azobenzenes have very large separations between the n - π^* bands of the *E* and *Z* isomers. Elements adapted with permission from Siewertsen *et al. J. Am. Chem. Soc.* 2009, **131**, 15594. Copyright 2009 American Chemical Society.

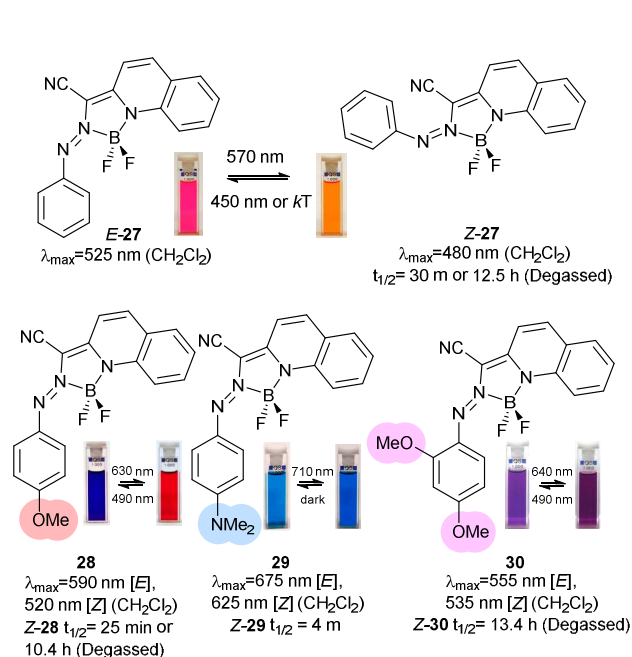


Figure 14. Borodifluoride bridged azobenzenes are strongly red shifted and the half-lives of the Z isomers are strongly affected by dissolved oxygen. Elements adapted with permission from Yang *et al. J. Am. Chem. Soc.* 2012, **134**, 15221 (Copyright 2012 American Chemical Society) and Yang *et al. J. Am. Chem. Soc.* 2014, **136**, 13190.

New Incorporation and Conjugation Approaches

Thiol-ene reactions are gaining popularity for bioorthogonal functionalisation of peptides and proteins.⁶³ The vinyl azobenzene amino acid **31** was incorporated into peptides by conventional solid phase synthesis and, when an *i*+4 cysteine residue and an appropriate photoinitiator were present, irradiation with 360 nm light caused isomerisation and a thiol-ene click reaction between the cysteine sulfur and the pendant vinyl group.⁶⁴ Consequent oxidation resulted in intramolecular crosslinking via a sulfone. Genetic code expansion can be used to introduce **31** into proteins expressed in *E. coli* or mammalian cells.⁶⁵ Azobenzene **32** can also be incorporated into expressed proteins; its pentafluoro ring is electrophilic enough to undergo nucleophilic aromatic substitution by a proximal cysteine.⁶⁶ The fluorine atoms of **32** provide sufficient $n-\pi^*$ band separation for bidirectional switching with 540 nm and 470 nm light to generate ~70% of the desired isomer in the photostationary state. This switch was incorporated into calmodulin with an orthogonal tRNA synthetase. Whilst a degree of side reaction with glutathione was noted, cyclisation proceeded smoothly to give a visible light responsive calmodulin with distinct binding affinities for neuronal nitric oxide synthase under green and blue light irradiation. Thiol-ene chemistry was also investigated to introduce a bifunctional glycosylated azobenzene photoswitch.⁶⁷ Using 2,2-dimethoxy-2-phenylacetophenone as a photoinitiator caused **33** to react with glutathione in 70% yield.

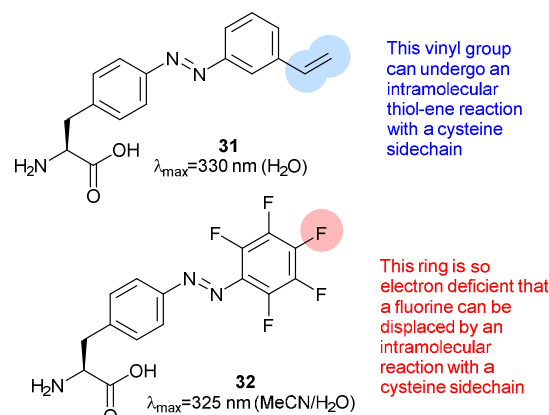


Figure 15. A vinyl azobenzene (**31**) capable of intramolecular thiol-ene cyclisation with an appropriately placed thiol-ene sidechain and a pentafluoroazobenzene (**32**) whose *para* fluoro group can be displaced by a thiolate anion.

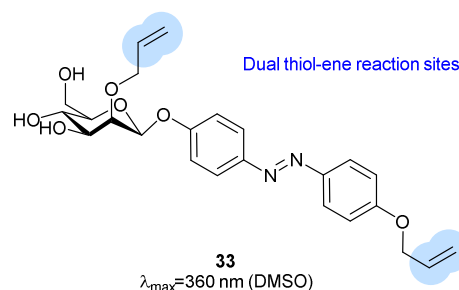


Figure 16. Glycoazobenzene derivative for conjugation thiol-ene reactions.

Staudinger ligations can be used to selectively functionalise synthetic biomolecules containing azide groups incorporated for *in situ* bioorthogonal labelling. The diphenylphosphoryl azobenzene **34** performs a variation of the Staudinger ligation, intramolecularly capturing the phosphazene intermediate formed by reaction of the phosphine with an azide-containing peptide. The resulting conjugate has a $\pi-\pi^*$ absorption maximum at 500 nm, but the reaction is relatively slow and the Z-isomer is short lived (0.5 ms).⁶⁸

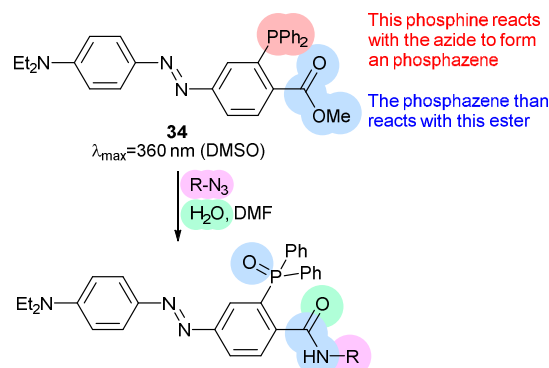


Figure 17. Diphenylphosphoryl azobenzene **34** can be used to introduce a pendant 'push-pull' azobenzene into peptides by Staudinger ligation chemistry.

In vitro translation and expanded genetic code techniques have been used to create a library of peptides containing the azobenzyl-lysine **35**.⁶⁹ By suppressing the stop codon and conducting the reaction at low temperature, neither the nascent peptide nor the coding mRNA dissociate from the ribosome. Streptavidin-coated beads were placed in the solution in the dark and the beads were washed and exposed to UV light, which resulted in dissociation of the peptide-ribosome-mRNA complex. The complexes were recovered, the mRNA dissociated and subjected to amplification by reverse transcriptase polymerase chain reaction; this identified peptide sequences that bound streptavidin 2.5-fold weaker in their photostationary state than in the dark. Libraries of peptides have also been generated using phage display.⁷⁰ Peptides containing *i,i+7* spaced cysteines were expressed in *E. coli*, reduced with resin-bound tris-carboxyethylphosphine and treated with **2**. Although the overall yield of modification with **2** was only 50%, peptides were identified with up to 22-fold decreases in affinity for streptavidin in their pure dark and light states. A later study also used *i,i+7* spaced cysteine residues alkylated with **2** and discovered peptides that had a greater affinity in for streptavidin in the light state with the best example found to bind 3-fold more tightly.⁷¹

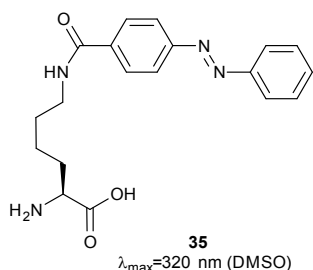


Figure 18. Azobenzyl-lysine, a readily synthesised amino acid with a pendant azobenzene in its sidechain.

Selected Applications

Control of Nucleic Acid Binding Peptides

Transcription factors act as 'master switches' in cells for the expression of genes in response to environmental cues. As such they are key targets to modulate downstream processes. Many common DNA binding motifs contain α -helices, which make direct and indirect contacts with the bases in the major groove of DNA. Such helices are often unstructured in the absence of DNA or when isolated from the rest of the protein, but are stabilised through interaction with other structural elements. Recognition helices in homeodomains, for instance, rely on a helix-turn-helix motif for stability (Figure 19). The DNA recognition helix from the engrailed homeodomain has been modified to incorporate two *i,i+11* spaced cysteine residues to accommodate **2**.⁷² When the peptide is folded into an α -helix, the intra-cysteine spacing matches the length of **E-2** so that the cross-linked peptide is strongly helical and binds to target DNA 25 times more strongly than a control peptide without **2**. Irradiation to form **Z-2** diminished the DNA binding affinity 18-fold.

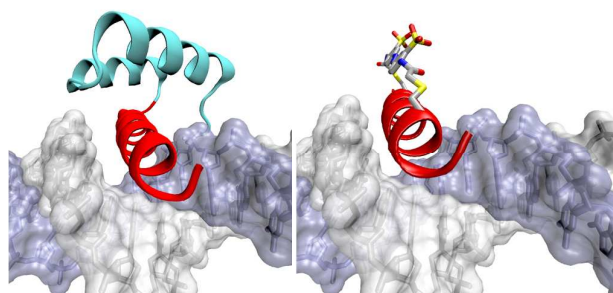


Figure 19. Cartoon of the engrailed homeodomain DNA binding helix in the major groove of its target DNA and its N-terminal extension (red), which contacts the minor groove of the DNA (left) with two short α -helices (cyan) that stabilise it. The stabilising role of these α -helices can be replaced by introducing a pair of cysteine residues whose sidechains project opposite the DNA binding face to react with **2**.

Basic helix-loop-helix proteins such as the transcription factor MyoD bind to DNA as homodimers (Figure 20). The DNA-binding region is unstructured in the absence of DNA but adopts an α -helical conformation when it contacts the DNA. The recognition helix was modified to include *i,i+7* spaced cysteine residues and reacted with **2**. The photo-activated state of the cross-linked protein not only bound DNA with an apparent dissociation constant for the complex three orders of magnitude higher than for the uncrosslinked protein, equivalent to a 60-fold decrease in the half-maximal binding concentration, but also displayed a significant increase in the specificity for its cellular target DNA.⁷³

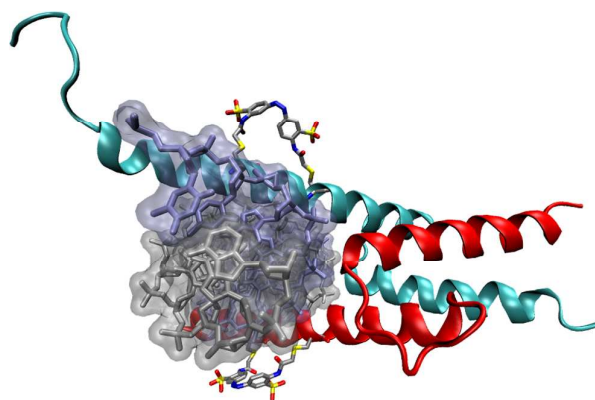


Figure 20. Cartoon illustrating the use of **2** to control the conformation of the DNA recognition α -helix of MyoD.

bZIP proteins form a different family of transcription factors that rely on α -helical structures. They contain discrete DNA-binding and leucine zipper regions. DNA binding requires dimerization through the formation of a coiled coil (Figure 21) leaving the DNA-binding region unchanged. *i,i+7* spaced cysteines were used to create a GCN4-bZIP peptide, which reacted with **2** to give a photoswitchable DNA-binding

peptide with increased α -helicity and a 20-fold decrease in dissociation constant towards a DNA target after irradiation.⁷⁴

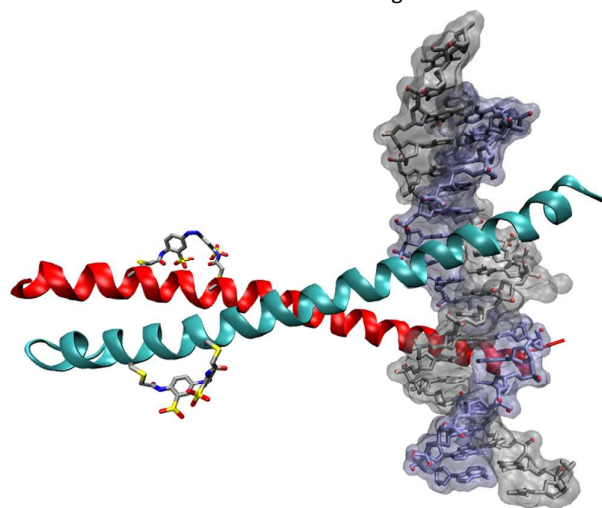


Figure 21. Cartoon of a crosslinked bZIP DNA complex. The crosslinker **2** controls the α -helicity of the coiled-coil dimerization domains, which in turn controls the DNA binding avidity of the complex.

Similar methodology was also effective in controlling of the cyclic AMP response element-binding protein (CREB).⁷⁵ Photocontrol of transcription in live cells was achieved by a modified bZIP helix with $i,i+7$ spaced cysteines and a pair of additional acidic heptad repeats at its *N*-terminus.⁷⁶ Once crosslinked with **2** and introduced into HEK293T cells alongside reporter DNA this molecule (0.1 to 1.2 μ M) was able to alter the expression level 2-fold upon irradiation (Figure 22).

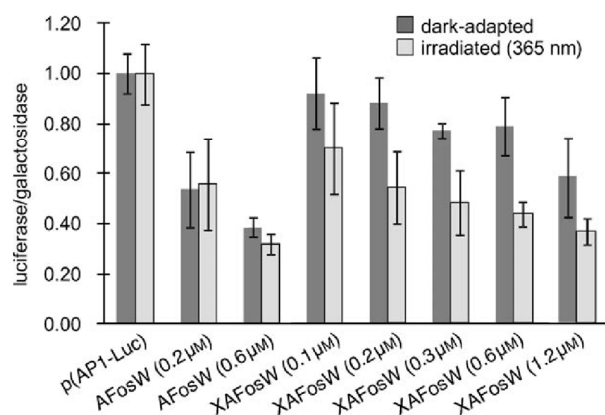


Figure 22. 293T cells were transfected with luciferase reporter plasmid p(AP1-Luc) and β -galactosidase plasmid pRSV-Gal. The ratio of the activities of the two enzymes was used to detect specific effects on AP1 promoter driven expression. Addition of non-crosslinked AFosW had a dose dependent effect as AFosW displaced native Fos from the Fos/Jun/AP-1 DNA complex. Addition of crosslinked XAFosW has less of an effect as the peptide is biased away from its active conformation by the crosslinker, UV irradiation removes this constraint and yields a 2-fold light-dependant inhibition of protein synthesis. Reproduced with permission from Zheng *et al.*, *Angew. Chem. Int. Ed.*, 2010, **49**, 3943. Copyright 2010 Wiley-VCH Verlag GmbH & Co. KGaA.

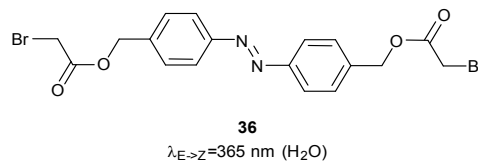


Figure 23. A bromoacetate ester azobenzene used to control the relative orientation of two basic DNA-binding helices.

In a second approach to controlling the bZIP family, an azobenzene linker (**36**) was bound to two DNA-binding helices by forming thioethers with cysteine residue side chains.⁷⁷ When the azobenzene was switched into the Z-conformation these helices aligned scissor-like into the major groove of a DNA strand and bound DNA 60-70 times more strongly than in the dark state. Single stranded RNA presents a greater variety of binding geometries than double stranded DNA because of its propensity to adopt different knot and loop structures. Nevertheless, good photocontrol of RNA binding was achieved using a peptide derived from the HIV1 Rev response element (RRE)⁷⁸ crosslinked with **1** for both $i,i+7$ and $i,i+11$ spacings.

Control of Protein-Protein Interactions

Although many examples of controlling the structure of β -hairpins⁷⁹⁻⁸⁶ and cyclic peptides⁸⁷⁻⁹⁴ are known, light is most frequently used to control protein-protein interactions by influencing the conformation of tethering loops or α -helices or by influencing the interactions of subunits (Figure 24).

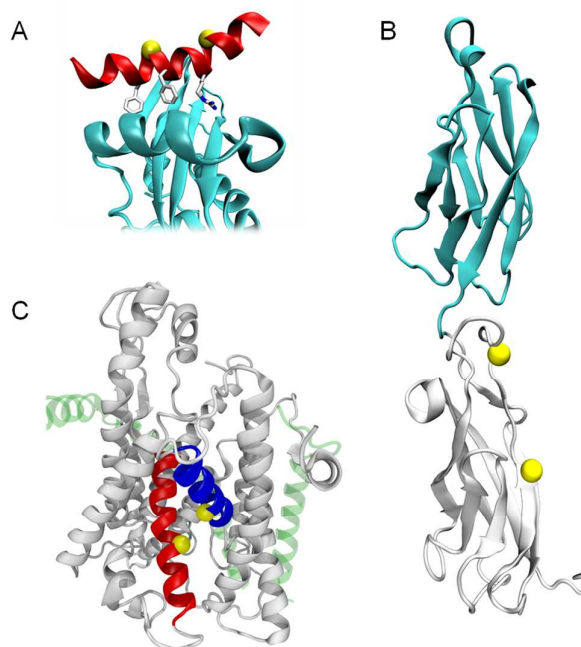


Figure 24. Cartoons illustrating the positions changed to cysteines to accept azobenzenes in A) AP2 β -appendage (PDB 2IV8). B) Loop regions of cadherin (PDB 1FF5). C) Trans-membrane helices 2 (blue) and 7 (red) of the SecYEG complex (PDB 1HRZ).

A photoswitchable peptide was designed so that its α -helical conformation bound to the adaptor protein AP2 (Figure 24A).^{95, 96} This protein is important for the generation of clathrin-coated pits for internalisation into cells. Peptides with $i,i+7$ and $i,i+11$ spaced cysteine residues, whose side chains were alkylated with **2**, were able to enter cells and allowed photocontrol of fluorescent transferrin uptake. The $i,i+7$ peptide was the most effective with a 3-fold decrease in intracellular fluorescence upon irradiation (Figure 25). Peptide concentrations from 20 μ M to 100 μ M were employed depending on the cell type.

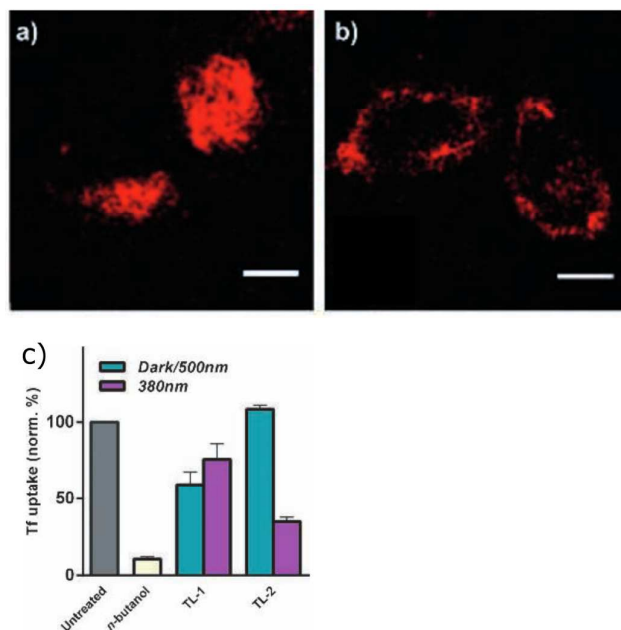


Figure 25. HeLa cells incubated with Alexa647 labelled transferrin and acid-washed show internalisation in (a) the absence of peptide, but (b) remain surface bound when the cells are treated with TL-1, an inhibitor peptide with $i,i+11$ spaced cysteines crosslinked with **2**. (c) Irradiation of TL-1 with 380 nm light decreases uptake inhibition, but the degree of photocontrol is significantly greater for TL-2 which has $i,i+7$ spaced cysteines crosslinked by **2**. The different cysteine spacing reverses the direction of the photocontrol with respect to TL-1. Reproduced with permission from Nevola *et al.*, *Angew. Chem. Int. Ed.*, 2013, **52**, 7704. Copyright 2013 Wiley-VCH Verlag GmbH & Co. KGaA.

A light-responsive cadherin molecule was created by introducing a pair of cysteine residues in the EC1 and EC2 domains (Figure 24B) and crosslinking them with **2**.⁹⁷ The resulting protein had different affinities for calcium ions under 365 nm or white light illumination and the half-life of the light state was strongly affected by the calcium ion concentrations. Experiments showed a light-dependent weakening of interactions between surface bound cadherin and irradiated crosslinked cadherins that indicated a degree of control over dimerisation. The B-cell leukemia-2 (Bcl-2) family of proteins, a vital part of the intrinsic apoptosis pathway, interact through conserved helical motifs. Photoswitchable peptides were designed based on the interacting helices of Bak and Bid.⁹⁸ These peptides were cross-linked with **2** and their affinity to

Bcl- x_L was increased by 23-fold ($i,i+4$), 20-fold ($i,i+7$) and decreased by 7-fold ($i,i+11$) upon irradiation. An NMR structure of Bak $i,i+11$ in complex with Bcl- x_L showed that the photoswitchable peptide bound to the canonical BH3 binding pocket.^{99, 100} When introduced to permeabilised SU-DHL-4 cancer cells, the Bak $i,i+7$ peptide (7.5 μ M) proved capable of inducing light-dependent mitochondrial membrane depolarisation and cytochrome release.¹⁰¹ The SecYEG complex (Figure 24c) translocates substrates from the cytosol to the lipid bilayer.¹⁰² Introduction of cysteine residues and a short dibromoalkyl azobenzene crosslinker (**37**) facilitated control of the opening of a lateral gate from the translocation pore to the interior of the lipid membrane. Inner membrane vesicles containing this modified complex transported the proOmpA pre-protein in a reversible, light-responsive manner.

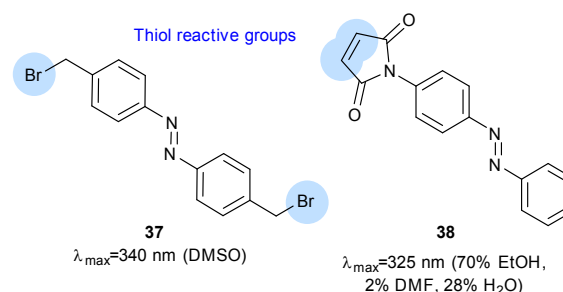


Figure 26. A bisbromoalkyl azobenzene (**37**) used to control the SecYEG complex of *E. coli* and a maleimidoazobenzene (**38**) used to modify calmodulin to insert a pendant photoswitchable group.

Calmodulin moderates many calcium-dependent cellular signalling events. Ten calmodulin mutants incorporating cysteines at positions that might interfere with calcium binding sites were constructed and then reacted with **38**.¹⁰³ The most successful allowed 4-fold photocontrol of its binding affinity for its binding partner presented in a fluorescent fusion-protein. Control of subunit interactions was demonstrated using restriction endonucleases. Bis-maleimidoazobenzene **39** was used to photo-regulate the restriction enzyme *PvuII* with a 16-fold change in endonuclease activity upon irradiation; this was achieved through regulation of the turnover number rather than a change of the Michaelis constant.^{104, 105}

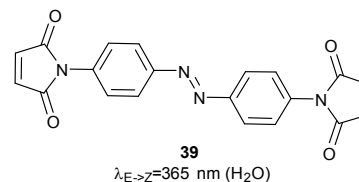


Figure 27. A bis-maleimide used to control homodimeric *PvuII* complexes and mixed P2X receptor complexes.

The DNA methyltransferase *SsoII* was also modified to include cysteine residues and reacted with **39**, which led to a 2-fold change of activity in response to light due to a 4-fold reduction in K_M .¹⁰⁶ The more subunits that are present in a complex, the greater the changes caused by irradiation; control of the relative orientations of the subunits of a group II chaperonin

using **39** resulted in structural changes that were large enough to be detected by cryo-electron microscopy (Figure 28).¹⁰⁷

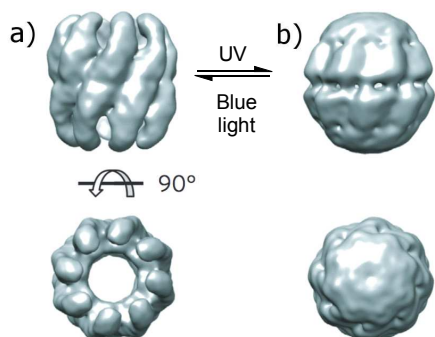


Figure 28. 15–19 Å single-particle cryo-electron microscopy reconstructions of an a homooligomeric chaparonin whose subunits are crosslinker with **39**. Adapted by permission from Macmillan Publishers Ltd: Nature Nanotechnology, 2013, **8**, 928, copyright 2013.

Control of Cellular Uptake

Photocontrol over the uptake of a peptide by cells was demonstrated by sandwiching azobenzene **40** between oligo-arginine and oligo-glutamate sequences.¹⁰⁸ *Z*-**40** allows the glutamates to mask the cationic polyarginine cell transduction sequence while *E*-**40** forces a conformational change that reveals the cell-penetrating sequence, resulting in a 10-fold increase in cellular uptake by HeLa cells incubated with the fluorescently labelled peptide (1 μM) (Figure 30).

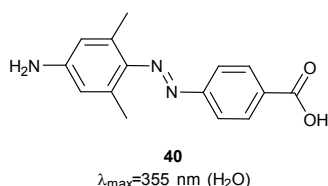


Figure 29. A sterically hindered push-pull azobenzene used to photocontrol cell uptake of a fluorescently labelled peptide.

The *para*-alkylthio azobenzene **41** was included between two nuclear localising signal peptides during solid phase synthesis.¹⁰⁹ The *para*-sulphur atom red-shifted the π - π^* transition to 377 nm, which is higher than the wavelength observed with an equivalent ether (360 nm) but lower than for a *para* secondary amine substituent (429 nm). A polyproline spacer and fluorophore were added and when the resulting molecule (120 nM) was incubated with HeLa cells the dark state was taken up more efficiently than after light activation.

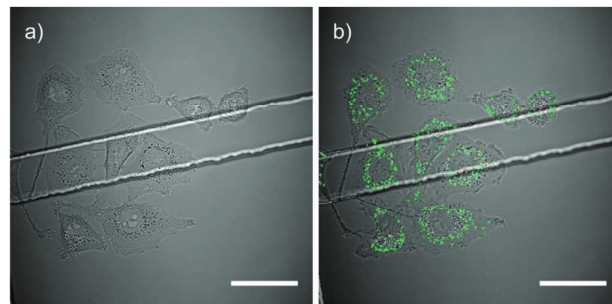


Figure 30. HeLa cells incubated with rhodamine B (Asp)₉-**40**-(Arg)₉ after a) UV light irradiation (1 μM) and b) blue light irradiation. Adapted under CC-BY 3.0 from Prestel and Möller, *Chem. Commun.*, 2016, **52**, 701.

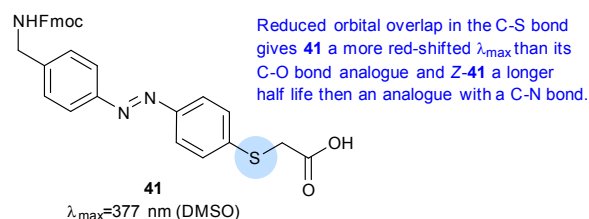


Figure 31. Azobenzene amino acid derived from thioacetic acid. The *para*-alkylthio substituent red-shifts the azobenzene absorbance to 377 nm.

Photopharmacology

Much work in photopharmacology is currently focused on creating molecules that can produce changes in cell behaviour.¹¹⁰ Such photopharmaceuticals allow increased activity at locations specified by irradiation to increase the effective therapeutic concentration range. For example, incretin glucagon-like peptide 1 can have beneficial effects on both the brain and the pancreas but can also cause numerous side effects elsewhere in the body. Introducing azobenzene **42** between two helical structural elements resulted in a peptide capable of controlling calm ion fluxes in pancreatic beta cells and moderating insulin secretion in response to light (Figure 33).¹¹¹ Despite the power of modifying natural systems for photocontrol by identifying critical elements and controlling the conformation of the corresponding peptides, the properties of peptides are not always ideal for their use as therapeutics. An increasing number of small molecule drugs have been modified to include azobenzenes to provide photoswitching functionality. Small molecule **43** targets the same glucagon-like peptide 1 pathway as the peptide modified with **42**¹¹¹ and yields similar levels of photocontrol of insulin secretion without a being linked to a peptide.¹¹²

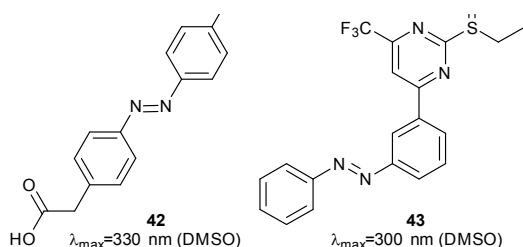


Figure 32. Azobenzene amino acid that when incorporated into the backbone of an incretin peptide could control β -turns formation and insulin secretion and a small molecule capable of the same feat without a peptide component.

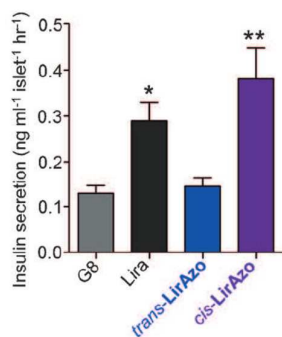


Figure 33. (a) Insulin secretion from pancreatic beta cells in response to glucose (8 mM, G8), a liraglutide-derived peptide (Lira) and the same peptide with **42** incorporated in its dark (*trans*-LirAzo) and UV light irradiated (*cis*-LirAzo) states. Adapted under CC-BY from Broichhagen *et al.*, *Angew. Chem. Int. Ed.*, 2015, **454**, 15565.

The stilbene element of combrestatin A4 (**44**), a cytotoxic experimental drug to treat advanced cancers, was replaced by azobenzene to yield **45**.^{113–115} While *E*-**45** is an extremely poor inhibitor of tubulin polymerisation (effective growth arrest at 25 μ M in a cell proliferation assay), *Z*-**45** (100 nM) is only tenfold less effective than **44** (10 nM) such that irradiation with 400 nm light causes effective photoswitching and a 200-fold change in cytotoxicity of the drug (Figure 35).¹¹⁴ **45** and similar photostatins can induce mitotic arrest and cell death in Jurkat, HEK, HeLa and MDA-MB-231 cells.¹¹³ ‘Mitotic catastrophe’ could be selectively induced on an individual cell level in an impressive display of non-selective drug delivery coupled with light-driven activation. The same strategy was successfully applied to the synthesis of a range of histone deacetylase inhibitors exemplified by **46**¹¹⁶ and **47**¹¹⁷ and a photoresponsive version of the antibiotic ciprofloxacin **48**.¹¹⁸

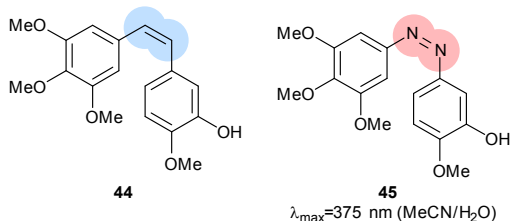


Figure 34. The antitumour natural product combrestatin A4 and an azobenzene containing analogue.

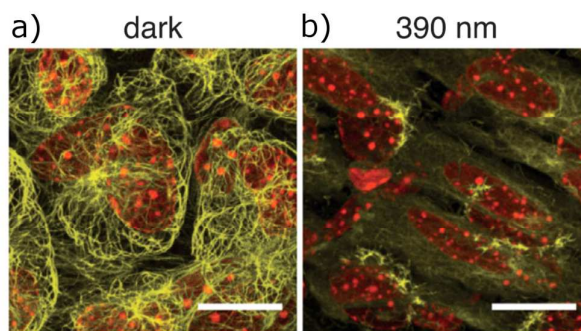


Figure 35. Cremaster muscle cells of live C57BL/6 mice superfused for 40 minutes with 50 μ M **45**-O-phosphate (a soluble prodrug) a) in the dark (left) or b) under 390 nm light irradiation. Samples were then excised and stained for α -tubulin (yellow) and nuclei (red). Scale bars 15 μ M. Adapted with permission from Borowiak *et al.*, *Cell*, 2015, **162**, 403. Copyright 2015 Cell Press.

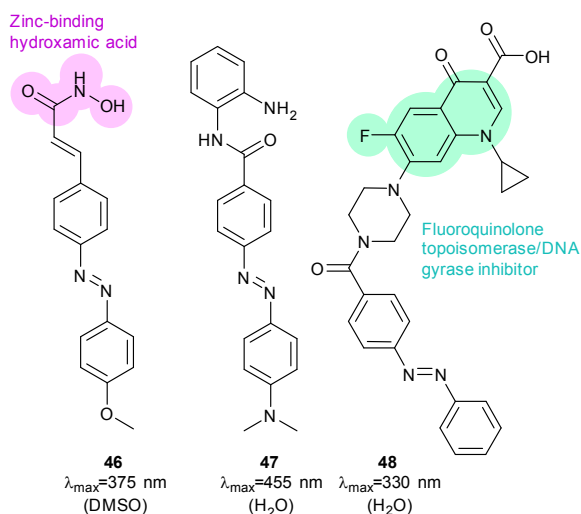


Figure 36. An azobenzene containing histone deacetylase inhibitor and a peripherally azobenzene-modified derivative of ciprofloxacin.

Azobenzene was also incorporated into acyl homoserine lactones to create a photoswitchable quorum sensing signalling molecule (**49**).¹¹⁹ When irradiated, **49** is five-fold more effective at activating the promoter of virulence genes in *P. aeruginosa*; its light-state has a half-life of 2.5 hours at 37 °C.

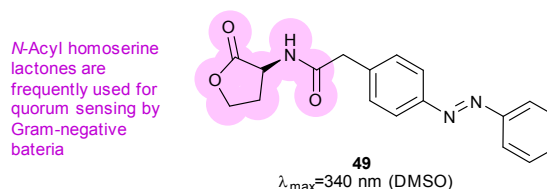


Figure 37. Azobenzene-containing molecule used to mimic naturally occurring homoserine lactone molecules used by bacteria for quorum sensing.

Control of Ion Channels

The application of photoswitchable ligands to control ion channels is a remarkable success story.¹²⁰ The first examples of photoswitchable channels were synthetic gramicidin derivatives,¹²¹⁻¹²⁶ but attention soon turned to the many ligand-gated ion channels that adorn the outer membrane of cells. Initial experiments showed that azobenzene-containing molecules could mimic the natural ligands for such channels,¹²⁷ leading to the development of two distinct methodologies. In the first a channel blocking ligand mimic group is tethered to an ion channel by an azobenzene-containing linker so that irradiation either leads to insertion or removal of a blocking group from the channel^{128, 129} or binding at a distal site that induces structural changes in the channel. The second approach uses less specific freely diffusible ligand mimics. Tethering is most conveniently achieved through the introduction of cysteine residues at solvent exposed positions and reacting their side chains with maleimides where the use of different cysteine positions allows the tuning of the response to photoswitching.¹³⁰ The genes for individual subunits of heteromeric pores can be altered and heterologously expressed to create hybrid receptors. The function of individual receptor subtypes can then be mapped by their selective activation with light.^{5, 131-135}

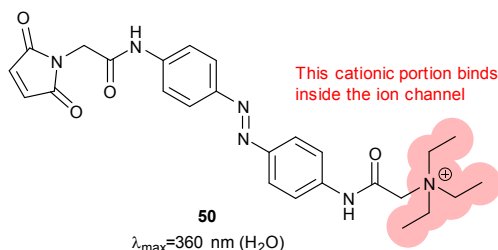


Figure 38. A photochromic, tethering ligand optimised to control a mutant Shaker ion channel with a carefully positioned cysteine residue near the binding pocket for positively charged groups.

The tethered photoswitchable ligand **50** was applied to the voltage-gated potassium ion channel Shaker by selectively conjugating it to a loop on the rim of the pore. Whilst the extended *E*-**50** form is long enough to allow the pendant ethylammonium group to block the pore, the photo-activated *Z*-**50** is too short (Figure 39). *Z*-**50** leaves the pore open allowing action potentials to be silenced by blocking potassium ion transport¹³⁶ or, with further mutations to allow sodium influx, action potentials to be triggered by irradiation.¹³⁷ Use of less selective acrylamide, chloroacetate or epoxide agents allowed the non-selective labelling of endogenously expressed Shaker channels to create UV responsive cultured neurons.¹³⁸ Viral co-transduction of DNA encoding mutant channels and GFP into the hippocampal region of mice caused no visible behavioural changes.¹³⁹

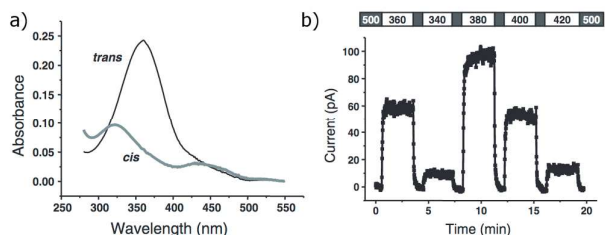


Figure 39. a) UV/visible absorbance spectra of **50** conjugated to glutathione. b) Action spectra of inside-out membrane patches containing Shaker ion channels conjugated with **50**. Adapted by permission from Macmillan Publishers Ltd: Nature Neuroscience, 2004, 7, 1381, copyright 2004.

To complement diffusible azobenzene-containing glutamate mimics,^{140, 141} the photoswitchable tethered ligand **51** was applied to ionotropic glutamate receptors, which feature extracellular binding sites that cause conformational changes when populated by glutamate.¹⁴²⁻¹⁴⁶ Zebrafish larvae engineered to express modified channels were bathed in a solution of **51** for 30 minutes. Touching the treated larvae revealed reversibly altered behaviour according to the irradiation they had received; 365 nm light deactivated their escape reflex and 488 nm light reactivated it.^{147, 148} Channels in five-day-old zebrafish were modified with **51** to elucidate the function of Kolmer-Agduhr cells, demonstrating the potential of this technique as an exquisitely selective investigative tool.¹⁴⁹ In addition to pure ligand gating approaches, bis-maleimide (**39**) has been used to control the trimeric proteins that comprise P2X2/3 receptors moving the subunits apart to open the pore.¹⁵⁰

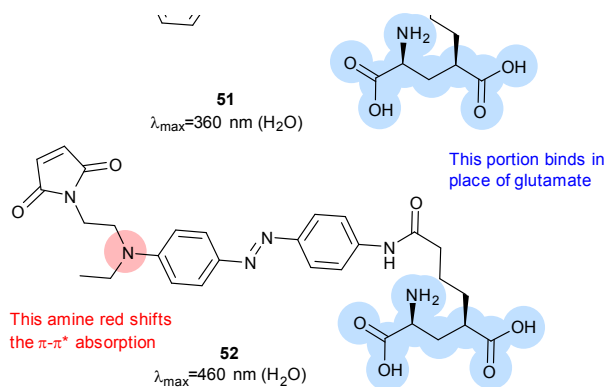


Figure 40. Azobenzene-containing photoswitchable tethering ligands for ionic glutamate receptors. Replacing the amido substituent with an amino substituent creates a red-shifted, fast-relaxing switch.

An exciting long-term research goal is the use of these channels to achieve a degree of sight restoration for vision-impaired patients, which requires rapid relaxation and absorption at distinct wavelengths in the visible spectrum. A modified azobenzene (**52**) where one ring bears an electron-releasing amine substituent increases the channel activation wavelength from 380 nm to 445 nm, a wavelength better suited for vision restoration research. Channels modified with **52** returned to their ground state 2200-fold faster than channels modified with **51**, allowing much higher temporal resolution control of neuron firing¹⁵¹ Azobenzene **52** has a much-improved two-photon absorption cross-section that is comparable to that in fluorescent proteins.¹⁵² A tethered tetra-*ortho*-chloroazobenzene glutamate ligand (**53**) allowed triggering of a modified GluK2 kainate receptor with 560 nm light.¹⁵³ High intensity 625 nm light could also be used, albeit at lower efficiency, but 440 nm irradiation induced no response so that **53** and **51** can work orthogonally with narrow wavelength light sources (Figure 41).

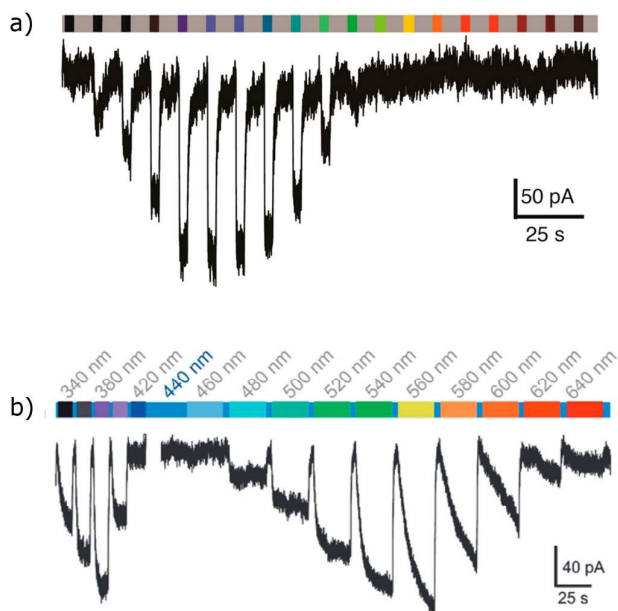


Figure 41. a) Representative action spectrum showing activation of the homotetrameric kainate receptor GluK2(439C) with **52** conjugated to its ligand binding domain. b) an action spectrum of the same protein conjugated with **53** showing how these two tethered ligands might be used in parallel. a) Adapted with permission from Kienzler *et al.* *J. Am. Chem. Soc.* 2013, **135**, 17683 and Copyright 2013 American Chemical Society. b) Adapted under CC-BY from Rullo *et al.*, *Chem. Commun.*, 2014, **50**, 14613.

A recent report describes late stage incorporation of *ortho*-chloro substituents to preformed azo compounds, making such molecules more accessible.¹⁵⁴ Even longer wavelengths can be used to switch **54** (880 nm) and **55** (900 nm), which were designed for improved two-photon absorption.¹⁵⁵

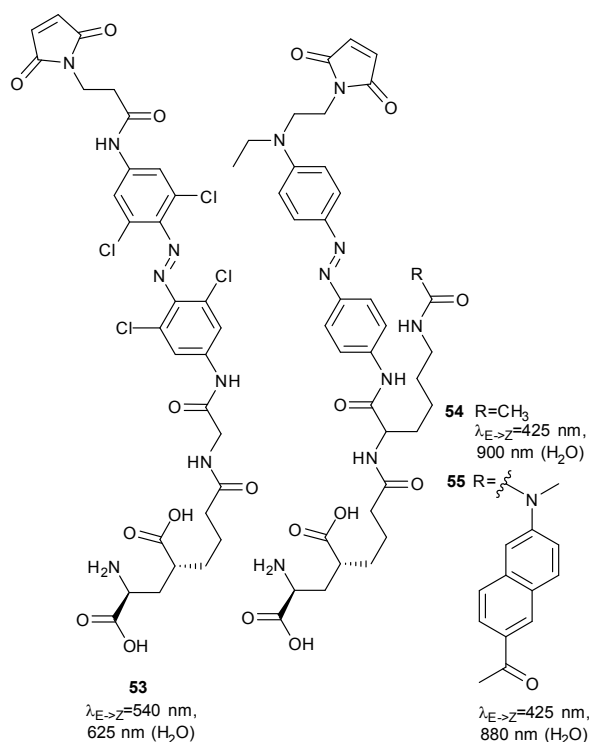


Figure 42. A tetra-*ortho*-chloroazobenzene photoswitchable tethered ligand. The $n-\pi^*$ absorbance band separation allows this molecule to be switched with light up to 625 nm at high intensities.

Non-tethered, diffusible azobenzene ligands can also be used to control a wide variety of channels. Successful endeavours include analysing the behaviour of Purkinje cells using **56**,¹⁵⁶ modulating insulin secretion *via* ATP-sensitive potassium channels with **57** and **58**,^{157, 158} transient receptor potential ion channel TRPV1{Mourou, 2011 #57}¹⁵⁹, that respond to compounds including azo-capsaicin **59**,¹⁶⁰ metabotropic glutamate receptors (class C GPCR receptors) controlled by **60**,¹⁶¹ trimeric amiloride-sensitive sodium channels gated by **61**,¹⁶² G-protein coupled inwardly rectifying potassium channels regulated by **62**,¹⁶³ and endowment of retinal function on blind mice using **63**.¹⁶⁴

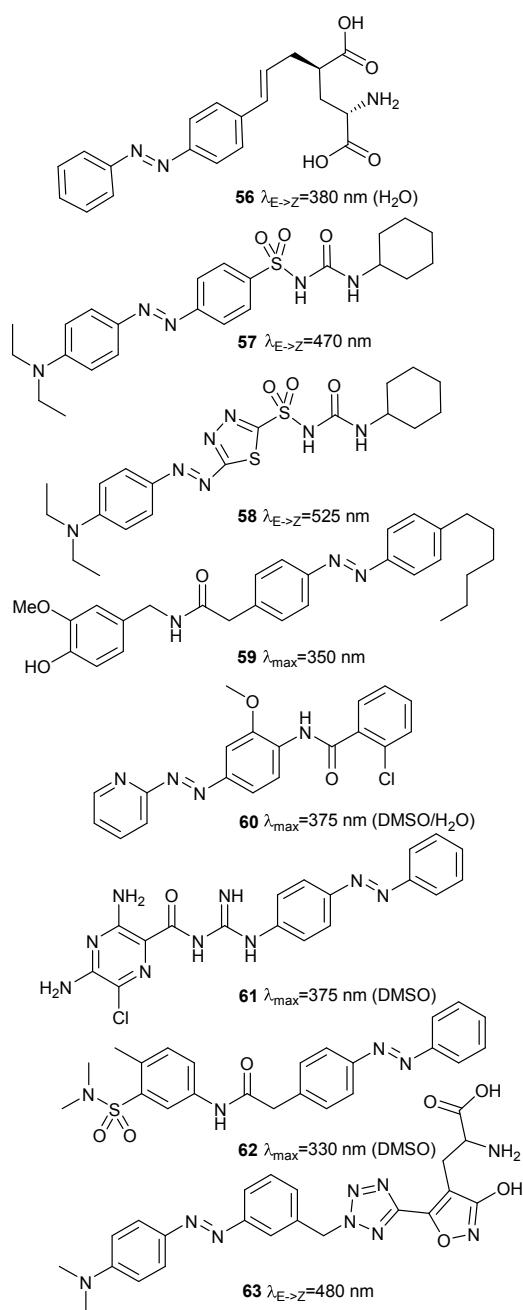


Figure 43. Non-tethered azobenzene ligand for controlling various ion channels.

Conclusions

Recent breakthroughs in tuning the physical properties of photoswitches have improved the scope of existing tools, but have also opened up new opportunities. Azobenzenes with extended Z-isomer half-lives now cover the timescales of more complex biological events and allow *in vivo* kinetics experiments to be conducted. Control over the lifetime of

active states is complemented by an expanded range of photoswitching wavelength and offers the possibility to combine two or more dyes to create multiplexed biophotonic systems. Such combinations of probes can lead to powerful tools to study thresholds in signalling events, as differentially timed light pulses could be used to simulate concentration gradients. The ever-growing number of photopharmaceutical tools could also be envisaged being used in pairs, with the effects of a pair of opposed molecules activated in different locations providing proportionally greater effects.

Future challenges include improving the mechanisms researchers use to translate photoisomerisation energy into structural change.¹⁶⁵⁻¹⁶⁸ Natural systems have evolved to be delicately balanced to change conformation when a ligand or binding partner is present, but directed evolution is presently impractical for proteins containing non-proteinogenic amino acids. Computational studies may therefore offer the best chance of successfully (re)designing systems to achieve improved photocontrol over protein structure and function.

Acknowledgements

Financial support from the UK Biotechnology and Biological Research Council through grants B16000, BB/I021396/1 and BB/M006158/1), the UK Engineering and Physical Sciences Research Council through grants EP/F040954/1 and EP/L027240/1), the Royal Society (IIF-2007/R1) and Cardiff University is gratefully acknowledged.

References

1. A. Fihey, A. Perrier, W. R. Browne and D. Jacquemin, *Chem. Soc. Rev.*, 2015, **44**, 3719.
2. D. Blegler and S. Hecht, *Angew. Chem., Int. Ed.*, 2015, **54**, 11338.
3. J. Zhang, J. Wang and H. Tian, *Mater. Horiz.*, 2014, **1**, 169.
4. A. Priimagi, C. J. Barrett and A. Shishido, *J. Mater. Chem. C*, 2014, **2**, 7155.
5. T. Fehrentz, M. Schonberger and D. Trauner, *Angew. Chem., Int. Ed.*, 2011, **50**, 12156.
6. A. Gautier, C. Gauron, M. Volovitch, D. Bensimon, L. Jullien and S. Vriz, *Nat. Chem. Biol.*, 2014, **10**, 533.
7. P. R. O'Neill and N. Gautam, *Photochem. Photobiol. Sci.*, 2015, **14**, 1578.
8. W. Szymanski, J. M. Beierle, H. A. Kistemaker, W. A. Velema and B. L. Feringa, *Chem. Rev.*, 2013, **113**, 6114.
9. M. Erdelyi, M. Varedian, C. Skold, I. B. Niklasson, J. Nurbo, A. Persson, J. Bergquist and A. Gogoll, *Org. Biomol. Chem.*, 2008, **6**, 4356.
10. M. Varedian, V. Langer, J. Bergquist and A. Gogoll, *Tetrahedron Lett.*, 2008, **49**, 6033.
11. N. J. Lindgren, M. Varedian and A. Gogoll, *Chem. Eur. J.*, 2009, **15**, 501.
12. M. Varedian, M. Erdelyi, A. Persson and A. Gogoll, *J. Pept. Sci.*, 2009, **15**, 107.

13. C. Karlsson, M. Blom, M. Johansson, A. M. Jansson, E. Scifo, A. Karlen, T. Govender and A. Gogoll, *Org. Biomol. Chem.*, 2015, **13**, 2612.
14. S. Herre, W. Steinle and K. Ruck-Braun, *Synthesis*, 2005, 3297.
15. T. Cordes, D. Weinrich, S. Kempa, K. Riesselmann, S. Herre, C. Hoppmann, K. Ruck-Braun and W. Zinth, *Chem. Phys. Lett.*, 2006, **428**, 167.
16. S. Herre, T. Schadendorf, I. Ivanov, C. Herrberger, W. Steinle, K. Ruck-Braun, R. Preissner and H. Kuhn, *ChemBioChem*, 2006, **7**, 1089.
17. M. Fullbeck, E. Michalsky, I. S. Jaeger, P. Henklein, H. Kuhn, K. Ruck-Braun and R. Preissner, *Genome Inform.*, 2006, **17**, 141.
18. T. Schadendorf, C. Hoppmann and K. Ruck-Braun, *Tetrahedron Lett.*, 2007, **48**, 9044.
19. T. Cordes, T. Schadendorf, B. Prieswisch, K. Ruck-Braun and W. Zinth, *J. Phys. Chem. A*, 2008, **112**, 581.
20. T. Cordes, T. Schadendorf, K. Ruck-Braun and W. Zinth, *Chem. Phys. Lett.*, 2008, **455**, 197.
21. T. Cordes, C. Elsner, T. T. Herzog, C. Hoppmann, T. Schadendorf, W. Summerer, K. Ruck-Braun and W. Zinth, *Chem. Phys.*, 2009, **358**, 103.
22. J. Hemphill, C. Chou, J. W. Chin and A. Deiters, *J. Am. Chem. Soc.*, 2013.
23. S. Wiedbrauk and H. Dube, *Tetrahedron Lett.*, 2015, **56**, 4266.
24. J. Z. Zhao, D. Wildemann, M. Jakob, C. Vargas and C. Schiene-Fischer, *Chem. Commun.*, 2003, 2810.
25. J. Helbing, H. Bregy, J. Bredenbeck, R. Pfister, P. Hamm, R. Huber, J. Wachtveitl, L. De Vico and M. Olivucci, *J. Am. Chem. Soc.*, 2004, **126**, 8823.
26. Y. Huang, Z. Y. Cong, L. F. Yang and S. L. Dong, *J. Pept. Sci.*, 2008, **14**, 1062.
27. Z. Y. Cong, L. F. Yang, L. Jiang, D. Ye and S. L. Dong, *Chin. Chem. Lett.*, 2010, **21**, 476.
28. J. M. Goldberg, S. Batjargal and E. J. Petersson, *J. Am. Chem. Soc.*, 2010, **132**, 14718.
29. Y. Huang, G. Jahreis, C. Lucke, D. Wildemann and G. Fischer, *J. Am. Chem. Soc.*, 2010, **132**, 7578.
30. M. Blanco-Lomas, S. Samanta, P. J. Campos, G. A. Woolley and D. Sampedro, *J. Am. Chem. Soc.*, 2012, **134**, 6960.
31. K. Fujimoto, T. Maruyama, Y. Okada, T. Itou and M. Inouye, *Tetrahedron*, 2013, **69**, 6170.
32. O. Babii, S. Afonin, M. Berditsch, S. Reibetaer, P. K. Mykhailiuk, V. S. Kubyshekin, T. Steinbrecher, A. S. Ulrich and I. V. Komarov, *Angew. Chem., Int. Ed.*, 2014, **53**, 3392.
33. D. J. van Dijken, P. Kovaricek, S. P. Ihrig and S. Hecht, *J. Am. Chem. Soc.*, 2015.
34. A. Sinicropi, C. Bernini, R. Basosi and M. Olivucci, *Photochem. Photobiol. Sci.*, 2009, **8**, 1639.
35. G. A. Woolley, *Acc. Chem. Res.*, 2005, **38**, 486.
36. C. Renner and L. Moroder, *ChemBioChem*, 2006, **7**, 869.
37. D. G. Flint, J. R. Kumita, O. S. Smart and G. A. Woolley, *Chem. Biol.*, 2002, **9**, 391.
38. S. H. Xia, G. Cui, W. H. Fang and W. Thiel, *Angew. Chem., Int. Ed.*, 2016, **55**, 2067.
39. Z. H. Zhang, D. C. Burns, J. R. Kumita, O. S. Smart and G. A. Woolley, *Bioconjugate Chem.*, 2003, **14**, 824.
40. D. C. Burns, F. Z. Zhang and G. A. Woolley, *Nat. Protoc.*, 2007, **2**, 251.
41. M. Dong, A. Babalhavaeji, S. Samanta, A. A. Beharry and G. A. Woolley, *Acc. Chem. Res.*, 2015, **48**, 2662.
42. O. Sadovski, A. A. Beharry, F. Z. Zhang and G. A. Woolley, *Angew. Chem., Int. Ed.*, 2009, **48**, 1484.
43. A. A. Beharry, O. Sadovski and G. A. Woolley, *J. Am. Chem. Soc.*, 2011, **133**, 19684.
44. S. Samanta, T. M. McCormick, S. K. Schmidt, D. S. Seferos and G. A. Woolley, *Chem. Commun.*, 2013, **49**, 10314.
45. S. Samanta, A. A. Beharry, O. Sadovski, T. M. McCormick, A. Babalhavaeji, V. Tropepe and G. A. Woolley, *J. Am. Chem. Soc.*, 2013, **135**, 9777.
46. A. A. Beharry, O. Sadovski and G. A. Woolley, *Org. Biomol. Chem.*, 2008, **6**, 4323.
47. S. Samanta, A. Babalhavaeji, M. X. Dong and G. A. Woolley, *Angew. Chem., Int. Ed.*, 2013, **52**, 14127.
48. M. Dong, A. Babalhavaeji, M. J. Hansen, L. Kalman and G. A. Woolley, *Chem. Commun.*, 2015, **51**, 12981.
49. D. Bleger, J. Schwarz, A. M. Brouwer and S. Hecht, *J. Am. Chem. Soc.*, 2012, **134**, 20597.
50. C. Knie, M. Utecht, F. Zhao, H. Kulla, S. Kovalenko, A. M. Brouwer, P. Saalfrank, S. Hecht and D. Bleger, *Chem. Eur. J.*, 2014, **20**, 16492.
51. A. A. Beharry, L. Wong, V. Tropepe and G. A. Woolley, *Angew. Chem., Int. Ed.*, 2011, **50**, 1325.
52. Y. X. Zhang, F. Erdmann and G. Fischer, *Nat. Chem. Biol.*, 2009, **5**, 724.
53. J. Moreno, M. Gerecke, A. L. Dobryakov, I. N. Ioffe, A. A. Granovsky, D. Bleger, S. Hecht and S. A. Kovalenko, *J. Phys. Chem. B*, 2015, **119**, 12281.
54. J. Moreno, M. Gerecke, L. Grubert, S. A. Kovalenko and S. Hecht, *Angew. Chem., Int. Ed.*, 2016, **55**, 1544.
55. R. Siewertsen, H. Neumann, B. Buchheim-Stehm, R. Herges, C. Nather, F. Renth and F. Temps, *J. Am. Chem. Soc.*, 2009, **131**, 15594.
56. O. Carstensen, J. Sielk, J. B. Schonborn, G. Granucci and B. Hartke, *J. Chem. Phys.*, 2010, **133**.
57. M. Boeckmann, N. L. Doltsinis and D. Marx, *Angew. Chem., Int. Ed.*, 2010, **49**, 3382.
58. C. W. Jiang, R. H. Xie, F. L. Li and R. E. Allen, *J. Phys. Chem. A*, 2011, **115**, 244.
59. S. Samanta, C. G. Qin, A. J. Lough and G. A. Woolley, *Angew. Chem., Int. Ed.*, 2012, **51**, 6452.
60. H. Sell, C. Nather and R. Herges, *Beilstein J. Org. Chem.*, 2013, **9**, 1.
61. Y. Yang, R. P. Hughes and I. Aprahamian, *J. Am. Chem. Soc.*, 2012, **134**, 15221.
62. Y. Yang, R. P. Hughes and I. Aprahamian, *J. Am. Chem. Soc.*, 2014, **136**, 13190.
63. Y. Wang and D. H. Chou, *Angew. Chem., Int. Ed.*, 2015, **54**, 10931.
64. C. Hoppmann, P. Schmieder, N. Heinrich and M. Beyermann, *ChemBioChem*, 2011, **12**, 2555.
65. C. Hoppmann, V. K. Lacey, G. V. Louie, J. Wei, J. P. Noel and L. Wang, *Angew. Chem., Int. Ed.*, 2014, **53**, 3932.

66. C. Hoppmann, I. Maslennikov, S. Choe and L. Wang, *J. Am. Chem. Soc.*, 2015, **137**, 11218.
67. A. Muller, H. Kobarg, V. Chandrasekaran, J. Gronow, F. D. Sonnichsen and T. K. Lindhorst, *Chem. Eur. J.*, 2015, **21**, 13723.
68. C. Poloni, W. Szymanski, L. Hou, W. R. Browne and B. L. Feringa, *Chem. Eur. J.*, 2014, **20**, 946.
69. M. Liu, S. Tada, M. Ito, H. Abe and Y. Ito, *Chem. Commun.*, 2012, **48**, 11871.
70. M. R. Jafari, L. Deng, P. I. Kitov, S. Ng, W. L. Matochko, K. F. Tjhung, A. Zeberoff, A. Elias, J. S. Klassen and R. Derda, *ACS Chem. Biol.*, 2014, **9**, 443.
71. S. Bellotto, S. Chen, I. Rentero Rebollo, H. A. Wegner and C. Heinis, *J. Am. Chem. Soc.*, 2014, **136**, 5880.
72. L. Guerrero, O. S. Smart, G. A. Woolley and R. K. Allemann, *J. Am. Chem. Soc.*, 2005, **127**, 15624.
73. L. Guerrero, O. S. Smart, C. J. Weston, D. C. Burns, G. A. Woolley and R. K. Allemann, *Angew. Chem., Int. Ed.*, 2005, **44**, 7778.
74. G. A. Woolley, A. S. I. Jaikaran, M. Berezovski, J. P. Calarco, S. N. Krylov, O. S. Smart and J. R. Kumita, *Biochemistry*, 2006, **45**, 6075.
75. A. M. Ali, M. W. Forbes and G. A. Woolley, *ChemBioChem*, 2015, **16**, 1757.
76. F. Z. Zhang, K. A. Timm, K. M. Arndt and G. A. Woolley, *Angew. Chem., Int. Ed.*, 2010, **49**, 3943.
77. A. M. Caamano, M. E. Vazquez, J. Martinez-Costas, L. Castedo and J. L. Mascarenas, *Angew. Chem., Int. Ed.*, 2000, **39**, 3104.
78. R. J. Mart, P. Wysoczanski, S. Kneissl, A. Ricci, A. Brancale and R. K. Allemann, *ChemBioChem*, 2012, **13**, 515.
79. L. Ulysse and J. Chmielewski, *Bioorg. Med. Chem. Lett.*, 1994, **4**, 2145.
80. L. Ulysse, J. Cubillos and J. Chmielewski, *J. Am. Chem. Soc.*, 1995, **117**, 8466.
81. L. G. Ulysse and J. Chmielewski, *Chem. Biol. Drug Des.*, 2006, **67**, 127.
82. A. Aemissegger, V. Krautler, W. F. van Gunsteren and D. Hilvert, *J. Am. Chem. Soc.*, 2005, **127**, 2929.
83. S. L. Dong, M. Loweneck, T. E. Schrader, W. J. Schreier, W. Zinth, L. Moroder and C. Renner, *Chem. Eur. J.*, 2006, **12**, 1114.
84. T. E. Schrader, W. J. Schreier, T. Cordes, F. O. Koller, G. Babitzki, R. Denschlag, C. Renner, M. Loweneck, S. L. Dong, L. Moroder, P. Tavan and W. Zinth, *Proc. Natl. Acad. Sci. U. S. A.*, 2007, **104**, 15729.
85. T. E. Schrader, T. Cordes, W. J. Schreier, F. O. Koller, S. L. Dong, L. Moroder and W. Zinth, *J. Phys. Chem. B*, 2011, **115**, 5219.
86. T. Podewin, M. S. Rampp, I. Turkanovic, K. L. Karaghiosoff, W. Zinth and A. Hoffmann-Roder, *Chem. Commun.*, 2015, **51**, 4001.
87. R. Behrendt, M. Schenk, H. J. Musiol and L. Moroder, *J. Pept. Sci.*, 1999, **5**, 519.
88. C. Renner, J. Cramer, R. Behrendt and L. Moroder, *Biopolymers*, 2000, **54**, 501.
89. C. Renner, R. Behrendt, S. Sporlein, J. Wachtveitl and L. Moroder, *Biopolymers*, 2000, **54**, 489.
90. R. Behrendt, C. Renner, M. Schenk, F. Q. Wang, J. Wachtveitl, D. Oesterheld and L. Moroder, *Angew. Chem., Int. Ed.*, 1999, **38**, 2771.
91. C. Renner, R. Behrendt, N. Heim and L. Moroder, *Biopolymers*, 2002, **63**, 382.
92. M. Schutt, S. S. Krupka, A. G. Milbradt, S. Deindl, E. K. Sinner, D. Oesterheld, C. Renner and L. Moroder, *Chem. Biol.*, 2003, **10**, 487.
93. A. G. Milbradt, M. Loweneck, S. S. Krupka, M. Reif, E. K. Sinner, L. Moroder and C. Renner, *Biopolymers*, 2005, **77**, 304.
94. C. Hoppmann, S. Seedorff, A. Richter, H. Fabian, P. Schmieder, K. Ruck-Braun and M. Beyermann, *Angew. Chem., Int. Ed.*, 2009, **48**, 6636.
95. L. Nevola, A. Martin-Quiros, K. Eckelt, N. Camarero, S. Tosi, A. Llobet, E. Giralt and P. Gorostiza, *Angew. Chem., Int. Ed.*, 2013, **52**, 7704.
96. A. Martin-Quiros, L. Nevola, K. Eckelt, S. Madurga, P. Gorostiza and E. Giralt, *Chem. Biol.*, 2015, **22**, 31.
97. R. S. Ritterson, K. M. Kuchenbecker, M. Michalik and T. Kortemme, *J. Am. Chem. Soc.*, 2013, **135**, 12516.
98. S. Kneissl, E. J. Loveridge, C. Williams, M. P. Crump and R. K. Allemann, *ChemBioChem*, 2008, **9**, 3046.
99. P. Wysoczanski, R. J. Mart, E. J. Loveridge, C. Williams, S. B. M. Whittaker, M. P. Crump and R. K. Allemann, *J. Am. Chem. Soc.*, 2012, **134**, 7644.
100. P. Wysoczanski, R. J. Mart, E. J. Loveridge, C. Williams, S. B. M. Whittaker, M. P. Crump and R. K. Allemann, *Biomol. NMR Assign.*, 2013, **7**, 187.
101. R. J. Mart, R. J. Errington, C. L. Watkins, S. C. Chappell, M. Wiltshire, A. T. Jones, P. J. Smith and R. K. Allemann, *Mol. BioSyst.*, 2013, **9**, 2597.
102. F. Bonardi, G. London, N. Nouwen, B. Feringa and A. J. M. Driessen, *Angew. Chem., Int. Ed.*, 2010, **49**, 7234.
103. H. Shishido, M. D. Yamada, K. Kondo and S. Maruta, *J. Biochem.*, 2009, **146**, 581.
104. B. Schierling, A. J. Noel, W. Wende, L. T. Hien, E. Volkov, E. Kubareva, T. Oretskaya, M. Kokkinidis, A. Rompp, B. Spengler and A. Pingoud, *Proc. Natl. Acad. Sci. U. S. A.*, 2010, **107**, 1361.
105. B. Schierling, N. Dannemann, L. Gabsalilow, W. Wende, T. Cathomen and A. Pingoud, *Nucleic Acids Res.*, 2012, **40**, 2623.
106. L. T. Hien, T. S. Zatsepina, B. Schierling, E. M. Volkov, W. Wende, A. Pingoud, E. A. Kubareva and T. S. Oretskaya, *Bioconjugate Chem.*, 2011, **22**, 1366.
107. D. Hoersch, S. H. Roh, W. Chiu and T. Kortemme, *Nat. Nanotechnol.*, 2013, **8**, 928.
108. A. Prestel and H. M. Moller, *Chem. Commun.*, 2015, **52**, 701.
109. S. Sawada, N. Kato and K. Kaihatsu, *Curr. Pharm. Biotechnol.*, 2012, **13**, 2642.
110. W. A. Velema, W. Szymanski and B. L. Feringa, *J. Am. Chem. Soc.*, 2014, **136**, 2178.
111. J. Broichhagen, T. Podewin, H. Meyer-Berg, Y. von Ohlen, N. R. Johnston, B. J. Jones, S. R. Bloom, G. A. Rutter, A. Hoffmann-Roder, D. J. Hodson and D. Trauner, *Angew. Chem., Int. Ed.*, 2015, **54**, 15565.
112. J. Broichhagen, N. R. Johnston, Y. von Ohlen, H. Meyer-Berg, B. J. Jones, S. R. Bloom, G. A. Rutter, D. Trauner and D. J. Hodson, *Angew. Chem., Int. Ed.*, 2016, **55**, 5865.
113. M. Borowiak, W. Nahaboo, M. Reynders, K. Nekolla, P. Jalinot, J. Hasserodt, M. Rehberg, M. Delattre, S. Zahler, A. Vollmar, D. Trauner and O. Thorn-Seshold, *Cell*, 2015, **162**, 403.
114. A. J. Engdahl, E. A. Torres, S. E. Lock, T. B. Engdahl, P. S. Mertz and C. N. Streu, *Org. Lett.*, 2015, **17**, 4546.

115. J. E. Sheldon, M. M. Dcona, C. E. Lyons, J. C. Hackett and M. C. Hartman, *Org. Biomol. Chem.*, 2016, **14**, 40.
116. W. Szymanski, M. E. Ourailidou, W. A. Velema, F. J. Dekker and B. L. Feringa, *Chem. Eur. J.*, 2015, **21**, 16517.
117. S. A. Reis, B. Ghosh, J. A. Hendricks, D. M. Szantai-Kis, L. Tork, K. N. Ross, J. Lamb, W. Read-Button, B. Zheng, H. Wang, C. Salthouse, S. J. Haggarty and R. Mazitschek, *Nat. Chem. Biol.*, 2016, **12**, 317.
118. W. A. Velema, M. J. Hansen, M. M. Lerch, A. J. Driessen, W. Szymanski and B. L. Feringa, *Bioconjug Chem*, 2015, **26**, 2592.
119. J. P. Van der Berg, W. A. Velema, W. Szymanski, A. J. M. Driessen and B. L. Feringa, *Chem. Sci.*, 2015, **6**, 3593.
120. A. Reiner, J. Levitz and E. Y. Isacoff, *Curr. Opin. Pharmacol.*, 2015, **20**, 135.
121. C. J. Stankovic, S. H. Heinemann and S. L. Schreiber, *Biochim. Biophys. Acta*, 1991, **1061**, 163.
122. G. A. Woolley, A. S. I. Jaikaran, Z. H. Zhang and S. Y. Peng, *J. Am. Chem. Soc.*, 1995, **117**, 4448.
123. L. Lien, D. C. J. Jaikaran, Z. H. Zhang and G. A. Woolley, *J. Am. Chem. Soc.*, 1996, **118**, 12222.
124. S. Rudolph-Bohner, M. Kruger, D. Oesterhelt, L. Moroder, T. Nagele and J. Wachtveitl, *J. Photochem. Photobiol. A*, 1997, **105**, 235.
125. V. Borisenko, D. C. Burns, Z. H. Zhang and G. A. Woolley, *J. Am. Chem. Soc.*, 2000, **122**, 6364.
126. W. Szymanski, B. Wu, C. Poloni, D. B. Janssen and B. L. Feringa, *Angew. Chem., Int. Ed.*, 2013, **52**, 2068.
127. E. Bartels, H. Wasserman and B. F. Erlanger, *Proc. Natl. Acad. Sci. U. S. A.*, 1971, **68**, 1820.
128. H. A. Lester, M. E. Krouse, M. M. Nass, N. H. Wassermann and B. F. Erlanger, *J. Gen. Physiol.*, 1980, **75**, 207.
129. M. E. Krouse, H. A. Lester, N. H. Wassermann and B. F. Erlanger, *J. Gen. Physiol.*, 1985, **86**, 235.
130. I. Tochitsky, M. R. Banghart, A. Mourot, J. Z. Yao, B. Gaub, R. H. Kramer and D. Trauner, *Nat Chem*, 2012, **4**, 105.
131. R. H. Kramer, J. J. Chambers and D. Trauner, *Nat. Chem. Biol.*, 2005, **1**, 360.
132. P. Gorostiza and E. Isacoff, *Mol. Biosyst.*, 2007, **3**, 686.
133. P. Gorostiza and E. Y. Isacoff, *Science*, 2008, **322**, 395.
134. R. H. Kramer, D. L. Fortin and D. Trauner, *Curr. Opin. Neurobiol.*, 2009, **19**, 544.
135. G. A. Woolley, *Nat Chem*, 2012, **4**, 75.
136. M. Banghart, K. Borges, E. Isacoff, D. Trauner and R. H. Kramer, *Nat. Neurosci.*, 2004, **7**, 1381.
137. J. J. Chambers, M. R. Banghart, D. Trauner and R. H. Kramer, *J. Neurophysiol.*, 2006, **96**, 2792.
138. D. L. Fortin, M. R. Banghart, T. W. Dunn, K. Borges, D. A. Wagenaar, Q. Gaudry, M. H. Karakossian, T. S. Otis, W. B. Kristan, D. Trauner and R. H. Kramer, *Nat. Methods*, 2008, **5**, 331.
139. D. L. Fortin, T. W. Dunn, A. Fedorchak, D. Allen, R. Montpetit, M. R. Banghart, D. Trauner, J. P. Adelman and R. H. Kramer, *J. Neurophysiol.*, 2011, **106**, 488.
140. M. Volgraf, P. Gorostiza, S. Szobota, M. R. Helix, E. Y. Isacoff and D. Trauner, *J. Am. Chem. Soc.*, 2007, **129**, 260.
141. L. Laprell, E. Repak, V. Franckevicius, F. Hartrampf, J. Terhag, M. Hollmann, M. Sumser, N. Rebola, D. A. DiGregorio and D. Trauner, *Nat. Commun.*, 2015, **6**, 8076.
142. M. Volgraf, P. Gorostiza, R. Numano, R. H. Kramer, E. Y. Isacoff and D. Trauner, *Nat. Chem. Biol.*, 2006, **2**, 47.
143. P. Gorostiza, M. Volgraf, R. Numano, S. Szobota, D. Trauner and E. Y. Isacoff, *Proc. Natl. Acad. Sci. U. S. A.*, 2007, **104**, 10865.
144. R. Numano, S. Szobota, A. Y. Lau, P. Gorostiza, M. Volgraf, B. Roux, D. Trauner and E. Y. Isacoff, *Proc. Natl. Acad. Sci. U. S. A.*, 2009, **106**, 6814.
145. J. Levitz, A. T. Popescu, A. Reiner and E. Y. Isacoff, *Frontiers in molecular neuroscience*, 2016, **9**, 2.
146. S. Berlin, S. Szobota, A. Reiner, E. C. Carroll, M. A. Kienzler, A. Guyon, T. Xiao, D. Tauner and E. Y. Isacoff, *eLife*, 2016, **5**.
147. S. Szobota, P. Gorostiza, F. Del Bene, C. Wyart, D. L. Fortin, K. D. Kolstad, O. Tulyathan, M. Volgraf, R. Numano, H. L. Aaron, E. K. Scott, R. H. Kramer, J. Flannery, H. Baier, D. Trauner and E. Y. Isacoff, *Neuron*, 2007, **54**, 535.
148. H. Janovjak, S. Szobota, C. Wyart, D. Trauner and E. Y. Isacoff, *Nat. Neurosci.*, 2010, **13**, 1027.
149. C. Wyart, F. Del Bene, E. Warp, E. K. Scott, D. Trauner, H. Baier and E. Y. Isacoff, *Nature*, 2009, **461**, 407.
150. L. E. Browne, J. P. Nunes, J. A. Sim, V. Chudasama, L. Bragg, S. Caddick and R. A. North, *Proc. Natl. Acad. Sci. U. S. A.*, 2014, **111**, 521.
151. M. A. Kienzler, A. Reiner, E. Trautman, S. Yoo, D. Trauner and E. Y. Isacoff, *J. Am. Chem. Soc.*, 2013, **135**, 17683.
152. E. C. Carroll, S. Berlin, J. Levitz, M. A. Kienzler, Z. Yuan, D. Madsen, D. S. Larsen and E. Y. Isacoff, *Proc. Natl. Acad. Sci. U. S. A.*, 2015, **112**, E776.
153. A. Rullo, A. Reiner, A. Reiter, D. Trauner, E. Y. Isacoff and G. A. Woolley, *Chem. Commun.*, 2014, **50**, 14613.
154. D. B. Konrad, J. A. Frank and D. Trauner, *Chem. Eur. J.*, 2016, **22**, 4364.
155. M. Izquierdo-Serra, M. Gascon-Moya, J. J. Hirtz, S. Pittolo, K. E. Poskanzer, E. Ferrer, R. Alibes, F. Busque, R. Yuste, J. Hernando and P. Gorostiza, *J. Am. Chem. Soc.*, 2014, **136**, 8693.
156. Z. R. Abrams, A. Warrier, D. Trauner and X. A. Zhang, *Frontiers in Neural Circuits*, 2010, **4**.
157. J. Broichhagen, M. Schonberger, S. C. Cork, J. A. Frank, P. Marchetti, M. Bugliani, A. M. Shapiro, S. Trapp, G. A. Rutter, D. J. Hodson and D. Trauner, *Nat. Commun.*, 2014, **5**, 5116.
158. J. Broichhagen, J. A. Frank, N. R. Johnston, R. K. Mitchell, K. Smid, P. Marchetti, M. Bugliani, G. A. Rutter, D. Trauner and D. J. Hodson, *Chem. Commun.*, 2015, **51**, 6018.
159. A. Mourot, T. Fehrentz, Y. Le Feuvre, C. M. Smith, C. Herold, D. Dalkara, F. Nagy, D. Trauner and R. H. Kramer, *Nat. Methods*, 2012, **9**, 396.
160. J. A. Frank, M. Moroni, R. Moshourab, M. Sumser, G. R. Lewin and D. Trauner, *Nat. Commun.*, 2015, **6**, 7118.
161. S. Pittolo, X. Gomez-Santacana, K. Eckelt, X. Rovira, J. Dalton, C. Goudet, J. P. Pin, A. Llobet, J. Giraldo, A. Llebaria and P. Gorostiza, *Nat. Chem. Biol.*, 2014, **10**, 813.
162. M. Schonberger, M. Althaus, M. Fronius, W. Claus and D. Trauner, *Nat Chem*, 2014, **6**, 712.

ARTICLE

Journal Name

163. D. M. Barber, M. Schonberger, J. Burgstaller, J. Levitz, C. D. Weaver, E. Y. Isacoff, H. Baieer and D. Trauner, *Chem. Sci.*, 2016, **7**, 2347.
164. L. Laprell, K. Hull, P. Stawski, C. Schon, S. Michalakis, M. Biel, M. P. Sumser and D. Trauner, *ACS Chem Neurosci*, 2016, **7**, 15.
165. F. Z. Zhang, A. Zarrine-Afsar, M. S. Al-Abdul-Wahid, R. S. Prosser, A. R. Davidson and G. A. Woolley, *J. Am. Chem. Soc.*, 2009, **131**, 2283.
166. A. A. Beharry, T. Chen, M. S. Al-Abdul-Wahid, S. Samanta, K. Davidov, O. Sadovski, A. M. Ali, S. B. Chen, R. S. Prosser, H. S. Chan and G. A. Woolley, *Biochemistry*, 2012, **51**, 6421.
167. B. Buchli, S. A. Waldauer, R. Walsler, M. L. Donten, R. Pfister, N. Blochliger, S. Steiner, A. Cafilisch, O. Zerbe and P. Hamm, *Proc. Natl. Acad. Sci. U. S. A.*, 2013, **110**, 11725.
168. Z. Yu and S. Hecht, *Chem. Commun.*, 2016, **52**, 6639.