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# A light-gated OmpG hybrid

# **Supporting Information**

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#### 1. Syntheses

Compounds 1<sup>1</sup>, 2<sup>2</sup>, 5<sup>2</sup> and 7<sup>1</sup> were prepared using reported procedures. All reactions were performed using dry solvents which were purchased from *Sigma Aldrich Co.* and *Acros*. Reagents were purchased from *Sigma Aldrich Co., Roth* and *TCI* and were used without further purification. For flash chromatography silica gel 60 by *Macherey-Nagel* was used. TLC analyses were performed on aluminum plates coated with silica gel 60 F 254 (*Merck*).

NMR spectra for the characterisation of the compounds were recorded on *Bruker* AVIII-HD500 (500 MHz), AV400 (400 MHz) and DRX600 (600MHz) instruments equipped with a N<sub>2</sub> cooled cryogenic probe head using DMSO-d<sup>6</sup> and CDCl<sub>3</sub> as solvent. HRMS spectra were recorded using a *Thermo Scientific MALDI LTQ Orbitrap*.

#### Synthesis of compound 3



7-(Diethylamino)-4-(hydroxymethyl)-2H-chromene-2-one<sup>1</sup> (2)

4.63 g (20.0 mmol, 1.0 eq.) 7-Diethylamino-4-methylcoumarin (**1**) were dissolved in 120 mL *p*-xylene and 3.33 g (30.0 mmol, 1.5 eq.) SeO<sub>2</sub> were slowly added. The reaction mixture was stirred at 155 °C overnight, filtered to remove SeO<sub>2</sub> and the solvent was removed *in vacuo*. To 130 mL EtOH 380 mg (10 mmol, 0.5 eq.) NaBH<sub>4</sub> were slowly added. The reaction mixture was stirred for 4 h at room temperature, acidified with 20 mL HCl (1.0 M) and 50 mL H<sub>2</sub>O were added and extracted three times with 50 mL CH<sub>2</sub>Cl<sub>2</sub>. The combined organic layers were washed with 50 mL brine, dried over Na<sub>2</sub>SO<sub>4</sub> and the solvent was removed *in vacuo*. Purification was accomplished via flash chromatography (CH<sub>2</sub>Cl<sub>2</sub>:acetone 5:1). The solvent was removed in vacuo to afford 7-(diethylamino)-4-(hydroxymethyl)-2*H*-chromene-2-one (**2**) (1.11 g, 4.5 mmol, 22%) as a light green-yellow solid.

C<sub>14</sub>H<sub>17</sub>NO<sub>3</sub> Mol. Wt.: 247.29 g/mol

<sup>1</sup>**H-NMR (500 MHz, DMSO-***d*<sub>6</sub>**) δ [ppm]:** 7.42 (d, 1H, *J* = 9.00 Hz, aromatic-*H*), 6.66 (dd, 1H, *J* = 9.05 Hz, aromatic-*H*), 6.52 (d, 1H, *J* = 2.55 Hz, aromatic-*H*), 6.06 (s, 1H, aromatic-*H*), 5.50 (t, 1H, *J* = 5.60 Hz, O*H*), 4.66 (dd, 2H, *J* = 5.60 Hz, C*H*<sub>2</sub>OH), 3.41 (q, 4H, *J* = 7.05 Hz, 2 x C*H*<sub>2</sub>CH<sub>3</sub>), 1.11 (t, 6H, *J* = 7.00 Hz, 2 x CH<sub>2</sub>C*H*<sub>3</sub>).

<sup>13</sup>**C-NMR (125 MHz, DMSO-***d***<sub>6</sub>) δ [ppm]:** 161.1, 156.9, 155.6, 150.2, 125.1, 108.5, 105.7, 103.9, 96.8, 59.0, 43.9, 12.3.

HRMS(MALDI): m/z calcd. for C<sub>14</sub>H<sub>18</sub>NO<sub>3</sub> [M+H]<sup>+</sup> 248.12812,

found 248.12866 (Δm = 0.00054, error 2.18 ppm).

(7-(Diethylamino)-2-oxo-2H-chromene-4-yl)methyl-2-bromoacetate<sup>2</sup> (3)



C<sub>16</sub>H<sub>18</sub>BrNO<sub>4</sub> Mol. Wt.: 368.22 g/mol

100 mg (0.40 mmol, 1.0 eq) 7-(Diethylamino)-4-(hydroxymethyl)-2*H*-chromen-2-on (**2**) were dissolved in 7 mL dry toluene. 50 μL (0.62 mmol, 1.5 eq.) dry pyridine and 54 μL (0.62 mmol, 1.5 eq.) bromoacetylbromide were added. The reaction mixture was stirred at room temperature overnight. The precipitate was filtered and washed with EtOAc. The organic solvent was removed *in vacuo*. Purification was accomplished via flash chromatography (EtOAc). The solvent was removed *in vacuo* to afford (7-(diethylamino)-2-oxo-2*H*-chromene-4-yl)methyl-2-bromoacetate (**3**) (136 mg, 0.37 mmol, 93%) as yellow solid.

<sup>1</sup>**H-NMR (400 MHz, DMSO-***d*<sub>6</sub>) δ [ppm]: 7.46 (d, 1H, *J* = 9.04 Hz, aromatic-*H*), 6.69 (dd, 1H, *J* = 9.04 Hz, aromatic-*H*), 6.55 (d, 1H, *J* = 2.52 Hz, aromatic-*H*), 6.07 (s, 1H, aromatic-*H*), 5.39 (s, 2H, C*H*<sub>2</sub>O), 4.34 (s, 2H, C*H*<sub>2</sub>Br), 3.43 (q, 4H, *J* = 8.00 Hz, 2 x C*H*<sub>2</sub>CH<sub>3</sub>), 1.12 (t, 6H, *J* = 8.00 Hz, 2 x CH<sub>2</sub>CH<sub>3</sub>).

<sup>13</sup>C-NMR (100 MHz, DMSO-*d*<sub>6</sub>) δ [ppm]: 166.8, 160.5, 155.8, 150.5, 149.9, 125.5, 108.7, 105.1, 105.0, 96.8, 62.5, 44.0, 26.8, 12.3.

HRMS(MALDI): m/z calcd. for C<sub>16</sub>H<sub>19</sub>BrNO<sub>4</sub> [M+H]<sup>+</sup> 368.04920,

found 368.04892 (Δm = 0.00028, error 0.76 ppm).

#### **Synthesis of Compound 5**

Compound **6** (7-(diethylamino)-2-oxo-2H-1-benzopyran-4-carboxaldehyde) was synthesized in analogy to compound **2** according to literature<sup>1</sup>.



Supporting Figure 1: reaction scheme for the synthesis of compound 5.

7-(Diethylamino)-4-(2-hydroxypentyl)-2H-chromene-2-one (7)



C<sub>18</sub>H<sub>25</sub>NO<sub>3</sub> Mol. Wt.: 303.40 g/mol

123 mg (0.50 mmol, 1.0 eq.) 7-(Diethylamino)-2-oxo-2H-1-benzopyran-4-carboxaldehyde (**6**) were dissolved in 2.0 mL THF and cooled to -80 °C. Then, 300  $\mu$ L (0.75 mmol, 1.5 eq., 2.5 M in *n*-hexane diluted in 3 mL THF) *n*-BuLi, prechilled to -80 °C, were added and the reaction mixture stirred for 60 minutes at -80 °C and at room temperature overnight. The reaction was stopped by addition of 3 mL saturated NH<sub>4</sub>Cl solution. After addition of EtOAc the organic layer was washed with water and brine, dried over MgSO<sub>4</sub> and the solvent evaporated under vacuum. Purification was accomplished via flash chromatography (CH<sub>2</sub>Cl<sub>2</sub>:acetone – 33:1) to yield 7-(diethylamino)-4-(2-hydroxypentyl)-2*H*-chromene-2-on (**7**)(50 mg, 0.17 mmol, 33%) as a yellow solid.

 $\mathbf{R}_{f}$ : 0.6 (CH<sub>2</sub>Cl<sub>2</sub>:acetone – 6:1).

<sup>1</sup>H-NMR (500 MHz, CDCl<sub>3</sub>)  $\delta$  [ppm]: 7.41 (d, 1H, *J* = 9.06 Hz, aromatic-*H*), 6.57 (dd, 1H, *J* = 9.06 Hz, *J* = 2.58 Hz, aromatic-*H*), 6.51 (d, 1H, *J* = 2.58 Hz, aromatic-*H*), 6.24 (s, 1H, aromatic-*H*), 5.02 – 4.94 (m, 1H, CHOH), 3.41 (q, 4H, *J* = 7.14 Hz, 2x CH<sub>2</sub>CH<sub>3</sub>), 2.07 (d, 1H, *J* = 3.84 Hz; OH); 1.89 – 1.83 (m, 1H, CHCH<sub>2,a</sub>), 1.77 – 1.70 (m, 1H, CHCH<sub>2,b</sub>), 1.56 – 1.48 (m, 1H, CH<sub>2</sub>CH<sub>2,a</sub>CH<sub>2</sub>), 1.48 – 1.43 (m, 1H, CH<sub>2</sub>CH<sub>2,b</sub>CH<sub>2</sub>), 1.42 – 1.32 (m, 2H, CH<sub>2</sub>CH<sub>2</sub>CH<sub>3</sub>), 1.21 (t, 6H, *J* = 7.08 Hz, 2x CH<sub>2</sub>CH<sub>3</sub>), 0.92 (t, 3H, *J* = 7,32 Hz, CH<sub>2</sub>CH<sub>2</sub>CH<sub>3</sub>).

<sup>13</sup>C-NMR (125 MHz, CDCl<sub>3</sub>) δ [ppm]: 162.7, 158.5, 156.5, 150.3, 125.0, 108.5, 106.3, 105.2, 98.0, 70.1,
44.7, 44.7, 37.0, 27.9, 22.5, 14.0, 12.5, 12.5.

HRMS(MALDI): m/z calcd. for C<sub>18</sub>H<sub>26</sub>NO<sub>3</sub> [M+H]<sup>+</sup> 304.19072, found 304.19076 (Δm=0.00004, 0.1 ppm).

(7-(Diethylamino)-2-oxo-2H-chromene-4-yl)pentyl-2-bromoacetate<sup>2</sup> (5)



C<sub>20</sub>H<sub>26</sub>BrNO<sub>4</sub> Mol. Wt.: 424.34 g/mol

50 mg (0.17 mmol, 1.0 eq.) 7-(diethylamino)-4-(2-hydroxypentyl)-2*H*-chromene-2-one (**7**) were dissolved in 2 mL toluene and 22 μL pyridine (0.25 mmol, 1.5 eq.) were added. After 5 min 0.20 mL (0.25 mmol, 1.5 eq.) bromoacetylbromide were added and the reaction mixture stirred at room temperature for 16 h. 15 mL EtOAc and 15 mL water were added and the organic phase washed with 10 mL water and 10 mL brine. Purification was accomplished via flash chromatography (CH:EtOAc – 5:1). The solvent was removed *in vacuo* to afford (7-(diethylamino)-2-oxo-2*H*-chromene-4-yl)pentyl-2-bromoacetate (**5**) (53 mg, 0.12 mmol, 74%) as yellow oil.

R<sub>f</sub>: 0.4 (CH:EE - 4:1).

<sup>1</sup>**H-NMR (600 MHz, CDCl<sub>3</sub>) δ [ppm]:** 7.40 (d, 1H, J = 9.06 Hz, aromatic-H), 6.60 (dd, 1H, J = 9.06 Hz, J = 1.98 Hz, aromatic-H), 6.52 (d, 1H, J = 2.12 Hz, aromatic-H), 6.10 (s, 1H, aromatic-H), 5.99 (t, 1H, J = 6.36 Hz, CH), 3.91 (d, 1H, J = 12.36 Hz,  $CH_{2,a}$ Br), 3.89 (d, 1H, J = 12.42 Hz,  $CH_{2,b}$ Br), 3.41 (q, 4H, J = 7.08 Hz, 2x  $CH_2$ CH<sub>3</sub>), 1.95 – 1.90 (m, 2H, CHC $H_2$ CH<sub>2</sub>), 1.46 – 1.31 (m, 4H, CH<sub>2</sub>C $H_2$ CH<sub>3</sub>), 1.21 (t, 6H, J = 7.08 Hz, 2x  $CH_2$ CH<sub>3</sub>), 0.90 (t, 3H, J = 7,20 Hz, CH<sub>2</sub>CH<sub>2</sub>CH<sub>3</sub>).

<sup>13</sup>**C-NMR DEPT (150 MHz, CDCl<sub>3</sub>) δ [ppm]:** 124.8, 108.8, 105.6, 98.2, 72.9, 44.9, 44.9, 34.5, 27.5, 25.4, 22.3, 13.9, 12.5, 12.5.

HRMS(MALDI): m/z calcd. for  $C_{20}H_{27}BrNO_4 [M+H]^+ 424.11180$ , found 424.11134 ( $\Delta m$ =0.00046, 1.1 ppm).

## 2. NMR-Spectra



Supporting Figure 3: <sup>13</sup>C-NMR of compound 2.



**Supporting Figure 4:** <sup>1</sup>H-NMR of compound **3**.







Supporting Figure 6: <sup>1</sup>H-NMR of compound 7.



Supporting Figure 7: <sup>13</sup>C-NMR of compound 7.



Supporting Figure 8: <sup>1</sup>H-NMR of compound 5.



Supporting Figure 9: <sup>13</sup>C-DEPT-NMR of compound 5.

## 3. Mutagenesis

OmpG/ $\Delta$ L6II constructs derived from *E. coli* K-12 *ompG* gene without signal sequence and C-terminal hexa-histidine tag as reported by Grosse *et al.*<sup>3</sup> served as template for the introduction of two cysteine residues. On the basis of OmpG/ $\Delta$ L6II plasmids C13 and C154 were introduced by site directed mutagenesis using primers 5'-CCGTGTCGAAACGTGTACAGGTCTGCAGTATACCTTCAACG-3' and 5'-CGTTGAAGGTATACTGCAGAACCTGTACACGTTTCGACACGG-3'.

Sequencing of the newly constructed primers confirmed introduction of cysteines at the desired positions:

Sequencing Data for OmpG/ $\Delta$ L6-M13C-E154C

Sequence translation was performed by using online tool *ExPASy translator*. Cysteine mutations highlighted in the amino acid sequence:

MEERNDWHFN IGACYEIENV EGYGEDMDGL AEPSVYFNAA NGPWRIALAY YQEGPVDYSA GKRGTWFDRP ELEVHYQFLE NDDFSFGLTG GFRNYGYHYV DEPGKDTANM QRWKIAPDWD VKLTDDLRFN GWLSMYKFAN DLNTTGYADT RVETCTGLQY TFNETVALRV NYYLERGFNM DDSRNNGEFS TQEIRAYLPL TLGNHSVTPY TRIGLDRAGH DFNRVGLFYG YDFQNGLSVS LEYAFEWQDH DEGDSDKFHY AGVGVNYSFH HHHHH

#### 4. Protein methods

#### **Expression and Purification**

OmpG mutant was produced as inclusion bodies after transformation in the porin-deficient *E.coli* BL21 (DE3)omp9 strain (B F<sup>-</sup> *ompT hsd*SB(rB<sup>-</sup> mB<sup>-</sup>) *gal dcm* (DE3)  $\Delta$ *lamB ompF*::Tn5  $\Delta$ *ompA*  $\Delta$ *ompC*  $\Delta$ *ompN* :: $\Omega$ )<sup>4</sup> as described by Grosse *et al.*<sup>3</sup> Cells were grown in 2YT medium with 100 µg/mL ampicillin and 1% glucose at 37 °C until OD<sub>595</sub>~0.7. Expression was induced by addition of IPTG (1 mM). Cells were grown for 4 h, harvested by centrifugation (4000 g, 4 °C, 30 min) and resuspended in TN buffer (50 mM Tris/HCl, pH 8.0, 100 mM NaCl). Before cell lysis in an *M-110P homogenizer (Microfluidics)*, DNase and 1mM PMSF were added. The resulting suspension was centrifuged (22000 g, 4 °C, 25 min) and supernatant discarded. Inclusion bodies were washed in Triton-TN buffer (50 mM Tris/HCl, pH 8.0, 100 mM NaCl) again for 3 times. Washed inclusion bodies were suspended in TN-buffer, aliquoted and stored at -20 °C until further usage.

#### Refolding

Purified inclusion bodies were resuspended in denaturation buffer (25 mM Tris/HCl, pH 8.0, 6 M urea, 1 mM TCEP), heated for 10 min at 60 °C and centrifuged at 20000 g for 3 min. The supernatant was moved to a fresh tube and the concentration determined. Refolding was performed by rapid dilution in refolding buffer (25 mM Tris/HCl, pH 8.0, 2.5 M urea, 1 mM TCEP, 3% OG) to a concentration of 1 mg/mL OmpG and mildly shaken overnight at 30 °C. Misfolded porin was removed by digestion with proteinase K (10  $\mu$ g/mL, 30 min, 25 °C) and the reaction stopped with 13 mM PMSF. Afterwards, refolded protein was concentrated and washed with IEC starting buffer prior to ion exchange chromatography (*HiTrap Q*, 10 mM Tris/HCl, pH 8.0, 10% glycerol, 0.05% LDAO, NaCl gradient, see supporting table 1 and figure 10) Purified OmpG was concentrated to at least 5 mg/mL and stored at -80 °C until further usage.

Supporting Table 1: NaCl gradient of ion exchange chromatography in IEC buffer.

Column volume	% 10 mM NaCl	% 1 M NaCl
0	100	0
6	100	0
16	50	50

#### Labelling

Refolded OmpG was diluted to a concentration of 3.0 mg/mL in labelling buffer (100 mM KP<sub>i</sub>, pH 7.5, 0.4% C<sub>8</sub>E<sub>4</sub>). Coumarin **3/5** was diluted in acetonitrile and added in a tenfold concentration to the protein together with 1 mM TCEP. After an incubation of about 30 minutes at 37 °C, the addition was repeated and the incubation extended overnight. OmpG hybrids were separated from excess coumarin by size exclusion PD-10 columns and another IEC chromatography (see supporting table 1 for gradient, figure 11). Afterwards, the product was concentrated to 5-10 mg/mL and dialyzed in storage buffer (25 mM Tris/HCl, pH 8.0, 10% glycerol, 100 mM NaCl, 0.4% C<sub>8</sub>E<sub>4</sub>). The final product was stored at -80 °C until further usage.



Supporting Figure 10: exemplary FPLC trace of OmpG-1. Absorbance at 390 nm (green) provides evidence for a successful labelling.

#### 5. Determination of labelling efficiency by absorption

The labelling efficiency of OmpG hybrids was determined by absorption measurements. Distinct absorption maxima of both OmpG and label were compared in order to determine the amount of label attached to the protein. Therefore, all molar extinction coefficients ( $\epsilon$ ) were determined for a calculation of concentrations. The molar extinction coefficient of OmpG was determined by its amino acid sequence applying the online tool *ExPASy ProtParam* (see supporting table 2). The molar extinction coefficients of the labels were determined by absorption measurements of a concentration series of the small molecule.

#### OmpG-2

In a first step, the molar extinction coefficient of DEACM was determined at its absorption maximum  $\epsilon_{max}(\lambda) = 390$  nm (see supporting figure 12). For the extinction coefficient of the doubly labelled hybrid the value of DEACM at 280 nm was added twice to the extinction coefficient of unlabelled **OmpG-1** (see supporting table 2).



Supporting Figure 11: left – graphical determination of ε. Right – absorption spectrum of DEACM.

	<b>E</b> 280 nm	<b>č</b> 390 nm
	L/mol·cm	L/mol∙cm
OmpG-1	71260	0
DEACM	3500	19900
OmpG-2	78260	19900

Supporting Table 2: Summary of extinction coefficient
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From the absorption spectrum in Supporting Figure 13 absorption at 280 and 390 nm can be drawn. In Supporting Table 3 the concentration is calculated from absorbance values. DEACM has approximatly the double concentration of OmpG which indicates full labelling.



Supporting Figure 12: absorption spectrum of OmpG-2.

Supporting Table 3: concentration of OmpG-2 determined at 280 and 390 nm.

OmpG-2	unit	280 nm	390 nm
ε	L/mol∙cm	78260	19900
Α	a.u.	0.965	0.495
с	μΜ	12.3	24.9

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#### OmpG-4

For *n-buty*/DEACM the same procedure was performed as above. The results are shown in Supporting Figure 14 and Table 4.



**Supporting Figure 13:** left – graphical determination of ε. Right – absorbance spectrum of *n*-buty/DEACM.

Supporting Table 4: Summary of extinction coefficients.

	<b>E</b> 280 nm	<b>8</b> 390 nm
	L/mol·cm	L/mol·cm
OmpG-1	71260	0
nbuty/DEACM	3500	20200
OmpG-4	78260	20200

From the absorption spectrum in Supporting Figure 15 absorption at 280 and 390 nm can be drawn. In Supporting Table 5 the concentration is calculated from absorbance values at the corrsponding wavelengths. Comparing the concentration of *nbuty*/DEACM and OmpG does not confer with the 2:1 ratio which results in a labelling efficiency of 87%. This is probably due to the hydrophobic nature of the label and the lower solubility in aqueous media.



Supporting Figure 14: absorption spectrum of OmpG-4.

OmpG-4	unit	280 nm	390 nm
3	L/mol∙cm	78260	20200
Α	a.u.	1.00	0.447
С	μM	12.8	22.2

Supporting Table 5: concentration of OmpG-2 determined at 280 and 390 nm.

### 6. Mass spectrometry

Depending on their concentration and the expected protein masses, 1-10  $\mu$ L of the buffered protein solutions were desalted online using a *Waters ACQUITY H-Class* HPLC-system equipped with a MassPrep column (*Waters*). Desalted proteins were eluted into the ESI source of a Synapt G2Si mass spectrometer (*Waters*) by the following gradient of buffer A (water/0.05% formic acid) and buffer B (acetonitrile/0.045% formic acid) at a column temperature of 60°C and a flow rate of 0.1 mL/min: Isocratic elution with 5% A for two minutes, followed by a linear gradient to 95% B within 8 min and holding 95% B for additional 4 minutes.

Positive ions within the mass range of 500-5000 m/z were detected. Glu-fibrinopeptide B was measured every 45 s for automatic mass drift correction. Averaged spectra were deconvoluted after baseline subtraction and eventually smoothing using MassLynx instrument software with MaxEnt1 extension.

Molecular weight of **OmpG-1** was calculated from its sequential data by using the online tool *ExPASy ProtParam*.

**OmpG-1:** OmpG/ΔL6-M13C-E154C



Supporting Figure 15: mass spectra of OmpG-1 (OmpG/ΔL6-M13C-E154C).

#### OmpG-2: OmpG/\DeltaL6-M13C-E154C/2xDEACM

Molecular weight of **OmpG-2** was calculated by addition of two DEACM residues (see below) to the confirmed mass of **OmpG-1**.



**Supporting Figure 16:** mass spectra of **OmpG-2** (OmpG/ΔL6-M13C-E154C/2xDEACM).

#### **OmpG-4**: OmpG/ΔL6-M13C-E154C/2xn-buty/DEACM

Molecular weight of **OmpG-4** was calculated by addition of two *n*-buty/DEACM residues (see below) to the confirmed mass of **OmpG-1**.

CH<sub>2</sub> O Ô Et<sub>2</sub>N ò

C<sub>20</sub>H<sub>26</sub>NO<sub>4</sub><sup>+</sup> Exact Mass: 344.18563 Mol. Wt.: 344.43045

calculated	<b>32675.37</b> g/mo
found	<b>32674.00</b> g/mo



180130\_IP\_081\_Ot\_2 502 (9.905) M1 [Ev-130033,lt16] (Gs,0.500,1110:1496,1.00,L33,R33); Sm (SG, 2x7.00); Sb (7,10.00 ); Cm (470:526) 1: TOF MS ES+ 100-9.3865

**Supporting Figure 17**: mass spectra of **OmpG-4** (OmpG/ $\Delta$ L6-M13C-E154C/2x*n*-buty/DEACM).

### 7. Spectroscopy

#### Absorption spectra

UV-vis absorption spectra were recorded on an *Evolution 300* Spectrometer (*Thermo Scientific*) using a 70 µL cuvette with 1 cm path length by *Hellma*.

#### **Fluorescence spectra**

Fluorescence spectra were recorded on a *Tecan Plate Reader infinite M200 PRO*. 50 µL of sample solution were pipetted into a black 96 well plate by *Corning*. Samples were excited at 390 nm, emission detected between 420 and 650 nm.

#### Irradiation

During Black Lipid Membrane measurements OmpG hybrids were irradiated by a *Thorlabs* fibercoupled LED M385F1 equipped with a *Thorlabs* LED driver DC2100 at 700 mA, 95 mW for 3-10 min (1000  $\mu$ m fiber).

Samples for SDS PAGE were irradiated with a custom made Multi-LED device equipped with LEDs by *Roithner Lasertechnik* at 385 nm for different time intervals (UVLED-385-310-SMD, 385 nm, 310 mW, 500 mA).

Read out of SDS PAGE fluorescent bands was performed on a *transilluminator BIO View UV light* (USDT-30ML-8R, *Biostep*) with excitation at 365 nm (5 mW).

#### Stability test

In order to proof stability of OmpG against irradiation and possible heating during irradiation a stability test was performed with unmodified **OmpG-1** (see supporting figure 18). In an SDS PAGE a read out was done comparing the different migration behavior of native and denatured OmpG. On the left side samples of **OmpG-1** were irradiated for 0, 1, 3, 5 and 10 min at 385 nm (details see above). On the right side samples were heated for 5 min at 37, 50, 60, 70, 80 and 90 °C. OmpG appears stable against irradiation at 385 nm for 10 min and heating up to 60 °C.



Supporting Figure 18: SDS PAGE of OmpG-1. Stability test against - left side: irradiation; - right side: temperature.

#### 8. Black Lipid Membrane Measurements

OmpG conductances were electrophysiologically monitored by the Black Lipid Membrane (BLM) method. Experiments were performed in 1,2-diphytanoyl-sn-glycero-3-phosphocholine (DPhPC, Avanti polar lipids) as previously described.<sup>3</sup> *N*-decane-solubilised lipid (25 mg/mL) was painted over a 200  $\mu$ m circular hole, separating two chamber compartments (polystyrene cuvette: CP2A, bilayer chamber: BCH-22A, *Warner Instruments*) filled equally with BLM buffer (10 mM Tris/HCl, pH 7.4, 1 M NaCl). The measurements were carried out under red light. Detergent refolded and purified porins (7.5–15  $\mu$ g) were added to one compartment next to the planar lipid layer and a voltage-gradient was imposed across the membrane whilst waiting for insertion of single channels. When a single channel was detected, the voltage was changed between + 60 mV and – 60 mV and the current recorded. The voltage was then set to 40 mV and the sample was exposed to blue light (ThorLabs DC2100, with LED M365L2: 365 nm) for up to 3 minutes. After that, voltage was changed as before and the currents recorded again.

Electric current was recorded using a Multipatch 700B patch-clamp amplifier connected to a Digidata 1440A A/D converter.

Current traces were analyzed using the *Clampex* 10.4 software (*Axon Instruments*), from a data collection frequency of 5 kHz and a sampling rate of 200 Hz. U-I-plots of five measurements where used to determine the conductivity.

## 9. Supporting References

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