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## Enhanced targeting of triple-negative breast carcinoma and malignant melanoma by photochemical internalization of CSPG4-targeting immunotoxins†

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## Abstract

Triple-negative breast cancer (TNBC) and malignant melanoma are highly aggressive cancers that widely express the cell surface chondroitin sulfate proteoglycan 4 (CSPG4/NG2). CSPG4 plays an important role in tumor cell growth and survival and promotes chemo- and radiotherapy resistance, suggesting that CSPG4 is an attractive target in cancer therapy. In the present work, we applied the drug delivery technology photochemical internalization (PCI) in combination with the novel CSPG4-targeting immunotoxin 225.28-saporin as an efficient and specific strategy to kill aggressive TNBC and amelanotic melanoma cells. Light-activation of the clinically relevant photosensitizer TPCS<sub>2a</sub> (fimaporfin) and 225.28-saporin was found to act in a synergistic manner, and was superior to both PCI of saporin and PCI-no-drug (TPCS<sub>2a</sub> + light only) in three TNBC cell lines (MDA-MB-231, MDA-MB-435 and SUM149) and two BRAFV600E mutated malignant melanoma cell lines (Melmet 1 and Melmet 5). The cytotoxic effect was highly dependent on the light dose and expression of CSPG4 since no enhanced cytotoxicity of PCI of 225.28-saporin compared to PCI of saporin was observed in the CSPG4-negative MCF-7 cells. The PCI of a smaller, and clinically relevant CSPG4-targeting toxin (scFvMEL-rGel) validated the CSPG4-

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Conflicts of interest

Dr Anders Høgset is the Chief Scientific Officer of PCI Biotech AS of which he also is a minority shareholder. The other authors have no conflicts to declare.

targeting concept *in vitro* and induced a strong inhibition of tumor growth in the amelanotic melanoma xenograft A-375 model. In conclusion, the combination of the drug delivery technology PCI and CSPG4-targeting immunotoxins is an efficient, specific and light-controlled strategy for the elimination of aggressive cells of TNBC and malignant melanoma origin. This study lays the foundation for further preclinical evaluation of PCI in combination with CSPG4-targeting.

### Introduction

Triple-negative breast cancer (TNBC) and malignant melanoma that do not respond to immune checkpoint inhibition are highly aggressive cancer types that have an unmet need for better treatment options. TNBC constitutes 10–20% of all breast cancers and is known for its particularly unfavourable clinical prognosis with high rates of metastasis.<sup>1-3</sup> TNBC is characterized as oestrogen receptor- (ER<sup>-</sup>), progesterone receptor-(PR<sup>-</sup>) and human epidermal growth factor receptor 2-negative (HER2<sup>-</sup>).<sup>3</sup> Due to the absence of ER and HER2, TNBC does not benefit from the otherwise effective anti-breast cancer drugs targeting these receptors, such as *tamoxifen* and *trastuzumab*, and surgery in combination with cytotoxic therapy remains the only clinically available treatment option.<sup>4</sup>

Pathological complete responses following neoadjuvant and adjuvant chemotherapy have been observed in 17 to 58% of patients.<sup>3</sup> However, patients who exhibit the chemoresistant subtype of TNBC have an exceptionally poor outcome, with a shorter overall survival compared to non-TNBC.<sup>5</sup>

Malignant melanoma, the most lethal form of skin cancer, is one of the deadliest forms of human cancers showing exceptional abilities to metastasize and develop drug resistance.<sup>6</sup> The conventional chemotherapeutic agent dacarbazine induces a response in very few patients (7–20%).<sup>7</sup> Important advances have been achieved with targeted drugs, such as the mutated BRAF inhibitor vemurafenib.<sup>8</sup> Although vemurafenib induces a strong response in melanoma patients with the BRAF<sup>V600E</sup> mutation, acquired resistance will develop leading to relapse over a few months.<sup>9</sup> Recently, immunotherapies with the anti-CTLA4 antibody ipilimumab or antibodies against the PD-1/PD-L1 axis have received considerable attention in malignant melanoma. These agents can lead to durable clinical responses, but only in a fraction of melanoma patients. In addition, severe adverse events including immune related toxicities, particularly for anti-CTL4 are often observed.<sup>10,11</sup> Thus, achieving prolonged therapeutic responses with limited adverse effects and in a larger fraction of patients remains to be a major challenge for malignant melanoma and especially for TNBC.

Chondroitin sulphate proteoglycan 4 (CSPG4, also known as neuron glia antigen-2 (NG2) or melanoma-associated chondroitin sulfate proteoglycan (MCSP)) is a transmembrane protein,<sup>12</sup> expressed in a majority of melanoma lesions and has been associated with melanoma therapy resistance and aggressiveness.<sup>13,14</sup> CSPG4 induces signalling cascades through FAK and MAPK, potentially leading to events promoting survival and chemoresistance, invasion, migration, proliferation and epithelial–mesenchymal transition.<sup>14</sup> CSPG4 has also been shown to be a novel marker of normal epidermal stem cells of the skin.<sup>15</sup> Recently, CSPG4 was shown to be expressed in triple-negative breast cancer, where the antigen was expressed in 32 of the 44 lesions tested, as well as in several cell lines and

tumour cells from pleural effusions.<sup>16</sup> Additionally, CSPG4 was strongly associated with the proposed cancer stem cell phenotype CD44<sup>high</sup>/CD24<sup>low</sup>.<sup>16,17</sup> Restricted distribution of CSPG4 in normal tissues makes CSPG4 a potentially valuable target for tailored TNBC and melanoma treatment.<sup>12</sup>

Photochemical internalization (PCI) is a drug delivery technology based on the use of photosensitizing compounds and targeted illumination, where the illumination induces the release of drugs from endo-lysosomal compartments into the cell cytosol.<sup>18,19</sup> Photosensitizers used with PCI accumulate in endocytic vesicle membranes and subsequent light exposure leads to the generation of reactive oxygen species (ROS), causing membrane disruption and release of the drug of interest into target cells. The PCI technology therefore enhances the endosomal and lysosomal escape of entrapped molecules before they are recycled or degraded following endocytosis. Examples of molecules unable to reach their target within the cell include proteins such as type I ribosome inactivating proteins (RIP), e.g. saporin<sup>20</sup> or gelonin, nucleic acids and nanoparticles.<sup>19</sup> The delivery of toxins linked to antibodies or growth factors using PCI may enhance the specificity of the PCI treatment and has previously been successfully demonstrated for targeting of EpCAM,<sup>21,22</sup> EGFR,<sup>23,24</sup> HER2,<sup>25</sup> VEGFR-1 and -2,<sup>26</sup> CD133,<sup>27-29</sup> CD44,<sup>30</sup> and CSPG4.<sup>31</sup> In addition, the PCI technology has been verified as an efficient strategy to kill therapy-resistant cancer cells.<sup>25,32-36</sup> The PCI technology is currently under evaluation in several clinical trials, where PCI of bleomycin for solid localized tumours was recently shown to be safe and encouraging with respect to clinical outcomes in a Phase I study.<sup>37</sup>

In the present study, we aimed to target and eliminate CSPG4-expressing TNBC and drug-resistant malignant melanoma cells by combining the PCI technology with the novel CSPG4-targeting immunotoxin 225.28-saporin. The efficacy and specificity of the PCI-based targeting of CSPG4 was found promising, and may therefore represent a rational strategy for the elimination of aggressive and therapy-resistant TNBCs and malignant melanoma.

## Materials and methods

#### **Cell culture**

The mammary cancer cell lines MDA-MB-231 (HTB-26), MDA-MB-435 (HTB-129), MCF-7 (HTB-22) and the non-pigmented malignant melanoma A-375 cells were all from ATCC (Manassas, VA, USA). MDA-MB-231 is described as a TNBC<sup>38</sup> and a mesenchymal stem-like cell line<sup>1</sup> with expression of putative cancer stem cell markers such as ALDH<sup>39,40</sup> and CD133.<sup>41</sup> SUM149, a TNBC cell line established from primary inflammatory breast cancer,<sup>42</sup> was obtained from the Department of Medical Genetics (Oslo University Hospital). The metastatic melanoma cell lines, Melmet 1 and Melmet 5, were established in-house as previously described.<sup>43</sup> All cell lines were routinely tested for *Mycoplasma* infections and cell ID. MDA-MB-231, MDA-MB-435, Melmet 1 and Melmet 5 were cultured in RPMI-1640 medium (Sigma-Aldrich, St Louis, MO, USA) supplied with L-glutamine, 10% fetal bovine serum (FBS) (PAA Laboratories, Pasching, Austria), 100 IU ml<sup>-1</sup> penicillin (Sigma-Aldrich) and 100 µg ml<sup>-1</sup> streptomycin (Sigma-Aldrich). MCF-7 was cultured in MEM with Earle's salts (PAA Laboratories), containing phenol red, supplied

with L-glutamine, 10% FBS, 100 IU ml<sup>-1</sup> penicillin/100  $\mu$ g ml<sup>-1</sup> streptomycin and 10  $\mu$ g ml<sup>-1</sup> recombinant human insulin (Sigma-Aldrich). SUM149 cells were cultured in a Ham's F12 nutrient mixture (Sigma-Aldrich) supplied with L-glutamine, 5% FBS, 100 IU ml<sup>-1</sup> penicillin/100  $\mu$ g ml<sup>-1</sup> streptomycin, 1  $\mu$ g ml<sup>-1</sup> hydrocortisone (H0888, Sigma-Aldrich) and 5  $\mu$ g ml<sup>-1</sup> insulin. A-375 cells were cultured in DMEM (Sigma-Aldrich) supplied with 10% FBS and 100 IU ml<sup>-1</sup> penicillin/100  $\mu$ g ml<sup>-1</sup> streptomycin. All cell lines were cultured as monolayers in Nunclon<sup>TM</sup> surface treated tissue culture flasks (NUNC A/S, Thermo Fisher Scientific, Roskilde, Denmark) in incubators at 37 °C with 5% (v/v) CO<sub>2</sub> in a humidified atmosphere.

We wish to stress that there is a controversy in the literature concerning the origin of the MDA-MB-435 cell line.<sup>38,44</sup> However, the focus of this work is the use of the PCI-technology to enhance the efficacy of CSPG4-targeting immunotoxins in both breast (TNBCs) and melanoma cancers and not on the biology of the included cell lines. Hence, independent of the origin of this cell line, the use of the MDA-MB-435 in this study as a CSPG4-expressing cell line is valid.

#### **Reagents and chemicals**

A stock solution of hydrocortisone for cell culture was prepared by dissolving 1 mg hydrocortisone powder per 1 ml absolute ethanol and stored as aliquots at -20 °C. TPCS<sub>2a</sub> (Fimaporfin) was provided by PCI Biotech AS (Lysaker, Norway) and stored protected from light as a stock solution (0.35 mg ml<sup>-1</sup> in 3% polysorbate 80, 2.8% mannitol, 50 mM Tris pH 8.5) at 4 °C. TPPS<sub>2a</sub> (LumiTrans) from PCI Biotech was dissolved in dimethyl sulfoxide to make a 0.35 mg ml<sup>-1</sup> stock solution. AlPcS<sub>2a</sub> (disulfonated aluminum phthalocyanine) provided by Frontier Scientific (Logan UT, USA) was dissolved in 0.1 M NaOH and thereafter diluted in phosphate-buffered saline, pH 7.5, to a final concentration of 5 mg ml<sup>-1</sup>. TPPS<sub>2a</sub> and AlPcS<sub>2a</sub> were stored protected from light as aliquots at -20 °C. LysoTracker Green (DND-26) was obtained from Life Technologies (Eugene, Oregon, USA) and stored as single-use aliquots at -20 °C. Streptavidin-conjugated Alexa Fluor 488 (Life Technologies) was stored at 4 °C according to the manufacturer's instructions.

## Preparation of 225.28-saporin and scFvMEL-rGel

CSPG4-specific mouse mAb 225.28, an IgG2a, was developed and characterized as described.<sup>45</sup> mAb was purified from ascitic fluid by affinity chromatography on a Protein G column. mAb was biotinylated as described.<sup>46</sup> To obtain the immunotoxin 225.28-saporin, streptavidin-saporin (Advanced Targeting Systems, San Diego, CA, USA), with an average of 2.5 molecules of saporin per molecule of streptavidin, was combined with biotinylated 225.28 mAbs at a biotin : streptavidin ratio of 4 : 1 by mixing and incubating at room temperature for 15–20 minutes. The reaction was assumed to be complete due to the high affinity of streptavidin to biotin. In all calculations and presentations, the molar concentration of the immunotoxin was set equal to the concentration of streptavidin-saporin in the mixture. The scFvMEL-rGel targeted toxin was expressed and purified as previously described.<sup>47</sup>

#### Flow cytometry investigation of antigen expression

CSPG4 expression was investigated using live cell flow cytometry. Monolayer cells were harvested using EDTA solution (0.54 mM) to avoid enzymatic degradation of the antigen. Subsequently, cells were incubated for 30 min in 1  $\mu$ g ml<sup>-1</sup> 225.28-biotin, followed by incubation in 0.5  $\mu$ g ml<sup>-1</sup> streptavidin-Alexa Fluor® 488 (Life Technologies) for 20 minutes, and were then thoroughly washed in PBS prior to examination. Samples were immediately analyzed on a BD LSR II flow cytometer (BD Biosciences) using a 488 nm laser, 505 nm dichroic filter and a 525/50 band pass excitation filter. Doublet cells, debris and unviable cells were excluded from the analyses based on forward/side scatter.

#### Intracellular localization studies by confocal and fluorescence microscopy

To validate the flow cytometry data, specific binding of the CSPG4-targeting antibody was compared by 4 h incubation of 225.28-Alexa488 in the CSPG4<sup>high</sup> MDA-MB-435 cells with the CSPG4<sup>low</sup> MCF-7 cells by *epi*-fluorescence microscopy. The cells were washed with cold  $1 \times$  PBS containing Ca<sup>2+</sup> and Mg<sup>2+</sup> prior to the fluorescence microscopy using a 63× oil-immersion objective with an Axioimager Z1 Microscope (Carl Zeiss, Oberkochen, Germany). A 450–490 nm band pass excitation filter, 495 nm dichroic mirror and a 500–550 band pass emission filter were used for the measurement of the Alexa-488-labeled 225.28 Ab. The micrographs were processed by use of the Axiovision Software (Carl Zeiss).

The intracellular localization of TPCS<sub>2a</sub> and the 225.28 mAb was investigated by confocal microscopy of live cells. MDA-MB-435 cells were seeded directly on 0.17 (±0.01) mm thick coverslips (Assistent, Sondheim, Germany) in 4-well dishes (NUNC A/S). To label biotinylated mAbs, 18 µg ml<sup>-1</sup> 225.28 mAb was mixed with 1.8 µg ml<sup>-1</sup> streptavidin-Cy3 (Jackson ImmunoResearch Europe, Newmarket, United Kingdom), followed by 15-20 minutes of incubation at room temperature as described in the Results section. For co-localization studies, cells were co-incubated with the Cy3-labelled 225.28 and 1  $\mu$ g ml<sup>-1</sup> TPCS<sub>2a</sub> for 18 h, followed by a 4-6 h chase prior to examination. LysoTracker® Green (Life Technologies) was added to a concentration of 0.5-5 µM, 30-120 minutes before examination as described in the Results section. The cover slips were washed briefly in PBS, and placed in 35 mm glass bottom MatTek dishes (MatTek Corp. Ashland, MA, USA) for live cell examination with a Zeiss LSM 710 confocal microscope (Carl Zeiss Microscopy GmbH, Jena, Germany), equipped with an Ar-Laser Multiline (458/488/514 nm), a DPSS-561 10 (561 nm), a Laser diode 405-30 CW (405 nm), and a HeNe-laser (633 nm). The objective used was a Zeiss C-Apochromat 40×/1.20 Water Imm DIC III. Image processing was performed with basic software ZEN 2009 (Carl Zeiss MicroImaging GmbH, Jena, Germany) and Photoshop CS2 (Adobe, Mountain View, CA).

#### Photochemical internalization (PCI) and cytotoxicity assays

Prior to each experiment the photochemical dose (light dose and concentration of TPCS2a) inducing around 50% reduction of cell viability was found for each cell line. Based on this, TPCS<sub>2a</sub> was diluted in medium to a final concentration of 0.05  $\mu$ g ml<sup>-1</sup> for SUM149 cells, 0.6  $\mu$ g ml<sup>-1</sup> for MCF-7 cells or 0.2  $\mu$ g ml<sup>-1</sup> for the additional cell lines. Cells were incubated in TPCS<sub>2a</sub> for 18 h, after which they were washed twice with fresh medium and chased for 4 h to allow the photosensitizer to be internalized and cleared from the

plasma membrane. Treatment with TPCS<sub>2a</sub> and light as described above, in the absence of toxin or immunotoxin, is referred to as PCI-no-drug (photochemical treatment alone = TPCS2a + light only). With PCI treatment, cells were co-incubated with TPCS<sub>2a</sub> and toxin or immunotoxin for 18 h prior to chase. The concentration of 225.28-saporin and streptavidin-saporin was 0.1 nM in MDA-MB-231, Melmet 1 and Melmet 5 cells and 1 nM in MDA-MB-435, SUM149 and MCF-7 cells. With an average of 2.5 molecules of saporin per complex, this is equivalent to 0.25 nM and 2.5 nM saporin, respectively. With the clonogenic assay, MDA-MB-435 cells were treated with 0.1 nM 225.28-saporin and streptavidin-saporin. The concentration of scFvMEL-rGel and rGelonin was 1 nM in the MDA-MB-231 and SUM149 cell lines, and 5 nM in the MCF-7 cell line. Cells were exposed to light using a LumiSource® light source (PCI Biotech AS), consisting of four light tubes with a peak emission at 435 nm, irradiance 12.6 mW cm<sup>-2</sup>. 1 J cm<sup>-2</sup> = 75 s light exposure.

The cell viability was assessed by the MTT assay as described earlier.<sup>32</sup> For MTT-based assays, cells were seeded on 96-well plates (NUNC A/S) at 13 000 cellsper well for MDA-MB-231 cells or 10 000 cellsper well for MDA-MB-435 cells 6 h prior to drug incubation, alternatively 8000 cellsper well for MCF-7, SUM149 and Melmet 5 cells, or 4000 cellsper well for MCF-7, SUM149 and Melmet 5 cells, or 4000 cellsper well for MCF-7, SUM149 and Melmet 5 cells, or 4000 cellsper well for McF-7, SUM149. The optical density at 570 nm was recorded using a Powerwave XS2 microplate spectrophotometer (BioTek, Winooski, VT, USA). Following normalization to blanks on each plate, the cell viability was calculated relative to the absorbance of untreated cells.

For the clonogenic survival assay, a modified version of Franken *et al.* was used.<sup>48</sup> In brief, the cells were seeded in 6-well plates (NUNC) at 1000 cellsper well for MDA-MB-231 and MDA-MB-435 cells, and 2000 cellsper well for SUM149 and Melmet 5 cells. In brief; 9–14 days post treatment, colonies were washed once in saline, fixed by absolute ethanol, stained by a methylene blue solution and counted manually. Colonies consisting of less than 50 cells were excluded.

#### Immunofluorescence

Melmet 1 and Melmet 5 cells grown on glass coverslips were fixed in 4% paraformaldehyde for 15 min on ice before staining with mouse anti-MITF (Thermo Scientific), or rabbit anti-Axl (Cell Signaling), both diluted 1 : 100 in PBS/0.05% saponin. After overnight incubation at 4 °C, the samples were stained for 1 h with donkey anti-mouse 488 or donkey anti-rabbit 549 (both Jackson ImmunoResearch) diluted 1 : 1500 or 1 : 400, respectively. The coverslips were mounted in Prolong Gold mounting medium (Life Technologies) containing DAPI and examined using a laser-scanning confocal microscope (LSM710; Carl Zeiss), equipped with a Plan-Apochromat ×63/1.4 Oil DICIII objective. Image processing and visualization were performed by using the ZEN Light 2011 software.

#### Drug response in melanoma cell lines

The growth inhibitory effect of dacarbazine (10, 50 and 100  $\mu$ g ml<sup>-1</sup>) and vemurafenib (0.33, 0.66, 3.3 and 6.6  $\mu$ M) was measured by using CellTiter 96 Aqueous One solution (MTS assay, Promega, Madison, WI, USA). Melmet 1 or Melmet 5 was seeded in 96-well plates and treated with the indicated drugs. Seventy two hours after treatment CellTiter 96

Aqueous One Solution was added to the wells and the absorbance was measured at 490 nm after approximately 2 hours using a micro plate reader (Victor2 1420 Multilabel Counter, PerkinElmer). The viability of the treated cells is reported as the percentage of viable cells relative to untreated control cells. Experiments were performed in four parallels and repeated at least in three independent experiments for each treatment condition.

#### In vivo PCI of scFv-rGel

Female athymic nude mice of the Hsd: Athymic nude Foxn1nu strain were bred at the Department of Comparative Medicine at Oslo University Hospital, Radiumhospitalet. All procedures involving mice were carried out in agreement with the protocols approved by the animal care unit at the Norwegian Radium Hospital—Oslo University Hospital, under control by the National Ethical Committee's guidelines on animal welfare. The mice were on average 20–25 g (6–8 weeks old) at the start of the experiment. The mice were kept under specific pathogen-free conditions. Water and food were given *ad libitum*. Exponentially growing A-375 cells were treated and transplanted s.c. as previously described.<sup>31</sup> Mice with 75 mm<sup>3</sup> (±25 mm<sup>3</sup>) A375 tumours were subjected to only one PCI of scFvMEL-rGel treatment and tumour growth was measured twice per week as previously described.<sup>31</sup>

#### Statistical analysis

Each experiment evaluating PCI-induced cytotoxicity was performed at least three times (*n* = 3-9 for each cell line) and the same trend in cytotoxic response was observed. Due to the multiple variables with the PCI protocol, such as photosensitizer concentration, toxin concentration, light source efficiency, the cytotoxicity data presented in Fig. 3, 4E-G and 5A (and Fig. ESI-4, 5 and 7<sup>+</sup>) stem from a single, representative experiment, while ESI-6<sup>+</sup> shows the average (3-9 replicated experiments) targeting ratio of PCI of saporin/PCI of 225.28-saporin for each individual experiment: briefly, we calculated the ratio between the time of light exposure needed to kill/reduce viability of 50% of the cells (LD50) after PCI of saporin (toxin) and the time of light exposure needed to kill/reduce viability of 50% of the cells after PCI of 225.28-saporin (immunotoxin) giving a targeting ratio for each cell line with standard error (SE). By using SigmaPlot 12.5 software, the two-sided Student's t-test was applied on technical replicates within each experiment, where the samples were found normally distributed by the Shapiro-Wilk Normality Test. Where normality test failed, a Mann–Whitney Rank Sum Test was used. The null hypothesis was rejected when P was <0.05. The synergistic toxicity in the 6 CSPG4 + cell lines was evaluated by comparing the theoretical additive effect of combining PCI (no drug) and IT with the observed combination (PCI of IT).<sup>49-51</sup> It was assumed that PCI(no drug) and the IT had distinct mechanisms of actions and the calculated additive effect was therefore assessed by the product of the survival fractions (SF) of the two treatments;

$$\begin{split} SF_{add} &= SF_{PCI(no\ drug)} \times SF_{IT} \text{ and } ln\ SF_{add} \\ &= ln\ SF_{PCI(no\ drug)} + ln\ SF_{IT} \,. \end{split}$$

The difference in logarithm (DL) of the SF<sub>add</sub> and SF<sub>PCI IT</sub> was calculated;

<sup>&</sup>lt;sup>†</sup>Electronic supplementary information (ESI) available. See DOI: 10.1039/c7pp00358g

 $DL = \ln SF_{add} - \ln SF_{PCI IT}$ 

and the standard error of DL was assessed by;

SE (DL) = 
$$\sqrt{(\text{SE}(\ln \text{SF}_{add})^2) + \text{SE}(\ln (\text{SF}_{PCLIT})^2)}$$

where

$$SE(\ln SF_{add}) = \sqrt{(SE(\ln SF_{PCI(no drug)})^2) + SE(\ln(SF_{IT})^2)}$$

A synergistic combination of the two treatments (PCI of IT) was defined as a significant positive DL as evaluated by a two-sided student *t* test

 $T = \frac{DL}{SE}(DL)$ df = 8

The data were considered significant when p = 0.05. All synergy calculations were established at a PCI(no drug)-dose reducing the viability to 20, 30 or 50% (LD<sub>50</sub> for the TNBC and LD<sub>30</sub> or LD<sub>30</sub> for Melmet 1 and Melmet 5, respectively) and extrapolation was used in cases where the LD<sub>50/30/20</sub> was not reached. The SF was set to 0.001 (99.9% toxicity) in experiments where no MTT signals or colonies were detected.

#### Results

#### CSPG4 expression in breast cancer and melanoma cell lines

The CSPG4 surface expression levels in four breast cancer and two melanoma cell lines were evaluated by flow cytometry (Fig. 1). The median fluorescence intensity (MFI) was determined in cells stained with Alexa488-225.28-mAbs, compared to the MFI in unstained control cells, and a fluorescence ratio was calculated (presented in Fig. 1). The MDA-MB-435 cells showed the highest expression of CSPG4 of all TNBC cell lines tested, with a fluorescence ratio of 42.0 (Fig. 1), and where 97% of the cells expressed the antigen (Fig. ESI-1†).

In the TNBC cell lines MDA-MB-231 and SUM149 the MFI ratio was 3.2 and 2.0, respectively. These cells showed a heterogeneous expression of CSPG4, where approximately 37% of MDA-MB-231 and 14% of SUM149 cells were strongly positive (Fig. ESI-1†). The luminal breast cancer cell line MCF-7 showed a negligible expression of CSPG4, with a fluorescence ratio of 1.3 and therefore further referred to as CSPG4-negative. The whole cell populations of the malignant melanoma cell lines Melmet 1 and Melmet 5 expressed high levels of CSPG4, with the fluorescence ratios of 26.4 and 241.7, respectively (Fig. 1).

The CSPG4-targeting mAb 225.28 binds to CSPG4-positive cells and is captured in endosomes/lysosomes together with the PCI photosensitizer TPCS2a.

To compare the binding of 225.28mAb to the cell surface of CSPG4-positive MDA-MB-435 cell line *versus* the CSPG4-negative MCF-7, the cells were incubated for 4 h with Alexa Fluor-labelled 225.28mAb (225.28-Alexa488) and observed by *epi*-fluorescence microscopy (Fig. 2A-C).

While a strong fluorescence signal was detected along the surface of MDA-MB-435 cells, CSPG4-targeting 225.28-Alexa488 could not be detected on the surface of the MCF-7 cell line (Fig. 2D-F). This indicates that the surface binding of 225.28-mAb was specific and depended on the CSPG4 level. The majority of the cell were completely negative, however, in some MCF-7 cells granular fluorescence was observed (Fig. 2D), which may be due to the aggregation and non-specific binding of the complex on the surface of the cells.

Drug delivery by PCI is dependent on co-localization of the photosensitizer and drug of interest within *endo*–lysosomal compartments. To study the intracellular localization of the 225.28 mAb-delivered moiety, 225.28 mAb was linked to streptavidin-Cyanine 3, referred to as 225.28-Cy3, and used as a surrogate for the immunotoxin. The intracellular localization of 225.28-Cy3 and the photosensitizer TPCS<sub>2a</sub> was investigated by confocal microscopy in the CSPG4-positive MDA-MB-435 cell line (Fig. 2G-J and ESI-2†). Following 18 h incubation and 4 h chase, TPCS<sub>2a</sub> and the 225.28-Cy3 complex was detected as intracellular fluorescent puncta. LysoTracker Green was used to stain endosomes and lysosomes (Fig. 2 and ESI-3†).

#### PCI of 225.28-saporin induces a specific and strong synergistic effect in TNBC cell lines

To generate the CSPG4-targeted immunotoxin, we combined biotinylated 225.28 with streptavidin-saporin, further referred to as 225.28-saporin.

It is important that the binding of the targeting moiety (in this respect: biotinylated 225.28) to streptavidin-saporin do not destroy the enzymatic activity (inhibition of protein synthesis) of saporin. Thus, we evaluated the ribosome-inactivating protein (RIP) activity of 225.28-saporin *versus* streptavidin-saporin in a cell-free system. No substantial difference or RIP activity between the targeted and non-targeted toxins was observed (Fig. ESI-4†).

To evaluate the potential of PCI to enhance the cytotoxicity of 225.28-saporin in TNBC cell lines, we combined light,  $TPCS_{2a}$  and the immunotoxin and measured cell viability by the MTT assay (Fig. 3A).

In the CSPG4-positive MDA-MB-435 cell line, PCI of 225.28-saporin with a light dose of 120 s resulted in a total elimination of the MTT signal (Fig. 3A). Incubation of the 225.28 mAb alone for 18 h (as in the PCI protocol) did not induce any effects on cell viability (Fig. ESI-5A†). Blocking CSPG4 with excess 225.28 mAb abolished the antigen-targeting effect of PCI with 225.28-saporin as the difference of cytotoxicity compared to PCI of saporin was not statistically significant (P = 0.62, Fig. ESI-5B†). In the MDA-MB-231 cell line, the PCI of 225.28-saporin resulted in a strong attenuation of cell viability despite the 10-fold reduction of the concentration of the 225.28-saporin immunotoxin compared with the concentration used for the PCI-based targeting of the MDA-MB-435 cells (Fig. 3B).

To evaluate the long-term effect on cell survival, we performed the clonogenic assay. In accordance with the MTT data, a reduction of colony-forming abilities by close to 99% was achieved in all three CSPG4-positive TNBC cell lines treated by PCI of 225.28-saporin (Fig. 3D-F). PCI of 225.28-saporin induced a synergistic cytotoxic effect in all three cell lines (Table ESI-1†).

In all three TNBC cell lines, there was a substantial and statistically significant difference when comparing PCI of the immunotoxin 225.28-saporin and PCI of the non-targeted toxin streptavidin-saporin (Fig. ESI-6†) or PCI-no-drug. The effect of the single treatments in the TNBC cell lines was minor compared to the effect of PCI of 225.28-saporin. PCI-no-drug induced a modest cytotoxic effect across the different cell lines (Fig. 3). There was no marked cytotoxic effect of the immunotoxin combined with TPCS<sub>2a</sub> in the absence of light (shown as 0 s. light exposure in Fig. 3), except for the MDA-MB-435 cell line, where the clonogenic assay revealed a 50% reduction in clonal cell survival (Fig. 3E). In the CSPG4-negative MCF-7 cell line, the PCI efficacy of 225.28-saporin *versus* streptavidin-saporin was indistinguishable (Fig. 3G). The observed difference between PCI of immunotoxin and non-targeted toxin in CSPG4-positive cell lines, but not the CSPG4-negative cell line, indicates that the PCI-enhanced toxicity of the immunotoxin depends on CSPG4 expression.

# PCI of 225.28-saporin is highly cytotoxic in BRAF<sup>V600E</sup> mutated MITF<sup>low</sup>/AXL<sup>high</sup> melanoma cells

To evaluate the PCI efficacy in another aggressive cancer type, we used two highly aggressive BRAFV600E mutated malignant melanoma cell lines, Melmet 1 and Melmet 5, originally derived from subcutaneous and lymph node metastases, respectively.<sup>43,52</sup> Melmet 5 exhibits a strong nuclear staining of Microphthalmia-associated transcription factor (MITF), the main regulator of melanocytic differentiation, while the MITF level in Melmet 1 was very low (Fig. 4A). The CSPG4 surface experiments including incubation with only secondary antibody-Alexa488 (resulting in no signal, data not shown) we ruled out that the low MITF signal of the Melmet 1 cells are neither unspecific staining nor autofluorescence or background staining. In contrast, the level of AXL receptor tyrosine kinase was high in Melmet 1 and negligible in Melmet 5 (Fig. 4B). This indicates that Melmet 1 and Melmet 5 represent two opposite phenotypes, known as MIT<sup>Flow</sup>/AXL<sup>high</sup> and MITF<sup>high</sup>/AXL<sup>low</sup>, respectively.<sup>53</sup> It was reported previously that the MITFlow/AXLhigh phenotype is associated with more aggressive, stem-like melanoma cells<sup>54,55</sup> showing higher resistance to therapy.<sup>53,56</sup> In accordance with these reports, we observed that Melmet 1 cells were more resistant to clinically used drugs, the chemotherapeutic agents dacarbazine, and the mutated BRAF inhibitor vemurafenib than Melmet 5 (Fig. 4C and D). Both Melmet 1 and Melmet 5 melanoma cell lines were sensitive to 225.28-saporin + TPSC<sub>2a</sub> in the absence of light. However, the cytotoxic efficacy was strongly enhanced by PCI in a light-dose dependent manner (Fig. 4E-G). The MTT assay revealed that the cell viability was below detectable levels following a 180 s and 300 s light exposure in Melmet 1 and Melmet 5, respectively (Fig. 4E and F). This suggests that the drug resistant Melmet 1 was more sensitive to PCI of 225.28-saporin than Melmet 5, even though the sensitivity to TPCS<sub>2a</sub> + light (PCI-no-drug) and PCI of streptavidin-saporin was comparable in both cell lines, and despite the fact that Melmet 1 expressed approximately 1/10 of the CSPG4

antigen compared to the Melmet 5 cells (Fig. 1). Furthermore, the efficacy of PCI of 225.28-saporin to inhibit colony formation was found superior to PCI-no-drug and PCI of streptavidin–saporin in Melmet 5 cells (Fig. 4G). Of notice, treatment with the immunotoxin alone resulted in relatively high cytotoxic responses in the Melmet cell lines (cells with very CSPG4 expression), which may explain that synergistic responses were not obtained when assessing synergy at  $LD_{50}$  doses. However, at higher light-doses ( $LD_{70-80}$ ) the PCI of 225.28-saporin resulted in strong synergistic responses (Table ESI-1†). The clonogenic assay could not be performed in Melmet 1, which shows a highly motile mesenchymal phenotype and is therefore not compatible with this assay. Collectively, these data indicate that 225.28-saporin combined with PCI is an effective strategy for targeting malignant melanoma cells that show resistance or only partial response to clinically-used drugs.

## PCI-based targeting of CSPG4 by the recombinant fusion toxin scFvMEL-rGel in TNBC cells and a malignant melanoma xenograft

To validate this PCI-based CSPG4 targeting concept and to ensure that the treatment response to 225.28-saporin was not influenced by the streptavidin-biotin linkage or the type of toxin we used; we tested the PCI of the fusion toxin scFvMEL-rGel. scFvMEL-rGel consists of the scFv fragment of the 225.28 mAb, recombinantly fused to the ribosome-inhibiting protein gelonin,<sup>47</sup> which belongs to the same RIP type I family as saporin. A similar cytotoxic response was observed with PCI of scFvMEL-rGel as with PCI of 225.28-saporin in the TNBC cell lines MDA-MB-231 and SUM149 and in the CSPG4-negative control cell line MCF-7 (Fig. ESI-7<sup>†</sup>).

The efficacy of PCI of scFvMEL-rGel was also validated in another malignant melanoma cell line, A-375 (Fig. 5A + ESI-8†), which represent the MITF<sup>low</sup>/AXL<sup>high</sup> phenotype as Melmet 1.<sup>56</sup> In these experiments we used the PCI photosensitizer AIPcS<sub>2a</sub>, which also localize to lysosomes.<sup>57</sup> Here we demonstrate that AIPcS<sub>2a</sub>, indeed is released to the cytosol of A375-GFP cells after light exposure (Fig. 5B and C).

In addition, the ability of PCI of scFvMEL-rGel (CSPG4-targeting toxin) to delay tumour growth was demonstrated in the A-375 xenograft model (Fig. 5F and G + Fig. ESI-9† for experimental design overview). While non-treated A-375 tumours (n = 14) had a >12-fold increase in tumour size at day 15, only one treatment with PCI of scFvMEL-rGel (n = 12) resulted in a substantial tumour growth delay where there was no statistical significant (P = 0.39) increase in the tumour size at day 15 compared to day 0. Altogether, the PCI of the scFvMEL-rGel group (n = 12) had a response rate of 83% of which 1/3 of the mice achieved complete responses (Fig. 5G), one mouse had a tumour that did not display aggressive growth (followed until day 91) and another long surviving mouse which reached the study endpoint at day 135. At day 20, the anti-tumour effect of PCI of scFvMEL-rGel was significantly (P < 0.001) better than PCI-no-drug (~7-fold increase at day 20, n = 14) or scFvMEL-rGel (~9-fold increase at day 20, n = 11) treatment groups.

## Discussion

CSPG4 is recognized as an attractive target in cancer therapy due to its very low or no expression in normal tissues and overexpression in several cancer types,<sup>12</sup> including

triple-negative breast cancer (TNBC),<sup>16</sup> malignant melanoma,<sup>13</sup> malignant mesothelioma,<sup>58</sup> sarcomas<sup>12</sup> and squamous cell carcinoma of the head and neck,<sup>16</sup> and known to correlate with radio- and chemoresistance and poor clinical outcome.<sup>59,60</sup> In this study, we developed the CSPG4-targeting model immunotoxin, 225.28-saporin and evaluated its use in combination with the clinically relevant photosensitizer TPCS2a (fimaporfin) for PCI-based targeting of TNBC and malignant melanoma. In addition, we studied the anti-cancer effect of CSPG4-targeting toxin, scFvMEL-rGel delivered by PCI both *in vitro* and *in vivo*.

PCI of 225.28-saporin was a highly cytotoxic treatment in all five tested CSPG4 expressing cancer cell lines, with no signs of resistance to the treatment. Interestingly, in malignant melanoma we observed a stronger response to PCI of 225.28-saporin in Melmet 1 cells than in the Melmet 5 cells, despite the lower sensitivity of Melmet 1 to conventional drugs used in the clinic and the fact that these cells have a 10-fold reduced expression of CSPG4 compared to Melmet 5. As shown in this study and reported previously,<sup>52</sup> these two malignant melanoma cell lines display opposite phenotypes: mesenchymal-like/invasive de-differentiated (Melmet 1 as well as A-375) and non-invasive differentiated (Melmet 5).<sup>52</sup> The mesenchymal-like phenotype has been linked to drug resistance, particularly to targeted therapies.<sup>53,56</sup> Hence, the efficient eradication of Melmet 1 and A-375 cells by PCI of CSPG4-targeting immunotoxins suggests that resistance mechanisms characteristic for this phenotype might not interfere with PCI-based therapies. The PCI approach might therefore be effective in targeting cancer cells in a mesenchymal-like de-differentiated state. This observation is also highly relevant in breast cancer, where malignant cells can acquire mesenchymal properties due to epithelial-mesenchymal transition (EMT), which is also linked with drug resistance.<sup>61</sup> The TNBC cell lines MDA-MB-231 and SUM149 have been shown to consist of stem cell-like subpopulations capable of self-renewal, increased tumourigenicity and resistance to radio- and chemotherapy.<sup>62-64</sup> Furthermore, MDA-MB-231 and MDA-MB-435 have been sub-classified as claudin-low, while the SUM149 cell line has been shown to have a claudin-low subpopulation.<sup>65</sup> Claudin-low TNBC cells are low in differentiation markers, high in EMT markers and show stem-cell like properties.<sup>65</sup> This is especially interesting, given that PCI of the CSPG4-targeting immunotoxin was effective even in MDA-MB-231 and SUM149, where only a fraction of cells were expressing CSPG4. This may indicate purging of a subpopulation important for cell-renewing and long-term survival. Taken together, these data underscores the high potential of targeting CSPG4 by PCI in aggressive cancers as triple-negative breast cancer and malignant melanoma, and implies a novel approach for targeting tumour cells with stem cell/-mesenchymal properties.

The PCI of 225.28-saporin induced a stronger cytotoxic effect compared to PCI of streptavidin–saporin. This was especially apparent in cell lines with a high CSPG4 expression, *i.e.* MDA-MB-435 and the Melmet cell lines. Also in the absence of light, 225.28-saporin exhibited a moderate to high cytotoxic effect in cell lines with high CSPG4 expression levels. The difference in cytotoxic responses to streptavidin–saporin *versus* 225.28-saporin is most likely explained by different cellular uptakes, as in the cell-free system no difference in the ribosome inhibiting activity was observed (Fig. ESI-4†). The immunotoxin 225.28-saporin is most probably taken up into cells by receptor-mediated endocytosis after binding to CSPG4. This conclusion is supported by

the cellular localization studies, showing that the 225.28-targeting moiety binds to the cell surface dependent on CSPG4-levels, and eventually accumulates in the *endo*–lysosomal compartment. In contrast, streptavidin–saporin, which is not known to bind to any cell surface receptor, is most likely taken up through the slower and less efficient process of pinocytosis. There might be also other reasons for differences in cytotoxicity between 225.28-saporin and streptavidin–saporin, *e.g.* a different route of intracellular trafficking following endocytosis and/or inhibition of FAK and ERK pathways by the 225.28 mAb itself.<sup>16</sup> The latter is, however, less likely as treatment with the 225.28 mAb did not influence the photochemical treatment alone (PCI-no-drug) induced cytotoxic response (Fig. ESI-5†). An elimination of the PCI based targeting of CSPG4 was obtained by blocking target cells with excess 225.28 mAb. In addition, no enhanced targeting effects of PCI of 225.28-saporin was achieved in the CSPG4-neg MCF-7 cells. Altogether, the survival data demonstrates the CSPG4-dependent specificity of 225.28-saporin and implies PCI of 225.28-saporin as a promising targeted treatment option.

The concept of TPCS2a-PCI-enhanced delivery of a CSPG4-targeting toxin should be highly relevant for clinical applications since it offers; (1) the advantage of a high tumour selectivity of the therapeutics due to preferential accumulation of the photosensitizer in tumour tissues, primarily by the enhanced permeability and retention phenomenon;<sup>66,67</sup> (2) a mAb or scFv-fragment, which is linked to the ribosome-inactivating toxin, ensures tumour cell targeting; (3) a spatiotemporal controlled laser-activation of the PCI-photosensitizer followed by endosomal escape and activation of the entrapped toxin. Furthermore, preclinical data indicates that a single injection of photosensitizer and toxin is sufficient for the anti-tumor effect with PCI,<sup>19</sup> reducing adverse effects and the formation of neutralizing antibodies, which is often a challenge with *in vivo* applications of immunotoxins.<sup>26</sup> Of relevance, also clinical data confirm that only one session with PCI-drug combination is sufficient to obtain strong anti-tumour responses.<sup>37</sup>

The molecular size of 225.28-saporin (>700 kDa) is most probably not optimal for targeting cancer cells in solid tumours, due to the limited transport across vascular endothelium and high interstitial fluid pressure within the tumours, as well as accumulation in organs presenting with a fenestrated vasculature. To overcome this obstacle, a smaller immunotoxin, *e.g.* scFvMEL-rGel (a 56 kDa) should be used in a preclinical/clinical setting. The *in vivo* studies with scFvMEL-rGel demonstrated the feasibility of the PCI-based targeting of CSPG4-expressing melanoma tumours. In immunocompromised mice lacking mature T-cells, PCI of scFvMEL-rGel delayed the tumour growth and was superior to all control treatments, including scFvMEL-rGel alone, PCI-no-drug or the combination of PCI-photosensitizer + scFvMEL-rGel without light. The superior anti-cancer effect of PCI was also reflected by enhanced animal survival in the PCI group compared to all other treatment groups as demonstrated previously.<sup>31</sup> Hence, the PCI- and CSPG4-based targeting approach may have great potential against CSPG4-expressing tumours.

Late stage TNBC and melanoma are often metastasized at presentation. This represents a problem for focal therapies. However, it has been demonstrated that by photodynamic therapy (PDT, photosensitizer + light) it is possible to stimulate naïve and memory T-cells for the attack on tumour cells distant from the original site of treatment.<sup>68</sup> As a proof-of-

concept we recently showed that PCI is able to potentiate ovalbumin-based model cancer vaccines,<sup>69,70</sup> and to enhance HPV peptide-based vaccines resulting in strong activation of tumour-suppressing CD8 killer cells, both in therapeutic and prophylactic settings.<sup>71</sup> The PCI method has therefore potential to be used in more advanced and metastasized cancers. Another challenge for light-based treatment strategies in melanotic melanoma is limited penetration of light into pigmented tissue.<sup>72</sup> However, as seen in this work, cytotoxic effects induced by PCI are substantially larger compared to PCI without drug-combination (PCI-no-drug), and requires a lower light dose to achieve anti-tumour effects in amelanotic melanoma. This indicates that the limited light-penetration should be less of a problem with PCI compared to photochemical treatment alone. In either case, localized non-pigmented melanomas should be a clinically relevant indication for PCI.

## Conclusions

PCI of CSPG4-based immunotoxins is a highly efficient and selective targeting approach for site-directed killing of drug-resistant cells of TNBC and malignant melanoma origin, as shown *in vitro* and *in vivo*. These results lay the foundation for further preclinical evaluation of this targeting approach for the treatment of aggressive melanomas and TNBCs.

### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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#### Fig. 1.

Cell surface expression of CSPG4 evaluated by flow cytometry. Cells stained with Alexa Fluor 488-linked 225.28 mAb are shown in blue. Red histogram represents unstained cells. The ratio of median fluorescence intensity of stained cells *versus* unstained cells is provided for each cell line. Figure shows representative histograms out of three independent experiments.



#### Fig. 2.

Specific binding of mAb 225.28 and co-localization of TPCS<sub>2a</sub>, mAb 225.28-Cy3 and LysoTracker. (A–F) MDA-MB-435 and MCF-7 cells were incubated with fluorochromelinked 225.28 for 4 h prior to examination by *epi*-fluorescence microscopy. (G–K) MDA-MB-435 cells incubated for 18 h in medium containing (G) TPCS2a (1  $\mu$ g ml<sup>-1</sup>) and (H) 225.28 mAbs labeled by cyanine 3 (Cy3) (30 nM), washed and chased in fresh medium for 4 h prior to examination by confocal microscopy. (I) LysoTracker Green (1  $\mu$ M) was added 1 h prior to investigation. (J) Merge of TPCS<sub>2a</sub>, 225.28-Cy3 and LysoTracker Green. (K) Corresponding DIC micrograph.



#### Fig. 3.

Specific and efficient cytotoxic effects induced by PCI of 225.28-saporin in CSPG4expressing TNBC cell lines. (A), MDA-MB-435, (B) MDA-MB-231 or (C) SUM149 cells were incubated with TPCS<sub>2a</sub> (PCI no drug), or TPCS2a in combination with streptavidin– saporin or 225.28-saporin (PCI), chased in drug-free medium for 4 h and illuminated as indicated on the *X*-axis. Relative cell viability was examined using the MTT assay 48 h post illumination. The concentration of TPCS<sub>2a</sub> was 0.2  $\mu$ g ml<sup>-1</sup> for MDA-MB-231 and MDA-MB-435 and 0.05  $\mu$ g ml<sup>-1</sup> for SUM149. The concentration of toxins was 0.1 nM for MDA-MB-231 and 1 nM for MDA-MB-435 and SUM149. (D–F) Clonogenic survival in (D) MDA-MB-435 (E) MDA-MB-231 or (F) SUM149 cells assessed 10–14 days post light exposure. (G) CSPG-negative MCF-7 control cells after treatments as described in A–C. Each figure shows a single representative experiment out of three independent experiments. Bars = SD of three technical replicates. (H) Experimental protocol of PCI of 225.8-saporin is the same as for PCI of saporin. For PCI no drug (TPCS<sub>2a</sub> + light only), TPCS<sub>2a</sub> was incubated alone.



#### Fig. 4.

PCI of 225.28-saporin is highly cytotoxic in drug-resistant melanoma cells that are BRAFV600E mutated and MITF<sup>low</sup>/AXL<sup>high</sup>. (A) MITF (green) or (B) AXL (red) expression was examined by immunofluorescence (left panels). Right panels, merge of MITF or AXL and nuclear staining with DAPI (blue). Melmet q and Melmet 5 cells were treated with either (C) the chemotherapeutic agent dacarbazine or (D) the BRAF inhibitor vemurafenib for 72 h prior to examination of cell viability by the MTS assay. Figures show average of three independent experiments with ±SE bars. (E) Melmet 1 and (F) Melmet 5 cells incubated with 0.2  $\mu$ g mL<sup>-1</sup> TPCS<sub>2a</sub> (PCI no drug), or TPCS<sub>2a</sub> in combination with 100 pM streptavidin-saporin (PCI saporin) or the CSPG4-targeting immunotoxin 225.28saporin (PCI 225.28-saporin) for 18 h, followed by a 4 h chase in drug-free medium prior to light exposure as indicated in the *X*-axis. Cell metabolic activity/viability was examined using the MTT assay 48 h post illumination. (G) Melmet 5 cells seeded at clonal density (1000 cells per 9.6 cm<sup>2</sup>) and treated as in F. Cells were assessed for clonogenicity 10–14 days post light exposure. Each figure shows a single representative experiment out of three independent experiments. Bars = SD of three technical replicates.



#### Fig. 5.

Potent in vitro and in vivo effects of the CSPG4-targeting recombinant immunotoxin scFvMEL-rGel. (A) A-375GFP cells were incubated with the PCI-photosensitizer AlPcS<sub>2a</sub> without (PCI no drug) or with 100 nM scFvMEL-rGel for 18 h and subsequently washed and incubated in drug-free medium for 4 h prior to red broad band light ( $\gamma_{max}$  around 670 nm) exposure. Cytotoxic responses were examined by the MTT assay 72 h post light (data are based on one representative experiment out of five independent experiments (the other 4 experiments are presented in ESI-8<sup>†</sup>)). Error bars are SD of triplets. (B) A-375 cells with granular (AlPcS2a (red) fluorescence prior to 30 s. microscopy light (650–690 nm) exposure and C) 5 min after PCI-induced cytosolic release of photosensitizer. (D) GFPand (E) Hoechst 33342 signal. (F) Strong delay of A-375 tumour growth after only one session with PCI of scFvMEL-rGel. These data are extracted from a Kaplan-Meier plot published previously Selbo PK et al. in PLoS One, 4(8), e6691. Athymic nude mice with subcutaneously growing A-375 tumours (approximately 75 mm<sup>3</sup>) were given AlPcS<sub>2a</sub> (5 mg kg<sup>-1</sup> i.p. 48 h before laser exposure) and scFvMEL-rGel (2 mg kg<sup>-1</sup> i.v, 24 h before laser exposure). Laser wavelength: 670 nm, Irradiance: 100 mW cm<sup>-2</sup>, Light dose: 20 J cm<sup>-2</sup>. n, number of animals in each group. (G) Individual tumour growth curves over time. The flat line up to day 110 in the PCI of the scFvMEL-rGel group represents 4 mice with complete tumour responses (33%). In the same group, one mouse had a tumour that started to grow slowly at day 100. Another PCI-treated mouse where the tumour size was reduced by 65%, gained a slow tumour growth between day 60 and 90, but had to be sacrificed due to a nose

infection. The number of long time survivors (animals with tumours less than 800 mm<sup>3</sup> at day 90) was n = 6 (50%) in the PCI group.