

1 **Transfection of pseudouridine-modified mRNA encoding CPD-photolyase leads to repair of**
2 **DNA damage in human keratinocytes: a new approach with future therapeutic potential**

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Abbreviations: NHEK – Normal Human Epidermal Keratinocyte; UVB – Ultraviolet B; CPD –
Cyclobutane Pyrimidine Dimer; Ψ – Pseudouridine-modified; CPD-PL – Cyclobutane Pyrimidine
Dimer-specific photolyase; eGFP – enhanced Green Fluorescent Protein; ELISA – Enzyme Linked
Immunosorbent Assay; IL-6 – Interleukin-6; SDHA – Succinate Dehydrogenase Complex, Subunit A;
PGK1 – Phosphoglycerate Kinase 1

24 **Abstract**

25 UVB irradiation induces harmful photochemical reactions, including formation of
26 cyclobutane pyrimidine dimers (CPDs) in DNA. Accumulation of unrepaired CPD lesions
27 causes inflammation, premature ageing and skin cancer. Photolyases are DNA repair enzymes
28 that can rapidly restore DNA integrity in a light-dependent process called photoreactivation,
29 but these enzymes are absent in humans. Here, we present a novel mRNA-based gene therapy
30 method that directs synthesis of a marsupial, *Potorous tridactylus*, CPD-photolyase in
31 cultured human keratinocytes. Pseudouridine was incorporated during *in vitro* transcription to
32 make the mRNA non-immunogenic and highly translatable. Keratinocytes transfected with
33 lipofectamine-complexed mRNA expressed photolyase in the nuclei for at least 2 days.
34 Exposing photolyase mRNA-transfected cells to UVB irradiation resulted in significantly less
35 CPD in those cells that were also treated with photoreactivating light, which is required for
36 photolyase activity. The functional photolyase also diminished other UVB-mediated effects,
37 including induction of IL-6 and inhibition of cell proliferation. These results demonstrate that
38 pseudouridine-containing photolyase mRNA is a powerful tool to repair UVB-induced DNA
39 lesions. The pseudouridine-modified mRNA approach has a strong potential to discern
40 cellular effects of CPD in UV-related cell biological studies. The mRNA-based transient
41 expression of proteins offers a number of opportunities for future application in medicine.

42 **Key words:** pseudouridine-modified mRNA – mRNA therapy – photolyase – human
43 keratinocyte – UVB – cyclobutane pyrimidine dimers

44 1. Introduction

45 Ultraviolet (UV) radiation, especially UVB (290-320 nm), is a common environmental
46 carcinogen that induces harmful photochemical reactions in the skin [1]. UVB photons
47 absorbed by the DNA cause a variety of photoproducts, including cyclobutane pyrimidine
48 dimers (CPD), which is the most frequent and deleterious photolesion. It forms in double-
49 stranded DNA when two adjacent pyrimidine bases in one strand fuse together [2].
50 Accumulated CPDs induce apoptosis, inflammation, immune suppression, and mutations that
51 may lead to melanoma and other types of skin cancer [3-6]. Timely repair of damaged DNA
52 is critical to prevent these adverse effects. In humans and rodents, photoproducts are removed
53 by nucleotide excision repair (NER), a versatile repair system for removal of bulky DNA
54 lesions. Genetic defects in NER result in photosensitive disorders, such as xeroderma
55 pigmentosum and Cockayne's syndrome, in which patients have UV sensitivity and display
56 skin cancer proneness and/or accelerated aging [7, 8]. NER excises CPDs at a very low rate,
57 requiring hours to days to restore DNA integrity [9-12]. Interestingly, in most organisms from
58 bacteria to marsupials, CPDs are rapidly removed by photolyases that specifically recognize
59 and split these dimers into monomers using energy from visible light [13-15]. Due to an
60 evolutionary loss, photolyases are absent in placental mammals, including humans, which
61 therefore solely rely on NER for removal of photolesions [16-19].

62 Studies have demonstrated that CPD photolyases from phylogenetically diverse organisms,
63 such as cyanobacteria (*Anacystis nidulans*) [20], yeast [21] or marsupials [22], can function in
64 mammalian cells. Liposome-encapsulated CPD photolyase enzyme from *Anacystis nidulans*
65 has been shown to protect cultured mammalian cells and human skin from the effects of UVB
66 [20, 23-26]. Superior resistance to UVB-induced sunburn, immune suppression and
67 carcinogenesis were demonstrated for transgenic mice ubiquitously expressing CPD
68 photolyase of the rat kangaroo (*Potorous tridactylus*) [22, 27-29]. Mammalian cell lines

69 stably expressing marsupial CPD photolyases also demonstrated reduced apoptosis and
70 mutation frequencies when exposed to UVB radiation [30-33]. To study sunlight-induced
71 DNA damage and their repair in primary cells, transient adenovirus-mediated CPD photolyase
72 expression was used [34, 35].

73 Gene therapy for the replacement of defective genes or for the expression of therapeutic
74 proteins has made great progress in the last decade. After exploring numerous systems for
75 expressing the encoded proteins [36], *in vitro* transcribed mRNA seems to be the most
76 suitable tool for transient protein expression [37]. It has many features that make mRNA-
77 mediated gene transfer especially valuable for functional characterization of encoded protein.
78 The transfected mRNA is translated with high efficiency in any cell, including primary and
79 non-dividing mammalian cells [37]. Importantly, when mRNA is delivered to the cell, only
80 the encoded protein of interest is generated, unlike other vectors, such as plasmids that
81 contain sequences for additional proteins, or viral vectors that not only code for but also carry
82 viral proteins into the cell. In the last several years, mRNA-mediated transfection technology
83 has improved tremendously [38]. It is now well documented that incorporation of
84 pseudouridine (Ψ), a naturally-occurring modified nucleoside, into mRNA makes it less
85 immunogenic by avoiding the activation of RNA sensors [39-41]. *In vitro*-transcribed mRNA
86 containing pseudouridine (Ψ -mRNA) is also translated more efficiently than mRNA
87 containing unmodified nucleosides [42]. We have demonstrated that the presence of Ψ
88 nucleosides improved the stability of RNA [43] and diminished both activation of RNA-
89 dependent protein kinase (PKR) and inhibition of translation [44]. A newly established
90 preparative HPLC purification procedure was critical to obtain Ψ -mRNA that is free of
91 aberrant transcription products, resulting in superior translational potential and no
92 immunogenicity [45, 46].

93 Here, we present a novel mRNA-based gene therapy method that directs functional

94 photolyase synthesis in human keratinocytes. Our study demonstrates that delivery of HPLC-
95 purified, Ψ -mRNA encoding CPD-photolyase into keratinocytes leads to the rapid repair of
96 UVB-induced CPDs and suggest that such mRNA has therapeutic potential to repair damage
97 caused by exposure to the sun and other sources of UV radiation.

98 **2. Material and methods**

99 **2.1 RNA synthesis**

100 A codon-optimized photolyase gene from *Potorous tridactylus* (rat kangaroo) containing GC-
101 rich codons for superior translation was synthesized by Entelechon (Bad Abbach, Germany).
102 The optimization increased the GC-content of the photolyase coding sequence (Accession:
103 D26020) from 51.8% to 65.0%. Messenger RNAs encoding CPD-photolyase (CPD-PL Ψ -
104 mRNA) and enhanced green fluorescent protein (eGFP Ψ -mRNA) were transcribed as
105 previously described [42] from linearized plasmids (pTEV-CPD-PL-A101 and pTEVeGFP-
106 A101), using the Megascript T7 RNA polymerase kit (Ambion, Austin, TX) in which UTP
107 was replaced with pseudouridine triphosphate (TriLink, San Diego, CA). Subsequently, the
108 mRNA was HPLC purified as described [45], and provided with a 5' cap using capping
109 enzyme and 2'-O-methyltransferase according to the manufacturer (CellScript, Madison, WI).
110 The RNA was transcribed to contain an encoded 101-nt long 3' poly(A) tail, which was
111 extended with ~ 300 nucleotides using poly(A) polymerase (USB, Cleveland OH). RNA
112 samples were analyzed by agarose gel electrophoresis for quality assurance. The mRNAs
113 were shown to be free of dsRNA contaminants using dsRNA-specific J2 mAb (English &
114 Scientific Consulting, Budapest, Hungary) in a dot-blot assay [45]. The mRNAs were stored
115 in siliconized tubes at -20°C.

116 **2.2 Cell cultures**

117 The human keratinocyte cell line HaCaT was obtained from the ATCC and grown in high
118 glucose DMEM (PAA, Traun, Austria) supplemented with 2 mM L-glutamine (PAA), 10%

119 heat-inactivated fetal bovine serum (Lonza, Verviers, Belgium) and 0.5%
120 antibiotic/antimycotic solution (Sigma-Aldrich, St. Louis, MO, USA) at 37°C in a 5% CO₂
121 atmosphere. Normal human epidermal keratinocytes (NHEK) were isolated from healthy
122 adult skin derived from plastic surgery and cultured in EpiLife serum-free, complete
123 keratinocyte growth medium (Life Technologies, Carlsbad, CA, USA). Ethics approval was
124 received from the National Scientific and Research Ethics Commission. HaCaT cells and
125 second passage NHEK were used at 70-80% confluency in each experiment.

126 **2.3 Transient transfection and treatments**

127 HaCaT cells and NHEK were seeded into 96-well plates at a density of 2×10^4 cells per well
128 one day prior to transfection. Aliquots of RNA samples (0.25 µg) were complexed with 0.8 µl
129 Lipofectamine LTX-PLUS (Life Technologies) in a final volume of 100 µl of EpiLife and the
130 complexed RNA was added to each well. After one hour the lipofectamine-RNA complex
131 was replaced with 100 µl culture medium. At 20 h post transfection, cells covered with 50 µl
132 Dulbecco's Phosphate Buffer Saline (DPBS) (Life Technologies) were subjected to 20
133 mJ/cm² UVB using two TL-20W/12 bulbs (Philips). Immediately after UVB treatment, cells
134 were either exposed to visible light using a F18W Daylight fluorescent bulb (Sylvania,) at a
135 distance of 16 cm and using a 4 mm thick glass filter as a shield or kept in the dark for one
136 hour. Cells were further cultured in complete medium for the indicated times. Cell viability
137 was determined by the EZ4U assay (Biomedica Gruppe, Vienna, Austria), according to the
138 manufacturer.

139 **2.4 Western blot analysis**

140 Cells were lysed in RIPA buffer (R0278, Sigma-Aldrich) containing protease inhibitor
141 cocktail (Sigma-Aldrich). Proteins were separated on 10% polyacrylamide gels and
142 transferred to nitrocellulose membranes. The membrane was blocked in 5% non-fat dry milk
143 for 1 h and incubated overnight at 4°C with primary CPD-photolyase antibody [27] diluted

144 (1:500) in PBGTNa made of PBS containing 0.5% bovine serum albumin (BSA, Amresco,
145 Solon, OH, USA), 0.05% gelatin (Sigma-Aldrich), 0.05% Tween-20 (Amresco) and 300 mM
146 NaCl (VWR, San Diego, CA, USA). HRP-conjugated anti-rabbit IgG was used as a
147 secondary antibody (1:3000; 1 h, at room temperature, BioRad, Berkeley, CA, USA). Proteins
148 were visualized with the Pierce ECL Plus Western blotting detection system (Thermo Fisher
149 Scientific, Rockford, IL, USA).

150 **2.5 Fluorescence microscopy**

151 Expression of eGFP in HaCaT cells was documented using an epifluorescent Zeiss Axiovert
152 100 microscope mounted with a Zeiss AxioCam MRc 5 digital camera using AxioVision 4.8
153 and 0.4-second exposure time.

154 In the expression studies, cells were fixed with 4% paraformaldehyde (Sigma-Aldrich) and
155 permeabilized with 1% Triton X-100 (Amresco). Before immunolabeling, cells were blocked
156 in 20% FBS and then probed overnight at 4°C with CPD-photolyase-specific antibody diluted
157 (1:100) in PBGTNa. FITC-conjugated secondary antibody (1:2000; anti-rabbit IgG;
158 Vectashield, Irvine, CA, USA) was added for 2 h at room temperature.

159 For the analysis of CPDs, experiments were carried out as described above, except, before
160 blocking, cellular DNA was denatured in 2 M HCl (VWR) at room temperature for 30 min
161 and then probed with primary anti-CPD antibody (TDM-2, Cosmo Bio, Tokyo, Japan) diluted
162 (1:1500) in PBS containing 5% FBS for 30 min. Alexa Fluor 568-conjugated anti-mouse IgG
163 was used as secondary antibody (1:2000; Life Technologies). In both studies, nuclei were
164 stained with DAPI (Vectashield) and cells were analyzed by fluorescence microscopy.

165 **2.6 Enzyme-linked immunosorbent assay (ELISA)**

166 Genomic DNA was extracted by the QIAamp DNA mini kit (Qiagen, Hilden, Germany),
167 according to the manufacturer. Flat-bottom 96-well plate were precoated with 0.003%
168 protamine sulfate (50 µl/well, Sigma-Aldrich) and incubated overnight with denatured DNA

169 (15 ng/well) at 37°C. The plates were washed with PBS containing 0.05% Tween-20 and
170 incubated with 2% FBS at 37°C for 30 min to prevent non-specific binding of the antibody
171 and washed again. Anti-CPD monoclonal antibody (TDM2) diluted in PBS (1:1000) was used
172 as primary antibody at 37°C for 30 min. The plates were washed and then incubated with
173 HRP conjugated anti-mouse IgG secondary antibody (1:3000, BioRad) at 37°C for 30 min.
174 After washing, samples were equilibrated with citrate-phosphate buffer (pH 5.0; 150 µl/well),
175 then substrate solution was added to each well (100 µl/well) containing 0.04% o-
176 phenylenediamine (Sigma-Aldrich) and 0.0063% H₂O₂ in citrate-phosphate buffer. Samples
177 were incubated at 37°C for 15 min, and 2 M sulfuric acid (VWR) was added (50 µl/well) to
178 stop the reaction and absorbance was measured at 492 nm using an Anthos 2020 microplate
179 reader (Biochrom Ltd; Cambridge; UK).

180 The concentration of IL-6 in the supernatants was determined by the human IL-6 Quantikine
181 ELISA kit (R&D Systems, Minneapolis, MN, USA), according to the manufacturer.

182 **2.7 RNA extraction and quantitative RT-PCR**

183 Total RNA was isolated using TRI reagent (MRC, Cincinnati, OH, USA) followed by DNase
184 I treatment (Fermentas, St. Leon-Rot, Germany). To quantitate IL-6, SDHA, and PGK1
185 mRNAs, reverse transcription was performed using the High-Capacity cDNA Reverse
186 Transcription Kit (Life Technologies), according to the manufacturer. Quantitative PCR was
187 performed on the ABI 7900 HT Sequence Detection System (Life Technologies) using a
188 reaction mixture containing 1x PCR buffer, 3 mM MgCl₂ and 0.1 mM dNTP mix
189 (Fermentas), 1x TaqMan Gene Expression assay (Life Technologies), 1x ROX reference dye
190 (Life Technologies), 0.0125 U Taq DNA polymerase (Fermentas) and 12.5 ng cDNA
191 template in a total volume of 10 µl. qPCR was performed in 384-well optical plates (Life
192 Technologies) and the PCR program used for amplification was: 94°C 1 min, followed by 40
193 cycles of 94°C for 12 sec and 60°C for 45 sec. The following TaqMan Gene Expression

194 assays were used: IL-6 (Hs00985639_m1), SDHA (Hs00188166_m1) and PGK1
195 (Hs00943178_g1), the sequences are proprietary and not released by the company. SDHA and
196 PGK1 mRNA levels were used for normalization [47].

197 **2.8 Statistical analysis**

198 All statistical analyses were performed using GraphPad Prism version 5.0 (GraphPad
199 Software Inc., San Diego, CA, USA). Values were compared between photoreactivated and
200 non-photoreactivated samples using two-tailed, paired *t*-test. A *p* value of less than 0.05 was
201 considered as statistically significant.

202 **3. Results**

203 **3.1 Synthesis of CPD-specific photolyase in human keratinocytes transfected with the** 204 **encoding pseudouridine-modified mRNA**

205 To test whether pseudouridine-modified mRNA (Ψ -mRNA) is translated in human
206 keratinocytes, eGFP-encoding Ψ -mRNA was complexed with lipofectamine LTX and
207 delivered to the cells. The expression of eGFP was monitored by fluorescence microscopy at
208 different time points after transfection. eGFP was detectable as early as 1 h, peaked between 6
209 and 20 h, and its level gradually declined, but remained detectable even at 48 h after mRNA
210 transfection (Fig. 1a). This eGFP mRNA served as a control for photolyase experiments.
211 Using western blotting, high levels of CPD-photolyase synthesized from the transfected CPD-
212 PL Ψ -mRNA could be detected in both HaCaT cells and normal human epidermal
213 keratinocytes (NHEK) at 20 h post transfection (Fig. 1b and c). The photolyase migrated with
214 the expected size of 62 kDa (Fig. 1c) and localized in the nuclei of keratinocytes, as
215 demonstrated by immunofluorescence imaging (Fig. 1d).

216 **3.2 Enhanced repair of UVB-induced CPDs in keratinocytes is mediated by CPD-** 217 **photolyase translated from the encoding Ψ -mRNA**

218 To investigate whether CPD-photolyase translated from the encoding Ψ -mRNA is functional
219 and capable of cleaving CPDs, HaCaT cells were first transfected with lipofectamine-
220 complexed CPD-PL Ψ -mRNA. Twenty hours later, the transfected cells were exposed to 20
221 mJ/cm^2 UVB. Immediately thereafter, cells were either exposed to photoreactivating light, or
222 left in the dark for 1 hour. Using CPD-specific antibodies, efficient repair of these
223 photolesions could be detected in CPD-PL Ψ -mRNA-transfected cells that were subjected to
224 photoreactivating light, while those deprived from this energy source or transfected with
225 eGFP Ψ -mRNA contained high levels of CPDs, when it was analyzed immediately after the
226 photoreactivation or incubation in the dark (Fig. 2).

227 To quantitate the amount of CPDs removed from the genomic DNA of the CPD-PL Ψ -
228 mRNA-transfected keratinocytes immediately after exposure to visible light, a CPD-specific
229 ELISA was used. We determined that photoreactivation of UVB-irradiated, CPD-PL Ψ -
230 mRNA-transfected cells eliminated more than 60% of the CPDs, as compared to cells kept in
231 the dark or transfected with control eGFP Ψ -mRNA (Fig. 3, left panel).

232 UVB treatment of keratinocytes results in the induction of apoptosis and reduction in cell
233 proliferation. Cell viability, determined 24 h after UVB irradiation, demonstrated that
234 photoreactivation significantly increased cell survival of CPD-PL Ψ -mRNA transfected
235 keratinocytes compared to those that were kept in the dark or transfected with eGFP Ψ -
236 mRNA (Fig. 3, right panel).

237 **3.3 Photorepair of CPDs in keratinocytes transfected with CPD-photolyase Ψ -mRNA** 238 **reduces UVB-induced IL-6 expression**

239 To determine whether repair of CPDs in photolyase mRNA-transfected cells can ameliorate
240 UVB-triggered cytokine induction, we measured the expression of IL-6 mRNA and protein
241 using qRT-PCR and ELISA. We found that the repair of CPDs in CPD-PL Ψ -mRNA
242 transfected keratinocytes significantly decreased the levels of IL-6 mRNA measured 24 h

243 after UVB irradiation (Fig. 4). Accordingly, the levels of IL-6 protein in the supernatants of
244 photolyase mRNA-transfected keratinocytes were also significantly reduced at 6 and 24 h
245 (Fig. 4).

246 **4. Discussion**

247 Intracellular delivery of therapeutic proteins has been performed using many different viral
248 and non-viral techniques. Recent advancements in the use of mRNA for expressing the
249 encoded proteins in mammalian cells made the present study possible. Here, we demonstrate
250 that HPLC-purified, lipofectamine-complexed mRNAs containing pseudouridine are
251 efficiently taken up and translated by primary human keratinocytes and an established
252 keratinocyte cell line, with greater than 95% of the cells expressing the protein encoded by the
253 mRNA (eGFP or photolyase). The CPD-photolyase of *Potorous tridactylus* encoded by the
254 mRNA is efficiently translated into a functional protein that correctly localized to the nucleus.
255 The CPD-photolyase mediated rapid repair of UVB-induced cyclobutane pyrimidine dimers.
256 This led to a significant survival advantage and reduction in IL-6 mRNA and protein
257 secretion.

258 This is the first report of transient expression of functional CPD-photolyase in human
259 cells from transfected mRNA. Several studies have reported successful protection from UVB-
260 induced damage using topical application of liposomes-encapsulating recombinant CPD-
261 photolyase protein, however, direct evidence for nuclear localization of this cyanobacteria-
262 derived protein has not been provided [20, 23, 26, 48-51]. Here, we found that nucleoside-
263 modified mRNA encoding CPD-photolyase correctly located to the nucleus and removed
264 more than 60% of CPDs within the first hour of photoreactivation. In placental mammals that
265 lack a functional CPD-photolyase, DNA repair of CPDs is performed by the NER complex
266 that in mammals contains 9 major proteins and over 20 associated genes. In UVB-irradiated
267 primary keratinocytes, NER-mediated repair removed only 30% of the CPDs in 24 hours [52].

268 If a cell containing CPD lesions divides before the repair can be completed, an alteration to
269 the DNA, i.e. mutation, can occur in the daughter cell and all of its progeny [6]. Such
270 mutations are the initial steps in the development of sun-induced skin cancers. As shown for
271 *Potorous tridactylus* CPD photolyase-expressing transgenic mice [27-29], very rapid
272 clearance of CPDs by photolyase produced from the transfected mRNA should result in a
273 large reduction of DNA mutations and other CPD-driven deleterious effects of UVB.

274 It is well established that UVB irradiation induces cell cycle arrest and inhibits cell
275 proliferation, which is partly mediated by CPD photolesions [53, 54]. Removal of CPDs after
276 photoreactivation of keratinocytes transfected with photolyase mRNA increased survival,
277 confirming that CPDs are major contributors to cell death after UVB irradiation [22, 31, 55,
278 56] and demonstrating that CPD-photolyase mRNA transfection is an effective tool for
279 photorepair.

280 UVB irradiation also induces erythema in the skin [57] and the involvement of the
281 proinflammatory cytokine IL-6 has been considered [58, 59]. Consistent with these
282 observations, we demonstrate that enhanced repair of CPDs in photolyase mRNA-transfected
283 keratinocytes reduced UVB-induced IL-6 expression when cells were subjected to
284 photoreactivation compared to those kept in dark. CPD-photolyase mediated repair of UVB-
285 induced damage did not return IL-6 levels to non-irradiated levels likely due to the presence
286 of modified host RNA acting through TLR3 [60] and other UVB mediated effects, including
287 the induction of reactive oxygen species, protein–DNA crosslinking, oxidative base damage
288 (e.g. 8-oxo-7,8-dihydroxyguanine), single-strand breaks and a 6-4 photoproduct (6-4PP).
289 Thus, these results demonstrate that CPDs play an important but not exclusive role in
290 triggering IL-6 release from keratinocytes, and transfection of Ψ -mRNA encoding CPD-
291 photolyase reduces production of IL-6 by UVB irradiated keratinocytes and supports their
292 survival.

293 In summary, we demonstrate that *in vitro*-synthesized, pseudouridine-containing mRNA
294 encoding *Potourus* photolyase is a powerful tool to rapidly repair UVB-induced CPD lesions
295 in human keratinocytes. This novel mRNA-based gene therapy method efficiently directs
296 functional protein synthesis providing number of opportunities for future dermatological
297 applications if the nucleic acid delivery to intact epidermis has been resolved. Concerning
298 photolyase, thus it could be used in UV-sensitive diseases [7, 8] and in the prevention of skin
299 cancer and tanning, which requires CPD formation to increase the production of melanin [61].
300 Ψ -mRNA encoding CPD-photolyase is also a new model system that uses primary human
301 keratinocytes to investigate the cellular and molecular effects of UVB irradiation with the
302 ability to distinguish between CPD-dependent and CPD-independent events and the potential
303 to deliver and study any protein of interest.

304 **Conflict of interests**

305 Katalin Karikó and Drew Weissman hold a patent for the use of nucleoside modified mRNA
306 as a therapeutic protein delivery platform. They have also formed a small biotech company,
307 RNARx, that receives funding from the National Institutes of Health (R42HL87688) to
308 explore the use of nucleoside-modified mRNA for gene therapy.

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483 **Figure captions**

484 **Fig. 1. Translation of CPD-photolyase-encoding, Ψ -modified mRNA in keratinocytes.**
485 HaCaT cells or normal human epidermal keratinocytes (NHEK) were transfected with
486 lipofectamine-complexed Ψ -mRNA encoding eGFP or CPD-photolyase. Translated eGFP
487 was detected at the indicated times post transfection in HaCaT cells transfected with eGFP Ψ -
488 mRNA by fluorescence microscopy (panel a) (original magnification $\times 20$). Cells were lysed

489 at the indicated times (panel b) or at 20 h post transfection (panel c) and analyzed by western
490 blot. Arrows mark positions of the 62 kDa CPD-photolyase and the loading reference, β -actin.
491 CPD-photolyase protein was observed in NHEK transfected with CPD-PL Ψ -mRNA using
492 immunofluorescent detection (panel d). Nuclei were visualized by DAPI staining (original
493 magnification $\times 40$).

494 **Fig. 2. Photorepair of CPDs in HaCaT cells transfected with CPD-PL Ψ -mRNA.** HaCaT
495 cells were transfected with lipofectamine-complexed Ψ -mRNA encoding CPD-photolyase or
496 eGFP, or left untransfected (none). Twenty hours later, cells were subjected to 20 mJ/cm^2
497 UVB and immediately exposed to photoreactivating light or left in the dark for 1 h.
498 Immediately after photoreactivation or incubation in the dark, CPDs were detected by
499 immunofluorescent labeling with CPD-specific antibody and Alexa-conjugated secondary
500 antibody (red). Nuclei were visualized by DAPI staining. Images are representatives of three
501 independently performed experiments (original magnification $\times 20$).

502 **Fig. 3. Reduced levels of CPDs and improved cell viability in photolyase mRNA-**
503 **transfected keratinocytes exposed to UVB and photoreactivating light.** HaCaT cells and
504 NHEK were irradiated with a physiological dose of UVB (20 mJ/cm^2) 20 h after delivery of
505 Ψ -mRNAs and immediately exposed to photoreactivating light (+Phr) or left in the dark (-
506 Phr) for 1 h. After photoreactivation or incubation in the dark, the amount of CPDs in the
507 DNA of keratinocytes transfected with Ψ -mRNA encoding CPD-PL or eGFP was determined
508 by ELISA (left panel). The OD values obtained were in the linear range of the ELISA. Cell
509 viability was determined by EZ4U assay at 24 h after UVB irradiation (right panel). The
510 values were calculated relative to those obtained with cells that were not UVB irradiated.
511 Significance was assessed by paired *t*-test, asterisk, $p < 0.05$. Error bars represent the standard
512 error of the mean. The results are means of three independent experiments.

513 **Fig. 4. Photorepair of CPDs reduces UVB-induced IL-6 expression in HaCaT cells.** Cells
514 were transfected with lipofectamine-complexed CPD-PL Ψ -mRNA, 20 h later irradiated with
515 20 mJ/cm² UVB and then immediately exposed to photoreactivating light (+Phr) or left in the
516 dark (-Phr) for 1 h. Total RNA was isolated and supernatants were collected 6 and 24 h after
517 UVB irradiation. Levels of IL-6 mRNA were determined by qRT-PCR. Values measured in
518 UVB irradiated -Phr or +Phr cells were normalized to those measured in the Ψ -mRNA-
519 transfected, non-UVB irradiated control cells. IL-6 was quantitated in the supernatant by
520 ELISA. Asterisks indicate significant differences (paired t-test; p<0.05) between
521 photoreactivated (+Phr) and non-photoreactivated (-Phr) samples. Error bars represent the
522 standard error of the mean. Three independent experiments were performed, each in triplicate.