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1 **Study of generated genetic polymorphisms during the photocatalytic elimination of**
2 ***Klebsiella pneumoniae* in water**

3

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16 **Abstract**

17 *Klebsiella pneumoniae* is considered an emerging pathogen persistent in extreme
18 environmental and stressed conditions. The aim of the present study is the investigation of
19 inactivation rates of this pathogen in water by means of heterogeneous photocatalytic
20 treatment under solar irradiation and the induced genetic variance applying RAPD-PCR as a
21 molecular typing tool. Novel Mn and Co-doped TiO₂ catalysts were assessed in terms of their
22 disinfection efficiency. The reference strain of *K. pneumoniae* proved to be readily
23 inactivated, since disinfection occurred rapidly (i.e. after only 10 min of treatment) and low
24 levels of bacterial regrowth were recorded in the dark and under natural sunlight. Binary
25 doped titania exhibited the best photocatalytic activity verifying the synergistic effect induced
26 by composite dopants. Applying RAPD analysis to viable cells after treatment we concluded
27 that increasing the treatment time led to a considerable alteration of RAPD profiles and
28 homology coefficient ranged almost between 35 and 60%. RAPD-PCR proved to be a useful
29 typing molecular tool, that upon standardized conditions exhibits highly reproducible results.
30 Genetic variation among isolates increased in relation to the period of treatment and
31 prolonged irradiation in each case affected the overall alteration in band patterns. RAPD
32 patterns were highly diverse between treated and untreated isolates when disinfection was
33 performed with the Co-doped titania. The broad spectrum of genetic variance and generated
34 polymorphisms has the potential to increase the already significant virulence of the species.

35

36 **Keywords:** *Klebsiella pneumoniae*; doped-TiO₂; solar photocatalysis; disinfection; RAPD

37

38 1. Introduction

39 Water may serve as the passive carrier of diverse bacterial groups and species, which have the
40 potential to act as infectious agents of waterborne diseases. In recent years considerable
41 attention has been drawn towards many opportunistic pathogens contained in water
42 distribution systems, among which *Klebsiella pneumoniae* is prominently included.^{1,2} This
43 bacillus is gram-negative, lactose-fermenting and it belongs to the broad family of
44 Enterobacteriaceae.³ *K. pneumoniae* accounts for a significant proportion of hospital-acquired
45 infections, like pneumonia, septicaemia, and soft tissue infections and is responsible for
46 numerous nosocomial outbreaks worldwide.^{4,5} The ability of this bacterial strain to spread
47 rapidly in the hospital, as well as in the aquatic environment and its resistant nature against
48 bactericidal factors enlist *K. pneumoniae* among the highly virulent and pathogenic
49 microorganisms, especially for susceptible population groups and immunocompromised
50 patients.^{2,6}

51 The frequency of reports concerns mainly the multi-drug resistance of this specific strain,
52 while equally significant is the resistance of *K. pneumoniae* isolates to environmental stressed
53 conditions and deleterious agents, such as disinfectants.⁶⁻¹⁰ The prominent polysaccharide
54 capsule that they possess increases their virulence through protection from phagocytosis and
55 prevents cellular destruction by bactericidal factors.^{3,11}

56 Given the persistence of this bacterium in the environment and the fact that it may be
57 transmitted through water consumption, it is imperative to explore effective and reliable
58 disinfection techniques, which would eventually inactivate it, providing appropriate control
59 measures of such pathogens. Although the beneficial effects of many treatment processes of
60 water and wastewater have been well addressed, *K. pneumoniae* has been merely mentioned
61 in research studies as far as its resistance against disinfection is concerned. Conventional
62 water disinfection involves chlorination, which is widely used to remove effectively an

63 extended variety of microorganisms.¹²⁻¹⁴ Yet, the diverse structural and physiological features
64 of the latter lead to different response in the course of treatment, raising some concerns
65 regarding the suitability of chlorination. Environmental isolates of *K. pneumoniae* display
66 considerable resistance to chlorination due to several factors, namely extracellular polymer
67 production, alteration of membrane lipids and increased cell aggregation.⁶ Another approach
68 towards efficient inactivation of persistent pathogens in water seems to be the application of
69 titanium dioxide (TiO₂) photocatalysis, which has already demonstrated high reduction rates
70 of viruses and bacteria in water/wastewater samples under ultraviolet (UV) irradiation.¹⁵⁻¹⁷
71 Further applications are based on the exploitation of total solar light spectrum and the use of
72 metal dopants, which have been explored for visible light-induced photocatalysis.¹⁸ The
73 exploitation of total solar light spectrum expands practical applications of TiO₂
74 photocatalysis, which has been successfully adopted as a disinfection technique.¹⁹⁻²² In our
75 previous work, *K. pneumoniae* was readily inactivated in aqueous samples under solar
76 irradiation with the use of Mn- and Co-doped titania and disinfection occurred rapidly (i.e.
77 after only 10 min of irradiation) with the dopant concentration affecting the overall process to
78 a certain extent.¹¹

79 The complexity of bacterial cells and their metabolic activity in the aquatic environment
80 make their inactivation rather complicated under solar irradiation. Generally, sunlight damage
81 involves the following principal mechanisms, which comprise direct DNA impairment and
82 indirect endogenous damage caused mainly by UVB wavelengths and indirect exogenous
83 damage involving UVB, UVA and visible wavelengths up to 550 nm.²³

84 Complete bacterial inactivation, which is mostly desired, is highly dependent upon
85 operating parameters of each water disinfection process that uses natural or simulated solar
86 irradiation. However, even in the case of residual cells post treatment, the extent of DNA
87 damage and the induced oxidative stress may result in loss of culturability or virulence of the

88 bacteria.²⁴ Efforts on delineating relationships among bacteria concerning alterations of their
89 genetic material focus on molecular typing techniques, among which PCR-based methods
90 predominate.^{3,5,8,10,25,26} Among various DNA-fingerprinting techniques, Randomly Amplified
91 Polymorphic DNA analysis (RAPD), offers a reliable, simple and cost effective tool to study
92 the genotypic relatedness of a limited number of isolates, highlighting their generated genetic
93 diversity through the course of a disinfection process.²⁵ RAPD typing has been a matter of
94 interest mainly because of its economical and practical merits.⁵ This technique is an arbitrarily
95 primed-PCR, with which strain-specific arrays of amplified DNA fragments are generated.²⁵
96 Upon meticulous optimization, RAPD could be a sensitive and reproducible assay for the
97 detection of DNA polymorphisms among bacterial isolates.^{5,10,26,27}

98 In the present work the main objective was the investigation of genetic diversity applying
99 RAPD analysis among isolates of *K. pneumoniae*, which were subject to solar photocatalysis
100 with the use of manganese- (Mn-), cobalt- (Co-), and binary(Mn/Co-)-doped titania catalysts,
101 prepared in previous work.¹¹ Also, inactivation rates and regrowth potential of the bacterium
102 after treatment were evaluated.

103

104 **2. Experimental**

105 **2.1. Metal-doped TiO₂ nanoparticles**

106 Commercially available titanium dioxide (TiO₂ P25) was purchased from Degussa - Evonik
107 Corp. (physicochemical characteristics are anatase:rutile 75:25, particle size of 21 nm and
108 BET area of 50 m² g⁻¹) and was used as benchmark. Mn-, Co- and Mn/Co binary-doped TiO₂
109 catalysts were prepared by a co-precipitation method with molar ratio in different
110 concentrations in the range of 0.02 to 0.3 wt%. Details regarding synthesis and
111 characterization of all catalysts used in the present study can be found in previous studies.^{11,28}
112 The crystal structure, particle size, and morphology were examined with powder X-ray

113 diffraction (XRD), scanning electron microscopy (SEM) and transmission electron
114 microscopy (TEM), respectfully. Powder X-ray diffraction patterns were collected on a
115 Rigaku D/MAX-2000H rotating anode diffractometer (CuK α radiation) equipped with the
116 secondary pyrolytic graphite monochromator operated at 40 kV and 80 mA over the 2θ
117 collection range of 10-80°. The scan rate was 0.05° s⁻¹. The UV- Visible diffuse reflectance
118 spectra of the final powders were measured on a Perkin Elmer LAMBDA 950 with BaSO₄, as
119 reference standard. Surface morphology and elemental analysis of the samples were carried
120 out using SEM and an energy dispersive spectrometer (EDS) on a JSM-6390LV instrument.

121

122 **2.2. Disinfection experiments**

123 The bacterial strain used in the present study was *K. pneumoniae* NCTC 5056 (Public Health
124 England Culture Collections). The reference strain was inoculated in 10 mL of nutrient broth
125 (HiMedia Laboratories) and grown overnight at 37 °C. The concentration of bacterial cells in
126 the suspension was estimated measuring its optical density at 600 nm (Shimadzu UV1240
127 spectrophotometer) where, according to McFarland scale, an absorbance of 0.132 corresponds
128 approximately to a cell density of 1.5×10⁸ CFU mL⁻¹. Plate counts were also performed for
129 accurate bacterial count. In each case, suspensions were properly diluted to achieve the
130 desired initial bacterial concentration, which was used for the subsequent experiments.

131 Photocatalytic experiments were conducted in batch type, laboratory scale photoreactor.
132 Solar irradiation experiments were carried out in a solar radiation simulator system (Newport,
133 model 96000) equipped with a 150 W xenon ozone-free lamp and an Air Mass 1.5 Global
134 Filter (Newport, model 81094), simulating solar radiation reaching the surface of the earth at
135 a zenith angle of 48.2°. The system is described in detail in our previous work.¹¹ Reactions
136 took place in an open, double-walled, cylindrical glass vessel under continuous stirring.

137 In a typical run, the bacteria suspension was spiked in sterile water of 200 mL which were
138 then loaded in the reaction vessel with the appropriate amount of catalyst. The solution was
139 left in the dark under stirring for 20 min to ensure complete equilibration of
140 adsorption/desorption of *K. pneumoniae* bacteria onto the catalyst surface and then exposed to
141 solar irradiation; this moment was taken as the starting point (time zero) of the disinfection
142 experiment. Temperature was maintained at 25 ± 2 °C with a temperature control unit. Catalyst
143 loading was 250 mg L^{-1} in all experiments, since this concentration proved to be optimum for
144 total elimination of *K. pneumoniae* under the current operating conditions. Disinfection rate
145 was measured in terms of *K. pneumoniae* inactivation, according to viable counts on M-FC
146 agar M1124 (HiMedia Laboratories) after incubation at 37 °C for 20-24 h.

147 Furthermore, disinfection durability experiments were carried out in the dark and at natural
148 sunlight irradiation under continuous stirring for 48 h, in order to evaluate the regrowth
149 potential of our strain.

150

151 **2.3. DNA isolation and RAPD-PCR analysis**

152 DNA was extracted from viable cells of *K. pneumoniae* after treatment using QIAamp DNA
153 mini kit (Qiagen) supplemented with lysozyme lysis buffer (100 mM NaCl, 500 mM Tris [pH
154 8], lysozyme 10 mg mL^{-1}). The quantity and purity of all DNA samples were determined
155 measuring their absorbance value at 260 nm and estimating the ratio of absorbance values at
156 260 nm and 280 nm, respectively.

157 For RAPD analyses different 10-nt primers of arbitrary sequence were tested and two of
158 them (RAPD4: 5'-AAGACGCCGT-3' and RAPD7: 5'-GTGGATGCGA-3') were chosen for
159 further study as they gave the most discriminating RAPD fingerprints.^{5,25} PCR was performed
160 with the use of peqlab thermal cycler (advanced primus 25) in 50µl reaction volumes
161 containing approximately 30 ng of bacterial DNA, MgCl_2 2.5 mM, 10 X PCR buffer, 20 pmol

162 of the primer, 1 U of AmpliTaq DNA polymerase and 250 μM of each dNTP. All data used in
163 the present analysis were generated in a cycling program of 45 cycles of 94 $^{\circ}\text{C}$, 1 min; 36 $^{\circ}\text{C}$,
164 1 min and 72 $^{\circ}\text{C}$, 1 min followed by a final extension at 72 $^{\circ}\text{C}$ for 9 min. PCR products were
165 electrophoretically separated by 2% (w/v) agarose gels, which contained 1 mg mL^{-1} ethidium
166 bromide staining. A molecular size marker (1 Kb) was used for reference in all gels, which
167 were further visualized and photographed on a UV transilluminator. Electrophoresed agarose
168 gels were analyzed visually and scored using a binary code. The UVibandmap software
169 (UVITEC Cambridge) was used to analyse the binary code, perform cluster analysis and
170 generate dendrogram based on Jaccard coefficient and unweighted-pair group method with
171 arithmetic mean clustering method (UPGMA).²⁷ The similarity index of RAPD profiles was
172 defined as the fraction of shared bands: $S_{xy} = 2n_{xy} / (n_x + n_y)$. Specifically, for individuals x and y,
173 it is the number of common bands in their RAPD profiles (n_{xy}) divided by the average number
174 of bands scored for both individuals. Similarity S_{xy} values range between 0 and 1. The highest
175 value represents completely similar RAPD profiles.²⁹

176 The reproducibility of the technique was assessed in the following ways: a) Independent
177 RAPD assays were applied in three runs using the same DNA template and b) RAPD assays
178 were applied to *K. pneumoniae* isolates obtained from different colonies of the reference
179 strain, grown on the same petri dish. Band repeatability was defined as $R_b = 2b_{12} / (b_1 + b_2)$,
180 where b_{12} is the number of individuals possessing band b in both replicates and b_1 and b_2 are
181 the numbers of individuals possessing that band in the first and the second replicate,
182 respectively. R_b can take values between 0 and 1.²⁹

183

184 **3. Results & Discussion**

185 **3.1 Inactivation of *K. pneumoniae* during photocatalysis**

186 Bacterial inactivation in water samples was recorded under solar irradiation in the presence of
187 the prepared doped catalysts. Metal-doped catalysts showed satisfactory inactivation, as may
188 be observed in Figure 1. Dopants improved the photocatalytic activity of pure titania, which
189 achieved a 2-Log reduction after 30 min of treatment. Moreover, increasing the dopant
190 concentration incorporated in titania nanoparticles, disinfection efficiency was improved as
191 reduction of *K. pneumoniae* density was 3 orders of magnitude within almost 10 min of
192 treatment. It is clearly shown that the 0.3 wt% Mn- and Co-doped TiO₂ catalysts showed
193 optimum photocatalytic performance when compared to the others and disinfection took place
194 after only 10 min, despite the fact that this specific emerging pathogen is considered
195 persistent during various treatments and disinfection techniques.⁶ These catalysts may
196 demonstrate equal performance even in the case of higher initial concentration of *K.*
197 *pneumoniae* (i.e. 10⁵ CFU mL⁻¹), as it has already been shown in our previous study.¹¹
198 Moreover, it can be noticed that in certain experimental runs and when Mn-doped catalysts
199 were employed, bacterial inactivation reached a plateau and became slow, leading in residual
200 cells at the end of the process. Similar findings were obtained in other study referred to *K.*
201 *pneumoniae* inactivation with the use of reactive plasma processed nanocrystalline TiO₂
202 powder.⁷ Vijay et al. noticed deceleration of the decay of this bacterial strain in the course of
203 treatment, attributing this outcome to the released metabolites from the killed bacteria, which
204 form a screen and protect the remaining active cells. The same conclusion was drawn in the
205 work of Yousef et al., who investigated the effect of CuO/TiO₂ nanofibers on *K. pneumoniae*
206 elimination under visible light.²

207 Further experiments were performed with the use of Mn/Co binary-doped TiO₂, which
208 showed enhanced disinfection efficiency, under the current experimental conditions.
209 Complete elimination of the pathogen was recorded in really short periods of time (Figure 1).
210 This finding verifies the general aspect that composite dopants induce a synergistic effect

211 such as an efficient charge separation and improvement of photostability, surpassing the
212 disadvantages of the individual components.³⁰ The superiority of binary-doped catalysts has
213 already been addressed in other cases with *E. coli* being the testing microorganism and an
214 average 4-Log reduction of its density after 90 min of exposure to visible light.^{31,32} The ratio
215 of doping levels (0.04 and 0.01 %wt of Mn/Co) proved to be optimum for rapid killing of *K.*
216 *pneumoniae* in water samples, although it should not be ignored that dopants have the
217 potential to cause photo-induced corrosion and promote charge recombination at some metal
218 sites.³³

219 Bacterial regrowth after treatment is considered an issue of great importance and many
220 concerns are raised for public health protection since microorganisms are capable of retaining
221 their virulent characteristics, which may be exhibited post disinfection process. Regrowth
222 potential of our strain was evaluated applying disinfection durability experiments. The vast
223 majority of treatment runs did not exhibit any regrowth of the bacterium. Only in three cases
224 was the strain reactivated during exposure under natural sunlight (Table 1). When *K.*
225 *pneumoniae* was processed with 0.02% Mn:TiO₂, 0.1% Mn:TiO₂ and 0.1% Co:TiO₂, there
226 was a photo-reactivation of the bacterium which did not exceed 0.2% of its initial population.
227 Recovery rates in the present study are very low, indicating that damage of cell membranes
228 during photocatalysis was extensive. According to the general observation cell death may be
229 achieved through membrane destruction, which in turn results in leakage of cellular
230 components. Conversely, a respective damage of the cell wall does not necessarily lead to cell
231 death, as bacteria possess a mechanism for its repair.³⁴ Moreover, the change in the
232 concentration of the cell wall components during illumination shows that, while the outer
233 membrane serves as a barrier, the peptidoglycan layer does not have a barrier function.² Our
234 results indicate that the detrimental effect of the oxidative species developed at the catalyst

235 surface was expanded towards the cytoplasmic membrane, increasing cell permeability which
236 finally caused cell death.¹¹

237

238 **3.2 RAPD-PCR analysis**

239 RAPD-PCR fingerprinting was employed as a typing method to highlight and discriminate
240 possible alterations in the genomic material of *K. pneumoniae* isolates, which remained viable
241 after treatment with the metal-doped catalysts. As a PCR-based technique, it was selected for
242 its credit classification results relying on genomic divergence of the tested bacterial isolates.
243 Generally, in RAPD analysis arbitrary oligonucleotides are used to promote DNA
244 amplification at low annealing temperatures determining genomic diversity. This generates
245 strain-specific arrays of amplified DNA fragments. The formation of these arrays does not
246 depend on prior knowledge of the nucleotide sequence, nor is it affected by DNA
247 modifications that complicate typing by restriction endonuclease digestion of genomic DNA.
248 This technique has been preferred in many studies because of its speed, simplicity and highly
249 discriminatory results.^{3-5,8,10,25-27}

250

251 **3.2.1 Reproducibility of RAPD analysis**

252 The reproducibility of the method was evaluated with preliminary trials using the same DNA
253 template and genetic material from different *K. pneumoniae* colonies. Despite the advantages
254 of RAPD there are some considerable concerns and criticism regarding the lack of
255 repeatability, due to low stringency conditions employed and variable factors that may affect
256 RAPD profiles.⁵ The reproducibility of the technique was assessed as mentioned before and
257 the average R_b (band repeatability) was estimated in profiles generated with the two selected
258 primers RAPD7 and RAPD4. When the same DNA template was used R_b ranged between
259 0.96 and 0.99 with both primers used. The corresponding values when DNA was isolated

260 from different colonies of the reference strain were in the range of 0.93-0.97. Generally, the
261 results were highly reproducible with both primers and under the specific experimental
262 conditions.

263 It is of great importance that reaction conditions, including DNA template concentration,
264 annealing temperature and the choice of the primers should be standardized to avoid
265 variations in RAPD patterns.²⁷ Upon optimization of conditions, RAPD offers the possibility
266 of studying, at least within a single run, the genotypic relatedness of a limited number of
267 isolates.²⁵ Ashayeri-panah et al., who worked with *Klebsiella* isolates and RAPD analysis,
268 concluded that the reproducibility of the technique may be achieved even when different
269 thermal cyclers are used under strictly defined conditions.⁵ In this sense, this method may
270 provide a stable and highly reproducible discrimination of bacterial isolates, as it has already
271 been documented in the literature.^{3,4,8}

272

273 **3.2.2 RAPD profiles of viable cells after treatment**

274 Several primers were tested for RAPD fingerprinting, among which RAPD7 and RAPD4
275 were the most discriminatory in relation to the number and intensity of generated bands, size
276 range and smear formation. Both primers were selected for detailed testing of sequence
277 divergence among isolates. Viable isolates after photocatalysis with the metal-doped titania
278 were subject to RAPD-PCR and all of them proved to be typeable with both primers,
279 developing a variety of amplification bands. RAPD profiles and the corresponding cluster
280 analysis of treated *K. pneumoniae* with Mn-, Co- and binary-doped catalysts are shown in
281 Figures 2, 3 and 4, respectively. The protocol performed in this study allowed the
282 amplification of 6-23 bands ranging in size from 200 to 2700 bp with primer RAPD7 and 7-
283 29 bands ranging from 200 to 4000 bp with primer RAPD4.

284 Reviewing the patterns obtained from isolates treated with Mn-doped TiO₂ in comparison
285 to that of intact cells, different combinations of amplification bands were obvious with both
286 primers resulting in their discrete grouping applying cluster analysis (Figure 2). Isolates were
287 divided into different groups, according to their homology with the intact *K. pneumoniae*. In
288 the course of treatment and increasing the period of irradiation a considerable alteration of
289 RAPD profiles occurred in viable bacterial cells. Homology of RAPD patterns ranged
290 between 27-88% and 38-90% with primers RAPD7 and RAPD4, respectively. Significant
291 discrimination among isolates was recorded after long treatment periods. For example when
292 bacteria were treated with 0.02% Mn:TiO₂, the homology of RAPD patterns between
293 untreated and viable cells after 30 min of irradiation was only 27%. The concentration of the
294 dopant did not affect significantly the profiles. Conversely, the period of irradiation in each
295 case affected the overall alteration in band patterns. Interestingly, although the band patterns
296 were different for each primer similar dendrograms and clusters were obtained but with
297 different similarity index S_{XY} (Table 2). Primer RAPD4 yielded patterns with lower values of
298 similarity index, compared with those recorded with primer RAPD7. The average S_{XY} of
299 profiles were 0.46 and 0.60 with RAPD4 and RAPD7, respectively, when Mn:TiO₂ was
300 applied. This finding indicates that the former primer exhibits higher discriminatory power, in
301 terms of highlighting the induced changes in genetic material of *K. pneumoniae* isolates under
302 solar irradiation with the use of the prepared metal-doped catalysts.

303 Screening results obtained when Co:TiO₂ was used it is evident that they follow the same
304 trend (Figure 3). Band patterns obtained with primer RAPD4 showed significant
305 heterogeneity among *K. pneumoniae* isolates, with the untreated one being grouped separately
306 from all others and the homology coefficient reaching a value of almost 35%. A period of 15
307 min of irradiation with the 0.3% Co:TiO₂ was sufficient to cause such alteration in the genetic
308 material of viable bacteria. The similarity index among profiles obtained with primer RAPD4

309 in the presence of Co-doped titania had even lower average value than that recorded with Mn-
310 doped titania (i.e. 0.23), as may be seen in table 2. The differences in the discriminatory
311 power of the tested primers became apparent after analysis of all viable isolates after
312 disinfection with the three sets of metal-doped catalysts (i.e. Mn-, Co- and Mn/Co binary-
313 doped). The use of two different primers to generate RAPD profiles gave an extended
314 flexibility and sensitivity to the typing method. The use of more than one primer seems to be
315 advantageous, since it adds to the consistency of the established groups and in many studies
316 applying RAPD it is preferred to employ a set of oligonucleotides for accurate and reliable
317 bacterial clustering.^{5,25}

318 In the case of binary-doped catalysts, lower homology coefficient among isolates was
319 achieved with primer RAPD4, reaching a value of almost 42%, whereas better discrimination
320 and grouping of patterns was generated with primer RAPD7 (Figure 4) but with higher
321 average value of similarity index (i.e. 0.69; Table 2). The demonstrated dendrogram showed
322 all treated isolates into one major group from a common lineage with further separation into
323 smaller sets, whose similarity level with the untreated *K. pneumoniae* was approximately
324 60%. It should be noted though that when treatment process was performed with binary-
325 doped titania, the recorded genetic variation was referred to isolates that were exposed to solar
326 irradiation for only 10 min, as complete removal of the bacteria occurred within this period of
327 time.

328 RAPD fingerprinting method could be used to rapidly identify major clones of this species,
329 including multi-drug resistant and particularly virulent clones. According to de Souza et al.,
330 pathogenic *K. pneumoniae* populations are highly heterogeneous.²⁷ The numerous serotypes
331 in this species could explain the relevant degree of genetic diversity highlighted by RAPD
332 analysis, as it was exhibited in our study. Polymorphisms detected throughout the entire
333 genome may include point mutation and genetic rearrangement, which have the potential to

334 increase the already significant virulence of the species. The occurrence of broad spectrum of
335 *K. pneumoniae* variants in stressed environments portrays the ability of this species to survive
336 abundantly under extreme conditions.¹⁰

337

338 **4. Conclusions**

339 The present study focused on the inactivation of *K. pneumoniae* in aqueous matrix under solar
340 irradiation with the use of metal-doped titania and the induced genetic variation among viable
341 cells after treatment. In this perspective RAPD-PCR was performed as a molecular typing
342 tool, that has the potential to highlight genetic diversity among bacterial isolates from various
343 sources. The main conclusions derived from this work are as follows:

- 344 - Inactivation rates of *K. pneumoniae* in water samples under simulated solar irradiation
345 were highly improved in the presence of metal-doped catalysts. Of all the catalysts used,
346 Mn/Co binary-doped titania exhibited the best photocatalytic activity and complete
347 bacterial removal was recorded in less than 10 min of treatment. This finding verifies the
348 synergistic effect induced by composite dopants.
- 349 - All catalysts were effective for considerable inactivation of *K. pneumoniae* with very low
350 levels of bacterial regrowth in the dark and under natural sunlight.
- 351 - RAPD-PCR proved to be a useful typing molecular tool, that upon standardized conditions
352 exhibits highly reproducible results.
- 353 - Genetic variation among isolates increased in relation to the period of treatment. The
354 period of irradiation in each case affected the overall alteration in band patterns.
- 355 - RAPD patterns were highly diverse between treated and untreated isolates when
356 disinfection was performed with the Co-doped titania.
- 357 - Generated polymorphisms may add to the virulence of the species.

358

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362

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- 423

424 **Table 1**

425 *K. pneumoniae* survival after the end of disinfection durability experiments either in the dark
 426 or under natural sunlight irradiation.

| Operating conditions | | | <i>K. pneumoniae</i> regrowth (CFU mL ⁻¹) | | | |
|----------------------|---------------------------|--------------------------------------|---|-------------------------------------|------------------------------------|-------------------------------------|
| Run | Catalyst | Initial bacterial concentration | Dark conditions (24 h of exposure) | Natural sunlight (24 h of exposure) | Dark conditions (48 h of exposure) | Natural sunlight (48 h of exposure) |
| 1 | 0.02% Mn:TiO ₂ | 10 ³ CFU mL ⁻¹ | 0 | 0 | 0 | 1 |
| 2 | 0.1% Mn:TiO ₂ | 10 ³ CFU mL ⁻¹ | 0 | 1 | 0 | 2 |
| 3 | 0.1% Co:TiO ₂ | 10 ³ CFU mL ⁻¹ | 0 | 0 | 0 | 1 |

427

428

429 **Table 2**

430 Average similarity index (S_{XY}) of generated RAPD profiles of viable isolates after treatment
431 with the metal-doped catalysts.

| Catalysts used during photocatalysis | Average similarity index (S_{XY}) of RAPD profiles | |
|---|--|----------------------|
| | <i>Primer RAPD 7</i> | <i>Primer RAPD 4</i> |
| Mn-doped TiO ₂ | 0.60 ± 0.14 | 0.46 ± 0.18 |
| Co-doped TiO ₂ | 0.68 ± 0.08 | 0.23 ± 0.07 |
| Mn/Co binary-doped TiO ₂ | 0.69 ± 0.05 | 0.39 ± 0.17 |

432

433 **Figure captions**

434 **Figure 1**

435 *K. pneumoniae* inactivation in the presence of different a) Mn- doped TiO₂ and the
436 commercially available TiO₂ (P25, Evonik) and b) Co- and Mn/Co binary-doped TiO₂.
437 Catalyst concentration is 250 mg L⁻¹.

438 **Figure 2**

439 RAPD patterns generated with a) primer RAPD7 and b) primer RAPD4 and corresponding
440 cluster analysis of *K. pneumoniae* isolates treated with Mn-doped TiO₂. Prefix NT indicates
441 the not treated isolate. Lane numbers correspond to *K. pneumoniae* treated isolates as follows,
442 1, 2: 0.02% Mn:TiO₂ after 6 and 30 min, respectively; 3, 4: 0.04% Mn:TiO₂ after 6 and 30
443 min, respectively; 5, 6: 0.1% Mn:TiO₂ after 3 and 10 min, respectively; 7, 8: 0.3% Mn:TiO₂
444 after 6 and 15 min, respectively. Dendrogram generated with Jaccard's coefficient and the
445 UPGMA clustering method.

446 **Figure 3**

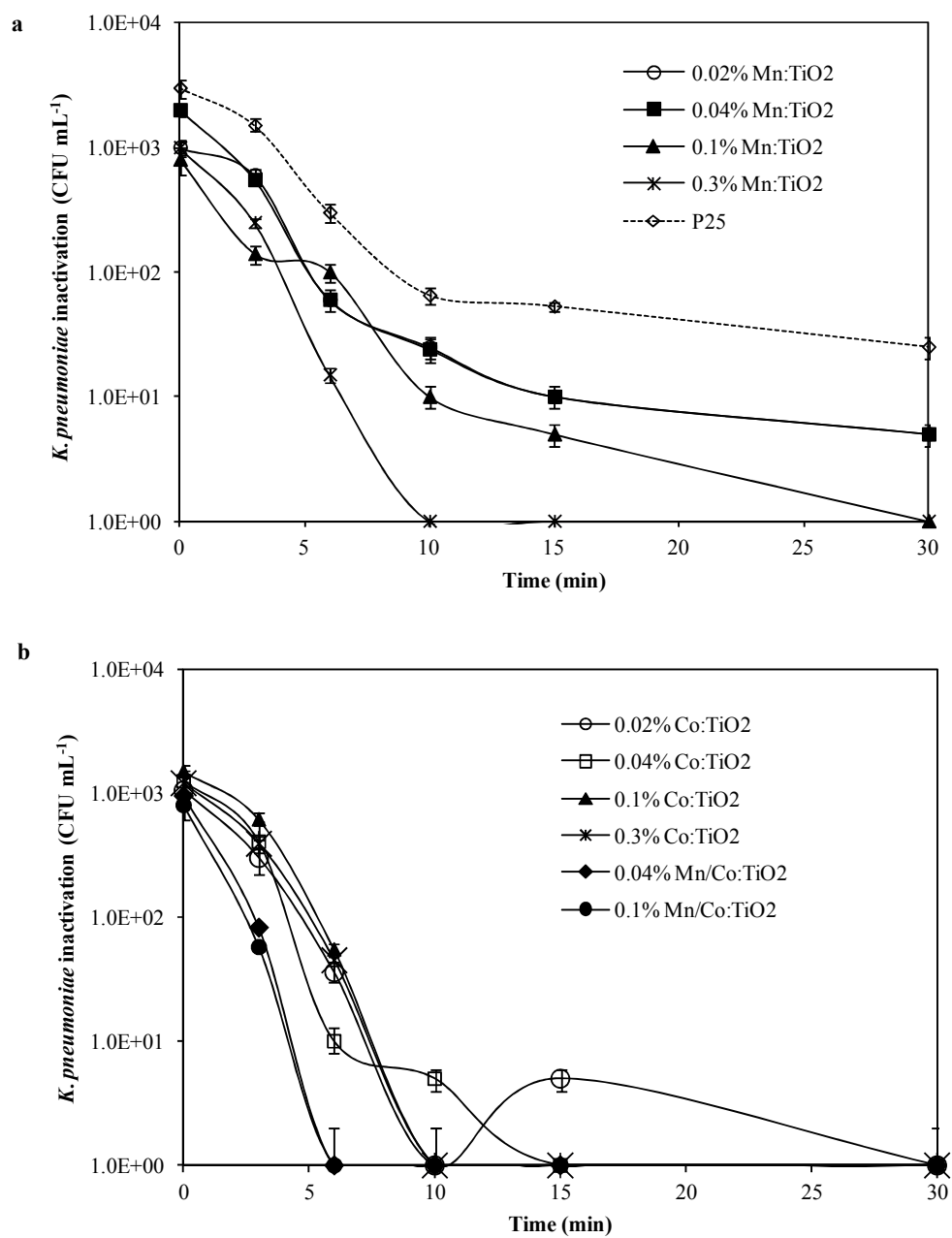
447 RAPD patterns generated with a) primer RAPD7 and b) primer RAPD4 and corresponding
448 cluster analysis of *K. pneumoniae* isolates treated with Co-doped TiO₂. Prefix NT indicates
449 the not treated isolate. Lane numbers correspond to *K. pneumoniae* treated isolates as follows,
450 1, 2: 0.02% Co:TiO₂ after 6 and 15 min, respectively; 3, 4: 0.04% Co:TiO₂ after 6 and 10
451 min, respectively; 5, 6: 0.1% Co:TiO₂ after 6 and 10 min, respectively; 7, 8: 0.3% Co:TiO₂
452 after 3 and 15 min, respectively. Dendrogram generated with Jaccard's coefficient and the
453 UPGMA clustering method.

454 **Figure 4**

455 RAPD patterns generated with a) primer RAPD7 and b) primer RAPD4 and corresponding
456 cluster analysis of *K. pneumoniae* isolates treated with Mn/Co binary-doped TiO₂. Prefix NT
457 indicates the not treated isolate. Lane numbers correspond to *K. pneumoniae* treated isolates

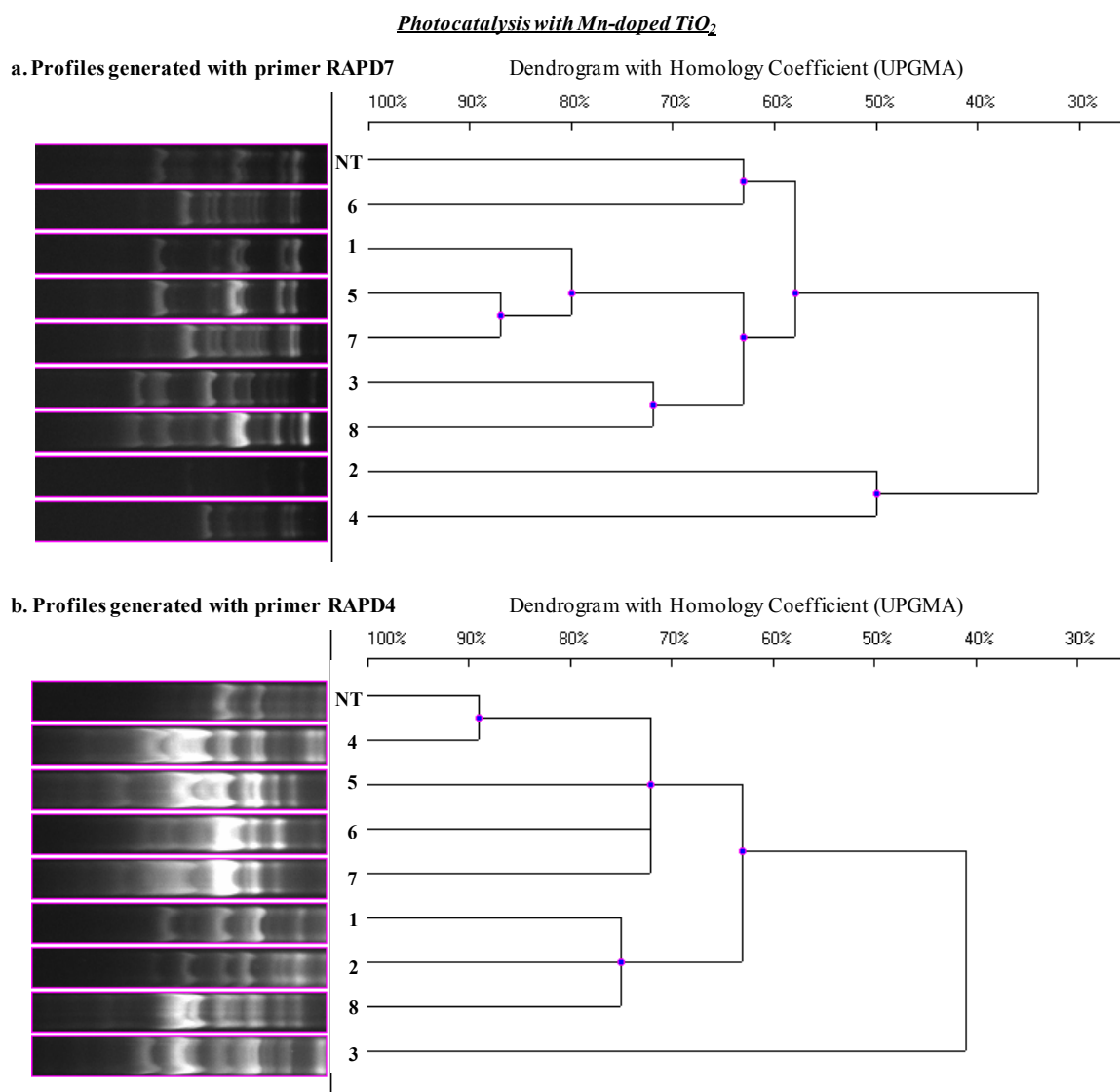
458 as follows, 1, 2: 0.04% Mn/Co:TiO₂ after 6 and 10 min, respectively; 3, 4: 0.1% Mn/Co:TiO₂
459 after 6 and 10 min, respectively. Dendrogram generated with Jaccard's coefficient and the
460 UPGMA clustering method.
461

462 Figure 1



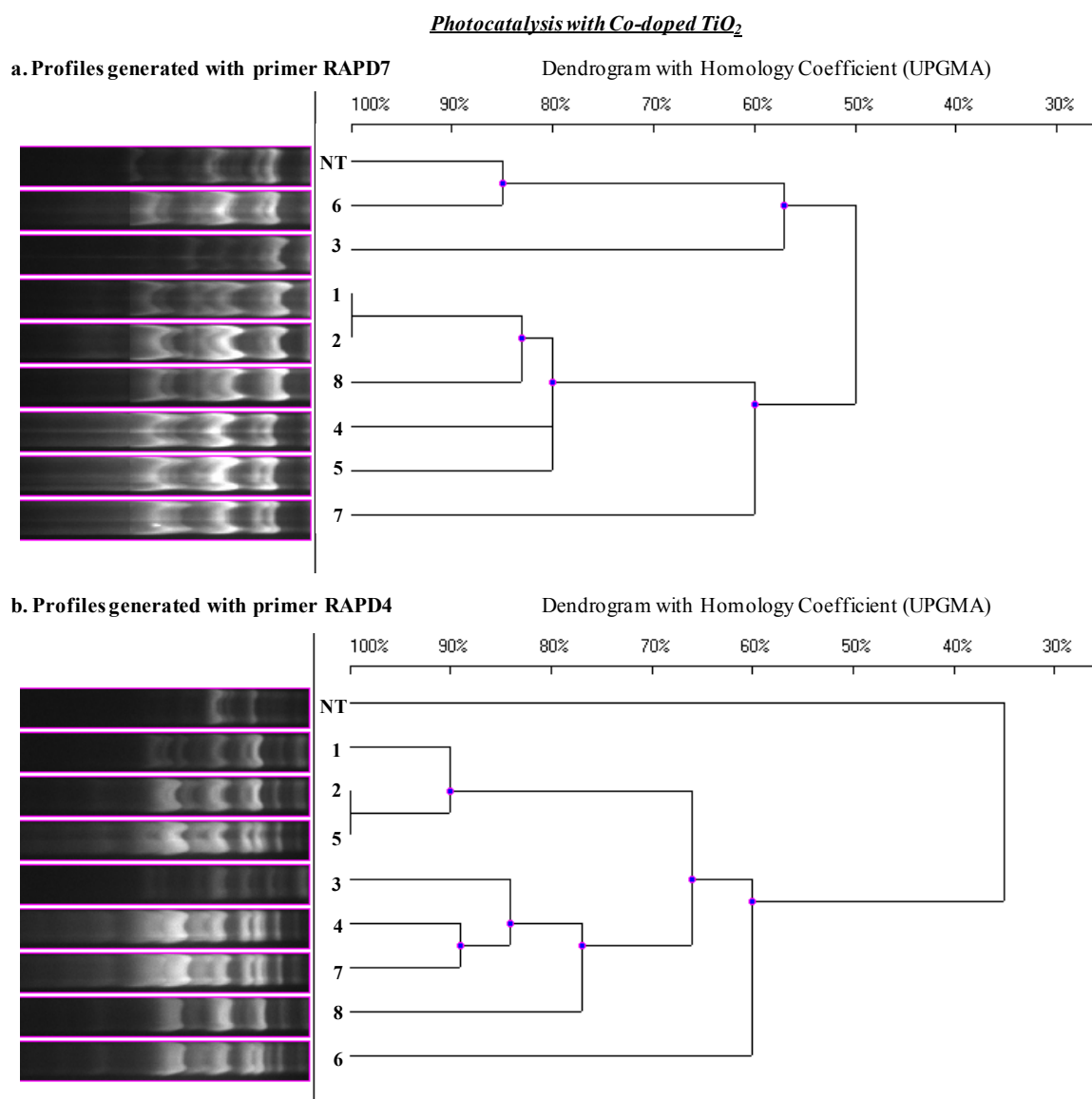
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464

465 **Figure 2**

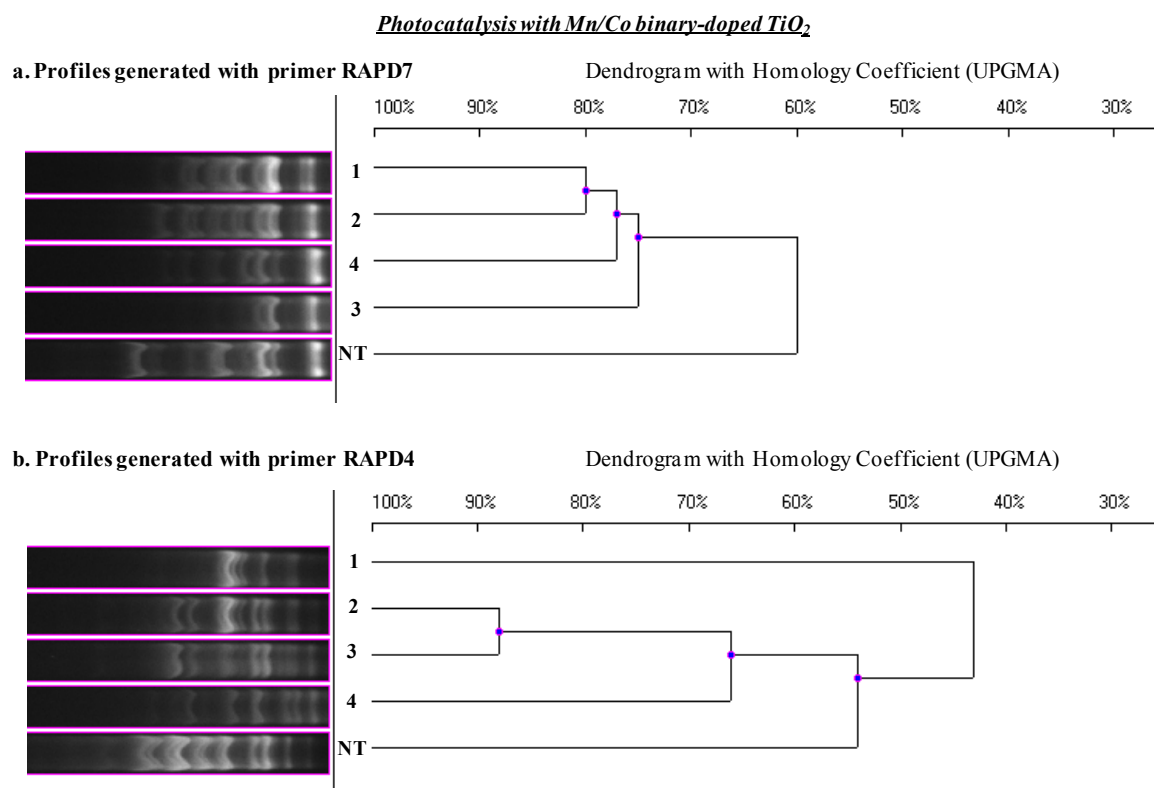
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468 **Figure 3**

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471 **Figure 4**

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