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1 **Exploring and strengthening the potential of R-phycoyanin from Nori flakes as**
2 **a food colourant**

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26 **Abstract**

27 This study aimed to purify, characterise and stabilise the natural food colourant, R-phycoyanin (R-
28 PC), from the red algae *Porphyra* spp. (Nori). We purified R-PC from dried Nori flakes with a high
29 purity ratio ($A_{618}/A_{280} \geq 3.4$) in native form (α -helix content 53%). SAXS measurements revealed
30 that R-PC is trimeric ($(\alpha\beta)_3$) in solution. The thermal denaturation of α -helix revealed one transition
31 (T_m at 52 °C), while the pH stability study showed R-PC is stable in the pH range 4 to 8. The
32 thermal treatment of R-PC at 60 °C has detrimental and irreversible effects on R-PC colour and
33 antioxidant capacity (22 % of residual capacity). However, immobilisation of R-PC within calcium
34 alginate beads completely preserves R-PC colour and mainly retains its antioxidant ability (78 % of
35 residual capacity). Results give new insights into the stability of R-PC and preservation of its purple
36 colour and bioactivity by encapsulation in calcium alginate beads.

37 **Keywords:** R-phycoyanin, Purification, Stability, SAXS, Immobilization, Calcium alginate.

38 Introduction

39 Alternative proteins have recently gained increasing attention due to substantial
40 environmental and health impacts compared to meat consumption (Onwezen, Bouwman, Reinders,
41 & Dagevos, 2021). Algae are a promising source of alternative proteins as they can meet health,
42 nutritional, environmental and production needs (Diaz et al., 2022). Red macroalgae *Porphyra* spp.,
43 commonly known as Nori, is a rich source of proteins, vitamins, minerals and antioxidants and has
44 been used as food for centuries in Asia, and its popularity has now spread to other parts of the world
45 (Bito, Teng, & Watanabe, 2017; Simovic, Combet, Cirkovic Velickovic, Nikolic, & Minic, 2022).
46 Consumed as dried Nori flakes, it is frequently used to prepare sushi, soup and cakes (Bito et al.,
47 2017). The most abundant proteins in *Porphyra* spp. are the phycobiliproteins (PBPs) R-
48 phycoerythrin (R-PE) and R-phycoerythrin (R-PC), contributing to approximately 3 % of the dry
49 mass of *Porphyra* spp. (Cao, Wang, Wang, & Xu, 2016). Considering that nearly one million tons
50 of *Porphyra* are produced annually (Baweja, Kumar, Sahoo, & Levine, 2016), R-PC and R-PE have
51 great potential to be used as alternatives to animal proteins. Besides R-PE and R-PC, *Porphyra* spp.
52 also contains the third phycobiliprotein allophycoerythrin (APC), the minor protein of the PBP
53 family (Liu et al., 1998). Compared to R-PE, which has been thoroughly studied, much less is
54 known about the properties of R-PC of *Porphyra* spp.

55 R-phycoerythrin is a strongly fluorescent and water-soluble molecule whose primary function
56 is to transfer excitation energy to reaction centres during photosynthesis (Glazer & Hixson, 1975;
57 Jiang, Zhang, Chang, & Liang, 2001). The intensive purple colour of R-PC arises from its
58 covalently attached (*via* thioether bond) linear tetrapyrrole chromophores, phycoerythrobilin (PEB)
59 and phycoerythrin (PCB). It is composed of two subunit types forming a trimer ($\alpha\beta$)₃ or hexamer
60 ($\alpha\beta$)₆, with α subunits of 18–20 kDa and β subunits of 19–21 kDa (Jian et al., 2001). One PCB
61 chromophore is bound to both α and β subunits, while one PEB chromophore is only attached to β
62 subunits.

63 R-phycoerythrin has potential bioactive properties due to the potent antioxidant,
64 immunomodulatory and anticancer activities of covalently bound tetrapyrrole chromophores (Guo
65 et al., 2022; Rashed, Hammad, Eldakak, Khalil, & Osman, 2023). The health benefits of R-PC, its
66 brilliant fluorescence, vivid colour, and the need to replace synthetic dyes give R-PC great potential
67 for food fortification and colouring. Nori seaweed represents an essential source of R-PC as natural
68 food colourant because it can be cultivated in vast amounts, as described above. Hence, tons of R-
69 PC could be produced annually to colour different foods and beverages. However, in this
70 perspective, R-PC's limited stability is the major obstacle. Indeed, generally, PBPs have moderate
71 pH stability (Munier et al., 2014; Yan et al., 2010), and their thermal stability is low: an increase in

72 temperature above 40 °C induces a substantial decrease in protein stability and colour intensity
73 (Faieta, Neri, Sacchetti, Di Michele, & Pittia, 2020; Munier et al., 2014; Simovic et al., 2022).

74 Therefore, developing R-PC as a natural food colourant and alternative protein requires a
75 deeper understanding of its structural and functional properties and improved stability. One
76 promising approach for stabilising R-PC and other PBPs could be encapsulation in solid food
77 matrices, which may decrease the mobility of protein molecules, making them more resistant
78 towards temperature and oxidation processes (Maghraby, El-Shabasy, Ibrahim, & Azzazy, 2023). In
79 this context, alginate immobilisation is a valuable strategy since it is widely used in biotechnology
80 and bioprocessing for the entrapment of living cells, proteins, and other biological molecules
81 (Andersen, Auk-Emblem, & Dornish, 2015). Alginate is a natural polysaccharide derived from
82 brown seaweed and has unique physical and chemical properties that make it ideal for
83 immobilisation. The process involves the formation of calcium-alginate beads, which can be easily
84 manipulated and controlled in terms of size, shape and mechanical stability (Segale, Giovannelli,
85 Mannina, & Pattarino, 2016).

86 In the present study, we aim to purify, characterise and stabilise the purple protein R-PC
87 from dried Nori flakes for the first time. We used precipitation and chromatography techniques to
88 obtain pure protein. Optical spectroscopy and SAXS measurements conclude that purified protein is
89 a typical R-PC, with α -helix as the dominant secondary structure and trimeric oligomerisation state
90 in solution. We determined the sequence of R-PC from Nori flakes by mass spectrometry and
91 compared it with phycocyanins from other sources. Further, we employed absorption and far-UV
92 CD spectroscopy to study the effects of temperature and pH on protein stability. Ultimately, we
93 tested if R-PC immobilisation within calcium alginate beads could improve protein thermal stability
94 and preserve its antioxidant capacity.

95 **2. Materials and Methods**

96 **2.1 Materials**

97 Dried Nori flakes (*Porphyra* spp.) were purchased from Fujian Friday Trading Co. LTD
98 (China). Hydroxyapatite (HA) and diethyl aminoethyl (DEAE) Sepharose resins were bought from
99 BioRad (CA, USA) and Sigma (MO, USA), respectively. Sodium alginate used for immobilisation
100 was of food grade, purchased from Health Leads (France). All other chemicals were of analytical
101 reagent grade.

102 **2.2 R-PC isolation and purification**

103 R-Phycocyanin was purified according to the previously published procedures with
104 modifications (Boussiba & Richmond, 1979; Simovic et al., 2022; Wang et al., 2014). Nori

105 seaweed leaves were frozen with liquid nitrogen and then pulverised using a mortar and pestle. The
106 powdered sample was then suspended in 20 mM phosphate buffer pH 7.0 (1 g of powdered sample
107 *per* 20 mL of buffer). Extraction of phycobiliproteins was performed overnight at 4 °C with
108 constant stirring. The resulting slurry was centrifuged at 6000 x g for 30 min at 6 °C. Proteins in the
109 supernatant were precipitated with ammonium sulfate (65 % final concentration). The pellet was
110 dissolved in 2.5 mM phosphate buffer pH 7.0 and was dialysed against the same buffer.

111 The dialysed solution was applied to the CHT ceramic hydroxyapatite (HA) column,
112 previously equilibrated with the same buffer. Non-bound proteins (mainly R-PE) were washed
113 away with the equilibration buffer, while fractions enriched with R-PC were obtained by stepwise
114 elution protocol employing increasing concentrations of sodium phosphate salt (pH 7.0; **Fig. 1A**).
115 R-PC containing fraction was applied onto the DEAE Sepharose CL-6B column, previously
116 equilibrated with 20 mM phosphate buffer pH 7.0. Unbound proteins were flushed with an
117 equilibration buffer. Bound proteins were eluted stepwise, increasing the ionic strength of NaCl
118 from 0.05 to 0.2 M (**Fig. 1B**). The R-PC-rich fractions were pulled, desalted and concentrated.
119 Purified R-PC was stored in 40 % (v/v) glycerol solution at -20 °C. We analysed the obtained
120 fractions by UV-Visible absorption spectroscopy (**Section 2.3**) and SDS-PAGE under reducing
121 conditions. Fluorescent PBP bands were first visualised under UV light, followed by staining with
122 Coomassie Brilliant Blue R-250.

123 **2.3 Determination of R-PC amino acid sequence by mass spectrometry**

124 Purified R-PC was digested by trypsin, and obtained peptides were analysed by nano Liquid
125 Chromatography-electron spray ionisation coupled to tandem mass spectrometry (nLC-ESI-
126 MS/MS). The protein identification was performed using PEAKS Suite X (Bioinformatics
127 Solutions Inc., Canada) software. Experimental details and mass spectrometry data analysis are
128 given in Supplementary Material 1.

129 **2.4 UV-Visible absorbance measurements**

130 UV-Visible absorption spectra were recorded on a NanoDrop 2000c spectrophotometer
131 (Thermo Scientific, USA) in a 1 cm pathlength quartz cell. The spectra were collected in the 250–
132 750 nm range at room temperature. R-PC concentration was determined using the following
133 equation (Nikolic, Minic, Macvanin, Stanic-Vucinic, & Velickovic, 2020):

$$134 \quad R - PC \left(\frac{mg}{mL} \right) = 0.154(A_{618} - A_{730}) \quad (1)$$

135 The effects of temperature and pH on visible absorption spectra of R-PC were studied using
136 a UV-1800 Shimadzu spectrophotometer (Japan). Temperature measurements were performed

137 using a Peltier element previously described (Simovic et al., 2022), and experimental details are
138 given in Supplementary Material 1.

139 The pH dependence of R-PC absorption spectra was studied using the same measurement
140 parameters described above for the thermal stability study. R-PC was incubated in the different
141 buffers (50 mM phosphate, citrate, Tris and glycine) in the pH range of 2–12 for 30 minutes,
142 followed by the acquisition of absorption spectra at room temperature.

143 **2.5 Fluorescence measurements**

144 Fluorescence spectra of 1.8 µg/mL of R-PC at pH 7.0 were recorded with a FluoroMax[®]-4
145 spectrofluorometer (HORIBA Scientific, Japan) under thermostable conditions (25 °C), using 5 nm
146 excitation and 5 nm emission slit widths. The PEB chromophore was excited at 488 and 590 nm,
147 and the emission spectra were recorded between 520 and 610 nm. Excitation of PCB chromophore
148 was performed at 590 nm, while the emission was measured in the range 610–680 nm.

149 The spectrofluorimetric oxygen radical absorbance capacity (ORAC) assay was performed
150 according to Ou et al. (Ou, Hampsch-Woodill, & Prior, 2001). Stock solutions of fluorescein (5
151 µM) as the substrate and 2,2'-azobis(2-amidinopropane) dihydrochloride (AAPH, 300 mM) as the
152 free radical generator were made in 75 mM sodium phosphate buffer, pH 7.4. The assay was
153 performed: 50 µL of 100 µg/mL R-PC solution (obtained as described in **section 2.7**) was mixed
154 with 1680 µL of buffer (75 mM phosphate, pH 7.4) and 20 µL of fluorescein solution. The reaction
155 was initiated by the addition of 250 µL of AAPH solution. Fluorescence was measured at the
156 excitation wavelength of 485 nm and an emission wavelength of 511 nm at 37 °C. Bandwidths of
157 slits were set to 2 nm. The fluorescence was recorded every 20 seconds for 30 min. Measurements
158 of blank samples (**section 2.7**) were performed in the same conditions. The blank sample's area
159 under the curve (AUC) was subtracted from the corresponding AUC of R-PC-containing samples,
160 and results were expressed as AUC per mg of R-PC. Measurements were performed at least in
161 duplicate, and results were presented as mean ± standard deviation. For the evaluation of
162 antioxidant capacity, an analysis of variance (ANOVA) of data was performed and means
163 comparisons were made using the Tukey test.

164 **2.6 CD spectroscopy measurements**

165 CD measurements were carried out on a Jasco J-815 spectropolarimeter (Jasco, Japan) under
166 constant nitrogen flow. All spectra were recorded at 25 °C. For the near-UV and visible region
167 (250–750 nm), the concentration and pH of R-PC were set to 0.3 g/L and 7.0, respectively. Scan
168 speed and the number of accumulations were set to 200 nm/min and two, respectively. A cell with
169 an optical pathlength of 1 cm was used.

170 Far-UV CD spectra of 100 µg/mL R-PC have been recorded in the range 190–260 nm at a
171 scan speed of 50 nm/min, using a cell with an optical pathlength of 1 mm and with an accumulation
172 of three scans. Spectra were measured in different buffers in the pH range of 2–12 (**section 2.3**).
173 The CD data analysis is given in Supplementary Material 1.

174 Thermal unfolding R-PC was performed as previously described (Simovic et al., 2022), and
175 experimental details and data analysis are given in Supplementary Material 1.

176 **2.7 SAXS experiments and analysis**

177 SAXS spectra were obtained on the SWING beamline at the French synchrotron facility
178 SOLEIL (St-Aubin, France) (Thureau, Roblin, & Pérez, 2021) using a wavelength of $\lambda = 0.764 \text{ \AA}$
179 and a sample-to-detector distance of 3 m. The achievable q -range was 0.0033 to 0.531 \AA^{-1} , where
180 $q = \frac{4\pi}{\lambda} \sin\theta$ is the modulus of the momentum transfer, and 2θ is the scattering angle. 40 µL of 5
181 mg/mL of R-PC at pH 5.7 (20 mM MES containing 150 mM NaCl) and buffer were injected into a
182 1.5 mm outside diameter quartz capillary thermalised at 15 °C.

183 The SAXS data analysis is given in Supplementary Material 1.

184 **2.8 Immobilisation of R-PC in sodium alginate**

185 The R-PC immobilisation in alginate hydrogel beads according to a previously published
186 procedure (Zhang, Zhang, Zou, & McClements, 2016) with modifications. A solution of R-PC (4
187 mg/mL) was prepared in 25 mM acetate buffer at pH 5. Subsequently, this solution was mixed with
188 a 2 % aqueous solution of sodium alginate in a volume ratio of 1:1. The resulting mixture was then
189 droplet-wise deposited into a 2 % calcium chloride solution of 50 mM acetate buffer at pH 5. The
190 formed beads were stored at 4 °C overnight. The beads containing immobilised proteins were
191 thoroughly washed several times with 50 mM acetate buffer pH 5. The beads were then transferred
192 into a microtube and subjected to thermal treatment at 60 °C for 10 minutes. Beads were dissolved
193 using 100 mM Tris-citrate buffer pH 7.3. Obtained solutions were subjected to an ORAC test
194 (**section 2.5**) to assess the effects of immobilisation and thermal treatment on the antioxidative
195 capacity of R-PC. The dissolved beads without R-PC served as a blank in the ORAC assay. The
196 control experiment comprises heating 2 mg/mL R-PC in solution (non-immobilised) at the same
197 conditions as those used for the thermal treatment of the R-PC containing alginate beads.

198 **3. Results**

199 **3.1 Purification of R-PC from Nori seaweed**

200 We combined precipitation and chromatographic techniques to purify R-PC from Nori
201 flakes. Phycobiliproteins from Nori seaweed exhibit the characteristic absorption peaks: R-PE

202 (peaks at 498 and 560 nm, and shoulder at 545 nm), R-PC (peaks at 550–560 and 618 nm) and APC
203 (peak at 652 nm with a shoulder at 620 nm) (Glazer et al., 1975; Simovic et al., 2022). Obtained
204 raw extract had a strong absorption in the region from 250 to 450 nm due to the presence of various
205 pigments (**Fig. 1C**). Precipitation with ammonium sulphate (65 %) increased the purity ratio
206 (A_{618}/A_{280}) from 0.31 to 0.61 (**Fig. 1C**) by separating most of the non-protein related pigments
207 absorbed in the region 250–450 nm. Different PBPs were not distinguished in this step since
208 ammonium sulphate induces non-selective precipitation of all PBPs presented in the extract (**Fig.**
209 **1C**). Separation of R-phycoerythrin from R-PE was achieved by applying HA chromatography (**Fig.**
210 **1A**). Most R-PE is eluted by equilibration buffer. At the same time, R-PC exhibits much stronger
211 binding to the HA column and 100 mM phosphate is required for its elution. This chromatography
212 step substantially increases A_{618}/A_{560} ratio (1.10) compared to the sample before chromatography
213 (0.67). Although this step removed a significant amount of R-PE, the ratio of A_{618}/A_{280} was still low
214 (1.32), suggesting the presence of other proteins in the R-PC enriched fraction. The same findings
215 are observed by SDS-PAGE after the fluorescence visualisation of PBPs and CBB staining (**Fig.**
216 **1D**). Hence, anion-exchange chromatography on DEAE-Sepharose was employed to purify R-PC
217 further. The chromatogram displayed two peaks, with the R-PC fraction eluting ionic strength of
218 150 mM NaCl. The fraction that was eluted with 200 mM NaCl corresponded to allophycoerythrin
219 and R-PE (high binding fraction).

220 Combining HA and ion-exchange chromatography resulted in the preparation of R-PC with
221 a purity index of A_{618}/A_{280} , around 3.4. The absence of a shoulder at 498 nm in absorption spectra
222 (Simovic et al., 2022), which indicates the complete removal of R-PE (**Fig. 1C**) and the presence of
223 solely two characteristic bands in the electrophoretic profile (**Fig. 1D**) served as confirmation that
224 R-PC of the high purity is obtained. The two observed bands at ~18 and 20 kDa arise from α and β
225 subunits of R-PC. Interestingly, visualisation of SDS-PAGE gel by UV lamp revealed that β
226 subunits exhibit much stronger fluorescence than α subunits. This result could be ascribed to the
227 higher quantum yield of PEB chromophore (Grabowski & Gantt, 1978; Saraswat et al., 2011),
228 exclusively bound to the R-PC β subunits, compared to the PCB chromophore, which is attached to
229 both α and β subunits.

230 **3.2 Spectroscopic and structural characterisation of purified R-phycoerythrin**

231 The purified R-PC was digested by trypsin, and we analysed obtained peptides by tandem
232 mass spectrometry. Based on this analysis (**Supplementary Material 2**), the R-PC sequence of
233 both subunits (α and β) was determined (**Fig. 2A**). Furthermore, we identified the binding sites for
234 tetrapyrrole chromophores: α subunit binds one chromophore *via* Cys84, while two chromophores
235 are attached to β subunit by Cys82 and Cys153 (**Fig. 2A**). Considering the same molecular mass of

236 PCB and PEB, we could not make a distinction between them in MS spectra, but only the position
237 of Cys residue involved in chromophore binding. However, based on the previously published
238 crystal structure of R-PC from *Polysiphonia urceolata* (Jiang et al., 2001), we were able to assign
239 PCB binding to Cys84 and Cys82 of α and β subunits, respectively, while PEB chromophore is
240 bound to the Cys153 of β subunits. BLAST analysis revealed the high sequence identity (around
241 90% or higher) between obtained R-PC from *Porphyra* spp. and other phycocyanins belonging to
242 the different classes of *Rhodophyta* division (**Tables S1 and S2**). We also detected about 70% of
243 sequence homology with the C-phycocyanin from cyanobacteria *Spirulina* with the identical
244 position of Cys residues involved in the chromophore binding sites (Minic et al., 2016).

245 As previously described, the VIS absorption spectrum of purified R-PC exhibited two
246 characteristic peaks at 552 and 618 nm, originating from PEB and PCB chromophores, respectively
247 (**Fig. 1C**). The presence of two PCB chromophores per $\alpha\beta$ heterodimer of R-PC gives higher
248 absorption of the peak at 618 nm compared to the peak intensity at 552 nm, because only one PEB
249 chromophore is bound to β subunit. Excitation of PEB and PCB chromophores in R-PC at 488 and
250 590 nm gives two intensive emission peaks at 580 and 640 nm, respectively (**Fig. 2B**). In
251 accordance with the absorption spectra, the higher content of PCB compared to PEB chromophores
252 in R-PC (2 vs 1, respectively) gives stronger emission at 640 nm compared to 580 nm. PEB and
253 PCB chromophores are in the asymmetric environment within the R-PC structure, which induces
254 strong signals in the near-UV/visible CD spectrum of this protein. In the visible region, two peaks
255 appear that originate from PEB (540 nm) and PCB (634 nm) chromophores (**Fig. 2C**). We also
256 observed two peaks at 306 and 341 nm, arising from a higher excitation state of the PEB and PCB
257 chromophores, respectively (**Fig. 2C**) (Glazer et al., 1975). Interestingly, we observed the
258 differences in the shape and the position of peaks in the CD spectra of PEB and PCB chromophores
259 (10–20 nm) compared to absorption spectra, especially in terms that the relative intensity of PEB
260 band is stronger in CD compared to the absorption spectra. This finding indicates the strong
261 interactions of tetrapyrrole chromophores between different subunits within R-PC oligomers
262 (Glazer et al., 1975).

263 Far-UV CD spectra were recorded to characterise the secondary structure of purified R-PC.
264 As shown in **Fig. 2D**, the far-UV CD spectra of R-PC have negative ellipticity between 202 and
265 240 nm, with minima at 209 and 222 nm and also show a positive ellipticity at 192 nm, which
266 corresponds to the signals of α -helical proteins. Indeed, α -helix is the main secondary structure in
267 R-PC (53 %).

268 SAXS measurements were performed to determine the R-PC shape and oligomerisation
269 state in solution at pH 5.7. A well-defined plateau characterises the SAXS profile of R-PC at $Q <$
270 0.01 \AA^{-1} (**Fig. 2E**), confirming the absence of aggregates in the purified protein. We also observed

271 the large shoulder at 0.1 \AA^{-1} , arising from the central hole in the R-PC structure (**Fig. 2E**). Indeed,
272 we demonstrated by CRY SOL software that our experimental SAXS curve fits well with the
273 theoretical curve of the crystal structure of trimeric R-PC (**Figs. 2E and 2F**) from *Polysiphonia*
274 *urceolata* (Chang et al., 1996), which resembles the shape of the hollow cylinder. Indeed, SAXS
275 intensities are well fitted to a hollow cylinder analytical model (**Fig. S1**), enabling us to determine
276 the R-PC dimensions (**Fig. 2F**). We also performed the *ab initio* modelling by GASBOR software
277 (**Figs. 2G and 2H**) without making any assumption of R-PC structure and obtained a model that
278 closely resembles the R-PC crystal structure and hollow cylinder analytical model (**Fig. 2F**). Radii
279 of gyration value ($R_g = 39.3 \text{ \AA}$), obtained from Guinier plots, is in very good agreement with the
280 previously determined R_g values for trimeric phycocyanin (**Saxena, 1988**).

281 **3.3 R-phycocyanin thermal stability**

282 The thermal stability of the R-PC has been determined using CD and UV/VIS absorption
283 spectroscopy. CD spectroscopy has allowed us to follow the temperature variation of the ellipticity
284 at 222 nm, indicating the loss of the α -helical structure. The resulting melting curve, shown in **Fig.**
285 **3A**, exhibits one steep transition with a melting temperature of 52 °C (325 K) and an apparent
286 unfolding enthalpy of 385.5 kJ/mol.

287 In an alternative approach, we evaluate R-PC's thermal stability by studying the absorption
288 spectra changes. The results, shown in **Fig. 3B**, indicate that up to 40 °C, no significant differences
289 in the spectral characteristics of the protein are observed. However, as the temperature is increased
290 further (above 45 °C), a substantial decrease in absorption intensity is observed in the region
291 between 500 and 700 nm, followed at 55 °C and above, by the disappearance of the characteristic
292 bands associated with PCB and PEB. Additionally, at temperatures above 50 °C, a substantial
293 increase of absorbances at lower wavelengths (below 500 nm) indicates the aggregation of unfolded
294 proteins. These findings suggest that R-phycocyanin undergoes significant structural changes at
295 higher temperatures, leading to the loss of its characteristic spectral features. Reducing the
296 temperature to 20 °C did not lead to the recovery of distinct absorption bands, indicating an
297 irreversible protein denaturation (**Fig. 4A**).

298 **3.4 R-phycocyanin pH stability**

299 The pH stability of R-PC was assessed using UV/VIS absorption spectrometry. Spectral
300 analysis of R-PC in the pH range 4–8 revealed well-defined absorption maxima. A noteworthy
301 absorbance increase at 552 nm was observed for pH between 4 and 5. In contrast, the peak at 618
302 nm displayed similar absorbance intensity throughout the pH range 4–8 (**Fig. 3C**), indicating that
303 moderate acidic conditions do not induce conformational changes in the vicinity of the PCB
304 chromophore. Further increase or decrease of pH (9 and 3, respectively) triggered a significant

305 reduction in the absorption intensities of both PEB and PCB chromophores, but without pronounced
306 changes in the peak shapes. In extreme pH conditions (pH 2 and pH 10–12), both spectral bands of
307 R-PC were substantially disturbed, indicating protein unfolding (**Fig. 3C**). At pH 2, a significant red
308 shift in the spectrum occurs, suggesting the protonation of pyrrole rings within tetrapyrrole
309 chromophores (Dietzek et al., 2004), while, at alkaline conditions, a blue shift in the R-PC spectrum
310 indicates chromophores oxidation (Minic et al., 2018).

311 We examined the effects of pH on R-PC's secondary structures by far-UV CD spectroscopy
312 (**Fig. 3D**). No significant change in the far-UV CD spectra shape was observed for pH values
313 ranging from 3 to 9. Although at pH 3 and 9, the α -helical bands at 209 and 222 nm are still
314 pronounced, a significant decrease in ellipticity (absolute values) is observed at these pH values
315 (**Fig. 3D**), confirming the reduction in R-PC α -helical content. Furthermore, at both low acidic
316 conditions (pH 2) and high alkaline conditions (pH above 9), we note the absence of the α -helix
317 characteristic bands (**Fig. 3D**), suggesting a complete denaturation of the protein.

318 **3.5 Thermal stability of R-phycoyanin immobilised in calcium alginate beads**

319 We immobilised R-PC in calcium alginate beads, and obtained product was subjected to
320 thermal treatment at 60 °C for 10 minutes. Our results demonstrate that the colour of R-PC
321 immobilised in alginate beads does not change upon heating (**Fig. 4B**). On the other hand, as
322 described above, heating of R-PC induces the detrimental loss of the protein colour (**Fig. 4A**).
323 Therefore, immobilised R-PC exhibit greater thermal stability (**Fig. 4B**) compared to a protein that
324 was not immobilised (**Fig 4A**). Furthermore, the absorption spectrum of the R-PC extracted from
325 the calcium alginate matrix after thermal treatment showed no significant change, except a small
326 increase in absorbance at wavelengths below 500 nm, which indicates a slight protein aggregation
327 within alginate beads upon thermal treatment. Therefore, our findings provide strong evidence of
328 the substantial thermal stabilisation of protein in the calcium alginate matrix.

329 We also employed an ORAC assay to test the effects of immobilisation and heating on the
330 antioxidative capacity of R-PC. We did not observe any significant changes in the ORAC values
331 between R-PC and immobilised R-PC protein before thermal treatment (**Fig. 5**). Heating R-PC has
332 detrimental effects on the protein's antioxidant activity (**Fig. 5**). On the other hand, heating the
333 immobilised R-PC preserved the antioxidant action in comparison to the heating R-PC in solution.
334 However, the heat treatment decreased for ~20 % the antioxidant activity of immobilised R-PC
335 compared to unheated immobilisate (**Fig. 5**), indicating that the immobilisation approach could not
336 preserve the antioxidant capacity of R-PC upon heating completely.

337 **4. Discussion**

338 In this study, we purified and characterised the R-phycoyanin (R-PC) from dried Nori
339 flakes. Further, we tested the stability of purified, food-derived R-PC using the combination of
340 several experimental approaches. We observed the low-temperature stability of R-PC since a
341 moderate thermal treatment induces irreversible loss of protein colour and antioxidant capacity. On
342 the other hand, protein immobilisation inside alginate beads significantly improves R-PC thermal
343 stability.

344 In the cells of algae, R-phycoyanin and homologous PBPs, like R-PE and APC, form large
345 photosynthetic complexes called phycobilisomes (Saluri, Kaldmäe, & Tuvikene, 2019). Therefore,
346 an important obstacle in R-PC purification is removing the other PBPs, especially R-PE, the most
347 abundant protein in *Porphyra*. We tackled this challenge with a combination of HA and IEC
348 chromatography. HA chromatography removed the majority of R-PE, while IEC separated R-PC
349 from other proteins, including APC. The achieved R-PC purity ratio of 3.4 (A_{618}/A_{280}) was
350 significantly higher than the purity of R-PC ($A_{618}/A_{280} < 3.0$) isolated from red macroalgae
351 *Polysiphonia urceolata* (Wang et al., 2014). However, the commonly accepted criterion for the
352 analytical grade purity of C-phycoyanin (C-PC) is 4 (Nikolic et al., 2020). C-PC and R-PC have
353 three and two PCB chromophores, respectively. Hence, obtained R-PC purity between 3 and 4
354 could not be ascribed to impurities but to a lower number of PCB chromophores bounds compared
355 to C-PC, which induces a lower A_{618}/A_{280} ratio. Additionally, SDS-PAGE justified the R-PC
356 purity and its heteromeric structure with two bands arising from α and β subunits. The heteromeric
357 structure of purified R-PC has also been confirmed by mass spectrometry (MS). Moreover, we
358 utilised MS to determine the sequence of R-PC from Nori flakes, and high sequence homology with
359 phycoyanins from different classes of red algae and cyanobacteria has been detected.

360 Overall, we successfully optimised a protocol for producing high-purity vibrant purple-
361 coloured R-PC. Although, to the best of our knowledge, there is no analysis on the costs of R-PC
362 extraction and purification, the purity grade of phycoyanins, in general, strongly influences its
363 commercial value. For example, the prices of C-PC from *Spirulina* range from \$0.35 to \$135 per
364 gram in the food and cosmetics industries and up to \$4600 per gram for therapeutic and diagnostic
365 applications (Zittelli, Lauceri, Faraloni, Benavides, & Torzillo, 2023). However, the
366 commercialisation of R-PC and other phycoyanins faces challenges due to their chemical
367 instability and the high costs and time-consuming nature of extraction and purification methods
368 (Fernandes et al., 2023, Zittelli, Lauceri, Faraloni, Benavides, & Torzillo, 2023). Therefore, there is
369 a pressing need to scale R-PC production for industrial purposes to enable its sustainable production
370 and meet market demands, representing essential aspects for future studies.

371 Optical spectroscopy and SAXS curves show typical features of R-PC: well-defined
372 absorption, CD bands in the near-UV-VIS region and intensive fluorescence detectable at low nM

373 concentrations (Wang et al., 2014). The specific feature of R-PC is the presence of additional bands
374 in absorption, fluorescence and CD spectra, arising from the PEB chromophore, not presented in
375 the cyanobacterial phycocyanins (Yan et al., 2010). The different peak positions between near-UV-
376 VIS CD and absorption spectra of R-PC indicate strong exciton-type interactions between
377 tetrapyrrole chromophores, confirming the oligomeric structure of the protein (Glazer et al., 1975;
378 Li et al., 2020). Additionally, the stronger intensity of the PEB band compared to the PCB band in
379 CD spectra, which is opposed to the intensity ratio of these two bands in absorption spectra, could
380 be ascribed to the deviation of the PEB chromophore D ring from BC rings conjugate plane, as
381 observed in the crystal structure of R-PC from *Polysiphonia urceolata* (Jiang et al., 2001; Pescitelli
382 et al., 2003). In this regard, it was previously shown that dihedral angles within and between
383 chromophores significantly contribute to the intensities in CD spectra (Berova, Di Bari, &
384 Pescitelli, 2007). Such an effect could be especially relevant within oligomeric PBPs where
385 chromophores of different subunits could interact with each other. Indeed, our SAXS experimental
386 curve of R-PC in solution (pH 5.7), purified from Nori (*Porphyra*), fits well with the theoretical
387 curve of R-PC crystal structure from *Polysiphonia urceolata* in trimeric oligomer state, indicating
388 conservation of R-PC 3D structure between different classes of organisms. Indeed, this finding is in
389 accordance with the above-stated high sequence homology between *Porphyra* R-PC and
390 phycocyanin from other sources. We also elucidated the structure of R-PC by the *ab initio* approach
391 and found very good agreement with the model obtained by crystallography (Jiang et al., 2001).
392 However, we observed that R-PC from red algae and C-PC from cyanobacteria do not have the
393 same propensity towards oligomerisation. According to Saxena (Saxena, 1988), C-PC from the
394 cyanobacterium *Synechocystis* sp. has a hexameric structure at pH 5.7, whereas R-PC is in a
395 trimeric state here. Moreover, our SAXS results (data not shown) demonstrated that C-PC from
396 *Spirulina*, the most cultivated microalgae for food purposes, also has a hexameric form at pH 5.7.
397 This difference indicates that the unique presence of PEB chromophore in R-PC, but not in C-PC,
398 could provoke the subtle conformational changes that influence the oligomerisation trend.

399 The native structure of R-PC was preserved in dried Nori flakes, as confirmed by high α -
400 helical content (>50 %). Although (to the best of our knowledge) no previous studies are focusing
401 on the far-UV CD spectra of R-PC, our results are in good agreement with the obtained α -helical
402 content for C-PC from *Spirulina* and other cyanobacteria (Chen, Liu, MacColl, & Berns, 1983; Chi
403 et al., 2020; Edwards, Hauer, Stack, Eisele, & MacColl, 1997; Li, Gillilan, & Abbaspourrad, 2021).
404 However, R-PC has a substantially lower amount of α -helices in comparison to R-PE (>70 %)
405 (Simovic et al., 2022), suggesting the presence of γ subunit is the main contributor to the higher
406 percentage of α -helix in R-PE.

407 Temperature increase induced a steep reduction in α -helix ellipticity, showing a one-
408 transition unfolding of R-PC at $T_m \sim 52^\circ\text{C}$. Although (to the best of our knowledge) there is no data
409 about the thermal stability of R-PC, the studies conducted on phycocyanins from *Spirulina* and
410 other cyanobacteria (C-PC) obtained similar thermodynamic data with the melting temperature of
411 C-PC in the range of $50\text{--}60^\circ\text{C}$ (Chen et al., 1983; Faieta et al., 2020). On the other hand, our
412 previous CD study demonstrated higher thermal stability of the R-PE from *Porphyra* spp. due to a
413 more stable γ subunit, a unique feature of the R-PE oligomer (Simovic et al., 2022). Absorption
414 measurements revealed that the R-PC colour is unstable above 40°C . At the same time, the thermal
415 treatment at 55°C or above induces the complete and irreversible loss of R-PC-specific bands in the
416 absorption spectra. Although previous studies also confirmed that R-PE colour from Nori flakes
417 starts to diminish above 40°C (Simovic et al., 2022), the R-PE specific absorption peaks, especially
418 the band arising from γ subunit, could be observed up to 75°C . Therefore, the presence of γ subunit
419 is R-PE's key structural feature, making its structure substantially more stable than R-PC. In this
420 context, the previous studies demonstrated that phycocyanin from *Spirulina*, with the same subunit
421 composition as R-PC, also has unstable colour at higher temperatures with the onset of colour loss
422 above 40°C (Chaiklahan, Chirasuwan, & Bunnag, 2012; Faieta et al., 2020).

423 The colour of R-PC exhibits good stability in the pH range of 4–8, previously shown for
424 phycocyanin from *Spirulina* (Li et al., 2021; Yan et al., 2010). The secondary structures (α -helices)
425 of R-PC are also well preserved in the same pH range, confirming that tetrapyrrole chromophores
426 are good indicators of the conformational state of PBPs. R-PE from red macroalgae has a slightly
427 wider pH stability range (Munier et al., 2014), which is in accordance with the higher thermal
428 stability of R-PE. Shifting the pH to more acidic (pH 3) or alkaline conditions (pH 9) will induce
429 loss in α -helical structure and colour intensity of R-PC. However, the main spectral feature of R-PC
430 is still preserved, indicating the absence of complete protein denaturation. A similar observation
431 was made on C-PC from *Spirulina*, which was not entirely unfolded at pH 3 and 9 (Li et al., 2021;
432 Zhang, Cho, Dadmohammadi, Li, & Abbaspourrad, 2021). Moreover, in contrast to R-PC, the α -
433 helical structure in C-PC is mostly preserved at pH 9 (Li et al., 2021). Still, C-PC becomes more
434 flexible, and dissociation from hexamers/trimers to monomers was observed at pH 9 (Li et al.,
435 2021). Increasing the pH above 10 or acidification to pH 2 entirely unfolds R-PC, while tetrapyrrole
436 chromophores oxidation at high pH was also observed. Considering that moderate acidic conditions
437 (pH 3–4) do not have detrimental effects on R-PC colour and most fruit juices display pH in this
438 range (Reddy, Norris, Momeni, Waldo, & Ruby, 2016), R-PC have a promising potential for
439 applications as a colouring agent for soft beverages, even if its properties can be improved.

440 *Porphyra* spp. proteins have a high essential amino acid index, representing a sustainable
441 alternative to animal proteins (Rawiwan, Peng, Paramayuda, & Quek, 2022). Further, the added

442 value of *Porphyra* spp., as the source of alternative proteins, is the high abundance of PBPs
443 characterised by the vivid colours and strong bioactive properties (Venkatraman & Mehta, 2019).
444 These specific properties of PBPs substantially strengthen the consumer acceptance of *Porphyra*
445 spp. as an alternative source of proteins. However, preserving the PBPs' colour and bioactivities
446 during the food treatment is a significant challenge for enhancing their application in the food
447 industry. Alginate hydrogel beads represent a promising system for encapsulating, protecting, and
448 delivering food proteins (Zhang et al., 2016). Our study unambiguously confirmed that
449 encapsulation of R-PC into alginate beads completely preserves the protein colour upon thermal
450 treatment at 60 °C. Moreover, the antioxidative capacity of encapsulated R-PC is maintained largely
451 upon thermal treatment but not wholly, probably due to slight protein aggregation within alginate
452 beads, which could mask R-PC antioxidant activity. On the other hand, heating of R-PC that was
453 not immobilised induces detrimental loss of protein colour and antioxidative activity, followed by
454 aggregation of unfolded R-PC subunits. Indeed, previous studies demonstrated the ability of
455 alginate-encapsulated C-PC from *Spirulina* to preserve its antioxidant capacity upon thermal
456 treatment (Qiao et al., 2022). Therefore, encapsulating R-PC or other temperature-sensitive, food-
457 derived PBPs into alginate beads could be an excellent approach to preserve and modulate its
458 techno-functional properties, such as colour, aggregation and antioxidant capacity. Furthermore,
459 this approach could be a good strategy for controlling the release of bioactive PBPs in the
460 gastrointestinal tract (GIT) (Qiao et al., 2022). Alginate immobilisation in the presence of calcium
461 ions represents the non-covalent entrapment of the protein of choice by making a gel-like network
462 that surrounds the protein, thus entrapping it within (Zhang et al., 2016). Factors like pH and
463 mechanical treatment could significantly affect the system's stability in this context. By having R-
464 PC non-covalently immobilised, easier protein leakage is expected. This could be a double-edged
465 sword. While the risk of leakage limits conditions at which alginate: R-PC beads can be stored and
466 prevents rigorous mechanical treatments, this could be beneficial during their potential consumption
467 since during the mechanical eating process of chewing and different pH values of GIT could easily
468 release bioactive R-PC. On the other hand, the utilisation of composite gels which combine alginate
469 with other biopolymers, such as chitosan, gelatin, and various proteins, improves the physical and
470 mechanical properties of the gel, enhancing encapsulation efficiency, resulting in better protection
471 and controlled release of bioactive compounds (McClements, 2018; Ramdhan, Ching, Prakash, &
472 Bhandari, 2020).

473 **5. Conclusions**

474 We successfully purified and characterised the R-PC from commercially dried Nori flakes.
475 The purified protein exhibited optical and structural characteristics of typical R-PC in the

476 oligomeric (trimeric) α -helical structure. R-PC has moderate pH stability while being more stable in
477 acidic than in alkaline conditions. It also has relatively low-temperature stability, and an irreversible
478 colour change and antioxidant capacity loss were observed upon thermal treatment at 60 °C.
479 However, immobilisation of the R-PC inside calcium alginate beads significantly improved protein
480 thermal stability and preserved its antioxidant capacity and colour.

481 Our results suggest that R-PC has a promising potential for colouring soft beverages at
482 slightly-acidic conditions. However, further studies are needed to further improve its pH stability.
483 Encapsulating R-PC into solid matrices could be an excellent approach to preserve protein colour
484 and bioactivities upon thermal processing, strengthening its potential for food colouring and
485 positioning it as an ideal alternative protein.

486 **Credit authorship contribution statement**

487 **Luka Velickovic:** Investigation, Conceptualisation, Formal analysis, Writing – original draft. **Ana**
488 **Simovic:** Investigation, Conceptualisation, Formal analysis, Writing – review and editing. **Nikola**
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490 **Aurélien Thureau:** Investigation, Formal analysis. **Milica Obradovic:** Investigation. **Tamara**
491 **Vasovic** – Investigation, Formal analysis. **Georgios Sotiroudis:** Investigation, Writing – review
492 and editing. **Maria Zoumpanioti:** Writing – review and editing. **Annie Brûlet:** Investigation,
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495 **Nikolic:** Conceptualisation, Funding acquisition, Supervision, Writing – review and editing.
496 **Simeon Minic:** Investigation, Conceptualisation, Formal analysis, Funding acquisition,
497 Supervision, Writing – review and editing.

498 **Declaration of Competing Interest**

499 The authors declare that they have no known competing financial interests or personal relationships
500 that could have appeared to influence the work reported in this paper.

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507 **Figure legends**

508 **Figure 1.** The chromatograms obtained after (A) hydroxyapatite and (B) ion-exchange
509 chromatography of R-PC; (C) Absorption spectra of R-PC after each purification step; (D) SDS-
510 PAGE electrophoresis under UV light (left) and after CBB staining (right) - lane 1: Purified RPC,
511 lane 2: hydroxyapatite eluate, lane 3: protein extract, M: molecular size markers.

512 **Figure 2.** (A) Amino acid sequences of R-PC α and β subunits determined by mass spectrometry. P
513 denotes the attachment site for PCB (Cys84 and Cys82 in α and β subunits, respectively) and PEB
514 chromophores (Cys153 in β subunit); (B) Fluorescence emission spectra of purified R-phycocyanin
515 (16 nM) after excitation at 488 nm (black line) and 590 nm (red line) at pH 7.0; (C) Near-
516 UV/Visible and (D) far-UV CD spectra of R-PC (pH 7.0); (E) Comparison of the experimental
517 SAXS curve of R-PC at pH 5.7 (dotted black line) with the theoretical curve (full red line)
518 calculated from R-PC crystal structure (PDB:1F99) by CRY SOL software (ATSAS); (F) Ribbon
519 model of the crystal structure of R-PC trimer superimposed with the hollow cylinder analytical
520 model; Dimensions are obtained by the best fit to this model (Table S3). Obtained χ^2 value was 2.8;
521 (G) Comparison of the same experimental SAXS curve with the curve obtained from *ab initio*
522 modelling (full red line) using GASBOR software (ATSAS). The obtained χ^2 value was 1.5; (H)
523 Ab initio envelope of the R-PC oligomer corresponding to the GASBOR model. T and S denote top
524 and side views, respectively.

525 **Figure 3.** (A) 20 $\mu\text{g/mL}$ (182 nM) R-PC thermal unfolding curve with the corresponding fit (full
526 red line), obtained by measuring ellipticity at 222 nm and pH 7.0; (B) The effects of temperature on
527 visible absorption spectra of 100 $\mu\text{g/mL}$ (0.9 μM) R-PC at pH 7.0 (optical pathlength 1 cm); (C)
528 The effects of pH on visible absorption spectra of 100 $\mu\text{g/mL}$ R-PC at 25 $^\circ\text{C}$; (D) The effects of pH
529 on far-UV CD spectra of 100 $\mu\text{g/mL}$ R-PC at 25 $^\circ\text{C}$.

530 **Figure 4.** Absorption spectra of (A) R-PC without immobilisation and (B) R-PC extracted from
531 calcium alginate beads before and after heating at 60 $^\circ\text{C}$ for 10 min; Insets show pictures of (A) R-
532 PC without immobilisation and (B) immobilised R-PC before (BH) and after (AH) heating.

533 **Figure 5.** Histogram comparing antioxidative activity (ORAC assay) of R-PC without
534 immobilisation (free R-PC) and R-PC immobilised in calcium alginate beads before and after
535 heating at 60 $^\circ\text{C}$. The data marked by different letters are significantly different ($p < 0.05$).

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