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### Title

Rapid adaptive evolution of colour vision in the threespine stickleback radiation

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1 Title: Rapid adaptive evolution of colour vision in the threespine stickleback radiation.

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31 **Abstract**

32 Vision is a sensory modality of fundamental importance for many animals,  
33 aiding in foraging, detection of predators, and mate choice. Adaptation to local  
34 ambient light conditions is thought to be commonplace, and a match between  
35 spectral sensitivity and light spectrum is predicted. We use opsin gene expression to  
36 test for local adaptation and matching of spectral sensitivity in multiple independent  
37 lake populations of threespine stickleback populations derived since the last ice age  
38 from an ancestral marine form. We show that sensitivity across the visual spectrum  
39 is shifted repeatedly towards longer wavelengths in freshwater compared with the  
40 ancestral marine form. Laboratory rearing suggests this shift is largely genetically  
41 based. Using a new metric, we found that the magnitude of shift in spectral  
42 sensitivity in each population corresponds strongly to the transition in the availability  
43 of different wavelengths of light between the marine and lake environment. We also  
44 found evidence of local adaptation by sympatric benthic and limnetic ecotypes to  
45 different light environments within lakes. Our findings indicate rapid parallel evolution  
46 of the visual system to altered light conditions. The changes have not, however,  
47 yielded a close matching of spectrum-wide sensitivity to wavelength availability, for  
48 reasons we discuss.

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52 Keywords: Visual Ecology, Local Adaptation, Evolution, Opsin, Gene Expression,  
53 *Gasterosteus aculeatus*.

54 **Background**

55           Sensory systems are often thought to be under strong natural selection [1], and  
56 are predicted to evolve to better correspond to signals in the local environment [2]. For  
57 example, sensitivity of the visual system to different wavelengths of light is expected to  
58 evolve to match roughly the availability of wavelengths [3,4], increasing ability to catch  
59 photons and detect contrast between objects and background [5,6,7]. However, few  
60 studies have tested the adaptive significance of spectral sensitivity across the whole  
61 visual spectrum. The degree of matching between spectral sensitivity of organisms and  
62 their light environment across the spectrum has not been quantified.

63           Aquatic organisms provide excellent opportunities to test for local adaptation and  
64 quantify matching [8]. This is because differential attenuation of wavelengths of light  
65 with water depth and by suspended particulates result in dramatic and predictable  
66 changes in local light spectra [9]. For example, the transition from marine to fresh  
67 waters is usually accompanied by a large reduction in the availability of ultraviolet (UV)  
68 wavelengths, largely because of an increase in the amount of dissolved organics [9,10].

69           We used threespine stickleback (*Gasterosteus aculeatus*), which inhabit both  
70 marine and freshwater habitats, to investigate predicted evolutionary changes in visual  
71 adaptations of populations to the different ambient light environments. Marine  
72 stickleback invaded and adapted to numerous lakes and streams at the end of the last  
73 ice age (~12,000 years ago) [11]. First, we tested for parallel evolution of opsin gene  
74 expression and spectral sensitivity over the visual light spectrum among these derived  
75 freshwater populations, which would represent strong evidence of natural selection [12].

76           Second, utilizing the extant marine form as a proxy for the ancestral state, we  
77 evaluated the extent to which shifts in the spectral sensitivity of freshwater populations

78 are correlated with shifts in the ambient light environment, and whether the outcome  
79 improves the match to local ambient light spectra. Finally, we tested for parallel  
80 divergence of spectral sensitivity of multiple pairs of sympatric limnetic and benthic  
81 stickleback ecotypes (or “species pairs”) to finer scale heterogeneity in the light  
82 environment within lakes. In each of the three species pairs analyzed here, the benthic  
83 stickleback forage in the vegetated littoral regions of the lake and deeper sediments,  
84 whereas limnetics are pelagic, found in the open water and near rocky cliffs [13]. The  
85 benthic environment contains relatively more long wavelengths than the open water  
86 [14].

87         We focused on expression of opsin genes, which encode the light sensitive G-  
88 protein coupled receptors that are expressed in retinal rod and cone cells. Opsins  
89 conjugate to vitamin A derived chromophores and play an important role in colour vision  
90 by mediating the conversion of photons into electrochemical signals, which initiate a  
91 neuronal response that is perceived by the brain [15]. The clear and well-characterized  
92 link between opsin genotype (coding sequence) and spectral phenotype (wavelength of  
93 maximal absorption,  $\lambda_{max}$ ) make opsins particularly useful for studying sensory  
94 adaptation [16]. Opsin mediated shifts in spectral sensitivity can be achieved by  
95 changes in opsin protein coding sequence (e.g. [17]) and by changes in levels of gene  
96 expression (e.g. [18]). We studied gene expression because analysis of whole genomes  
97 of marine and freshwater stickleback has not found consistent differences in opsin gene  
98 coding sequence between marine and freshwater populations [19]. Compared to other  
99 fish, stickleback have relatively few (four) opsins, with a single functional opsin gene in  
100 each of the four cone opsin subfamilies: short-wavelength sensitive 1 (*SWS1*), short-

101 wavelength sensitive 2 (*SWS2*), middle-wavelength sensitive (*RH2*), and long-  
102 wavelength sensitive (*LWS*) [20]. We measured expression levels of each of the four  
103 unique cone opsin genes in 11 stickleback populations. We also measured expression  
104 in fish from two populations raised in a common laboratory environment to test the  
105 extent to which it is genetically determined.

106         We used opsin gene expression levels to estimate spectrum-wide spectral  
107 sensitivity to evaluate two general expectations. First, the advantages of photon capture  
108 and contrast should result in spectral sensitivity evolving roughly to correspond with  
109 wavelength availability [3,4]. We measure this correspondence (“matching”) with the  
110 correlation across wavelengths between spectral sensitivity and two measures of light  
111 availability: irradiance (photons of each wavelength available at a specific water depth)  
112 and transmission (indicating the absorption of specific wavelengths by water). Large  
113 discrepancies between spectral sensitivity and light availability in specific regions of the  
114 visual spectrum might suggest specialized visual functions. Second, changes in  
115 wavelength availability from marine to fresh water should lead to similar shifts in  
116 spectral sensitivity (“local adaptation”). For example, as some wavelengths become  
117 scarce in the new environment and others common, relative to the ancestral  
118 environment, we expect spectral sensitivity to shift to correspond [2]. Throughout, we  
119 use the whole light spectrum to study association, rather than studying associations  
120 between summary measures such as the median. We introduce a new metric to  
121 quantify the correlation between shift in spectral sensitivity and the transition between  
122 light environments.

123 Shifts in spectral sensitivity can additionally be achieved by differential use of  
124 vitamin A derived chromophores [21, 22]. Conjugation of an opsin to an A<sub>1</sub>  
125 chromophore (11-*cis* retinal) leads to a shorter wavelength of maximal absorption ( $\lambda_{max}$ )  
126 than conjugation to an A<sub>2</sub> chromophore (3-dehydro 11-*cis* retinal) [21]. Switches in  
127 chromophore use have been shown to occur in fishes over ontogeny [23] and between  
128 habitats via phenotypic plasticity [22]. Fish in the ocean generally use A<sub>1</sub> chromophores,  
129 while freshwater fish have a mixture of A<sub>1</sub> and A<sub>2</sub> chromophores (varying from  
130 completely A<sub>1</sub> to completely A<sub>2</sub>) [22]. Complete use of A<sub>2</sub> is generally found in lakes  
131 whose waters are strongly stained with tannins [e.g. 24], and such lakes are not  
132 included in our study. To account for possible variation in chromophore use, we model  
133 the effects of changes in chromophore and describe how this affects our measures of  
134 local adaptation and spectral matching of opsin expression in stickleback.

135

## 136 **Materials and Methods**

### 137 **Sampling**

138 Six gravid females were collected from each of 11 populations inhabiting different  
139 breeding environments in the Strait of Georgia region of British Columbia, Canada.  
140 Collections were made under the Species At Risk Act collection permit 236 and British  
141 Columbia Fish Collection permit NA-SU12-76311. The samples came from two marine  
142 locations, three lakes containing just a single species of stickleback, and three lakes  
143 containing stickleback species pairs (see Supp. Mat. Section 1 and Table S1 for site  
144 details). Fish were euthanized at the collection site and eye tissue was immediately

145 preserved in RNAlater® (Qiagen, Netherlands) and then kept at –20 °C for up to a  
146 month before RNA was extracted.

### 147 **Opsin Expression and Spectral Sensitivity**

148 The expression of each of the stickleback’s four unique cone opsin genes  
149 (*SWS1*, *SWS2A*, *RH2-1*, and *LWS* [20]) was measured using a standard reverse-  
150 transcriptase quantitative polymerase chain reaction protocol (details in the Supp. Mat.  
151 Section 2). We normalized the absolute number of transcripts for each gene from each  
152 individual by dividing the expression of a given opsin by the sum of the expression of all  
153 four opsins to get relative opsin expression. We also measured gene expression of a  
154 reference gene, *Beta actin*, and calculated the expression of each opsin gene relative to  
155 it.

156 All statistical analyses in the paper were conducted in R 3.0.2 [25]. To test for  
157 differences in mean expression of each opsin gene between marine and freshwater  
158 populations, we used a linear mixed-effects model (using the nlme package, [26]) with  
159 water type (marine or fresh) as fixed effect and location as a random effect. For this  
160 comparison, individuals from the benthic and limnetic species in a given location were  
161 combined and treated as a single population. Results were the same when only the  
162 benthics, or only the limnetics, were used instead. In separate analyses we tested for  
163 differences in gene expression between the sympatric benthic and limnetic species in  
164 three lakes, with lake as a random effect and species as a fixed effect in the model. We  
165 treat lake populations as independent replicates that require no phylogenetic correction.  
166 This is justified by the geological origins of lakes, which are in separate drainages and  
167 were accessible via the sea for a limited period of time. Previous studies show that



168 phylogenies of freshwater stickleback populations in British Columbia based on  
169 putatively neutral markers are well approximated by a star phylogeny (e.g. [27]).

170 We bred three families of one marine population (Oyster Lagoon) and three  
171 families of one benthic population (Priest Lake) by *in vitro* fertilization and reared them  
172 under laboratory conditions in stand-alone 100 L tanks with fluorescent lights. Animals  
173 were treated in accordance with University of British Columbia Animal Care protocols  
174 (Animal Care Permit A11-0402). A gravid adult female from each family was euthanized  
175 and her opsin expression was quantified as described above. We used linear models to  
176 test differences between lab-reared marine and freshwater fish and between lab- and  
177 wild-reared fish from the same populations.

178 Upon finding differences in mean opsin expression between marine and  
179 freshwater stickleback, and between sympatric benthic and limnetic stickleback, we  
180 estimated how they translated into differences in spectral sensitivity. We calculated a  
181 spectral sensitivity curve  $S_i$  (350 – 700 nm) for each individual  $i$  based on its relative  
182 expression of the four opsin genes, and using the absorbance templates from  
183 Govardovskii *et al.* [28] and estimates by Flamarique *et al.* [24] of the wavelength of  
184 maximum absorbance ( $\lambda_{\max}$ ) of each opsin gene (details in Section 3 of the Supp. Mat.).  
185 This model assumes that opsin expression contributes additively to spectral sensitivity;  
186 at this point in time it is a necessary simplification as we still lack empirical informed  
187 models that describe and generalize any potential inhibitory interactions among opsins  
188 during signal integration and interpretation.

189 Chromophore ( $A_1$  and  $A_2$ ) ratios in the surveyed freshwater populations are not  
190 known. Based on empirical observations [24] we assumed that marine stickleback used

191 100%  $A_1$  in the ocean. We estimated spectral sensitivity of stickleback in fresh water  
192 using three different chromophore ratios representing the extremes: 100%  $A_1$ ; 50%  $A_1$   
193 and 50%  $A_2$ ; and 100%  $A_2$ . We assumed that benthic and limnetic stickleback have the  
194 same  $A_1 : A_2$  ratio.

195

## 196 **Association between Spectral Sensitivity and Ambient Light**

197 We measured the spectral conditions of each location, with the exception of  
198 Cranby Lake and Little Quarry Lake. We used two measures to quantify the ambient  
199 light environment: irradiance and transmission. Irradiance measures the abundance of  
200 photons at each wavelength in the environment at a given point in time. Irradiance  
201 measurements of side-welling light ( $I_s$ ) were taken at 10 cm, 20 cm, 50 cm, 100 cm and  
202 200 cm depth at 10 or more sites within each sampling location using a cosine corrector  
203 attached to a spectrophotometer (Ocean Optics, USA). In subsequent analyses we  
204 used the irradiance at 50 cm. A limitation of using irradiance to quantify available light is  
205 that it varies with depth and with the weather and the angle of the sun. Transmission is  
206 the relative rate of loss of photons of a given wavelength per unit distance traveled  
207 through water. Transmission is a property of the body of water and may be less variable  
208 than irradiance, at least on short time scales. Transmission was measured as the light  
209 extinction coefficient with depth ( $K_s$ ) (method for calculation outlined in Supp. Mat.  
210 Section 5).

211 To test for local adaptation, we developed a statistic to quantify the association  
212 between the shift in spectral sensitivity and the transition in light environment, from  
213 marine to fresh water, across all wavelengths for each lake population. First, we chose

214 a marine population (Oyster Lagoon) to represent the ancestral phenotype and breeding  
215 environment. Next, we constructed transmission ( $K_s$ ) and irradiance ( $I_s$ ) curves by  
216 calculating at each wavelength ( $\lambda$ ) the median from all samples within a location. At  
217 each wavelength we then subtracted the median value of the reference marine location  
218 from the median value in each freshwater location. This yielded change in transmission  
219 ( $\Delta K_s$ ) and change in irradiance ( $\Delta I_s$ ) values at every wavelength ( $\lambda$ ) at each freshwater  
220 location. A positive value of  $\Delta I_s$  at a given wavelength indicates that there are more  
221 photons of that wavelength ( $\lambda$ ) present at the freshwater location relative to the marine  
222 environment. A positive value of  $\Delta K_s$  at a given wavelength ( $\lambda$ ) indicates greater light  
223 transmission (fewer photons lost as light travels through water) at the freshwater  
224 location than at the reference marine location.

225 Change in spectral sensitivity  $\Delta S$  was calculated similarly, as follows. We  
226 calculated the median sensitivity at each wavelength ( $\lambda$ ) of the sample of individuals  
227 from the reference marine population. Change in sensitivity was calculated for each  
228 freshwater individual as the difference between its sensitivity curve and the median  
229 marine curve. Finally, for each freshwater individual, we calculated the correlation  
230 coefficient ( $r$ ) of the change in sensitivity ( $\Delta S$ ) against the change in light environment  
231 ( $\Delta K_s$  or  $\Delta I_s$ ), with each wavelength yielding a data point for each freshwater individual. A  
232 positive  $r$  indicates that regions of the spectrum with increased irradiance (or  
233 transmission) are correlated with increased spectral sensitivity, and regions of the  
234 spectrum with a decrease in irradiance (or transmission) are correlated with decreased  
235 spectral sensitivity. We used a mixed-effects model (with population as a random effect)  
236 to test whether mean correlation coefficients ( $r$ ) differed significantly from zero.

237 We carried out a similar analysis of local adaptation of spectral sensitivity  
238 between the sympatric species in relation to differences in their local light environments.  
239 For each lake, we used the limnetic population and the pelagic environment as the  
240 reference. Other calculations were the same as described above for the marine and  
241 freshwater comparison (see Supp. Mat. Sections 5 and 6, Tables S3 and S4, and Supp.  
242 Figs. 1 and 2, for further details and justification of our reference populations).

243 To quantify the degree to which populations are matched to their native light  
244 environments we estimated the correlation, wavelength by wavelength, between each  
245 population's mean spectral sensitivity and the transmission and irradiance measured in  
246 its local environment. The significance of the mean correlation was tested separately for  
247 marine and freshwater populations using linear models.

248 Because analyses of local adaptation and matching involved a suite of tests that  
249 incorporated different measures of light environment and three chromophore scenarios,  
250 we adjusted the p-values for multiple testing in each table of results using the "BH" false  
251 discovery rate method [29] and the *p.adjust* function in R (Tables S3, S4, and S5). Raw  
252 p-values are reported in the main paper and adjusted p-values are reported in the  
253 statistics tables in the supplement. In all cases significant p-values remained significant  
254 after the correction for multiple testing.

255

## 256 **Results**

### 257 **Opsin Expression and Spectral Sensitivity**

258 Freshwater stickleback populations had significantly lower expression of the  
259 *SWS1* (UV) opsin gene than the marine populations (difference =  $-0.20 \pm 0.02$  SE,  $F_{1,6}$

260 = 145.2,  $p < 0.001$ ) and higher expression of the *RH2* (green) opsin gene (difference =  
261  $0.21 \pm 0.06$  SE,  $F_{1,6} = 18.1$ ,  $p = 0.005$ ). We did not detect a significant difference in the  
262 other two opsin genes, *LWS* (red) (difference =  $0.02 \pm 0.04$  SE,  $F_{1,6} = 0.2$ ,  $p = 0.68$ ) and  
263 *SWS2* (blue) (difference =  $-0.009 \pm 0.008$  SE,  $F_{1,6} = 1.2$ ,  $p = 0.31$ ) (Figure 1).  
264 Differences in *SWS1* and *RH2* remained significant if expression was calculated relative  
265 to the reference gene *Beta actin* (*SWS1* difference =  $2.1 \pm 0.3$  SE,  $F_{1,6} = 49.2$ ,  $p <$   
266  $0.001$ ; *RH2* difference =  $2.97 \pm 0.9$  SE,  $F_{1,6} = 10.7$ ,  $p = 0.017$ ). Thus we proceeded  
267 using cone opsin proportion as our metric of gene expression when modeling spectral  
268 sensitivity, as this has been shown to be best for making inferences about overall colour  
269 vision capacities [30].

270         These differences in overall expression translated into large differences in  
271 estimated spectral sensitivity in two portions of the spectrum (Figure 2). Freshwater fish  
272 had reduced sensitivity in the 350-375 nm (UV and violet) region of the spectrum, and  
273 they had greater sensitivity in the 450-600 nm (blue and green) region relative to both  
274 marine populations.

275         Within lakes we found that the limnetic stickleback populations had significantly  
276 greater *RH2* (green) expression than the benthics (difference =  $0.05 \pm 0.02$  SE,  $F_{1,31} =$   
277  $7$ ,  $p = 0.01$ ), and benthics had greater *LWS* (red) expression ( $0.04 \pm 0.02$  SE,  $F_{1,31} =$   
278  $4.3$ ,  $p = 0.05$ ). However, the magnitudes of the differences were small (Figure 3). The  
279 expression of *SWS1* and *SWS2* opsins did not differ significantly ( $p > 0.29$ ) between the  
280 two species (Figure 3). The difference in *RH2* expression between the species was still  
281 significant when expression was calculated relative to *Beta actin* gene expression  
282 (difference =  $1.3 \pm 0.6$  SE,  $F_{1,31} = 4.4$ ,  $p = 0.04$ ), but the difference in *LWS* was not

283 (difference =  $0.3 \pm 0.84$  SE,  $F_{1,31} = 0.12$ ,  $p = 0.70$ ). These differences in expression  
284 translate to reduced sensitivity in the 525-575 nm (green) region of the spectrum and  
285 increased sensitivity in the portion of the spectrum above 600 nm (red) in benthics  
286 compared to limnetics (Supp. Figure 3).

287

## 288 **Laboratory Rearing**

289 In the lab, Oyster Lagoon (marine) and Priest benthic fish (freshwater) had  
290 similar expression differences as in the wild (Figure 4). *SWS1* gene expression  
291 remained different between the marine and freshwater populations in the lab (difference  
292  $0.11 \pm 0.02$  SE,  $df = 1,4$ ,  $F = 27.1$ ,  $p = 0.01$ ) as did *RH2* (difference  $0.18 \pm 0.03$  SE,  $df =$   
293  $1,4$ ,  $F = 40.1$ ,  $p = 0.003$ ). The difference in *SWS1* was, however, greater in the wild  
294 samples, as indicated by an interaction between rearing condition (wild or lab) and  
295 population of origin (effect size =  $0.096 \pm 0.039$  SE,  $t_{1,4} = 2.486$ ,  $p = 0.03$ ). No other  
296 interactions were significant (all  $p > 0.17$ ). Finally, we also detected a small difference in  
297 *LWS* expression between the two populations in the lab only (Figure 4;  $0.06 \pm 0.02$  SE,  
298  $F_{1,4} = 11.5$ ,  $p = 0.03$ ). Additional tests examining changes in the gene expression of lab-  
299 reared fish from each population compared to their wild counterparts are outlined in the  
300 Supp. Mat. Section 4.

301

## 302 **Association between Shifts in Spectral Sensitivity and Ambient Light**

303 The shift in spectral sensitivity from marine to freshwater environments was  
304 positively correlated with the change in ambient light spectrum, when sensitivity was  
305 estimated assuming that both populations used only the A<sub>1</sub> chromophore. On average,

306 the correlation measured using transmission (mean  $r = 0.39$ ,  $\pm 0.12$  SE,  $t_{1,31} = 3.3$ ,  $p =$   
307  $0.002$ ; Figure 5A) was of similar magnitude when using irradiance (mean  $r = 0.32$ ,  $\pm$   
308  $0.06$  SE,  $t_{1,31} = 4.95$ ,  $p < 0.0001$ ; Figure 5B). These correlations arose primarily from  
309 shifts in the short- (UV-blue) and middle-wavelength (green) regions (Supp. Figure 4).  
310 Decreased transmission of UV (350-400 nm) and violet (380-450 nm) in the freshwater  
311 environment (indicated by values below the dashed line in Supp. Figure 4) correspond  
312 with decreased sensitivity to these wavelengths in freshwater populations. Increased  
313 transmission of blue (450-495 nm) and green (495-570 nm) wavelengths in fresh water  
314 is correlated with increased sensitivity to these wavelengths. Freshwater populations  
315 varied considerably in the strength of the correlation (Figure 5).

316         These results isolate the effects of shifts in spectral sensitivity caused by  
317 changes in opsin gene expression in freshwater, when controlling for chromophore. We  
318 also measured the effects of these expression changes if combined with a hypothetical  
319 increase in the use of the A<sub>2</sub> chromophore in these freshwater populations. The  
320 correlation between shifts in spectral sensitivity and transmission weakens slightly when  
321 a 50:50 mix of A<sub>1</sub> and A<sub>2</sub> chromophores is projected (mean  $r = 0.22$ ,  $\pm 0.12$  SE,  $t_{1,31} =$   
322  $1.85$ ,  $p = 0.07$ ). When 100% A<sub>2</sub> chromophore is used, the correlation between shifts in  
323 sensitivity and transmission weaken further (mean  $r = 0.14$ ,  $\pm 0.09$  SE,  $t_{1,31} = 1.53$ ,  $p =$   
324  $0.14$ ) and the correlation between shifts in sensitivity and irradiance becomes negative  
325 (mean  $r = -0.48$ ,  $\pm 0.05$  SE,  $t_{1,31} = -9.3$ ,  $p = <0.0001$ ) (see Table S3 for details, including  
326 adjusted p-values).

327         Within species pair lakes, there was a moderate, although not quite significant,  
328 correlation between divergence in spectral sensitivity and the difference in transmission

329 (modeled using the A<sub>1</sub> chromophore; Figure 5C; mean  $r = 0.27 \pm 0.13$  SE,  $t_{1,10} = 1.97$ ,  $p$   
330  $= 0.077$ ). This correlation was not significant for the difference in irradiance (Figure 5D;  
331 mean  $r = 0.18 \pm 0.18$  SE,  $t_{1,10} = 1.00$ ,  $p = 0.339$ ). The results were similar when other  
332 chromophore ratios were used to estimate spectral sensitivity, assuming that ratios  
333 were the same in both sympatric forms (see Table S4 for details, including adjusted  $p$ -  
334 values).

335

### 336 **Match of Spectral Sensitivity to Ambient Light**

337         Despite strong correlations between shifts in spectral sensitivity and changes in  
338 the distribution of available wavelengths, spectral sensitivity is not closely matched to  
339 wavelength availability in either marine or freshwater environments. The mean  
340 correlation between spectral sensitivity of freshwater fish and ambient light in lakes,  
341 while statistically significant, was small ( $0.07 \pm 0.03$  for transmission and  $0.12 \pm 0.02$  for  
342 irradiance). This low level of matching has arisen multiple times in parallel in lake  
343 stickleback, which suggests that natural selection favors it. Substituting the  
344 chromophore did little to alter the mean correlation for transmission (although it became  
345 statistically insignificant) and slightly changed the strength for irradiance (See Supp.  
346 Mat. Section 7 and Table S5 for details, including adjusted  $p$ -values). In the marine  
347 environment the mean correlations between marine spectral sensitivity and  
348 transmission or irradiance are negative ( $r = -0.66 \pm 0.16$  SE and  $r = -0.11 \pm 0.07$  SE,  
349 respectively). The main cause of the strong negative correlation in marine waters is the  
350 excessive UV sensitivity compared with UV light availability. Nevertheless, UV  
351 expression declines in fresh water, where these wavelengths are even more scarce,



352 contributing to the observed correlation between shifts in sensitivity and the change in  
353 wavelength distribution.

354

## 355 **Discussion**

356 Our findings indicate that there has been rapid parallel evolution of opsin gene  
357 expression and spectral sensitivity across the light spectrum in freshwater stickleback  
358 populations. All surveyed freshwater populations have their spectral sensitivity shifted  
359 towards blue and green wavelengths, and away from ultraviolet and violet, relative to  
360 the marine populations. This has been accomplished entirely by shifts in opsin gene  
361 expression rather than protein sequence changes. We provide evidence that this  
362 difference has a genetic basis, as the main differences in expression were maintained in  
363 two lab-reared populations. Our analyses also reveal a strong association between  
364 shifts in spectral sensitivity and changes in light transmission from marine to fresh water  
365 environments, suggesting that these shifts are in an adaptive direction. On a smaller  
366 scale, we also find support for parallel adaptive divergence of gene expression and  
367 spectral sensitivity within lakes, between sympatric limnetic and benthic species. The  
368 evolution of the visual system in stickleback has been rapid, as these freshwater  
369 populations have evolved within the last 12,000 after the last glacial maxima [11].

370 The degree of phenotypic parallelism in opsin expression and spectral sensitivity  
371 that we describe is unprecedented over such a short time span. Nine independently  
372 derived populations exhibit the same direction of shift in opsin expression following the  
373 colonization of freshwater. In East African cichlids, parallel evolutionary divergence of  
374 opsin expression has been detected between species within two of the three major lake

375 cichlid radiations [31], but these radiations are much older than the freshwater  
376 stickleback populations studied here. Our findings are in line with previous work in  
377 stickleback, which has found extensive parallel evolution of morphological traits and  
378 patterns of genomic divergence among freshwater populations [19,32,33]. Some but not  
379 all of this morphological parallelism involves changes at the same underlying genes,  
380 which frequently represents adaptation from a common ancestral pool of standing  
381 genetic variation [32]. Possibly, the parallelism we observe in spectral sensitivity also  
382 represents adaptation from a common pool of standing genetic variation, which would  
383 help to explain the speed of evolution in this trait in stickleback. Further genetics work is  
384 required to test this idea.

385         The result from our lab rearing experiment suggests a substantial genetic  
386 component to the population differences in opsin expression. This contrasts with many  
387 other systems in which differential opsin gene expression and/or spectral sensitivity is  
388 largely phenotypically plastic [e.g. 34]. For example, wild Bluefin Killifish (*Lucania*  
389 *goodie*) living in clear springs and tannin stained waters exhibit large differences in their  
390 opsin gene expression [34]; however, light treatment and rearing experiments in the lab  
391 have shown that most of these differences are due to environmental effects [34].

392         Smaller but detectable differences in opsin expression and sensitivity between  
393 limnetic and benthic stickleback inhabiting the same lake were repeated in multiple  
394 lakes, suggesting a role for natural selection in divergence of visual systems on a small,  
395 within-lake scale. Benthics had slightly higher estimated sensitivity to red wavelengths  
396 than did limnetics, in accord with a more red-shifted local light environment. Previous  
397 work using optomotor behavioural response assays indicated that limnetic stickleback in

398 Enos Lake have higher red wavelength sensitivity than the benthic population from the  
399 same lake, and similar red wavelength sensitivity to the benthic in Paxton Lake [14]. In  
400 contrast, we found higher expression of long wavelength opsins in benthics compared  
401 to limnetics. Future work is required to determine how these differences in opsin  
402 expression affect foraging and mate choice in stickleback, as has been suggested in  
403 Lake Victoria cichlids [35].

404         Early work in the field of visual ecology focused on the hypothesis that spectral  
405 sensitivity should evolve to maximize an individual's photon catch [5,6,7]. Tests of this  
406 hypothesis have examined the relationship between the  $\lambda_{\max}$  of visual pigments (opsins)  
407 and the wavelengths most prevalent in ambient environment and have often found a  
408 strong relationship (e.g. [36,37]). However, detection of contrast and colour  
409 discrimination also likely shapes the evolution of spectral sensitivity. With multiple  
410 functions, it may be difficult to predict *a priori* the evolved degree of spectrum-wide  
411 matching of spectral sensitivity to the available light spectrum. We did not find a close  
412 match in freshwater populations, and indeed, the correlation was negative in marine  
413 populations. The low match in marines is driven by their high estimated sensitivity to  
414 short wavelengths such as UV, despite the relative rarity of these light wavelengths in  
415 the marine environment compared to mid-wavelengths. The low degree of matching  
416 suggests that increasing photon capture alone is unlikely to explain the evolution of  
417 spectral sensitivity. Predicting a shift in sensitivity with change in light spectrum may be  
418 more straightforward: reduced investment in capturing specific wavelengths that are  
419 increasingly rare is expected. For example in the deep sea, long-wavelength light is  
420 rare, and some deep-sea fish have lost long-wave sensitive opsins and shifted their

421 sensitivity towards shorter wavelengths [37]. Similarly, we found that freshwater  
422 stickleback have reduced expression of short wavelengths, which are even scarcer in  
423 freshwater than in the sea. Nevertheless, freshwater fish retain relatively high sensitivity  
424 to UV light compared to background irradiance.

425         One possible explanation for the low match between sensitivity and ambient  
426 wavelengths is that high expression of pigments whose sensitivity is offset from the  
427 dominant wavelengths of the environment could play an important role in contrast  
428 detection under low light conditions [36]. For example, in stickleback UV wavelengths  
429 are important for detection of zooplankton prey against the background light [38]. This  
430 idea is consistent with the observed trend toward reduced UV opsin expression in  
431 freshwater stickleback populations, since most are less zooplanktivorous than marine  
432 stickleback [39]. Experimental work in other fish species has also shown that reduced  
433 UV sensitivity coincides with reduced zooplanktivory and zooplankton foraging ability  
434 [40,41] A second possible explanation for the low match between spectral sensitivity  
435 and ambient light is that detection of specific wavelengths might be important for mate  
436 choice and intraspecific signaling. Short (UV-blue) and long wavelengths (yellow-red)  
437 are important signals for mate choice in stickleback [42], as male nuptial colouration  
438 often involves blue and red pigmentation [43], as well as reflection in the UV [44].  
439 Tuning of perception towards these nuptial signals and detection of contrast among  
440 them could also contribute to the mismatch of sensitivity to available light. It is also  
441 conceivable that our estimates of sensitivity, which do not account for non-additive  
442 signal integration during neuronal processing, underestimate the environmental  
443 correlation.

444           A<sub>2</sub> opsin chromophore complexes do not necessarily act synergistically with  
445 changes in opsin expression to produce adaptive shifts in spectral sensitivity. In the  
446 populations surveyed substitution of A<sub>1</sub> chromophores with A<sub>2</sub> chromophores weakens  
447 the relationship between shifts in spectral sensitivity and shifts in ambient light. While  
448 the empirical ratios of A<sub>1</sub> and A<sub>2</sub> in the wild are unknown for these freshwater  
449 populations, our analyses suggest that A<sub>2</sub> domination would be unlikely. A<sub>2</sub> dominated  
450 retinas result in shifts in spectral sensitivity that do not correlate to shifts in these  
451 environments, and thus are unlikely to be in an adaptive direction. This was a somewhat  
452 surprising result as A<sub>2</sub> chromophores are commonly used by many species of fish found  
453 in freshwater lakes or streams [22]. The potentially maladaptive shifts seen when  
454 substituting to A<sub>2</sub> are a result of overshooting long-wavelength sensitivity relative to the  
455 prevalence of these wavelengths in the surveyed freshwater lakes. This finding is  
456 consistent with work suggesting A<sub>2</sub> dominated retinas are common for threespine  
457 stickleback from dystrophic lakes that are strongly red-shifted relative to the marine  
458 environment, as A<sub>2</sub> use in such an environment would likely result in shifts in an  
459 adaptive direction [24].

460           In this study we provide three lines of evidence to suggest that observed shifts in  
461 spectral sensitivity are adaptive: we show that they have evolved repeatedly, are  
462 genetically based and that regions of the spectrum that differ between marine and  
463 freshwater locations are largely the same regions that exhibit differences in spectral  
464 sensitivity between populations. The methods used in this study help to understand the  
465 direction of evolution of spectral sensitivity, and its relationship with ambient light.  
466 However, our approach does not allow us to disentangle the relative contribution of

467 selection on colour discrimination, contrast detection and photon capture to shifts in  
468 spectral sensitivity. Future experimental and theoretical work will be required to  
469 determine the importance of selection on each of these functions.

470

471 **Author's Contributions:** D.J.R, D.S & T.V conceived of and designed the study; D.J.R,  
472 G.L.O, & T.V collected the data; D.J.R, T.V & N.H analyzed the data; T.V and N.H  
473 developed the statistical method; D.J.R wrote the manuscript with help from T.V and  
474 D.S. All authors contributed to the editing.

475

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485

#### 486 **Data Accessibility**

487 The raw data and accompanying meta data are archived in the dryad database,  
488 doi: 10.5061/dryad.1mc01

489

490 **Competing Interests**

491 The authors have no competing interests.

492

493 **Figure Captions**

494 Figure 1. Normalized cone opsin gene expression of marine and freshwater  
495 populations. Marine populations are indicated in black, freshwater populations in grey.

496 Horizontal lines indicate the mean of all populations; circles indicate individual fish.

497 Location abbreviations: Oyster Lagoon (O), Little Campbell River (LC), Priest Lake (Pr),  
498 Paxton Lake (Pa), Little Quarry Lake (LQ), Trout Lake (T), Kirk Lake (K), and Cranby  
499 Lake (C).

500

501 Figure 2. Estimated spectral sensitivity of marine and freshwater populations assuming  
502 both only use the A<sub>1</sub> chromophore. Marine populations are indicated in black, freshwater  
503 in grey. The thin lines are the fitted values of spectral sensitivity from the mixed-effects  
504 model. The shaded regions are one standard error above and below the fitted values,  
505 with standard errors also derived from the mixed-effects model.

506

507 Figure 3. Normalized cone opsin gene expression of benthic and limnetic populations.

508 The benthic populations are in black, limnetic populations in grey. Horizontal lines  
509 indicate the mean of all populations; triangles indicate individual fish. Location names  
510 abbreviated as: Priest Lake (Pr), Paxton Lake (Pa), Little Quarry Lake (LQ).

511

512 Figure 4. Opsin expression in wild and lab reared fish from a marine (Oyster Lagoon  
513 (O)) and freshwater location (Priest Lake (Pr)). Wild fish are indicated in black, lab  
514 reared fish in grey. Horizontal lines indicate the mean of the population, and points  
515 indicate individual fish.

516

517 Figure 5. (A) Correlations between shifts in spectral sensitivity of individuals from  
518 freshwater populations and differences in local light transmission relative to the  
519 reference marine population, Oyster Bay. (B) As in (A) but using irradiance to measure  
520 light environment shift. (C) Correlations between shifts in spectral sensitivity between  
521 sympatric benthic and limnetic stickleback species and differences in local light  
522 transmission. (D) As in (C) but using irradiance to compare light environments.

523

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