

SI GUIDE

File Name: Supplementary Information

Description: Supplementary Figures, Supplementary Tables and Supplementary References.

File Name: Supplementary Data 1

Description: Differentially expressed genes in transition from SC to LW conditions.

Black and blue characters indicate long day/warm temperature (LW)-up-regulated and -down-regulated genes, respectively. Genes highlighted in yellow are those involved in phototransduction.

File Name: Supplementary Movie 1

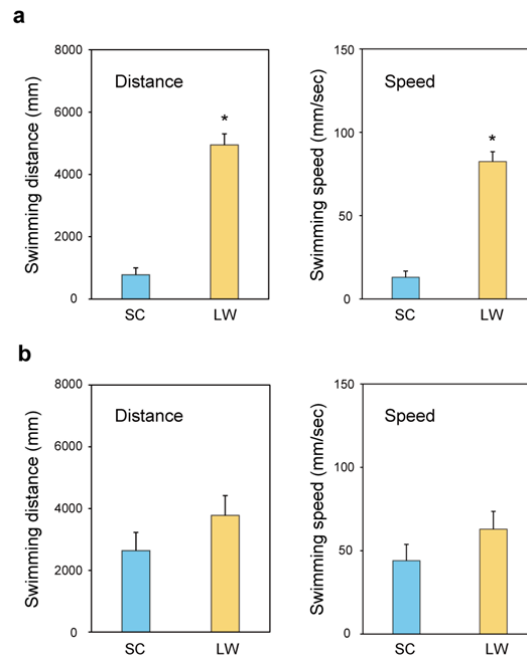
Description: Virtual medaka generated by three-dimensional computer graphics (3D-CG).

File Name: Peer Review File

Description:

1 **Supplementary Figures**

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5 **Supplementary Figure 1. Although activity level of short day/cool**

6 **temperature (SC) fish is low, they have the capacity to move. (a)** Significant

7 differences were observed in the swimming distance (left; $P < 0.05$, t -test, mean

8 + SEM, $n = 10-12$) and swimming speed (right; $P < 0.05$, t -test, mean + SEM, $n =$

9 10-12) between SC fish and long day/warm temperature (LW) fish in an open

10 field tank. Swimming distance and speed were calculated from the data shown in

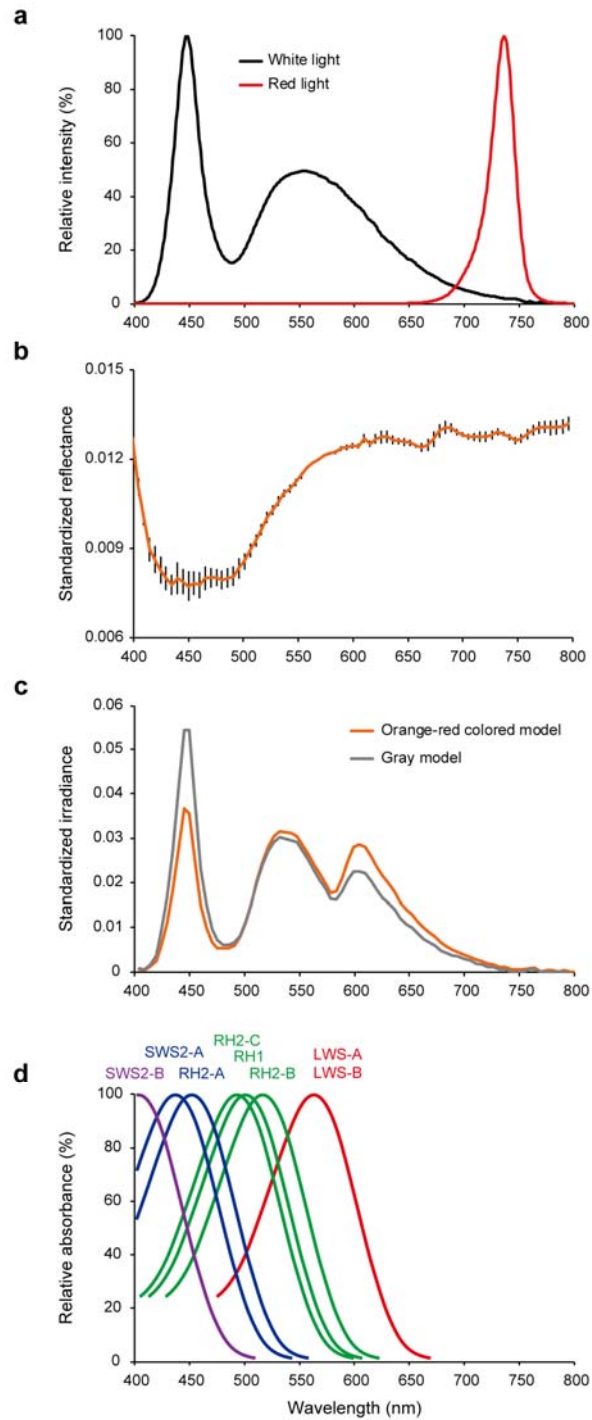
11 Figure 1a. **(b)** No significant differences were observed in the swimming

12 distance (left; $P > 0.05$, t -test, mean + SEM, $n = 5-9$) and swimming speed (right;

13 $P > 0.05$, t -test, mean + SEM, $n = 5-9$) between SC and LW fish in a novel test

14 tank. Swimming distance and speed were calculated from the data shown in

15 Figure 1b.

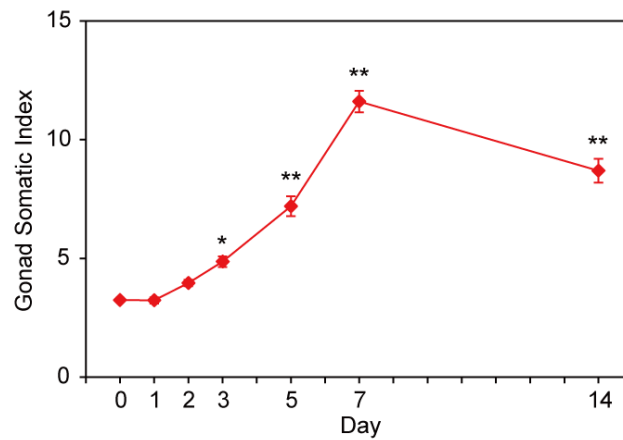


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 17 **Supplementary Figure 2. Spectral curves for light sources, body color and**
 18 **visual pigments. (a) Relative radiant output of white and monochromatic red**
 19 **light. (b) Spectral reflectance from the caudal fin of medaka kept under long**

20 day/warm temperature (LW) conditions (mean $\pm$ SEM, n = 4) (c) Spectral power
21 distribution of orange-red colored model fish and gray model fish on the screen.
22 Orange-red-colored model had high spectral power at long wavelengths covered
23 by LWS. Note that gray model fish also contained spectral power in the red
24 range. This is because gray is produced by mixing three-additive primary color
25 beams (RGB). (d) Schematic drawing of absorption spectra of medaka
26 photopigments. These absorption spectra were drawn by using a spectral
27 template¹ according to the reported values for absorption maxima². Note that
28 although the λ_{max} of LWS is about 560 nm, LWS has been demonstrated to cover
29 an unexpectedly wide range of wavelengths (up to 830 nm) under light-adapted
30 conditions³.

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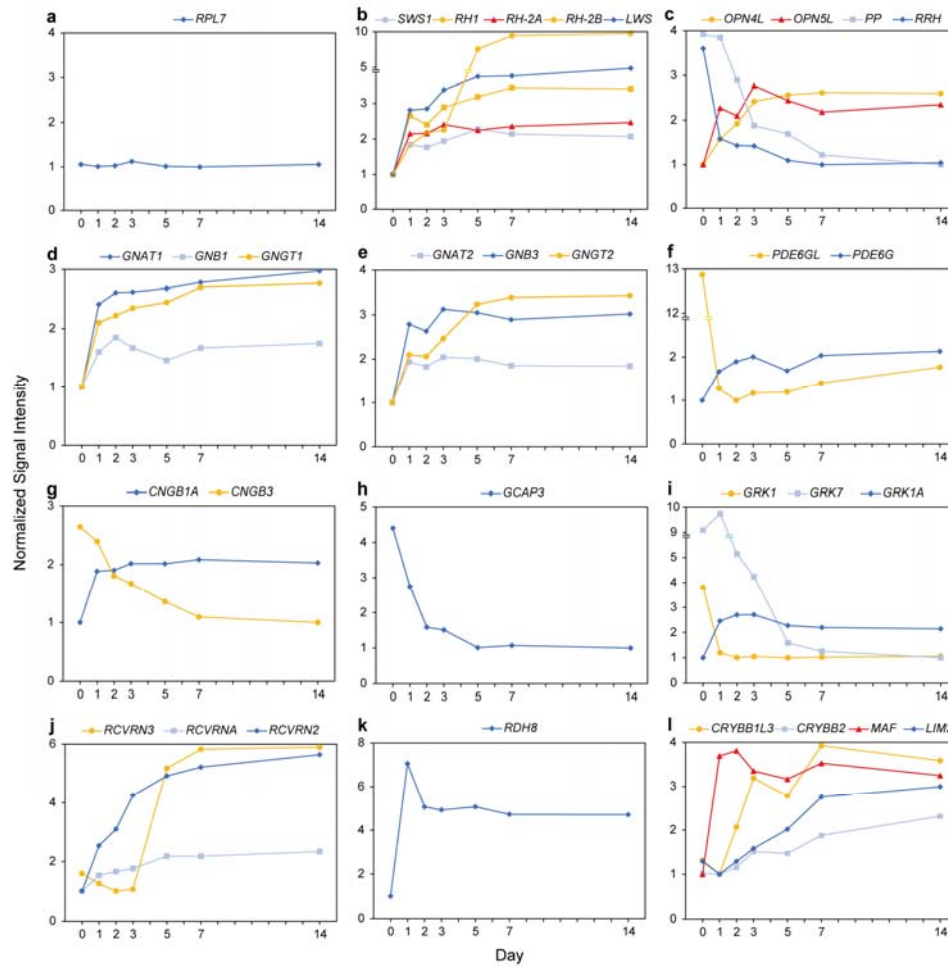
32



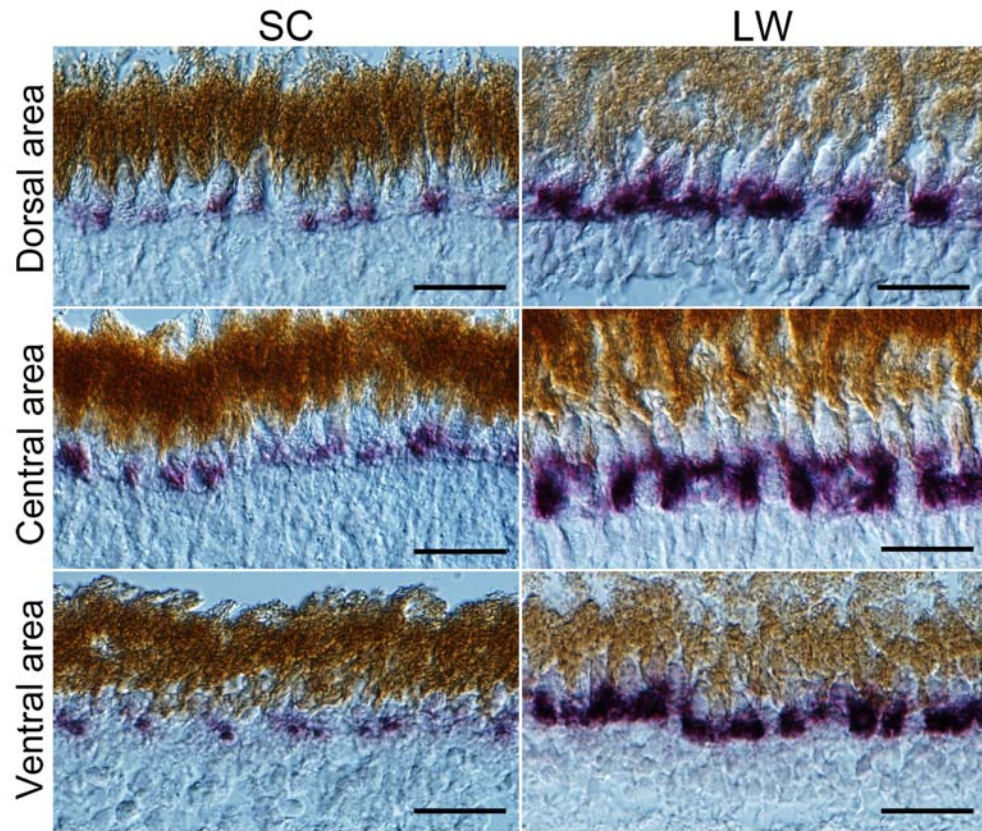
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34 **Supplementary Figure 3. Induction of gonadal development by LW.** When
 35 fish kept under SC conditions were transferred to LW, a significant increase in
 36 the gonad somatic index (gonadal weight / body weight $\times$ 100) was first detected
 37 on day 3, and ovaries reached their full size on day 7 ($F_{6,35} = 73.8$, $P < 0.01$,
 38 ANOVA; * $P < 0.05$, ** $P < 0.01$, vs. day 0, Dunnett's test; mean $\pm$ SEM, n = 6).

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Supplementary Figure 4. Seasonal changes in the expression of genes involved in phototransduction determined by microarray analysis. (a) House-keeping gene. (b) Image-forming visual pigment genes. (c) Non-image-forming visual pigment genes. (d) Rod transducin genes. (e) Cone transducin genes. (f) Phosphodiesterase genes. (g) Cyclic-nucleotide-gated cation channel genes. (h) Guanylate-cyclase-activating protein gene. (i) Rhodopsin kinase genes. (j) Recoverin genes. (k) Retinol dehydrogenase gene. (l) Genes involved in lens formation. Data were normalized such that the minimum expression values over all time points for each gene is 1.



Supplementary Figure 6. Long day/warm temperature (LW)-induction of *LWS* is observed in whole retina. Representative high-resolution images from the dorsal, central and ventral areas of the retina taken by Differential Interference Contrast (DIC) microscope (Olympus BX63). Scale bars: 20 μ m.

65 **Supplementary Tables**66 **Supplementary Table 1. Primer sequences used in the Q-PCR**

Gene symbol	Primer sequence (sense)	Primer sequence (antisense)
<i>RPL7</i>	5' - aactggcctttgtcatcagg - 3'	5' - aaagtctgcagcttctccg - 3'
<i>SWS1</i>	5' - ggagtcggtacatccctgaa - 3'	5' - agttcagttgcgccagctat - 3'
<i>RH1</i>	5' - ttgaaggcttcttggcact - 3'	5' - aatgaccaacttccgctttg - 3'
<i>RH-2A</i>	5' - gagggattcatggcaacact - 3'	5' - ttcaaattcactgcaacca - 3'
<i>RH-2B</i>	5' - gctggagtccttctgacctg - 3'	5' - tacactctggctcctggctt - 3'
<i>LWS</i>	5' - gggtagtcgtgtgcaaacct - 3'	5' - ttggtggaatcgtcttctcc - 3'
<i>OPN4L</i>	5' - accacatggacatcaagcaa - 3'	5' - acctggacacttcgttttgg - 3'
<i>OPN5L</i>	5' - gatcacgctgaaggcttttc - 3'	5' - cagcactctgcactcagagc - 3'
<i>PP</i>	5' - ggatgtttgtggtgtgcaag - 3'	5' - ctgtggtcccttacctggaa - 3'
<i>RRH</i>	5' - gcatccttgggtgaccctaa - 3'	5' - ctcttggccaagtctccac - 3'
<i>GNAT1</i>	5' - ggagctagcgtgaagagaa - 3'	5' - ctggagaatcaggcaaaagc - 3'
<i>GNB1</i>	5' - cctcctgggtcatgactgt - 3'	5' - aacctgaaaacacgtgaggg - 3'
<i>GNGT1</i>	5' - tcgacgacctcacagacaag - 3'	5' - ggaagaggacacctgtga - 3'
<i>GNAT2</i>	5' - tgaagaaggccagaaagaa - 3'	5' - agatcgtgttcaacgctgtg - 3'
<i>GNB3</i>	5' - tgaagcaggcgtgattgag - 3'	5' - aaaaccggagacaattgtgc - 3'
<i>GNGT2</i>	5' - tgaagaaggccagaaagaa - 3'	5' - agatcgtgttcaacgctgtg - 3'
<i>PDE6GL</i>	5' - acgctgcacgtgtatcttg - 3'	5' - caactgtcagcgtgtgtt - 3'
<i>PDE6G</i>	5' - cctgctcttttctcgtttgc - 3'	5' - gaaacttacgcaagcaaggc - 3'
<i>CNGB1A</i>	5' - gctggcgagaactactaccg - 3'	5' - gaagtccagaaccggatcaa - 3'
<i>CNGB3</i>	5' - ttctgccgttggactgttg - 3'	5' - ttgagttcagtgatgcctg - 3'
<i>GCAP3</i>	5' - cgtcaaaggagaaggtgagc - 3'	5' - tcatggaaatgctcaccaa - 3'
<i>GRK1</i>	5' - gtcagtgtacgagctgagga - 3'	5' - aactgaatgcaggatcctg - 3'
<i>GRK7</i>	5' - acgagatggtggtggtgatac - 3'	5' - tctcaacgaggagccaaagt - 3'
<i>GRK1A</i>	5' - tatctcgatctgggtctgg - 3'	5' - ctaagggttatgccgggaca - 3'
<i>RCVRN3</i>	5' - tgatggcacttggatttca - 3'	5' - aaaactggagtgggcctttt - 3'
<i>RCVRNA</i>	5' - aacatggggaacacgaagag - 3'	5' - caaatactcagaggccgagc - 3'
<i>RCVRN2</i>	5' - cgatgatgggactctggact - 3'	5' - aaccactcgcaaactcgagt - 3'
<i>RDH8</i>	5' - cagcaaagctgacaaaccaa - 3'	5' - agacatcgagagcacacac - 3'
<i>CRYBB1L3</i>	5' - aactttcagggccgttaggat - 3'	5' - attattgtcagtgctggacc - 3'
<i>CRYBB2</i>	5' - accagctgtaaggagagca - 3'	5' - acagccaggagcacaagatt - 3'
<i>MAF</i>	5' - cagactgcacaaaagcaca - 3'	5' - ccaagagctgatcgtcatca - 3'
<i>LIM2</i>	5' - ctgtgctgtttgtctggtg - 3'	5' - gtcctgtcagccttcttgc - 3'

67 **Supplementary References**

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- 73 3. Homma, N., Harada, Y., Uchiyama, T., Kamei, Y. & Fukamachi, S. Protanopia (Red
74 color-blindness) in medaka: A simple system for making color-blind fish and testing
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