Channelrhodopsin-1: A Light-Gated Proton Channel in Green Algae

Georg Nagel,^{1*} Doris Ollig,¹ Markus Fuhrmann,² Suneel Kateriya,² Anna Maria Musti,³ Ernst Bamberg,¹ Peter Hegemann²

Phototaxis and photophobic responses of green algae are mediated by rhodopsins with microbial-type chromophores. We report a complementary DNA sequence in the green alga *Chlamydomonas reinhardtii* that encodes a microbial opsin-related protein, which we term Channelopsin-1. The hydrophobic core region of the protein shows homology to the light-activated proton pump bacteriorhodopsin. Expression of Channelopsin-1, or only the hydrophobic core, in *Xenopus laevis* oocytes in the presence of all-*trans* retinal produces a lightgated conductance that shows characteristics of a channel selectively permeable for protons. We suggest that Channelrhodopsins are involved in phototaxis of green algae.

The earliest detectable rhodopsin-mediated responses in phototactic green algae are electrical currents within the eye spot. These currents are carried by Ca^{2+} and $H^+(1, 2)$. In Chlamydomonas reinhardtii or Haematococcus pluvialis, photoreceptor currents occur with a delay of $<30 \,\mu s$ after flash stimulation (3, 4), suggesting an intimate link between photoreceptor and electrical conductance. The two dominant retinal-binding proteins in the eye spot, Cop1 and Cop2, are not the photoreceptors that mediate photomovement responses (5). Searching a C. reinhardtii EST (expressed sequence tag) database (6) revealed overlapping cDNA sequences that encode a 76.4-kD opsin-related protein. We have named this protein Channelopsin-1 (Chop1). The core region (amino acid residues 76 to 309 out of 712 total) comprises seven hypothetical transmembrane segments with sequence similarity (15 to 20%) to the archaeal sensory rhodopsins (SRs), the ion transporters bacteriorhodopsin (BR) and halorhodopsin (HR), and rhodopsin from Neurospora (nop1) (7). Although the overall sequence homology is low, several amino acids are conserved that define the retinalbinding site and the H⁺-transporting network in BR (8, 9) (Fig. 1 and Supporting Online Material). The consensus motif LDxxxKxxW (10) suggests that K296 is the retinal-binding amino acid. Thirteen of the 22 amino acids that are in contact with the retinal in BR are identical (9 amino acids) or conservatively

*To whom correspondence should be addressed. Email: nagel@mpibp-frankfurt.mpg.de exchanged (4 amino acids) in Chop1. The conserved residues are close to the protonated Schiff-base site in the archaeal rhodopsins.

We expressed cRNA encoding Chop1 in oocytes of *Xenopus laevis*, in the presence of all-*trans* retinal, to study ion transport under voltage-clamp conditions, as demonstrated for BR and sensory rhodopsin II (11–13). Illumination by green light, but not red light, induced inward currents in *Chop1* RNA-injected oocytes at a membrane potential of -100 mV (Fig. 2A). Similar results were obtained with truncated *Chop1* RNAs encoding amino acids 1 to 346 or 1 to 517. At an external pH (pH_o) of 7.5, the inward current induced by green light reversed at a voltage near -15 mV (Fig. 2B) with clearly visible outward photocurrents at positive membrane potentials. The dependence of photocurrent direction on the applied potential suggests that the reconstituted Channelrhodopsin-1 (ChR1) mediates a light-induced passive ion conductance. A reversal potential of -15 mV is close to the Nernst potential for Cl⁻ [-20 mV, as deduced by expressing CFTR (cystic fibrosis transmembrane conductance regulator) Cl⁻ channels in oocytes (14)] or H⁺ [-12 mV, at an intracellular pH (pH_i) of 7.3, as measured with microelectrodes by us and others (15)], but far from Nernst potentials for Na^+ , K^+ , or Ca^{2+} .

To investigate the ionic specificity of the light-induced permeability, we systematically changed the composition of the bath solution. Lowering the pH_o to 6.0 increased inward photocurrents and shifted the reversal potential to >+40 mV (Fig. 3, A and B). Replacing Cl^- by aspartate (pH₀ = 6) had no discernible effect on the photocurrent amplitude or its current-voltage (I-V) relation (Fig. 3, A and B), thus excluding Cl⁻ as the conducted ion. Similarly, Na⁺ and Ca²⁺ were excluded: Photocurrents were not changed either by replacing Na⁺ by N-methyl-D-glucamine (NMG) or by replacing Ca^{2+} with Mg²⁺ (Fig. 3, A and B). However, increasing the H⁺ concentration of the bath solution, $[H^+]_o$, to 10 μM (pH_o = 5) enhanced the light-induced inward currents, and changing

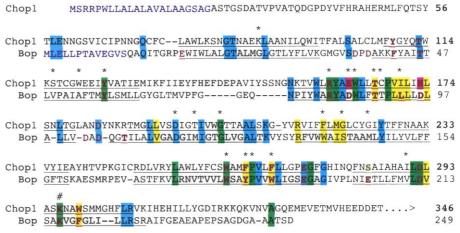
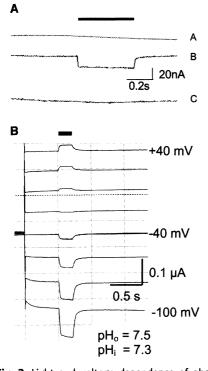


Fig. 1. Comparison of the truncated Chop1 (GenBank accession no. AF385748) sequence (10), Chop1-346 (encoding amino acids 1 to 346), with bacterioopsin (Bop) from *Halobacterium salinarum*. Amino acids that are known from the bacteriorhodopsin (BR) structure to interact directly with the retinal (8, 9) are indicated with an asterisk (*). Amino acids that are identical in most microbial opsins are highlighted in green; those that are functionally homologous in microbial opsin sequences are in yellow; and those that are identical in Chop1, BR, and only some other microbial opsins are inblue. Amino acids that contribute to the H⁺-conducting network in BR (8, 9) are shown in red. Residues that are part of the transmembrane H⁺-network are in bold type. The key substitutions, D⁸⁵ and D⁹⁶ in BR to E¹⁶² and H¹⁷³ in Chop1, are shown in red. Underlined amino acids indicate identified or hypothetical transmembrane helices. Amino acids labeled in blue are leader sequences. The numbering of bacterioopsin begins after the leader sequence for historical reasons. The hypothetical retinal-binding Lys is labeled with "#". For further details, see supporting online material.

¹Max-Planck-Institut für Biophysik, Kennedyallee 70, 60596 Frankfurt am Main, Germany. ²Institut für Biochemie, Universität Regensburg, Universitätsstrasse 31, 93040 Regensburg, Germany. ³Dipartimento Farmaco-Biologico Università della Calabria, 87036 Rende, Italy.

 $[H^+]_o$ to 100 μM (pH_o = 4) further increased the photocurrents (Fig. 3, A to C). Figure 3C demonstrates the magnitude of light-induced H^+ currents at $pH_o = 4$ in an oocyte where light-induced H⁺ conductance exceeds the background conductance of the oocyte by more than fivefold. Lowering pH shifts the reversal potential of the photocurrent to more positive values (Fig. 3B). Taken together, these results indicate that the photocurrent is carried by protons.

The reversal potential of the photocurrent should be shifted to more negative potentials by increasing the cytosolic $[H^+]$. At $pH_0 =$ 7.4, a concentration of 40 mM Na-butyrate, which corresponds to 100 µM undissociated, membrane-permeable butyric acid, has been reported to increase the cytosolic [H⁺] of an oocyte from 50 nM ($pH_i = 7.3$) to 160 nM $(pH_i = 6.8)$ (15). We confirmed by pH measurements with ion-selective microelectrodes at $pH_0 = 7.4$ that 40 mM butyrate caused a



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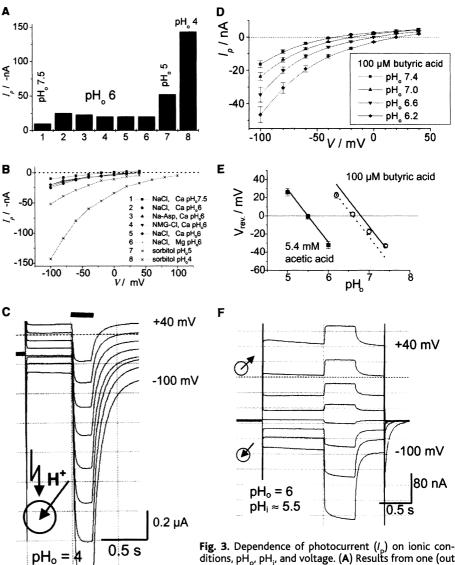
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0.5-unit decrease in pH_i . At $pH_o = 7.4$, 100 µM butyric acid caused a shift of the photo current reversal potential from $-5 \pm 2 \text{ mV}$ to -33 ± 2 mV (n = 5) (Fig. 3D). Photocurrents were measured at four pH_o values-7.4, 7.0, 6.6, and 6.2-with the undissociated butyric acid concentration of 100 µM to maintain the pH_i at ~ 6.8 (15). The direction of photocurrent reversed at a different potential for each pH_o (Fig. 3D). The dependence of



ditions, pH_o , pH_i , and voltage. (A) Results from one (out of five) characteristic Chop1 oocyte, plotted in the order of measurement (\sim 150-s interval); V = -100 mV, green light as in Fig. 2A. Solutions are buffered with 5

Fig. 2. Light and voltage dependence of photocurrents at pH = 7.5. (A) Photocurrents recorded during illumination of oocytes with green or red light (500 \pm 25 nm or 700 \pm 25 nm, respectively; 10²² photons m⁻² s⁻¹). Membrane potential (V) = -100 mV; the light pulse is indicated by a bar. Bath solution: 96 mM NaCl, 5 mM KCl, 2 mM CaCl₂, 1 mM MgCl₂ 5 mM MOPS (pH = 7.5). Trace A: A noninjected oocyte, green light. Trace B: A Chop1 oocyte, green light. Trace C: The same Chop1 oocyte as in trace B, irradiated with red light. (B) Current responses of a Chop1 oocyte to voltage steps from -100 mV to +40 mV (in 20-mV steps; holding potential $V_{\rm h} = -40$ mV), followed by green light pulses of 200-ms duration. Bath solution: 96 mM NaCl, 5 mM KCl, 2 mM CaCl₂, 1 mM MgCl_2 , 5 mM MOPS (pH = 7.5).

mM MOPS (pH = 7.5), MES (pH = 6), or citrate (pH = 5 and 4). Concentration (in mM): bar 1: 100 NaCl, 2 CaCl₂ (pH = 7.5); bar 2: 100 NaCl, 2 CaCl₂ (pH = 6.0) (reference condition); bar 3: 100 Na-aspartate, 2 CaCl₂ (pH = 6.0); bar 4: 100 NMG-Cl, 2 CaCl₂ (pH = 6.0); bar 5: same as bar 2; bar 6: 100 NaCl, 2 EGTA, 2 MgCl₂ (pH = 6.0); bar 7: 200 sorbitol, 5 EGTA (pH = 5.0); bar 8: 200 sorbitol, 5 EGTA (pH = 4.0). (B) Voltage dependence of photocurrents (l_p) from (A). Concentrations as in (A). (C) Current responses to voltage jumps from -100 to +40 mV (in 20-mV steps, $V_h = -40$ mV), followed by green light pulses of 200-ms duration. Bath solution: 200 mM sorbitol, 5 mM citrate (pH = 4.0) with NMG (same oocyte as in Fig. 2B). (D) Voltage dependence of photocurrent (*I*) in different bath solutions, containing 100 μ M undissociated butyric acid: (**I**) 60 mM NaCl + 40 mM Na-butyrate (pH = 7.4); (**A**) 84 mM NaCl + 16 mM Na-butyrate (pH = 7.0); (**V**) 93.6 mM NaCl + 6.4 mM Na-butyrate (pH = 6.6); (\blacklozenge) 97.4 mM NaCl + 2.6 mM Na-butyrate (pH = 6.2) (n = 5). (E) (O) pH_o dependence of reversal potentials, derived from (D). The lines show the theoretical relation for a proton-selective conductance (-58 mV/pH) at a constant pH_i of 6.6 (dotted) or 6.8 (solid). (\blacksquare) pH_o dependence of reversal potentials, derived from experiments (n =6) with 5.4 mM undissociated acetic acid, as demonstrated in (F). The line shows the theoretical relation for a constant pH, of 5.5 and -58 mV/pH. (F) Current responses to voltage steps from V = -100 mV to +40 mV, followed by green light pulses. Bath solution: 100 mM Na-acetate, 5 mM MES (pH = 6.0) with NaOH.

of protons during each ChR1 photocycle or

a light-gated proton channel in which an

intermediate of the ChR1 photocycle al-

lows permeation of many H⁺ ions. Howev-

er, further analysis supported the model of

a light-gated channel. The photocurrent de-

cays with two time constants after illumi-

nation, similar to bacteriorhodopsin (16)

(Fig. 4A). These time constants are strong-

ly temperature sensitive, indicating an activation energy for the photocycle of ~ 60

kJ/mol. The temperature sensitivity of the

photocurrent amplitude is much smaller, \sim 20 kJ/mol (Fig. 4, A and B). This sug-

gests that intermediates of the photocycle

serve as the proton-conducting channel.

The maximal ChR1 photocurrent exceeds

maximal BR photocurrents, providing an

additional argument against a stoichiomet-

ric coupling of photocycle and proton

transport. Large photocurrents of a strictly

coupled system would require either a pho-

tocycle that is faster than that of BR, which

is incompatible with the observed slow de-

cay time constants, or a high expression

level (\sim 100-fold greater than that of BR),

which is unlikely. If we assume that the

level of ChR1 expression in oocytes is

similar to that of BR [$\sim 2 \times 10^9$ molecules

per oocyte (11)], then we can calculate the

electrical current contribution of a single

ChR1 molecule. At $pH_0 = 4$ and -100 mV,

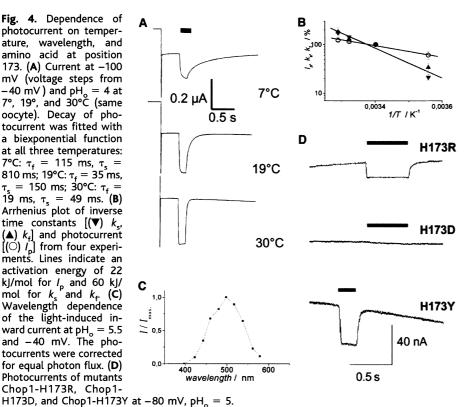
we measured a light-induced inward cur-

rent of $\sim 1.4 \mu A$, i.e., ~ 4000 protons/s or

reversal potential (V_{rev}) on pH_o is shown in Fig. 3E (open circles). The $V_{\rm rev}$ of -33 \pm 2 mV at $pH_o = 7.4$ and $pH_i = 6.8$ (15) is in close agreement with the H⁺ V_{rev} of -35 mV at $\Delta pH = 0.6$. This close fit shows that the light-activated currents are passive. In addition, the dependence of photocurrent V_{rev} on pH_o indicates a high selectivity for protons. A small change of pH_i with changing pH_o, as reported previously for oocytes (15), can explain the deviation from the ideal slope of -58 mV/pH unit. Indeed, a shift of pH_i from 6.8 to 6.6 (as shown by the two lines with ideal slope in Fig. 4C) would bring the observed dependence in accordance with theoretical expectation for a photoconductance that is selectively permeable for H⁺. When a high concentration (5.4 mM) of undissociated acetic acid ($pH_0 = 5$ to 6) was administered, large outward photocurrents at positive potentials (Fig. 3F) and an ideally pHo-dependent V_{rev} with a slope of -58 mV/pH unit were observed (Fig. 3E, filled circles). The $V_{\rm rev}$ observed in these experiments suggests that application of 5.4 mM acetic acid lowers the pH_i to \sim 5.5 (Fig. 3E). We confirmed this value using microelectrodes to measure the pH_i of oocytes in the presence of 5.4 mM acetic acid. Thus, the observed dependence of $V_{\rm rev}$ on pH_o implies that the ChR1-mediated light-sensitive conductance is passive and highly selective for protons.

The dependence of the direction of proton transport on the electrochemical potential for H⁺ is compatible with either facil-

Fig. 4. Dependence of photocurrent on temperature, wavelength, and amino acid at position 173. (A) Current at -100 mV (voltage steps from -40 mV) and pH = 4 at 7°, 19°, and 30°C (same oocyte). Decay of photocurrent was fitted with a biexponential function at all three temperatures: 7°C: $\tau_f = 115$ ms, $\tau_s =$ 810 ms; 19°C: $\tau_f = 35$ ms, _ = 150 ms; 30°C: τ_f = 19 ms, $\tau_s = 49$ ms. (B) Arrhenius plot of inverse time constants $[(\mathbf{\nabla}) k_s]$ (**(**) $k_{\rm f}$] and photocurrent [(\bigcirc) $I_{\rm p}$] from four experiments. Lines indicate an activation energy of 22 kJ/mol for I_p and 60 kJ/ mol for k_s and k_f . (C) Wavelength dependence of the light-induced inward current at $pH_o = 5.5$ and -40 mV. The photocurrents were corrected for equal photon flux. (D) Photocurrents of mutants Chop1-H173R, Chop1-



0.7 fA per molecule, which is much too small for single-channel resolution.

The wavelength dependence of the inward photocurrent was determined at $pH_o = 5.5$ and -40 mV (Fig. 4C). The maximum wavelength of \sim 500 nm closely resembles that observed in the action spectra for photoreceptor currents (1), phototaxis (17), and photoshock responses (18) of intact C. reinhardtii cells. The pH_o-dependent photocurrent, I_{P2} , recorded from intact C. reinhardtii cells (2) was found to dominate the stationary current in continuous light at low pH_o (2). We speculate that I_{P2} might be carried by ChR1.

During the BR photocycle, a proton is transferred from the Schiff base to Asp⁸⁵ (corresponding to Glu¹⁶² in Chop1) and released to the surface by way of Arg⁸², Glu¹⁹⁴, and Glu²⁰⁴. The corresponding residues in Chop1 are Arg¹⁶¹, Glu²⁷⁴, and Ser²⁸⁴. The key amino acid for the reprotonation of the retinal Schiff base in BR is $Asp^{96}(8, 19)$. The corresponding amino acid in Chop1 is His¹⁷³ (Fig. 1), which we exchanged with three different amino acids: Asp, Arg, or Tyr. The substitution $His^{173} \rightarrow Asp$ (H173D) resulted in a complete loss of light-gated conductance, whereas H173R or H173Y were still functional (Fig. 4D). These results indicate that His¹⁷³ does not function as a proton donor of a deprotonated Schiff base. Therefore, we suggest that in ChR1 the retinal Schiff base is not deprotonated during the photocycle. This claim is corroborated by the observation that blue light, which is absorbed by deprotonated retinylidene, did not quench the stationary currents (20). It is conceivable that isomerization of retinal or a conformational change, tightly coupled to isomerization, gates the ChR1 proton channel.

In conclusion, ChR1 is an ion channel that opens in response to absorption of light, i.e., a combined photoreceptor and ion channel. It is not unlikely that such directly light-sensitive ion channels are widely distributed in other phototactic microalgae, as well as in gametes and zoospores of macroalgae, or even in fungi (e.g., nop1) (7). Moreover, the ability of ChR1 to mediate a large lightswitched H⁺ conductance in oocytes holds promise for the use of ChR1 as a tool for measuring and/or manipulating electrical and proton gradients across cell membranes, simply by illumination.

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A Low Genomic Number of Recessive Lethals in Natural Populations of **Bluefin Killifish and Zebrafish**

Amy R. McCune, ¹*† Rebecca C. Fuller, ²*† Allisan A. Aquilina, ¹‡ Robert M. Dawley,³ James M. Fadool,² David Houle,² Joseph Travis,² Alexey S. Kondrashov⁴

Despite the importance of selection against deleterious mutations in natural populations, reliable estimates of the genomic numbers of mutant alleles in wild populations are scarce. We found that, in wild-caught bluefin killifish Lucania goodei (Fundulidae) and wild-caught zebrafish Danio rerio (Cyprinidae), the average numbers of recessive lethal alleles per individual are 1.9 (95% confidence limits 1.3 to 2.6) and 1.4 (95% confidence limits 1.0 to 2.0), respectively. These results, together with data on several Drosophila species and on Xenopus laevis, show that phylogenetically distant animals with different genome sizes and numbers of genes carry similar numbers of lethal mutations.

One of the key genetic processes in any population is the dynamic equilibrium between the mutational origin of deleterious alleles and their eventual elimination by selection. Deleterious alleles maintained by mutation-selection balance create mutation load and contribute to genetic polymorphism and inbreeding depression (1, 2). There are very few estimates of the genomic numbers of even those kinds of deleterious alleles that can individually produce drastic phenotypes and are readily detected by simple genetic assays for lethal and visible recessives. In several Drosophila species, the average number of visible alleles per individual ranges from 0.3 to 1.3 (3-5), and the number of lethals R is 1.0 to 3.0 (6-11). In Xenopus

†These authors contributed equally to this work. ‡Present address: Department of Biology, Case West-

ern Reserve University, Cleveland, OH 44106, USA.

laevis, the only vertebrate for which there are published data, $R \sim 1.9$ (12).

Here, we report data on R in natural populations of two phylogenetically distant teleost fishes, Lucania goodei and Danio rerio. Both L. goodei and D. rerio were sampled from large wild populations in their native ranges, and a classical design (3) was used to detect their recessive alleles. We produced F₁ sibships by individual matings between different wild-caught parents. Within each sibship, individual brother-sister crosses yielded F₂ offspring, which were scored for mutant phenotypes, identified by abnormal morphology. Twenty-five percent of the lethals present in the wild-caught parents will be exposed as homozygotes in offspring from each F_1 cross. When this happens, 25% of the F_2 offspring show the effects of the recessive allele.

For the L. goodei experiment, we produced 20 F_1 sibships (13) and performed between one and four brother-sister crosses within each sibship. A total of 43 F_1 sib crosses was performed, and 30 to 90 offspring were examined from each cross, resulting in a total of 2569 F₂ offspring, 465 (18%) of which expressed mutant lethal phenotypes. No recessive lethals were found in 17 crosses, and 39 recessive lethals with

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Supporting Online Material

www.sciencemag.org/cgi/content/full/1072068/DC1 Materials and Methods Fig. S1

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clear-cut abnormal morphology were observed in the remaining 26 crosses (14). The fraction of abnormal F₂ embryos or fry was consistent with a 3:1 Mendelian ratio in all crosses except three, where it was too high. In two of these cases, the proportion of mutant phenotypes was explained better by hypothesizing that two unlinked loci act to produce the same effect. In contrast, only 8 (1.6%) out of 499 F₂ offspring, produced in nine control crosses between F₁ individuals from different sibships, were morphologically abnormal.

Thirty-one different lethals (18 before hatching and 13 after hatching) were found in 20 parental pairs (Table 1). Seventeen prehatching lethals could be subdivided into four clear phenotypic classes. We called them "early death," "worm," "pinhead" (which had several variants), and "bulbous head with beady eyes." They were clearly manifested (14) at 1 to 2, 4 to 7, 6 to 9, and 7 to 10 days after fertilization, respectively (Fig. 1). None of these mutants hatched. Twelve lethals that acted after hatching (12 to 14 days after fertilization) could be subdivided into four clear phenotypic classes called "humped," "curly," "uptail," and "no body pigment."

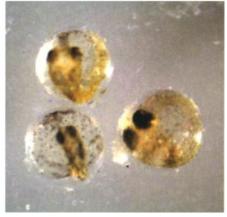


Fig. 1. Mutant and normal phenotypes for Lucania goodei. The phenotypes shown were obtained in cross 15b at 8 days after fertilization. The embryo on top is categorized as bulbous head with beady eyes. The embryo on the bottom left is a pinhead with bulbous ventricles. The embryo on the bottom right is wild type.

¹Department of Ecology and Evolutionary Biology, Cornell University, Ithaca, NY 14853, USA. ²Department of Biological Sciences, Florida State University, Tallahassee, FL 32306-4340, USA. ³Department of Biology, Ursinus College, Main Street, Collegeville, PA 19426-1000, USA. ⁴National Center for Biotechnology Information, National Institutes of Health, 45 Center Drive, Bethesda, MD 20892-6510, USA.

^{*}To whom correspondence should be addressed. E-mail: arm2@cornell.edu (A.R.M.) and fuller@ neuro.fsu.edu (R.C.F.)