

	$J_{-60\text{mV}}$ (pA/pF)	τ_{off} (-60 mV) RT
Chronos (a)	24.01 ± 7.46 (n = 14) # b***, c**, d(ns), e(ns), f(ns), g(ns), h(ns), i(ns)	3.72 ± 0.67 (n = 10) § b(ns), c(ns), d(ns), e(ns), f*, g(ns), h****, i****
f-Chronos (b)	4.66 ± 2.98 (n = 13) # a***, c(ns), d(ns), e(ns), f(ns), g****, h****, i***	1.72 ± 0.12 ms (n = 4) § a(ns), c(ns), d(ns), e(ns), f*, g(ns), h****, i****
f-Chronos LC (c)	4.55 ± 3.09 (n = 9) # a**, b (ns), d(ns), e(ns), f(ns), g****, h****, i***	3.56 ± 0.94 ms (n = 8) § a(ns), b(ns), d(ns), e(ns), f*, g(ns), h****, i****
Chronos LC (d)	17 ± 6.85 (n = 11) # a(ns), b(ns), (ns), e(ns), f(ns), g(ns), h(ns), i(ns)	8.22 ± 1.73 ms (n = 11) § a(ns), b(ns), c(ns), e(ns), f(ns), g(ns), h**, i*
ChR2 (e)	19.33 ± 7.57 (n = 11) # a(ns), b(ns), c(ns), d(ns), f(ns), g(ns), h(ns), i (ns)	10.54 ± 2.34 ms (n = 9) § a(ns), b(ns), c(ns), d(ns), f(ns), g(ns), h(ns), i(ns)
ChR2 ET/TC (f)	14.15 ± 5.37 (n = 11) # a(ns), b(ns), c(ns), d(ns), e (ns), g(ns), h**, i(ns)	10.99 ± 2.23 ms (n = 11) § a*, b*, c*, d(ns), e(ns), g(ns), h(ns), i(ns)
f-ChR2 TC (g)	28.42 ± 10.36 (n = 16) # a (ns), b****, c****, d(ns), e(ns), f(ns), h(ns), i(ns)	9.73 ± 1 ms (n = 9) § a(ns), b(ns), c(ns), d(ns), e(ns), f(ns), h(ns), i(ns)
CatCh (h)	37.05 ± 12.46 (n = 11) # a(ns), b****, c****, d(ns), e(ns), f**, g(ns), i(ns)	33.09 ± 5.72 ms (n = 9) § a****, b****, c****, d**, e(ns), f(ns), g(ns), i (ns)
ChR2 TC (i)	31.39 ± 17.53 (n = 10) # a(ns), b***, c***, d(ns), e(ns), f(ns), g(ns), h(ns)	28.22 ± 6.52 ms (n = 10) # a****, b****, c****, d*, e(ns), f(ns), g(ns), h(ns)

Legend:

Table EV1. Stationary current densities [$J_{-60\text{mV}}$ (pA/pF)] and closing kinetics (τ_{off} values) of blue light activated ChRs. Stationary current densities were obtained from the quotient of the mean stationary photocurrent upon 500 ms light stimulation (saturating intensity of $\sim 30 \text{ mW/mm}^2$, $\lambda = 473 \text{ nm}$) and the capacitance of the cell. [#] Significantly different current densities compared to a) Chronos, b) f-Chronos, c) f-Chronos LC, d) Chronos LC, e) ChR2, f) ChR2 ET/TC, g) f-ChR2 TC, h) CatCh, and i) ChR2 TC. Closing kinetics were determined at RT by a monoexponential fit of the decaying photocurrent after 3 ms light pulse (saturating intensity of $\sim 30 \text{ mW/mm}^2$, $\lambda = 473 \text{ nm}$). f-Chronos closing kinetics at RT were obtained from photocurrents elicited by 7 ns light pulse ($\lambda = 500 \text{ nm}$, 1020 photons/m^2) to avoid interference in the off-kinetics due to shutter opening/closing time ($\sim 700 \mu\text{s}$) using the Opolette 355 tunable laser system (Opotek Inc, Carlsbad, USA). [§] Significantly different closing kinetics compared to a) Chronos, b) f-Chronos, c) f-Chronos LC, d) Chronos LC, e) ChR2, f) ChR2 ET/TC, g) f-ChR2 TC, h) CatCh, and i) ChR2 TC. Kruskal-Wallis followed by Dunn's test: $p > 0.05$ (ns); * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$; **** $p < 0.0001$. All measurements were performed in NG108-15 cells transiently transfected with the specified ChR variants by whole-cell patch clamp at membrane potential of -60 mV . Data are

	$\tau_{\text{off}} (-60 \text{ mV})$ RT	$\tau_{\text{off}} (-60 \text{ mV})$ 33 to 34 °C	EC50 (mW/mm ²)	Desensitization (Stat/Peak)
Chronos (a)	3.72 ± 0.67 (n = 10)	1.875 ± 0.47 ms (n = 6) § b(ns), d(ns), g(ns), h**	1.25 ± 0.64 (n = 6) \$ b(ns), d(ns), e(ns), f(ns), g(ns), h(ns)	0.35 ± 0.085 (n = 10) # b****, c(ns), d****, e(ns), f(ns), g(ns), h****, i(ns)
f-Chronos (b)	1.72 ± 0.12 ms (n = 4)	0.80 ± 0.099 ms (n = 4) § a(ns), d(ns), g*, h***	5.12 ± 0.91 (n = 6) \$ a(ns), d(ns), e(ns), f(ns), g(ns), h**	0.19 ± 0.044 (n = 14) # a****, c(ns), d****, e(ns), f*, g****, h****, i(ns)
f-Chronos LC (c)	3.56 ± 0.94 ms (n = 8)			0.29 ± 0.026 (n = 3) # a(ns), b(ns), d****, e(ns), f(ns), g(ns), h****, i(ns)
Chronos LC (d)	8.22 ± 1.73 ms (n = 11)	3.93 ± 0.19 ms (n = 4) § a(ns), b(ns), g(ns), h(ns)	1.02 ± 0.14 (n = 4) \$ a(ns), b(ns), e(ns), f(ns), g(ns), h(ns)	0.62 ± 0.077 (n = 6) # a****, b****, c****, e****, f****, g****, h(ns), i****
ChR2 (e)	10.54 ± 2.34 ms (n = 9)		1.29 ± 0.57 (n = 3) \$ a(ns), b(ns), d(ns), f(ns), g(ns), h(ns)	0.26 ± 0.064 (n = 6) # a(ns), b(ns), c(ns), d****, f(ns), g(ns), h****, i(ns)
ChR2 ET/TC (f)	10.99 ± 2.23 ms (n = 11)		1.7 ± 0.33 (n = 3) \$ a(ns), b(ns), d(ns), e(ns), g(ns), h(ns)	0.298 ± 0.063 (n = 9) # a(ns), b*, c(ns), d****, e(ns), g(ns), h****, i(ns)
f-ChR2 TC (g)	9.73 ± 1 ms (n = 9)	4.1 ± 0.96 ms (n = 12) § a**, b****, d(ns), h(ns)	1.29 ± 0.315 (n = 3) \$ a(ns), b(ns), d(ns), e(ns), f(ns), h(ns)	0.37 ± 0.073 (n = 7) # a(ns), b****, c(ns), d****, e(ns), f(ns), h****, i(ns)
CatCh (h)	33.09 ± 5.72 ms (n = 9)	15.71 ± 2.17 ms (n = 5)	0.54 ± 0.12 (n = 3)	0.74 ± 0.058 (n = 8)

		§ a**, b***, d(ns), g(ns)	§ a(ns), b**, d(ns), e(ns), f(ns), g(ns)	# a****, b****, c****, d(ns), e****, f****, g****, i****
ChR2 TC (i)	28.22 ± 6.52 ms (n = 10)		0.637 (n = 1)	0.28 ± 0.079 (n = 6) # a(ns), b(ns), c(ns), d****, e(ns), f(ns), g(ns), i****

Legend:

Table EV2. Closing kinetics (τ_{off} values), EC50 values and stationary-peak-ratios of blue light activated ChRs. Closing kinetics were determined at RT by a monoexponential fit of the decaying photocurrent after 3 ms light pulse (saturating intensity of 30 to 40 mW/mm², λ = 473 nm), and f-Chronos closing kinetics were obtained from photocurrents elicited by 7 ns light pulse (λ = 500 nm, 1,020 photons/m²), or more physiological temperature (33 to 34 °C; 1 ms, saturating intensity of 30 to 40 mW/mm², λ = 488 nm). The half maximal activation value (effective power density for 50 % activation, analogous to an EC₅₀) was determined by hyperbolic fitting of the stationary photocurrent amplitude obtained upon 0.5 or 1 s light pulses at different irradiances (mW/mm²). Desensitization values were determined by the quotient of the mean stationary photocurrent of 0.5 to 1 s light pulse and their respective peak photocurrent. All measurements were performed in NG108-15 cells transiently transfected with the specified ChR variants by whole-cell patch clamp at membrane potential of -60 mV. Data are presented as mean \pm SD. All values are derived from the data shown in Figure 1. § Significantly different closing kinetics compared to a) Chronos, b) f-Chronos, d) Chronos LC, g) f-ChR2 TC, and h) CatCh. \$ Significantly different EC50 compared to a) Chronos, b) f-Chronos, d) Chronos LC, e) ChR2, f) ChR2 ET/TC, g) f-ChR2 TC, and h) CatCh. # Significantly different desensitization to a) Chronos, b) f-Chronos, c) f-Chronos LC, d) Chronos LC, e) ChR2, f) ChR2 ET/TC, g) f-ChR2 TC, h) CatCh, and i) ChR2 TC. τ_{off} (-60 mV; at 33 to 34 °C) and EC50 were tested by Kruskal-Wallis t-test and post-hoc Dunn's test: p > 0.05 (ns); * p < 0.05; ** p < 0.01; *** p < 0.001; **** p < 0.0001. Desensitization was tested by ANOVA Bonferroni: p

	Frequency (Hz)	Stationary/peak	Fluctuation/peak
f-ChR2 TC (a)	5	0.54 ± 0.06, n = 4 # b(ns), c(ns), d**	0.99 ± 0.002, n = 4 \$ b(ns), c(ns), d(ns)
	10	0.58 ± 0.06, n = 4 # b(ns), c(ns), d***	0.99 ± 0.009, n = 4 \$ b*, c(ns), d(ns)
	20	0.57 ± 0.07, n = 4 # b(ns), c***, d***	0.98 ± 0.01, n = 4 \$ b**, c(ns), d(ns)
	30	0.54 ± 0.06, n = 4 # b*, c***, d***	0.98 ± 0.01, n = 4 \$ b***, c(ns), d(ns)
	40	0.52 ± 0.05, n = 4 # b***, c****, d****	0.98 ± 0.01, n = 4 \$ b****, c(ns), d(ns)
	50	0.49 ± 0.04, n = 4 # b****, c****, d****	0.97 ± 0.01, n = 4 \$ b****, c(ns), d(ns)
	125	0.38 ± 0.05, n = 4 # b****, c**, d****	0.84 ± 0.05, n = 4 \$ b****, c(ns), d(ns)
	200	0.37 ± 0.05, n = 4 # b****, c**, d****	0.68 ± 0.07, n = 4 \$ b****, c**, d*
	333	0.36 ± 0.05, n = 4 # b****, c**, d***	0.46 ± 0.09, n = 4 \$ b****, c**, d*
	500	0.38 ± 0.06, n = 4 # b****, c***, d***	0.27 ± 0.09, n = 4 \$ b***, c**, d(ns)
CatCh (b)	5	0.6 ± 0.04, n = 4 # a(ns), c(ns), d*	0.98 ± 0.006, n = 4 \$ a(ns), c(ns), d(ns)
	10	0.64 ± 0.02, n = 4 # a(ns), c**, d**	0.97 ± 0.0064, n = 4 \$ a*, c*, d**
	20	0.63 ± 0.008, n = 4 # a(ns), c****, d**	0.89 ± 0.04, n = 4 \$ a**, c**, d**
	30	0.64 ± 0.01, n = 4 # a*, c****, d*	0.73 ± 0.08, n = 4 \$ a***, c***, d***
	40	0.66 ± 0.02, n = 4 # a***, c****, d(ns)	0.63 ± 0.1, n = 4 \$ a****, c***, d***
	50	0.68 ± 0.02, n = 4 # a****, c****, d(ns)	0.54 ± 0.12, n = 4 \$ a****, c***, d***
	125	0.74 ± 0.02, n = 4 # a****, c****, d***	0.21 ± 0.08, n = 4 \$ a****, c****, d****
	200	0.78 ± 0.03, n = 4 # a****, c****, d****	0.11 ± 0.04, n = 4 \$ a****, c****, d****
	333	0.79 ± 0.04, n = 4 # a****, c****, d***	0.05 ± 0.02, n = 4 \$ a****, c****, d**
	500	0.82 ± 0.03, n = 4 # a****, c****, d***	0.02 ± 0.01, n = 4 \$ a***, c****, d(ns)
f-Chronos (c)	5	0.6 ± 0.009, n = 3 # a(ns), b(ns), d*	1 ± 0.01, n = 3 \$ a(ns), b(ns), d(ns)
	10	0.5 ± 0.02, n = 3 # a(ns), b**, d****	0.99 ± 0.003, n = 3 \$ a(ns), b*, d(ns)
	20	0.37 ± 0.01, n = 3 # a***, b****, d****	0.98 ± 0.01, n = 3 \$ a(ns), b**, d(ns)
	30	0.32 ± 0.01, n = 3 # a***, b****, d****	0.98 ± 0.01, n = 3 \$ a(ns), b***, d(ns)
	40	0.32 ± 0.01, n = 3 # a****, b****, d****	0.96 ± 0.01, n = 3 \$ a(ns), b***, d(ns)

	50	0.3 ± 0.01, n = 3 # a****, b****, d****	0.94 ± 0.05, n = 3 \$ a(ns), b***, d(ns)
	125	0.25 ± 0.008, n = 3 # a**, b****, d****	0.89 ± 0.01, n = 3 \$ a(ns), b****, d*
	200	0.22 ± 0.01, n = 3 # a**, b****, d****	0.89 ± 0.05, n = 3 \$ a**, b****, d****
	333	0.21 ± 0.01, n = 3 # a**, b****, d****	0.7 ± 0.06, n = 3 \$ a**, b****, d****
	500	0.18 ± 0.02, n = 3 # a****, b****, d****	0.52 ± 0.04, n = 3 \$ a**, b****, d****
Chronos LC (d)	5	0.75 ± 0.03, n = 3 # a**, b*, c*	1 ± 0.004, n = 3 \$ a(ns), b(ns), c(ns)
	10	0.82 ± 0.01, n = 3 # a****, b**, c****	0.99 ± 0.001, n = 3 \$ a(ns), b**, c(ns)
	20	0.78 ± 0.01, n = 3 # a****, b**, c****	0.99 ± 0.004, n = 3 \$ a(ns), b**, c(ns)
	30	0.74 ± 0.01, n = 3 # a****, b*, c****	0.98 ± 0.0008, n = 3 \$ a(ns), b****, c(ns)
	40	0.73 ± 0.01, n = 3 # a****, b(ns), c****	0.97 ± 0.005, n = 3 \$ a(ns), b****, c(ns)
	50	0.7 ± 0.01, n = 3 # a****, b(ns), c****	0.96 ± 0.005, n = 3 \$ a(ns), b****, c(ns)
	125	0.6 ± 0.01, n = 3 # a****, b***, c****	0.74 ± 0.02, n = 3 \$ a(ns), b****, c*
	200	0.57 ± 0.01, n = 3 # a****, b****, c****	0.53 ± 0.03, n = 3 \$ a*, b****, c***
	333	0.57 ± 0.006, n = 3 # a****, b****, c****	0.29 ± 0.02, n = 3 \$ a*, b**, c****
	500	0.58 ± 0.01, n = 3 # a****, b****, c****	0.14 ± 0.01, n = 3 \$ a(ns), b(ns), c****

Legend:

Table EV3. Frequency dependencies of stationary-peak-ratios and photocurrent fluctuations at physiological temperature. Statistical comparison of data shown in Figure EV1. Whole-cell patch-clamp measurements at ~ 34 °C in NG108-15 cells expressing ChRs 2 to 3 days after transient transfection. ChRs were light activated by 50 light pulses of 1 ms at 488 nm (39.43 mW/mm²) at different frequencies. Stationary photocurrent amplitudes of the different ChR variants were divided by the peak photocurrent amplitude to show how desensitization is influenced by the frequency of stimulation using short light pulses. Data are presented as mean ± SD. # Significantly different desensitization levels at the indicated frequency of light stimulation compared to a) f-ChR2 TC, b) CatCh, c) f-Chronos, and d) Chronos LC. Quantification of the photocurrent fluctuation at the stationary state (end of the pulse train) of different ChR variants normalized to the stationary current amplitude. \$ Significantly different fluctuation amplitudes at the indicated frequency of light stimulation compared to a) f-ChR2 TC, b) CatCh, c) f-Chronos, and d) Chronos LC. Multiple comparison results were obtained by ANOVA Bonferroni: p > 0.05 (ns); * p < 0.05; ** p < 0.01; *** p < 0.001; **** p < 0.0001.

Appendix for Channelrhodopsin variants for high-rate optogenetic neurostimulation at low light intensities

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Appendix Tables

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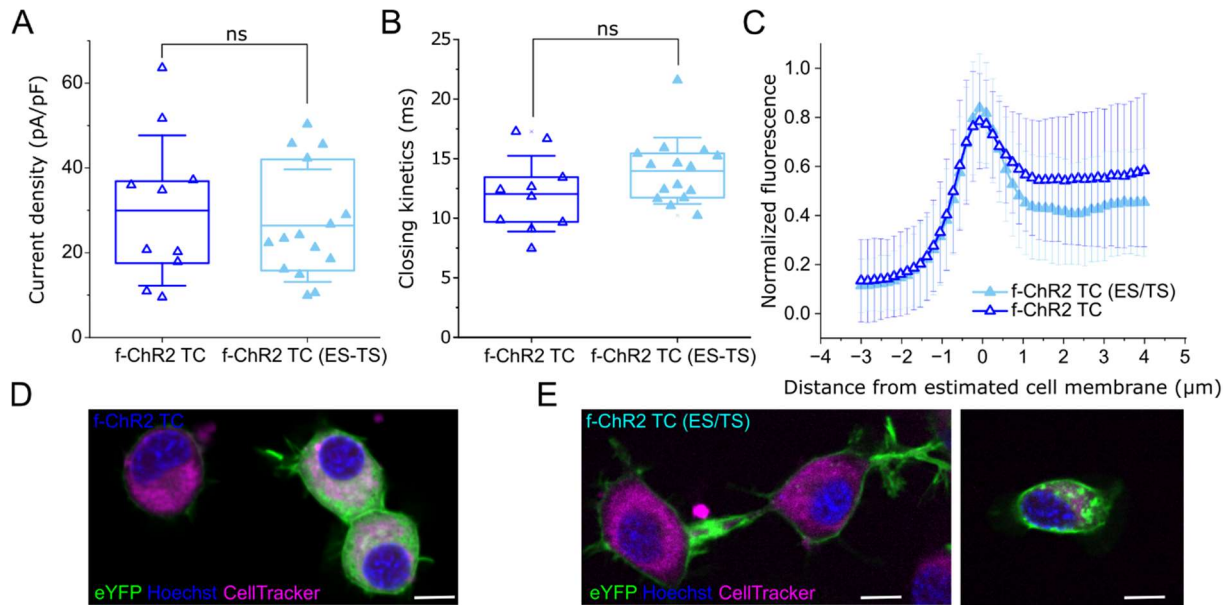
Appendix Table S2. Channelrhodopsin variants.

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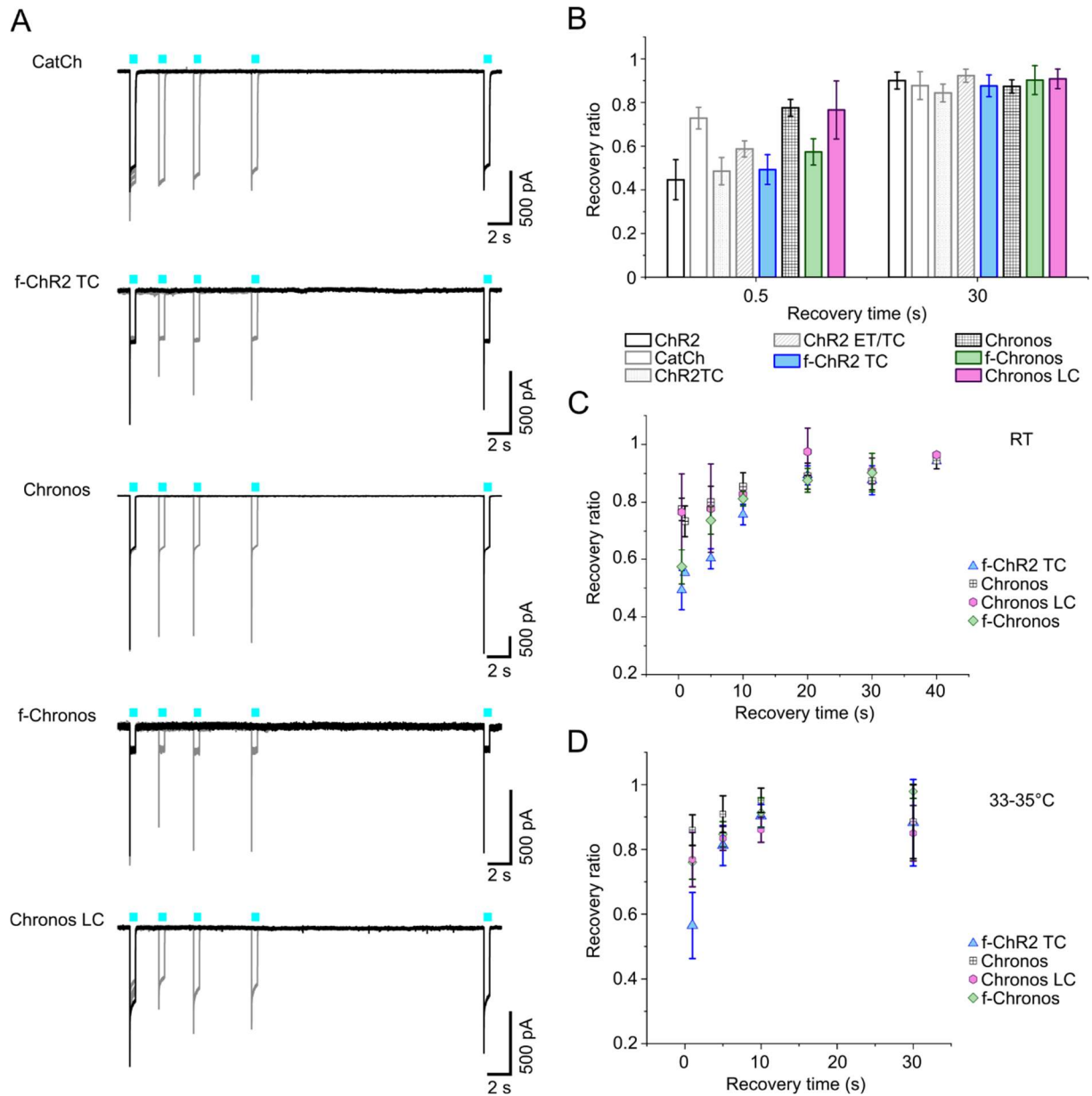
Appendix Table S3. List of primers used for ChR mutant generation.

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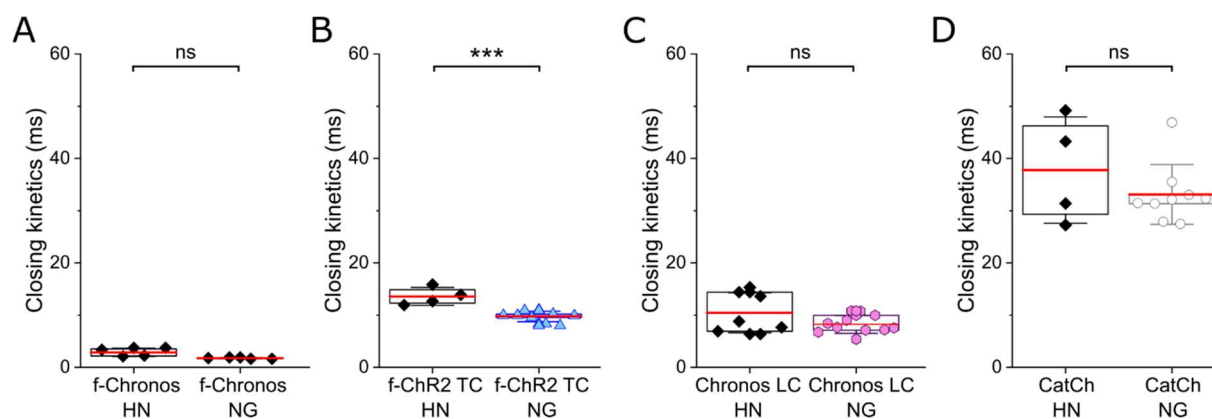
Appendix Figures



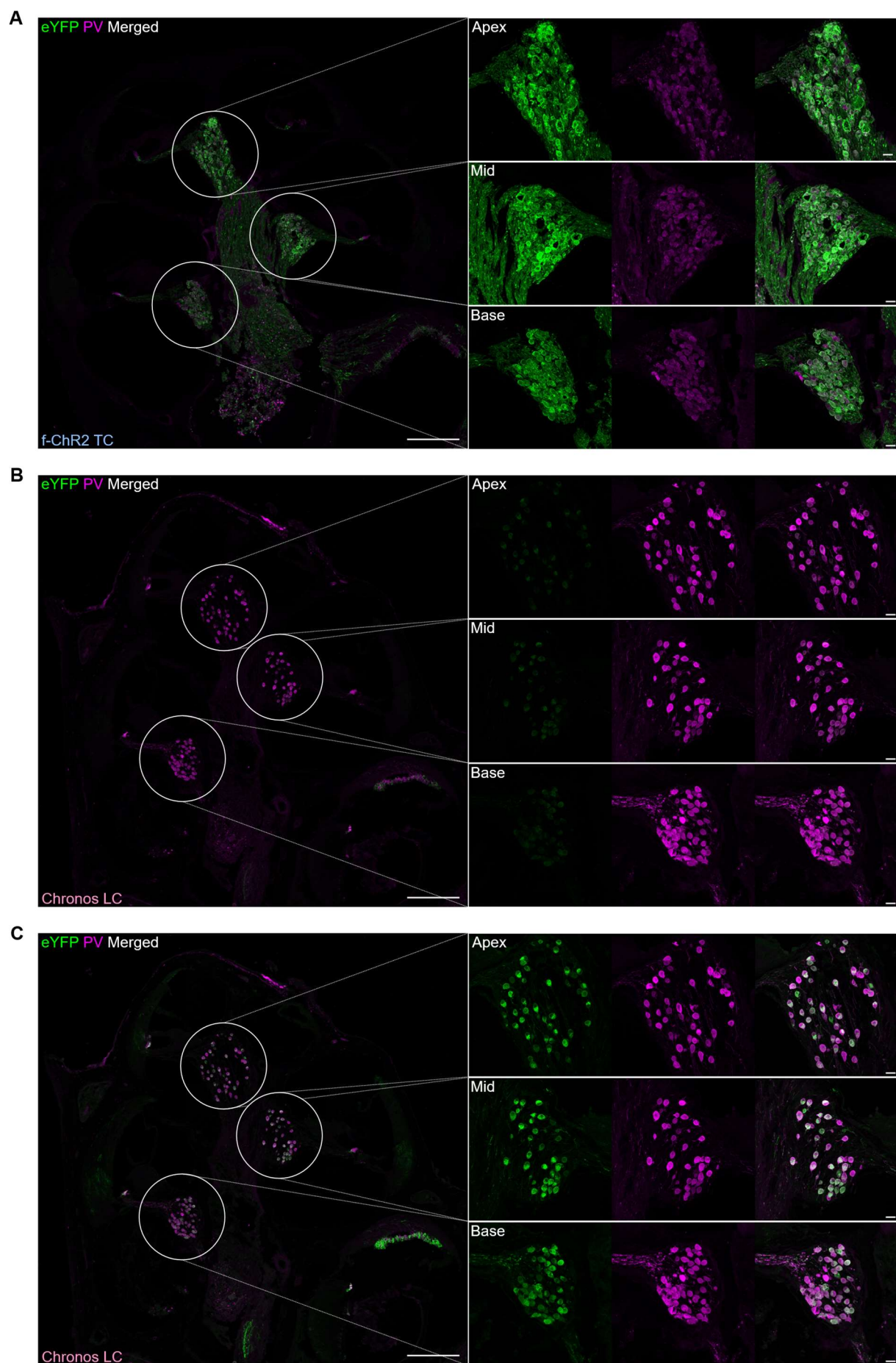
Appendix Figure S1. Effect of the C-terminal fusion of targeting motifs on the plasma membrane expression of f-ChR2 TC. **A-B**, Comparison of current densities (**A**) and closing kinetics (**B**) of f-ChR2 TC and f-ChR2 TC (ES/TS). The whole-cell patch-clamp measurements were performed at RT ($\sim 22^\circ\text{C}$) in NG cells 2 to 3 days after transient transfection. τ_{off} values were determined by monoexponential fits to photocurrents evoked by 3 ms light pulses ($\lambda = 473\text{ nm}$, $\sim 30\text{ mW/mm}^2$) at a membrane potential of -60 mV . The statistical comparison was performed by unpaired t-tests (f-ChR2 TC: $n = 10$ vs f-ChR2 TC (ES/TS): $n = 15$; ns: $p = 0.5719$, $p = 0.1231$). **C**, Line profile analysis of single cells from three different transfections. Plots show means \pm SDs for f-ChR2 TC: $n = 91$ and f-ChR2 TC (ES/TS): $n = 51$. **D-E**, Confocal images of exemplary transfected NG cells expressing f-ChR2 TC (**D**, panel D corresponds to Figure 1A) or f-ChR2 TC (ES/TS) (**E**) that are included in the quantification shown in **C**, eYFP fluorescence shown in green, nucleus stain (Hoechst, 1:2000) shown in blue, cytosol stain (CellTrackerTM, Invitrogen) shown in magenta. Single-plane images of the NG cells were collected by automated spinning disk confocal microscopy (CellVoyager CQ1, Yokogawa) at a magnification of 40 x. Scale bar 10 μm .



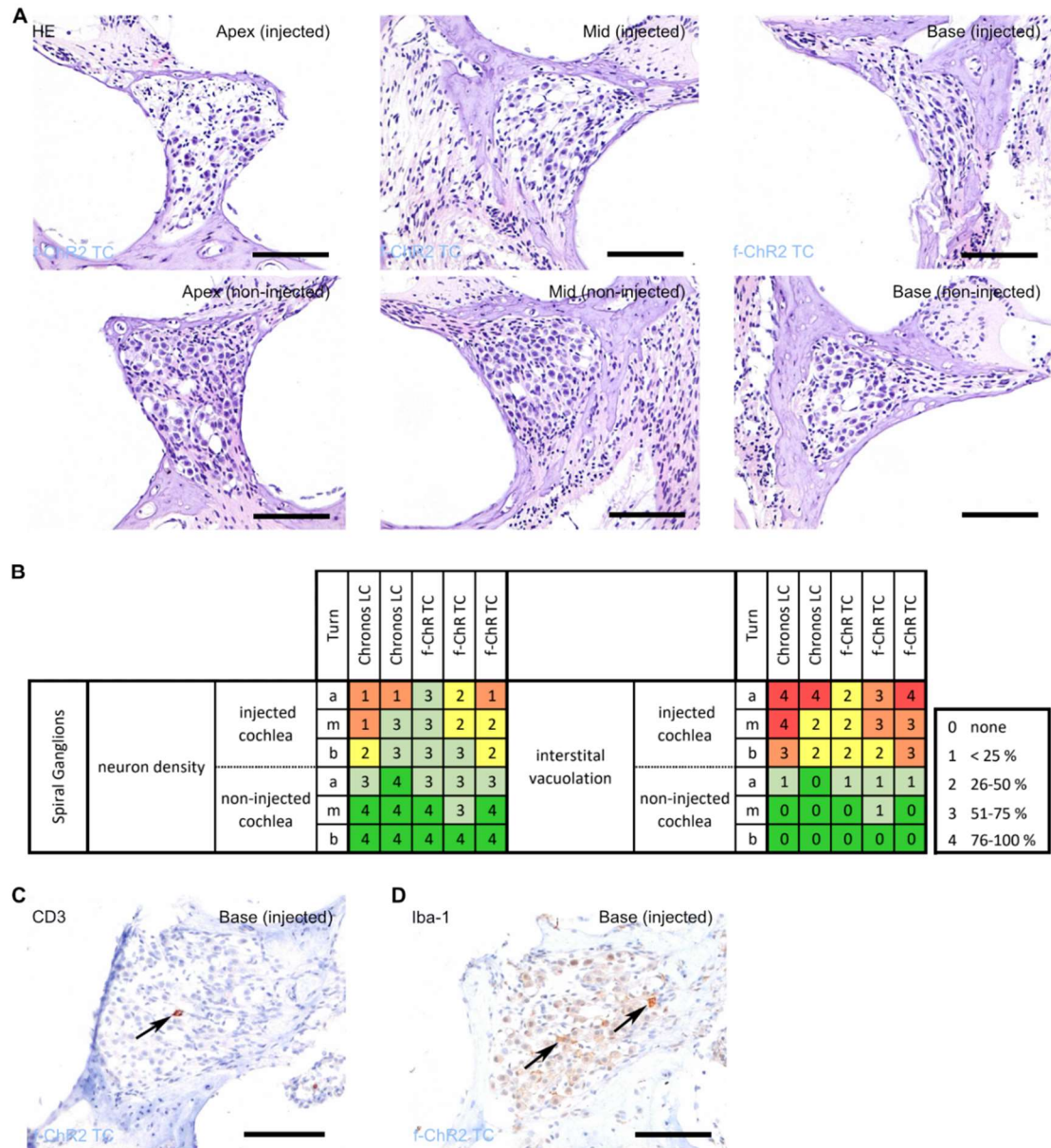
Appendix Figure S2. Recovery from photocurrent desensitization. **A**, Whole cell patch-clamp recordings from NG108-15 cells transiently expressing CatCh, f-ChR2 TC, Chronos, f-Chronos or Chronos LC. Overlaid representative traces of photocurrent peak recovery kinetics after applying two sequential 500 ms light pulses (488 nm; ~ 40 mW/mm²) at different dark time intervals (1, 5, 10, and 30 sec). **B**, Peak photocurrent recovery ratios at the indicated time points. Bars display the mean \pm SD. ChR2: n = 3; CatCh: n = 4; ChR2 TC: n = 4; ChR2 ET/TC: n = 4; f-ChR2 TC: n = 4 to 5; Chronos: n = 5; f-Chronos: n = 4; Chronos LC: n = 4. **C**, Peak photocurrent recovery ratios of selected variants were determined at different time intervals at RT. Dots display the mean \pm SD. f-ChR2 TC: n = 3 to 5; Chronos: n = 2 to 5; f-Chronos: n = 2 to 4; Chronos LC: n = 3 to 4. **D**, Peak photocurrent recovery ratios of selected variants determined after indicated time intervals at a temperature of 33°C to 35 °C. Dots display the mean \pm SD. f-ChR2 TC: n = 5 to 7; Chronos: n = 3 to 6; f-Chronos: n = 3; Chronos LC: n = 3.



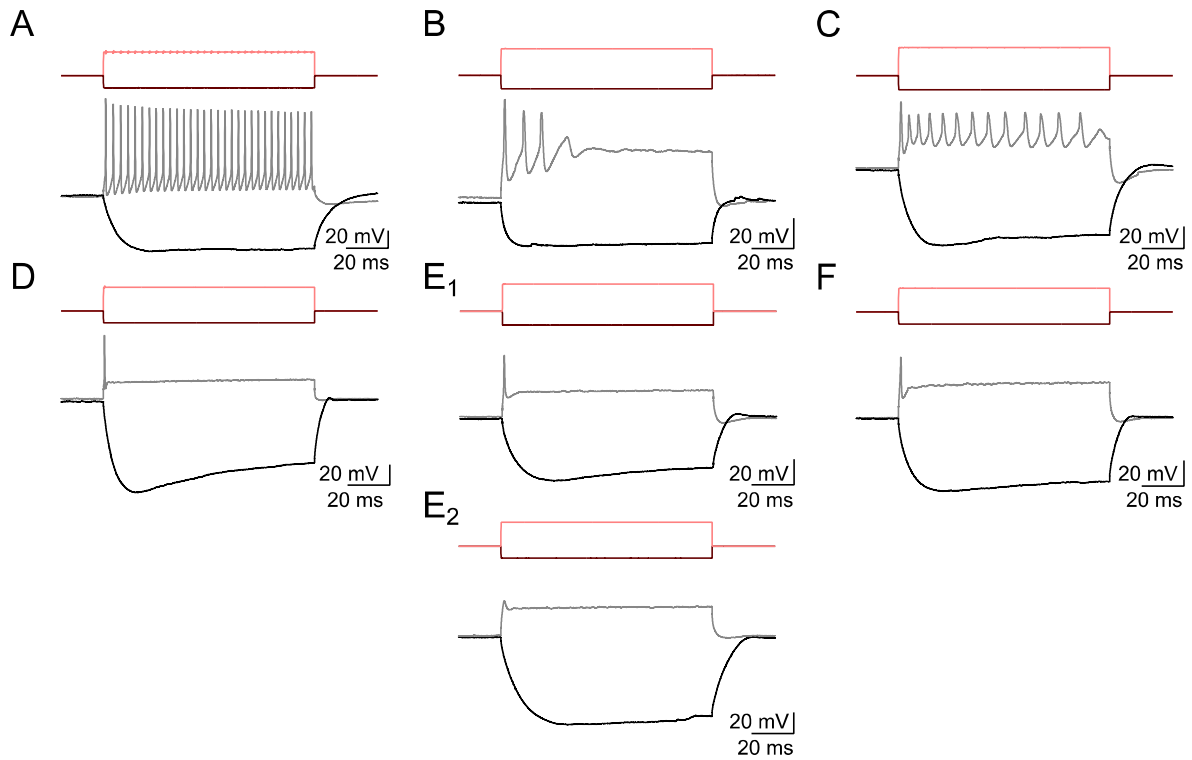
Appendix Figure S3. Comparison of ChR channel closing kinetics in rat hippocampal neurons (HN) and NG cells (NG). Shown is the statistical comparison of the channel closing kinetics of f-Chronos (A) by unpaired t-test with Welch's correction ($p = 0.0656$); f-ChR2 TC (B) by unpaired t-test ($p = 0.0003$); Chronos LC (C) by unpaired t-test with Welch's correction ($p = 0.1930$), and CatCh (D) by Mann-Whitney test ($p = 0.8252$). CatCh (HN): $n = 4$; CatCh (NG): $n = 9$; f-ChR2 TC (HN): $n = 4$; f-ChR2 TC (NG): $n = 9$; Chronos LC (HN): $n = 7$; Chronos LC (NG): $n = 11$; f-Chronos (HN): $n = 4$; f-Chronos (NG): $n = 4$. Data taken from Figure 1B (NG cells) and Figure 2 (hippocampal neurons).



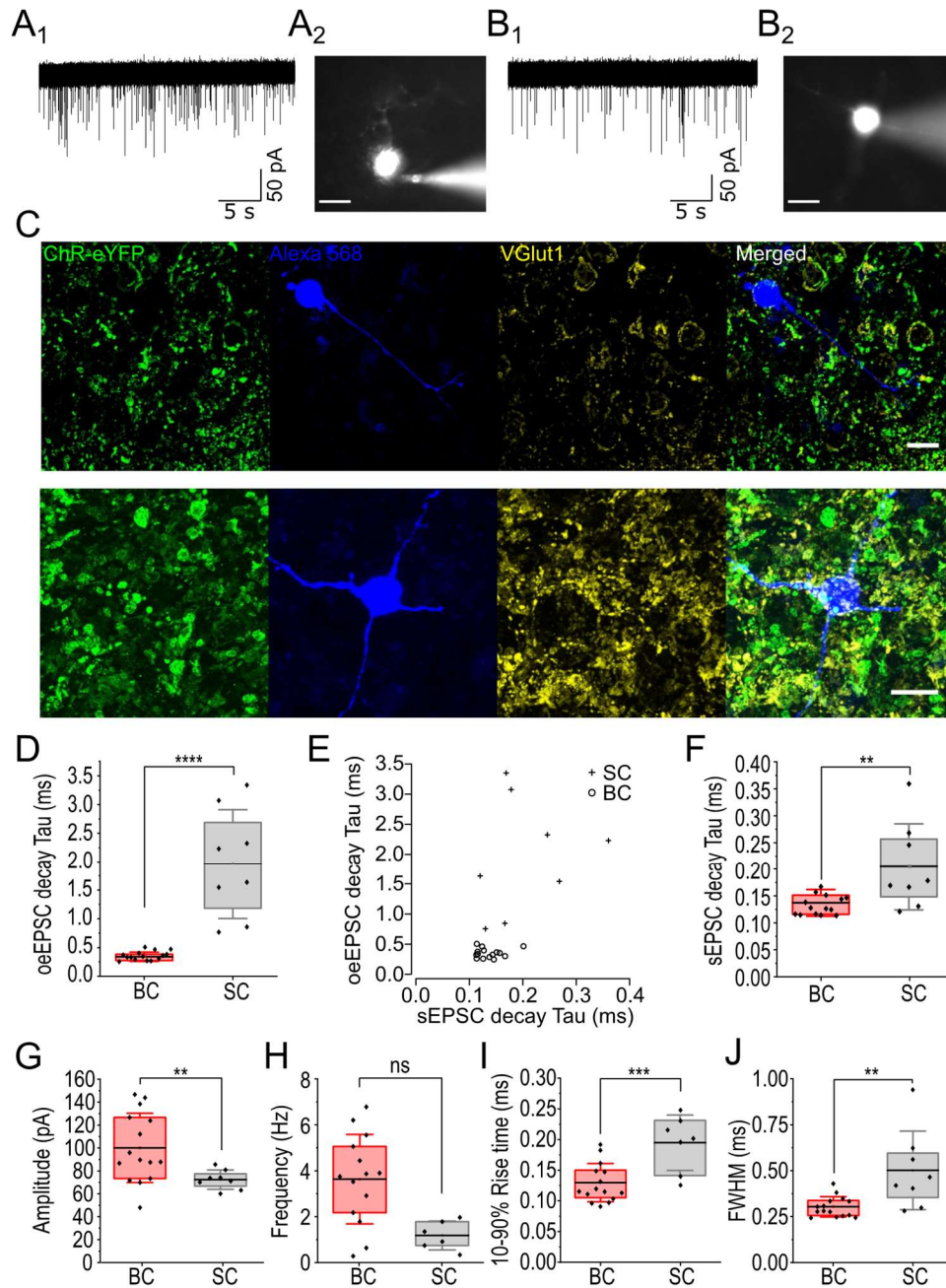
Appendix Figure S4. Immunohistochemical analysis of ChR expression in exemplary oABR-positive cochleae. **A-C**, 20x overview images of exemplary, mid-modiolar sliced cochleae, postnatally injected with AAV2/9_hSyn_ChR2(T159C/F219Y)-EYFP (**A**) and AAV2/9_hSyn_Chronos(L149C)-ES-EYFP-TS (**B-C**) shown on the left. 40 x magnifications of Apex, Mid, and Base shown on the right. Stainings: anti-parvalbumin (PV, SGN marker) and anti-GFP (ChR-EYFP). Scale bar at 20 x = 200µm and at 40 x = 20 µm.



Appendix Figure S5. Histological and immunohistochemical analysis of cochlear paraffin-sections. A, Exemplary mid-modiolar HE-stained sections of apical, medial and basal turns of an animal injected with f-ChR2 TC. **B,** Quantification of neuronal density and interstitial vacuolation (optically empty matrix) for left = injected and right = non-injected (contralateral) cochleae of animals from the f-ChR2 TC (n = 3) and Chronos LC (n = 2) groups. **C,** Staining for CD3 lymphocytes of a left, basal turn of an animal from the f-ChR2 TC group. Arrow indicates a CD3 positive lymphocyte. **D,** Staining for Iba-1 (macrophages) of a left, basal turn of the same animal from the f-ChR2 TC group as in **C**. Arrows indicate cell-associated signal enhancement of Iba-1 expression (not further quantified due to strong diffuse background staining). Scale bars = 100 μ m.

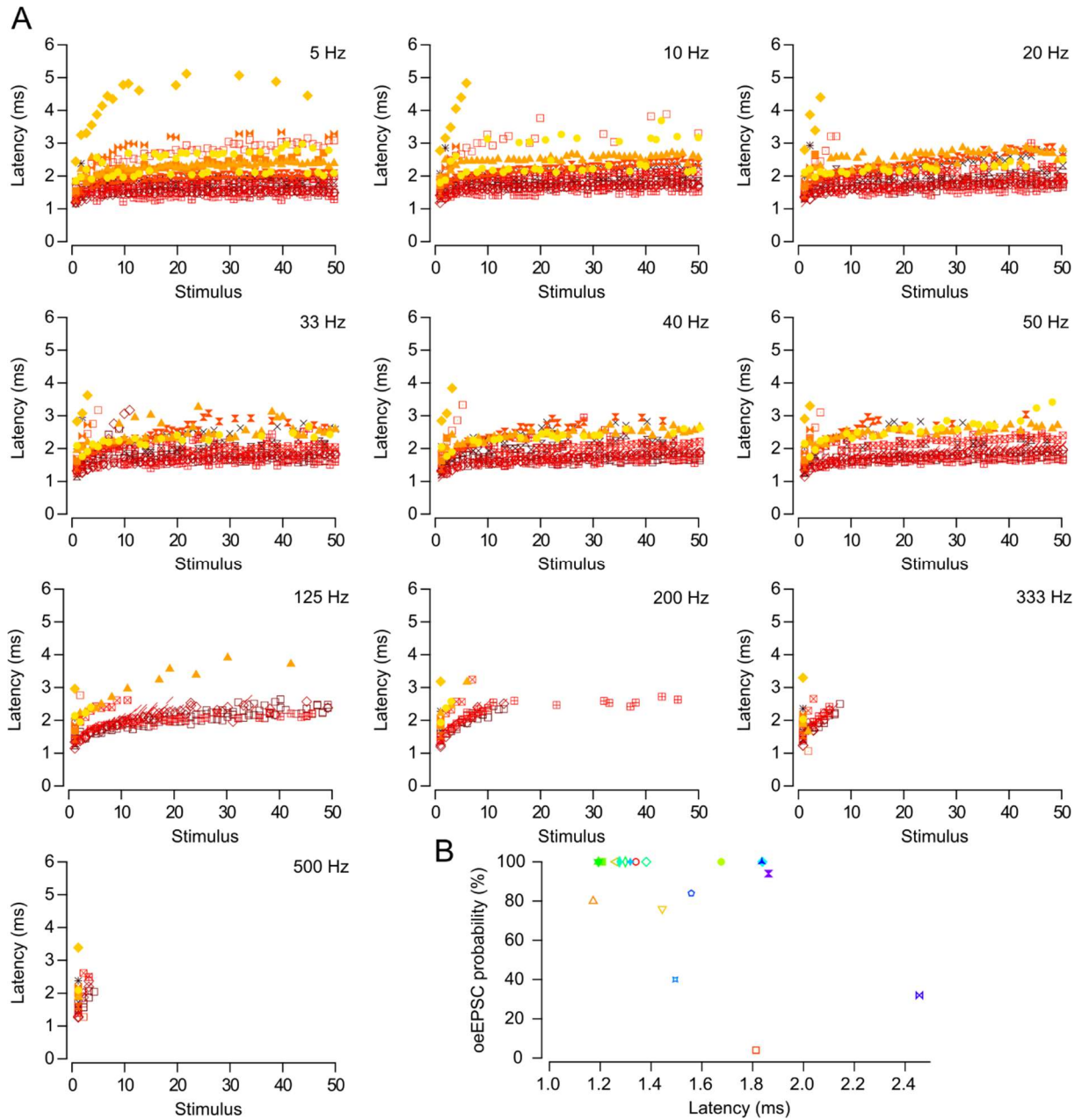


Appendix Figure S6. Characteristic firing patterns of AVCN neurons in response to 100 ms depolarizing current injection. Shown is tonic firing of a stellate cell (**A**) and phasic firing of a bushy cell (**D**) upon current injection. Measurements with an intracellular sodium channel blocker (1 mM QX-314) into the pipette solution immediately after establishing whole-cell configuration (**B-C**: stellate cells; **E-F**: bushy cells) and 30 min after establishing whole-cell configuration (**E₂**). Injection of hyperpolarizing currents results in the activation of hyperpolarization-activated (I_h) current (Cao et al. 2007; Rodrigues and Oertel 2006)

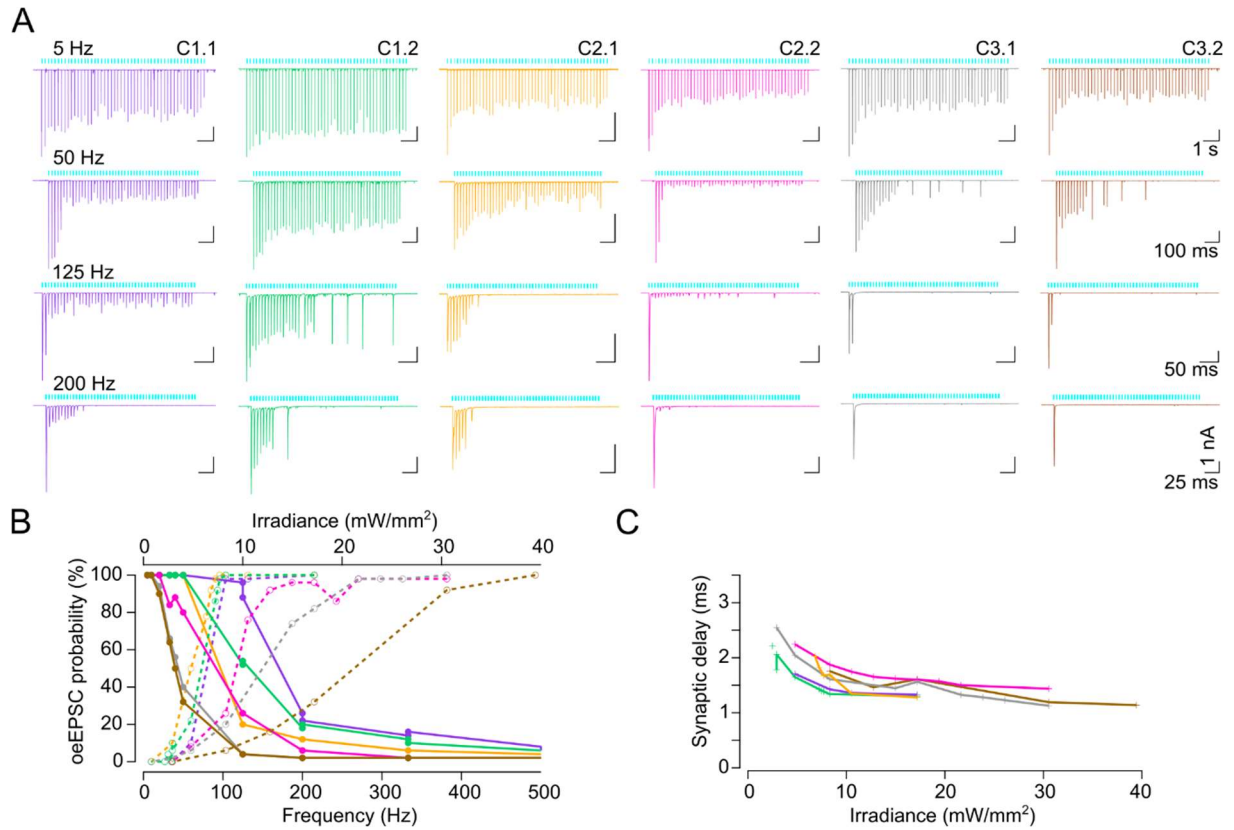


Appendix Figure S7. Classification of principal cells in the cochlear nucleus based on their spontaneous excitatory postsynaptic currents and cell morphology. **A-B**, Exemplary traces of spontaneous excitatory postsynaptic currents (sEPSCs) from a bushy cell (**A₁**) and from a stellate cell (**B₁**). Representative fluorescent image of a bushy cell (**A₂**) and a stellate cell (**B₂**) filled with Alexa 568. Bushy cells are characterized by short bushy-like dendrites branching next to the cell soma and are innervated by large endbulb of Held. Stellate cells have long dendrites (more than 2 usually) and receive bouton-like excitatory inputs from the auditory nerve terminals. **C**, Maximal z-projections of confocal image stacks of sagittal slices fixed and immunolabelled for EYFP (green) for f-ChR2 TC-EYFP localization, and context marker VGlut1 (yellow) as presynaptic vesicular transporter. Patched cells were filled with fixable Alexa 568 (blue). Scale bar = 20 μ m. **D**, Spontaneous and optically evoked EPSCs decay kinetics show two clear populations that we classified as bushy cells (oeEPSC decay lower than 0.5 ms) and stellate cells (oeEPSC decay higher than 0.5 ms), as previously reported in (Isaacson and Walmsley 1995; Cao and Oertel 2010). **E-I**, sEPSCs of previously classified cells as BC or SC (**D**) were analyzed in terms of amplitude (**E**), frequency of events (**F**), decay time (**G**), full width half maxima time

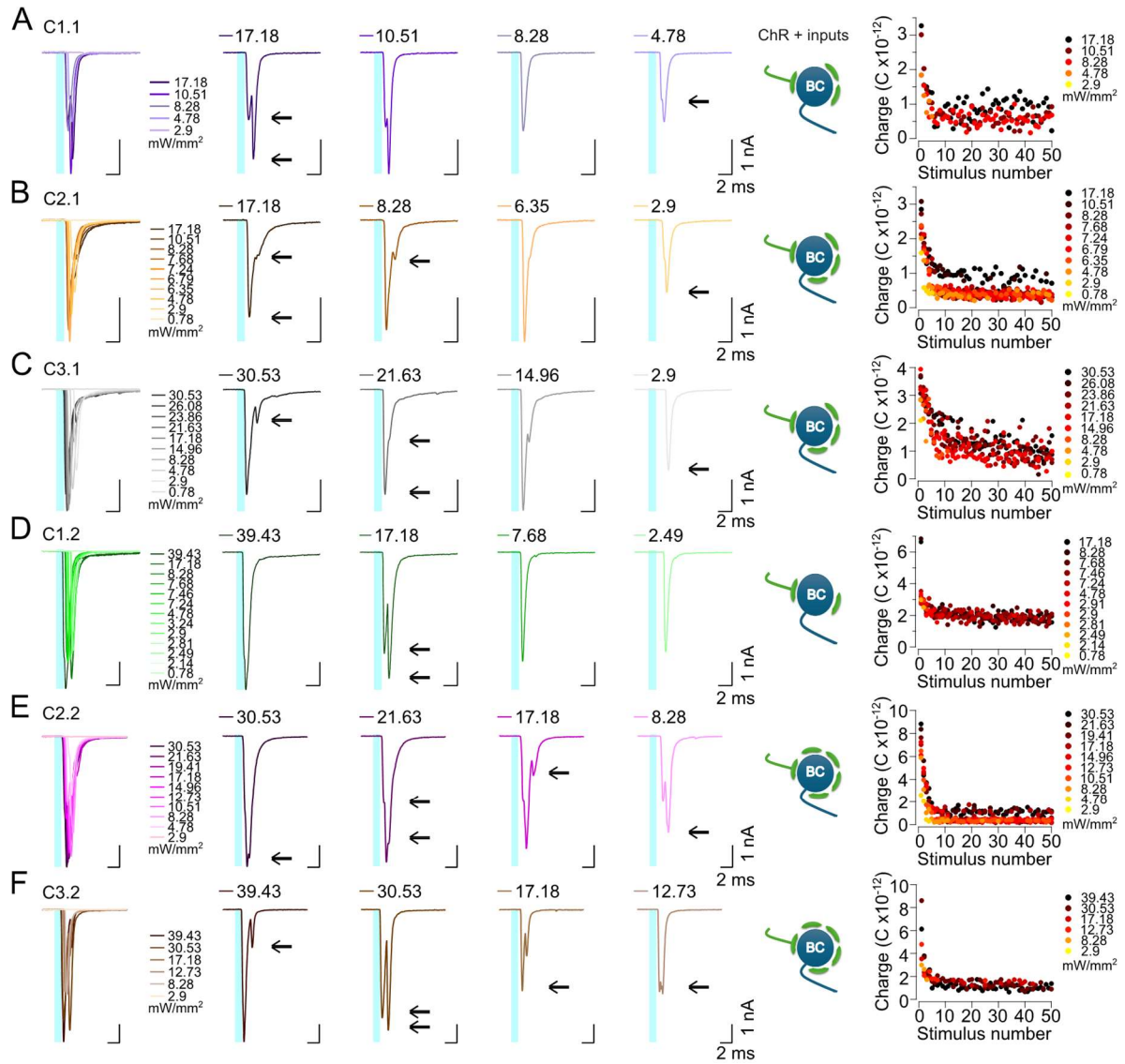
(H), and rise time taken from 10 % to 90 % of the event (I). Significant differences were detected between the two groups by Mann-Whitney test: **** $p < 0.0001$; ** $p = 0.0087$; ** $p = 0.0087$; $p = 0.2381$ (ns); *** $p = 0.0005$; ** $p = 0.0022$. Outliers were detected by ROUT test and excluded. Values obtained are in line with previous reports (Gardner et al. 1999; Lu et al. 2007; Cao and Oertel 2010). BC: number of animals = 9; number of cells = 16. SC number of animals = 6; number of cells = 8.



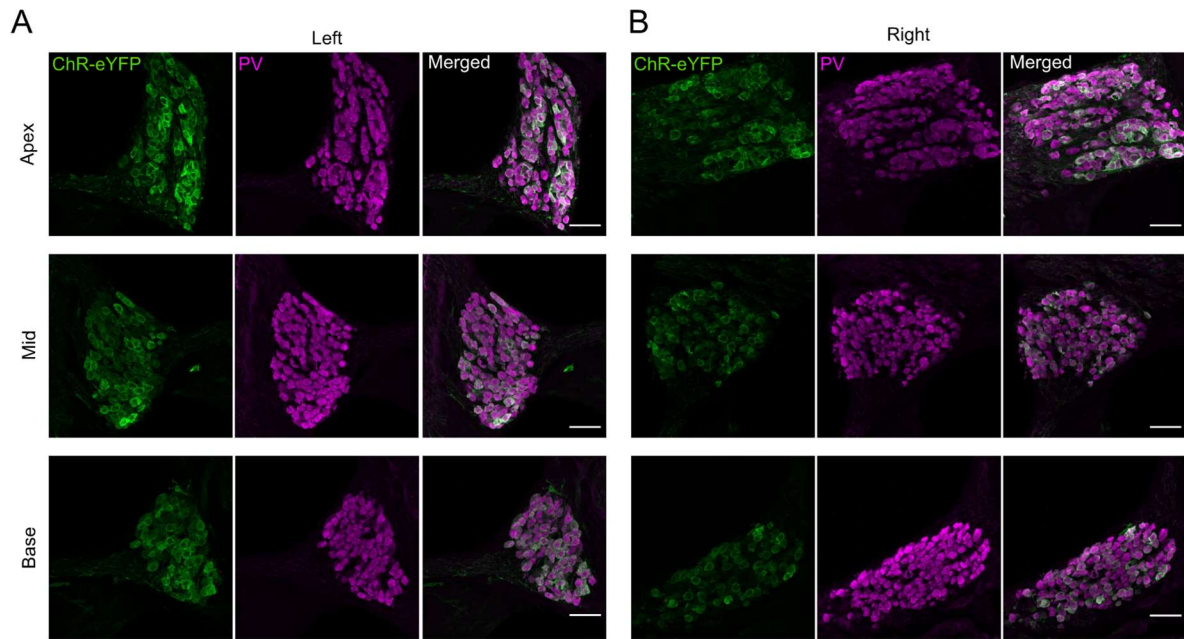
Appendix Figure S8. Spike latency increases over the train of light pulse stimulation. **A**, Quantification of latency by subtracting the time of stimulus onset from the time of the peak of the first oeEPSC ($n = 15$ BC; $N = 9$ slices from 9 mice). **B**, Cells with shorter latency show higher oeEPSC probabilities (frequency: 5 Hz, pulse length: 1 ms, $\lambda = 488$ nm, ~ 40 mW/mm²). Note that low oeEPSC probability was not always correlating to long latencies, as shown exemplarily for the endbulb synapses depicted in red square when compared to other synapses with similar latency (displayed by cyan rhombus, purple hourglass icon (X) and blue triangular star).



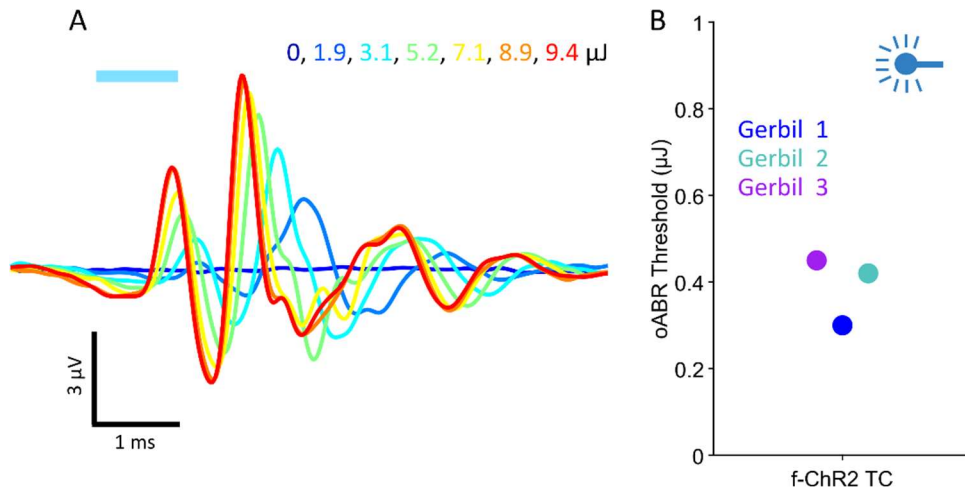
Appendix Figure S9. Bushy cells with greater light sensitivity can follow higher frequencies of stimulation. **A**, oeEPSCs in 6 different bushy cells, as indicated by the color code, receiving optogenetically evoked presynaptic inputs at different frequencies of light stimulation (pulse length: 1 ms, $\lambda = 488$ nm, ~ 40 mW/mm²). Cells are displayed from left to right from higher to lower success rate at different frequencies. **B**, Dependence of oeEPSC probability on irradiance (dashed line and empty circles) at a frequency of 10 Hz and a pulse length of 1 ms and dependence of oeEPSC probability on frequency (continuous line and filled circles) (pulse length: 1 ms, $\lambda = 488$ nm, ~ 40 mW/mm²). **C**, Quantification of synaptic delay dependent on frequency of stimulation (~ 40 mW/mm², 1 ms; continuous line). Cells plotted in **B-C** are the same.



Appendix Figure S10. oeEPSC measurements at different light intensities. Shown are exemplary traces recorded in 6 different bushy cells (A-F). Overlapped oeEPSCs showing the dependence of amplitude and synaptic delay on light intensity (panels on the left). For clarity, selected traces are shown separately (panels in the middle). Dependence of oeEPSC charge on stimulus number at different light intensities (panels on the right). The charge levels likely correlated to the number of synaptic inputs. Data taken from measurements shown in Appendix Fig. S9.

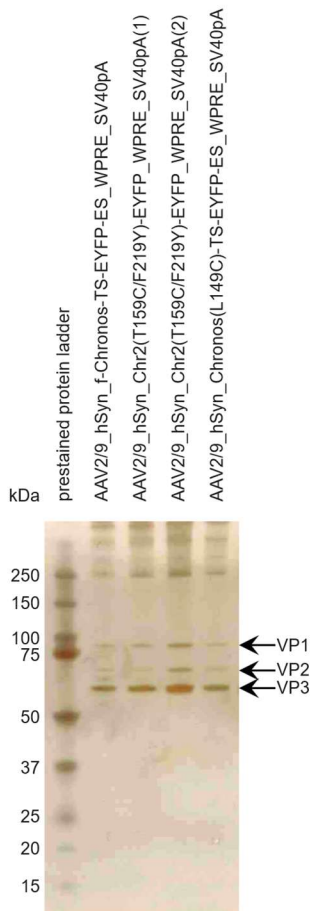
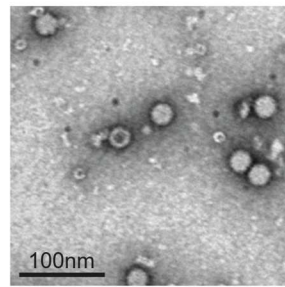


Appendix Figure S11. f-ChR2 TC expression in SGNs at P20 in postnatally injected mice. Projections of confocal mid-modiolar cryosections immunostained for GFP (green) showing ChR-EYFP expression, and parvalbumin (magenta) as context marker in SGNs in three cochlear regions. Scale bar: 50 μ m. Mice were injected unilaterally in the left side (**A**) but ChR expression was also detected in the right side (**B**) as previously reported in early postnatal injections (Bali et al. 2021).

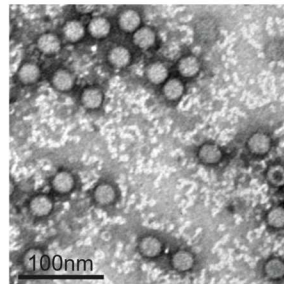


Appendix Figure S12. oABR shape and threshold for Mongolian gerbils expressing f-ChR2 TC in SGNs.

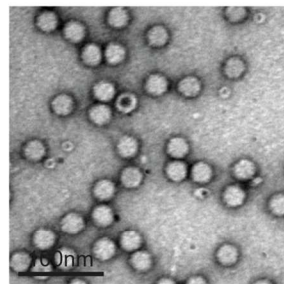
A, Exemplary oABRs evoked by a laser fiber placed into the round window of a Mongolian gerbil, emitting blue light at 488 nm. 1 ms pulses were presented at 17 Hz. Different colors resemble radiant energy in μJ . **B**, Radiant energy thresholds for occurrence of an oABR for stimulus conditions in **A**. $n = 3$ gerbils (indicated by different colors). One gerbil did not show oABR responses. Icon in B was created in BioRender. Albrecht, N. (2025) <https://BioRender.com/mz4pj4h>.

A**B**

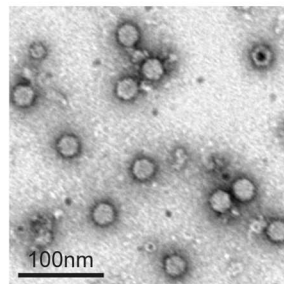
AAV2/9_hSyn_f-Chronos-TS-EYFP-ES_WPRE_SV40pA
92.8% filled capsids



AAV2/9_hSyn_ChR2(T159C/F219C)-EYFP_WPRE_SV40pA(1)
97.1% filled capsids



AAV2/9_hSyn_ChR2(T159C/F219C)-EYFP_WPRE_SV40pA(2)
95.1% filled capsids



AAV2/9_hSyn_Chronos(L149C)-TS-EYFP-ES_WPRE_SV40pA
90.1% filled capsids

Appendix Figure S13. Quality control of AAV vector preparations. **A**, SDS-PAGE of AAV vector preparations used in this study depicted by silver impregnation. For virus protein (VP) size comparison Bio-Rad Dual Protein plus was used. **B**, Visualization of AAV particles using negative stainings in electron microscopy of AAV preparations used in this study. Here, AAV particles, 25 nm in size, filled with ChR-constructs depicted by white shapes can be distinguished from empty AAV particles containing a dark, negatively stained core. Additional smaller structures originate from serum protein or host cell contaminations. Percentages of AAV particle filling is calculated based on manual count in 10 random images per of AAV vector.

Appendix Tables

	Input resistance (G Ω)	Mean capacitance (pF)	Mean resting membrane potential (mV)
f-Chronos (b)	0.38 \pm 0.27 (n = 15)	19.78 \pm 9.59 (n = 15)	-62.87 \pm 10.09 (n = 12)
Chronos LC (d)	0.22 \pm 0.11 (n = 13)	23.99 \pm 8.44 (n = 13)	-64.38 \pm 8.09 (n = 13)
f-ChR2 TC (g)	0.35 \pm 0.14 (n = 19)	21.38 \pm 12.14 (n = 19)	-66.87 \pm 9.94 (n = 18)
CatCh (h)	0.34 \pm 0.26 (n = 15)	23.75 \pm 9.75 (n = 15)	-64.4 \pm 7.62 (n = 15)

Appendix Table S1. Passive properties of hippocampal neurons transduced with blue light activated ChRs. The passive properties of patched neurons after AAV transduction were comparable among groups.

	Species	Mutation(s)	Effect	Reference
ChR2	<i>Chlamydomonas reinhardtii</i>			(Nagel et al. 2003)
CatCh	<i>Chlamydomonas reinhardtii</i>	ChR2 L132C	Increased photocurrent amplitude. Reduced desensitization. Slower closing kinetics. Increased relative calcium permeability.	(Kleinlogel et al. 2011)
ChR2 TC	<i>Chlamydomonas reinhardtii</i>	ChR2 T159C	Increased photocurrent amplitude. Slower closing kinetics.	(Berndt et al. 2011)
ChR2 ET/TC	<i>Chlamydomonas reinhardtii</i>	ChR2 E123T/T159C	Increased photocurrent amplitude with fast closing kinetics.	(Berndt et al. 2011)
f-ChR2	<i>Chlamydomonas reinhardtii</i>	ChR2 F219Y	Faster closing kinetics.	(Mager et al. 2018)
f-ChR2 TC	<i>Chlamydomonas reinhardtii</i>	ChR2 T159C/F219Y	Increased photocurrent amplitude with fast closing kinetics.	This study
Chronos	<i>Stigeoclonium helveticum</i>			(Klapoetke et al. 2014)
f-Chronos	<i>Stigeoclonium helveticum</i>	Chronos F236Y	Ultra fast closing kinetics.	This study
Chronos LC	<i>Stigeoclonium helveticum</i>	Chronos L149C	Reduced desensitization. Slower closing kinetics.	This study
f-Chronos LC	<i>Stigeoclonium helveticum</i>	Chronos L149C/F236Y	Reduced desensitization. Slower closing kinetics.	This study
Chrimson	<i>Chlamydomonas noctigama</i>			(Klapoetke et al. 2014)
f-Chrimson	<i>Chlamydomonas noctigama</i>	Y261F/S267M	Faster closing kinetics.	(Mager et al. 2018)
vf-Chrimson	<i>Chlamydomonas noctigama</i>	K176R/Y261F/S267M	Faster closing kinetics.	(Mager et al. 2018)

Appendix Table S2. Channelrhodopsin variants. The table shows the ChR optimization strategy by the combination of point mutations with the shown effects.

construct	template	sequence of forward primer	sequence of reverse primer
Chronos F236Y	humanized Chronos	5'- TCATGGCCTACGTCTACTTC TGCTCTTGGC -3'	5'- GCCAAGAGCAGAAGTAGA CGTAGGCCATGA -3'
Chronos L149C	humanized Chronos	5'- ACTTGCCCCGTGATCTGCA TCCACCTGAGCAAC -3'	5'- GTTGCTCAGGTGGATGCAG ATCACGGGGCAAGT -3'
Chronos L149C/F236Y	humanized Chronos F236Y	5'- ACTTGCCCCGTGATCTGCA TCCACCTGAGCAAC -3'	5'- GTTGCTCAGGTGGATGCAG ATCACGGGGCAAGT -3'
Chr2 T159C	humanized Chr2	5'-TCCGACAT CGGCTGTATCGTAT GGGG-3'	5'- CCCCAT ACGATACAGCCG ATGTCGGA-3'
Chr2 E123T/T159C	humanized Chr2 T159C	5'- CTGCGGTATGCCACTTGGC TGCTGAC -3'	5'- GTCAGCAGCCAAGTGGCAT ACCGCAG -3'
Chr2 F219Y/T159C	humanized Chr2 T159C	5'- TGGCTTGGCTGTACTTCGT GTCCTG -3'	5'- CAGGACACGAAGTACAGCC AAGCCA -3'

Appendix Table S3. List of primers used for Chr mutant generation.

Channelrhodopsin variants for high-rate optogenetic neurostimulation at low light intensities

Thomas Mager, Tobias Moser, Lennart Roos, Aida Garrido-Charles, Alexey Alekseev, Anupriya Thirumalai, Artur Mittring, Theodor Alvanos, Antoine Huet, Ernst Bamberg, Kathrin Kusch, Bettina Wolf, Niels Albrecht, Anna Vavakou, and Martina Bleyer

Corresponding authors: Thomas Mager (thomas.mager@med.uni-goettingen.de) , Tobias Moser (tmoser@gwdg.de)

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Revision Received:	14th Nov 25
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Editor: Jingyi Hou

Transaction Report:

(Note: With the exception of the correction of typographical or spelling errors that could be a source of ambiguity, letters and reports are not edited. Depending on transfer agreements, referee reports obtained elsewhere may or may not be included in this compilation. Referee reports are anonymous unless the Referee chooses to sign their reports.)

23rd Apr 2025

Dear Tobias,

Thank you again for submitting your work to EMBO Molecular Medicine. We have now received feedback from the two referees who agreed to evaluate your manuscript. As you will see in the reports below, the referees are overall supportive but have raised several concerns that will need to be thoroughly addressed in a major revision of the manuscript.

The referees' recommendations are clear, so I won't repeat the points listed below. In particular, the concerns regarding the long-term stability and safety of f-ChR2 TC need to be addressed. Additionally, we kindly ask that you revise your manuscript to place greater emphasis on the biomedical aspects, improve overall clarity, and make the content more accessible to the general audience of EMBO Mol Med. All other issues raised by the referees need to be addressed as well.

We would welcome the submission of a revised version within three months for further consideration. As you may already know, our editorial policy allows in principle a single round of major revision, and it is therefore essential to provide responses to the referees' comments that are as complete as possible.

EMBO Molecular Medicine has a "scooping protection" policy, whereby similar findings that are published by others during review or revision are not a criterion for rejection. Should you decide to submit a revised version, I do ask that you get in touch after three months if you have not completed it, to update us on the status.

Please also contact us as soon as possible if similar work is published elsewhere. If other work is published, we may not be able to extend the revision period beyond three months.

Please read below for important editorial formatting and consult our author's guidelines for proper formatting of your revised article for EMBO Molecular Medicine.

I look forward to receiving your revised manuscript.

Kind regards,
Jingyi

Jingyi Hou
Senior Editor
EMBO Molecular Medicine

When submitting your revised manuscript, please carefully review the instructions that follow below. We perform an initial quality control of all revised manuscripts before re-review; failure to include requested items will delay the evaluation of your revision.

We require:

- 1) A .docx formatted version of the manuscript text (including legends for main figures, EV figures and tables). Please make sure that the changes are highlighted to be clearly visible.
- 2) Individual production quality figure files as .eps, .tif, .jpg (one file per figure). For guidance, download the 'Figure Guide PDF': (<https://www.embopress.org/page/journal/17574684/authorguide#figureformat>).
- 3) A .docx formatted letter INCLUDING the reviewers' reports and your detailed point-by-point responses to their comments. As part of the EMBO Press transparent editorial process, the point-by-point response is part of the Review Process File (RPF), which will be published alongside your paper.
- 4) A complete author checklist, which you can download from our author guidelines (<https://www.embopress.org/page/journal/17574684/authorguide#submissionofrevisions>). Please insert information in the checklist that is also reflected in the manuscript. The completed author checklist will also be part of the RPF.

5) Please note that all corresponding authors are required to supply an ORCID ID for their name upon submission of a revised manuscript.

6) It is mandatory to include a 'Data Availability' section after the Materials and Methods. Before submitting your revision, primary datasets produced in this study need to be deposited in an appropriate public database, and the accession numbers and database listed under 'Data Availability'. Please remember to provide a reviewer password if the datasets are not yet public (see <https://www.embopress.org/page/journal/17574684/authorguide#dataavailability>).

In case you have no data that requires deposition in a public database, please state so in this section. Note that the Data Availability Section is restricted to new primary data that are part of this study.

7) For data quantification: please specify the name of the statistical test used to generate error bars and P values, the number (n) of independent experiments (specify technical or biological replicates) underlying each data point and the test used to calculate p-values in each figure legend. The figure legends should contain a basic description of n, P and the test applied. Graphs must include a description of the bars and the error bars (s.d., s.e.m.). See also 'Figure Legend' guidelines: <https://www.embopress.org/page/journal/17574684/authorguide#figureformat>

8) At EMBO Press we ask authors to provide source data for the main manuscript figures. You will receive a separate email with instructions for providing source data with your revised manuscript, including how to upload and organize the files.

9) Our journal encourages inclusion of *data citations in the reference list* to directly cite datasets that were re-used and obtained from public databases. Data citations in the article text are distinct from normal bibliographical citations and should directly link to the database records from which the data can be accessed. In the main text, data citations are formatted as follows: "Data ref: Smith et al, 2001" or "Data ref: NCBI Sequence Read Archive PRJNA342805, 2017". In the Reference list, data citations must be labeled with "[DATASET]". A data reference must provide the database name, accession number/identifiers and a resolvable link to the landing page from which the data can be accessed at the end of the reference. Further instructions are available at .

10) We replaced Supplementary Information with Expanded View (EV) Figures and Tables that are collapsible/expandable online. A maximum of 5 EV Figures can be typeset. EV Figures should be cited as 'Figure EV1, Figure EV2' etc... in the text and their respective legends should be included in the main text after the legends of regular figures.

- For the figures that you do NOT wish to display as Expanded View figures, they should be bundled together with their legends in a single PDF file called *Appendix*, which should start with a short Table of Content. Appendix figures should be referred to in the main text as: "Appendix Figure S1, Appendix Figure S2" etc.

- Additional Tables/Datasets should be labeled and referred to as Table EV1, Dataset EV1, etc. Legends have to be provided in a separate tab in case of .xls files. Alternatively, the legend can be supplied as a separate text file (README) and zipped together with the Table/Dataset file.

See detailed instructions here:

.

11) The paper explained: EMBO Molecular Medicine articles are accompanied by a summary of the articles to emphasize the major findings in the paper and their medical implications for the non-specialist reader. Please provide a draft summary of your article highlighting

- the medical issue you are addressing,
- the results obtained and
- their clinical impact.

This may be edited to ensure that readers understand the significance and context of the research. Please refer to any of our published articles for an example.

12) Author contributions: You will be asked to provide CRediT (Contributor Role Taxonomy) terms in the submission system. These replace a narrative author contribution section in the manuscript.

13) A Conflict of Interest statement should be provided in the main text.

14) Every published paper now includes a 'Synopsis' to further enhance discoverability. Synopses are displayed on the journal

webpage and are freely accessible to all readers. They include a short stand first (maximum of 300 characters, including space) as well as 2-5 one-sentences bullet points that summarizes the paper. Please write the bullet points to summarize the key NEW findings. They should be designed to be complementary to the abstract - i.e. not repeat the same text. We encourage inclusion of key acronyms and quantitative information (maximum of 30 words / bullet point). Please use the passive voice. Please attach these in a separate file or send them by email, we will incorporate them accordingly.

Please also suggest a visual abstract to illustrate your article as a PNG file 550 px wide x 300-600 px high.

15) All Materials and Methods need to be described in the main text using our 'Structured Methods' format. According to this format, the Methods section includes a Reagents and Tools Table (listing key reagents, experimental models, software and relevant equipment and including their sources and relevant identifiers) followed by a Methods and Protocols section describing the methods, ideally using a step-by-step protocol format. The aim is to facilitate adoption of the methodologies across labs.

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When submitting your revised manuscript, please DO NOT include the Reagents and Tools Table in the Methods section of the manuscript but upload it as a separate file choosing the file type "Reagent Table".

EMBO Molecular Medicine has a "scooping protection" policy, whereby similar findings that are published by others during review or revision are not a criterion for rejection. Should you decide to submit a revised version, I do ask that you get in touch after three months if you have not completed it, to update us on the status.

***** Reviewer's comments *****

Referee #1 (Comments on Novelty/Model System for Author):

Dear Dr Hou, the paper and the data are great and very convincing. However, it is written for a more biophysical audience, so I would recommend streamlining it and focusing on the biomedical aspects. Please see my comments to the authors for details.

Referee #1 (Remarks for Author):

The paper "Channelrhodopsin variants for high-rate optogenetic neurostimulation at low light intensities" by Roos et al. presents a sophisticated biophysical characterisation of an advanced ChR2 derivative called f-ChR2 TC. It is being tested in various cell lines and model systems with a view to its application in optogenetic hearing restoration. The depth of data provided is impressive, but also overwhelming. It might be helpful to reduce the data and figures provided and tailor them to the main messages/findings of the manuscript. The provision of expanded figures and, on top of this, appendix figures is an overload and reduces clarity and readability. There is also a huge imbalance towards the biophysical aspects and more biomedical impact would be desirable for the readership of this journal. Otherwise, the manuscript is very well written. Scientifically, the question remains as to why the authors now focus on a blue-shifted optogenetic tool for hearing restoration, having previously presented red-shifted versions. Blue light can cause phototoxicity. The manuscript would benefit greatly if the authors provided a comprehensive rationale for why the described enhanced f-ChR2-TC is better for biomedical applications than, for example, f-ChRimson. In this regard, it would be great if the authors could discuss or provide data on whether the enhanced expression of f-ChR2 TC and its sensitivity to blue light does not cause photo- or cytotoxicity over a meaningful period of time. In particular, both the in vivo and in vitro experiments were performed a few days after AAV delivery. It is unclear whether f-ChR2 TC is stable, functional and safe over time.

Minor points:

Line 67: "spectral coding" is not introduced - what is meant by this?

Line 173 (Figure 1F - right panel, x-axis label): is f-ChR2 TC meant?

Line 210: What is "ultrafast f-Chronos"?

Line 214: It would be great to show representative fluorescence images in the main figures.

Line 346: What is known about the cytotoxicity of optogenes after AAV overexpression?

Line 351: GFP or EYFP expression?

Line 418: Could you correlate optogene expression levels with better performance in bushy cells or could this be a reason for the differences observed?

Line 463: Which 'ChR' versions are meant?

Referee #2 (Remarks for Author):

General comments :

This study strength lies in the engineering and comprehensive characterization of a novel blue-light-sensitive channelrhodopsin variant, f-ChR2 TC, which uniquely combines fast closing kinetics with good plasma membrane expression and low light requirements, enabling efficient and safe high-rate neurostimulation. Through in vitro and in vivo experiments in the auditory pathway, the study demonstrates that f-ChR2 TC outperforms other blue-light activated ChRs and compares favorably to advanced red-light options, highlighting its potential for basic research and preclinical optogenetic therapies, particularly for hearing restoration.

The study is well designed, from careful characterisation of photocurrent kinetics in non-neuronal cells to quantification of the optogenetic auditory brainstem response. Interpretation of the results are deepened by the measurement made at these two levels, and the role of the introduced mutations at the molecular level, is elegantly outlined by the functional results obtained at the level of the full organism. The study also delves into the synaptic response driven by optogenetic stimulation, an important step into the comprehension of critical parameters for transmission and integration of a restored percept.

Specific comments:

1-

The large number of variants makes it difficult to clearly understand the relationships between the different constructs. Including a summary figure, such as a table or a phylogenetic-style tree, could help illustrate and organize the various optimization strategies explored in this study. Additionally, the use of "ChRs" as a generic term for channelrhodopsins and "ChR2" for the specific Channelrhodopsin-2 can be confusing—especially since some variants are derived from ChR2 mutations, while others originate from different microorganisms (e.g., *S. helveticum* for Chronos, *C. reinhardtii* for ChR2, or *C. noctigama* for Chrimson).

2-

Line 217 "... the success rate was low for f-Chronos expressing neurons reaching a spike probability of 80 to 100 % in only 3 out of 14 neurons (Fig. 2)."

The statement should be clarified, as a spike probability of 80~100% in only 3 out of 14 shows great limitation of the construct. It should be clear what type of stim was done here (it is clarified later, but should be stated here). It also appears misleading to display f-Chronos in Fig 2A if there is such a decreased efficacy.

Line 219 "i) suboptimal plasma membrane targeted expression, which is reflected in the comparatively low stationary photocurrent density values measured at saturating light intensities (Fig. 2)"

this refers, I presume, to the result in Fig 2B, then I'd recommend to display in Fig 2A the photocurrent quantified in Fig 2B rather than the spike trains (although, showing both is fine). As photocurrents are displayed in Fig 1E, I wonder if the recordings show similar kinetics in neurons and NG cells.

3-

Line 222 "ii) the limited charge transfer evoked by the short (1 ms) light pulses given the ultrafast channel closing kinetics (Fig. EV1, Fig. 1)."

For clarity, I suggest to separate the figure references "... by the short light pulses (Fig EV1) ... closing kinetics (Fig 1B)". It would also be easier to understand the sentence meaning if the charge transfer was quantified in Fig EV1. Alternatively, rewrite the sentence to clarify that charge transfer is illustrated by the normalized stationary current.

4-

Line 270 "oABRs mediated by f-ChR2 TC compared favorably to published results obtained with ultrafast ChR variants: Chronos and f-Chrimson (Fig. 3 E-G, replotted from Keppeler et al. 2018 and Zerche et al. 2023)."

The statement is a bit unfair, and leads to the notion that f-ChR2 TC outperforms the other constructs. While true for Chronos, f-Chrimson seems on par with f-ChR2 TC here. Adjust phrasing accordingly.

5- Major remark

My next point concerns the reported reduction in SGN density following injection. The authors should comment on the magnitude of this reduction, especially since the effect appears to vary between constructs. This variability weakens the argument that the observed SGN loss is solely due to injection pressure. To strengthen this hypothesis, it would be important to include a control group in which mice are injected with a saline solution or better yet an AAV with YFP alone. The authors could also discuss ways to adapt the injection protocol to reduce this unwanted effect.

Additionally, in Figure EV3, the rationale behind the left/right panel separation is unclear. If the right side corresponds to the injected cochlea and the left to the non-injected one, it is important to clarify what was injected in the control group, and why YFP-positive SGNs are visible on the right side in panel EV3-D.

It is also important for a potential therapeutic development to assess if the reduction is due to the pressure injection, a toxic overexpression of the transgene, or an immune response. As the reduction in the number of cells is important, it should be possible to localise the dead cells, or identify the lost volume?

It might be worth to relate the observation with what seems like a reduced PV expression in the f-ChR2 TC.

6-

I acknowledge the challenges involved in analyzing the subcellular localization of the optogene from cryosections, and the authors have made a commendable effort through careful quantification and appropriately cautious terminology in describing their findings. While the methods section outlines the procedure for the line analysis, I was unable to find a clear explanation of how the ratio presented in Figure 4-C was calculated. Furthermore, it would have been beneficial to perform membrane surface staining in at least a small subset of tissue to validate the quantification.

7-

line392 " injected (P5-P9) C57BL/6 wild-type mice after a standardized expression period of ~ 2 weeks"
it is weird to have a standard being approximatively 2 weeks, please rephrase or clarify.

8-

line 404 " Thus, by reducing the light irradiance, fewer inputs are recruited, which is reflected by smaller step sizes in the oeEPSC amplitude eventually arriving at monosynaptic input (single endbulb synapse)."
From my understanding, this method is not used here. If it is indeed the case, please consider rephrasing, as it may confuse the reader.

Line 409" Where positive," what is positive here? Typo?

9- Major point

To measure the maximal frequency driving synaptic inputs the authors use variable frequency and saturating light (@ 40mW/mm²). While it help normalize the variability in light sensitivity, isn't it counterproductive in term of maximal frequency responses? I would assume that saturating light might increase the oeEPSC probability reduction observed in the train of stimuli.

10- Major point

Line 416 "Quantification of oeEPSC probability versus irradiance and frequency of stimulation shows that bushy cells that are more light-sensitive in terms of higher oeEPSC probability and shorter synaptic delay also follow higher frequencies of stimulation (Fig. 5 E-G)."

This aspect is a crucial component of the synaptic findings and deserves a clearer representation. While the current data allow the reader to infer the conclusion by comparing individual cells, the current layout is suboptimal. Notably, the line colors in panels E/F and G are inconsistent. The authors should consider a more effective way to illustrate this relationship within a single panel. One possible solution would be to extract, for each cell, a single irradiance or frequency value corresponding to a 50% oEPSC probability, and plot these values against one another.

11-

Line 421 "Given the fast-closing kinetics of f-ChR2 TC at physiological temperature ($\tau_{off} \sim 5$ ms, Table EV2), we reason that limited expression and photocurrent desensitization of f-ChR2 TC as well as a potential depolarization block hinder endbulb synapses to follow stimulation frequencies of more than 100 Hz."

The rationale behind this hypothesis is not fully elaborated. While the limited expression aspect is relatively straightforward, the desensitization observed with f-ChR2 TC should be more thoroughly discussed. Additionally, the role of depolarization block remains unclear. Since the authors refer to results from cells receiving "optogenetic inputs" rather than directly expressing the opsin, does this imply that the depolarization block occurs at the presynaptic terminals? Typically, a depolarization block would be expected in cells with higher-than-average opsin expression, which might be reflected by a shorter synaptic delay. Is such a correlation observed in this study?

12-

Fig EV4: does the panels D and F relate to stellate cells and C and E to bushy cells? If so, it is not clearly indicated in the legend.

13-

Line 437 "However, low oeEPSC probability was not always predicted by long latency, as shown exemplarily for the endbulb synapses depicted in Appendix Fig. S5 B-C, and compared to other synapses with similar oeEPSC probability."

I might misunderstand the point, but the example referred to in the legend of Fig S5 B-C seems to have a low probability AND a long latency (green dots). It may be the phrasing that is unclear as the legend indicate:

"Note that low oeEPSC probability was not always predicted by long latency, as shown exemplarily for the endbulb synapses depicted in green dots, and compared to other synapses with similar oeEPSC probability (displayed by purple hourglass icon () and red square). »

The current phrasing suggests that it is specifically for the green dots that the oeEPSC probability fails to predict long latency.

14-

Fig S6 C legend: « Quantification of synaptic delay dependent on frequency of stimulation (~ 40 mW/mm², 1 ms; continuous line) and dependent on irradiance (10 Hz, 1 ms; dashed line). «

does not fit with the actual figure, i only have continuous line, and 1 X axis, rendering issue?

Discussion:

It would be valuable to further elaborate on strategies that could help minimize desensitization.

Since neurons can differ in their intrinsic properties-affecting, for instance, the maximal sustainable firing rate-the authors should also expand on how the results observed in hippocampal neurons relate to the later findings in SGNs or oABR recordings.

Line 551 : "Limitations explaining failure or sustained high rate..." should be "...OF sustained high rate"

Although, that sentence remain unclear even with the proposed modification

Methods:

Line 656: Toff is determined as a monoexponential fit of the current, but there is no quantification of the goodness of the fit?

Could a double exponential lead to better fit?

It is then mentioned line 796 that 10/15 cells have a biexponential fit, without much explanation.

Line 663: the authors mention peak recovery of ChRs, where is that data presented?

Line 858 : the sentence is uncomplete (no verb).

RESPONSE TO REVIEWER REPORT(S):

Referee: 1

Referee #1 (Comments on Novelty/Model System for Author):

Dear Dr Hou, the paper and the data are great and very convincing. However, it is written for a more biophysical audience, so I would recommend streamlining it and focusing on the biomedical aspects. Please see my comments to the authors for details.

We would like to thank the reviewer for the thorough review of our manuscript. We understand the Referee's assessment of the scope of our study. Thus, we have streamlined the manuscript towards its biomedical aspects and revised our manuscript substantially by including additional experiments in a new main figure highlighting translational transferability of our newly engineered channelrhodopsin. Here, we employed f-ChR2 TC in Mongolian gerbils, which serve as a translational rodent model in hearing research as they offer a larger cochlea and a more human-like low frequency hearing than mice and rats. These experiments demonstrated low light requirements for activating the auditory pathway by f-ChR2 TC mediated optogenetic stimulation of the auditory nerve. We then used LED-based multichannel optical cochlear implants and recorded the ensuing multi-unit activity in the inferior colliculus of these gerbils by multielectrode array recordings and confirmed sub- μ J light requirements and good temporal fidelity of coding.

Referee #1 (Remarks for Author):

The paper "Channelrhodopsin variants for high-rate optogenetic neurostimulation at low light intensities" by Roos et al. presents a sophisticated biophysical characterisation of an advanced ChR2 derivative called f-ChR2 TC. It is being tested in various cell lines and model systems with a view to its application in optogenetic hearing restoration. The depth of data provided is impressive, but also overwhelming. It might be helpful to reduce the data and figures provided and tailor them to the main messages/findings of the manuscript. The provision of expanded figures and, on top of this, appendix figures is an overload and reduces clarity and readability. There is also a huge imbalance towards the biophysical aspects and more biomedical impact would be desirable for the readership of this journal. Otherwise, the manuscript is very well written.

We would like to thank the reviewer for the appreciation of our work and the advice to further improve our manuscript. In response to the reviewer's comment, we have completely overhauled the manuscript taking the suggestions of the reviewer into account. In particular we have better aligned the balance of the manuscript toward the biomedical application in the auditory system, including in the Mongolian gerbil.

Scientifically, the question remains as to why the authors now focus on a blue-shifted optogenetic tool for hearing restoration, having previously presented red-shifted versions. Blue light can cause phototoxicity. The manuscript would benefit greatly if the authors provided a comprehensive rationale for why the described enhanced f-

ChR2-TC is better for biomedical applications than, for example, f-ChRimson. In this regard, it would be great if the authors could discuss or provide data on whether the enhanced expression of f-ChR2 TC and its sensitivity to blue light does not cause photo- or cytotoxicity over a meaningful period of time.

In particular, both the in vivo and in vitro experiments were performed a few days after AAV delivery. It is unclear whether f-ChR2 TC is stable, functional and safe over time.

We fully understand the reviewers concerns and appreciate the comment. While previous studies indeed described the benefits of red-shifted f-Chrimson, we now focused on a blue-shifted optogenetic tool for two reasons.

First, we aimed to engineer a well-balanced channelrhodopsin to combine robust current densities while maintaining fast closing kinetics to allow for optogenetic stimulation of the cochlea at low pulse energy thresholds and at near physiological temporal fidelity. Thus, we now have laid the basis for further preclinical assessment of optogenetic hearing restoration and its efficacy in gerbils and non-human primates employing μ LED oCIs and its comparisons to eCIs. To date, this is not possible using red-shifted ChRs as oCI prototypes emitting red-light are still in development.

Second, a blue-light-shifted ChR with similar properties to f-Chrimson, such as f-ChR2 TC, will ultimately help to expand the current dynamic range limitation of oCIs by facilitating SGN subtype specific activation using two non-overlapping spectra of visible light when combining blue and red-light activated ChRs for dual color stimulation.

However, we agree with the reviewer that cyto- and phototoxicity upon expression of blue-light activated ChRs needs to be carefully evaluated and tested over a meaningful period of time and is highly warranted for any clinical translation. In this regard the lab has already published previous work on stable, longevous expression of ChRs (f-Chrimson) in mice that we discussed in our manuscript and now highlighted to a greater extend (Bali et al. 2022). Further investigation on potential phototoxic effects may be out of the scope of this manuscript. Nevertheless, development of ChRs such as f-ChR2 TC, with low energy thresholds for activation will ultimately contribute in reducing phototoxicity when using blue light.

Minor points:

Line 67: "spectral coding" is not introduced - what is meant by this?

We thank the reviewer for the observation and replaced "spectral coding" by "sound frequency coding" for clarification.

Line 173 (Figure 1F - right panel, x-axis label): is f-ChR2 TC meant?

We thank the reviewer for the question and apologize for the confusion. The right panel of Fig. 1F, highlights the comparison of the current densities of ChR2 TC and CatCh, which are two other ChR2 variants. The current densities of f-ChR2 TC are shown in the middle panel of Fig. 1F. Further, the panels in Figure 1B and 1F are designed to compare channel closing kinetics (B) and current densities (F) for: very fast channelrhodopsins (left panel) with closing kinetics faster than 10 ms; medium-

fast channelrhodopsins (middle panel) with closing kinetics around 10 ms, and slow channelrhodopsins (right) with closing kinetics slower than 10 ms.

Line 210: What is "ultrafast f-Chronos"?

We apologize for the confusion cause and have now clarified the main text: f-Chronos is the ultrafast ChR variant Chronos F236Y (see the revised sections below):

Lines 107-108

"Toward the first strategy, we generated Chronos F236Y (fast Chronos: f-Chronos), which, to our knowledge, is the fastest closing ChR to date."

Lines 121-125

"Indeed, as shown by whole-cell patch clamp experiments in transiently transfected NG cells, the ultrafast ChR Chronos F236Y (f-Chronos, $\tau_{\text{off}} = 1.7 \pm 0.1$ ms, $n = 4$ at room temperature (RT), $\tau_{\text{off}} = 0.8 \pm 0.1$ ms, $n = 4$ at 33 °C) was considerably faster than Chronos ($\tau_{\text{off}} = 3.1 \pm 0.5$ ms, $n = 6$ at RT (** $p = 0.0009$, unpaired t-test with Welch's correction), $\tau_{\text{off}} = 1.9 \pm 0.5$ ms, $n = 6$ at 33 °C (** $p = 0.021$, unpaired t-test with Welch's correction); **Fig. 1 A-B; Table EV1-2**)"

Line 214: It would be great to show representative fluorescence images in the main figures.

We included the respective images.

Line 346: What is known about the cytotoxicity of optogenes after AAV overexpression?

Changes in neuronal biology up to toxicity and neural loss e.g. due to proteostatic stress following AAV-mediated overexpression of proteins present a risk also in optogenetic therapies (Kleinlogel *et al*, 2020; Miyashita *et al*, 2013; Stone *et al*, 2025). In response to this comment, we have performed additional histological analysis of AAV-injected left cochleae and compared them to the non-injected right cochlea, which also contained optogenetically modified SGNs. With the AAV doses used we found signs of neural pathology (vacuolization and neural loss) yet no evidence for innate or adaptive immune response (**Appendix Fig. S5**). We have now included these results in the manuscript and also discussed potential mechanisms of cytotoxicity.

Attached respective section of the manuscript:

Lines 351-357

"SGN loss might be an aftermath of the intracochlear pressure injection at young age. In addition, we cannot rule out potential cytotoxicity due to protein overload based proteostatic stress (Stone *et al*. 2025) or immune response following AAV-mediated overexpression. Further, histological and immunohistochemical investigation of cochlear paraffin sections can be found in **Appendix Fig. S5**. In brief, we observed neuropathological changes in SGNs but did not find evidence for a prevailing adaptive or innate immune response."

.”

Line 351: GFP or EYFP expression?

We thank the reviewer for noticing the inconsistency. Our transgene is composed of the respective ChR and if needed enhancing sequences as well as the reporter EYFP. Thus, it is EYFP expression. GFP antibodies were used for immunolabelling of EYFP in immunohistochemistry.

Line 418: Could you correlate optogene expression levels with better performance in bushy cells or could this be a reason for the differences observed?

We agree to the reviewer that ChR expression is a likely reason for the differences observed. As described in the main text, we observed a correlation between synaptic delay and the light sensitivity as well as the maximal frequency. Spike latency and therefore synaptic delay depends on photocurrent size, which in turn depends on the expression. We modified the following sentence, which is following the description of the correlation:

Lines 433-437

“Given the fast-closing kinetics of f-ChR2 TC at physiological temperature ($T_{\text{off}} \sim 5$ ms, **Table EV2**), we reason that limited expression and photocurrent desensitization of f-ChR2 TC as well as a potential depolarization block led to the observed heterogeneity of endbulb synaptic transmission to bushy cells and hindered most endbulb synapses to follow stimulation frequencies of more than 100 Hz.”

Line 463: Which 'ChR' versions are meant?

We thank the reviewer and corrected the legend of Figure 5 in order to specify the ChR version.

Referee #2 (Remarks for Author):

General comments :

This study strength lies in the engineering and comprehensive characterization of a novel blue-light-sensitive channelrhodopsin variant, f-ChR2 TC, which uniquely combines fast closing kinetics with good plasma membrane expression and low light requirements, enabling efficient and safe high-rate neurostimulation. Through in vitro and in vivo experiments in the auditory pathway, the study demonstrates that f-ChR2 TC outperforms other blue-light activated ChRs and compares favorably to advanced red-light options, highlighting its potential for basic research and preclinical optogenetic therapies, particularly for hearing restoration.

The study is well designed, from careful characterisation of photocurrent kinetics in non-neuronal cells to quantification of the optogenetic auditory brainstem response. Interpretation of the results are deepened by the measurement made at these two levels, and the role of the introduced mutations at the molecular level, is elegantly outlined by the functional results obtained at the level of the full organism. The study also delves into the synaptic response driven by optogenetic stimulation, an important step into the comprehension of critical parameters for transmission and integration of a restored percept.

We would like to thank the reviewer for the appreciation of our work and the advice to further improve our manuscript. We have completely overhauled the manuscript taking the reviewer's recommendations into account.

Specific comments:

1-

The large number of variants makes it difficult to clearly understand the relationships between the different constructs. Including a summary figure, such as a table or a phylogenetic-style tree, could help illustrate and organize the various optimization strategies explored in this study. Additionally, the use of "ChRs" as a generic term for channelrhodopsins and "ChR2" for the specific Channelrhodopsin-2 can be confusing—especially since some variants are derived from ChR2 mutations, while others originate from different microorganisms (e.g., *S. helveticum* for Chronos, *C. reinhardtii* for ChR2, or *C. noctigama* for Chrimson).

We thank the reviewer for the comment and completely understand that this can cause confusion. Following the reviewer's comment, we included "**Appendix Table S2**" to summarize all the variants and optimization strategies used in this study.

2-

Line 217 "... the success rate was low for f-Chronos expressing neurons reaching a spike probability of 80 to 100 % in only 3 out of 14 neurons (Fig. 2)."

The statement should be clarified, as a spike probability of 80~100% in only 3 out of 14 shows great limitation of the construct. It should be clear what type of stim was done here (it is clarified later, but should be stated here). It also appears misleading to display f-Chronos in the Fig2A if there is such a decreased efficacy.

Line 219 "i) suboptimal plasma membrane targeted expression, which is reflected in the comparatively low stationary photocurrent density values measured at saturating light intensities (Fig. 2)" this refer, I presume, to the result in Fig2B, then I'd recommend to display in Fig2A the photocurrent quantified in Fig2B rather than the spike trains (although, showing both is fine). As photocurrents are displayed in fig 1E, I wonder if the recordings show similar kinetics in neurons and NG cells.

We thank the reviewer for pointing this out and adapted the manuscript accordingly. We exchanged the f-Chronos trace in Fig. 2 A (now **Fig. 2 E**) for a more representative recording showing a reduced spike probability upon photostimulation by a train of 50 light pulses with a pulse length of 1ms. Moreover, we included **Fig. 2 B** showing representative photocurrent recordings (quantified in **Fig. 2 C**). The comparison of the channel closing kinetics in hippocampal neurons (**Fig. 2 D**) and NG cells (**Fig. 1 B**), which is provided in the new **Appendix Figure S2**, shows that the closing kinetics are similar in the different cell types.

Lines 215-225

"The photocurrent densities and τ_{off} values determined in hippocampal neurons were similar to the values obtained in NG cells (**Fig. 1, Fig. 2B, Appendix Fig. S3, Appendix Table S1**). Whereas Chronos LC, f-ChR2 TC and CatCh enabled reliable neuronal photostimulation by the short (1 ms) light pulses, spike probability for f-Chronos expressing neurons was low, with 20% of the neurons (3 out of 14) showing a spike probability higher than 80% (**Fig. 2C**). The low success rate in f-Chronos expressing neurons likely results from i) suboptimal plasma membrane targeted expression, which is reflected in the comparatively low stationary photocurrent density values measured at saturating light intensities (**Fig. 2D-E**) and ii) the limited charge transfer evoked by the short (1 ms) light pulses (**Fig. EV1D**) given the ultrafast channel closing kinetics (**Fig. 1B**)."

3-

Line 222 " ii) the limited charge transfer evoked by the short (1 ms) light pulses given the ultrafast channel closing kinetics (Fig. EV1, Fig. 1). "

For clarity, I suggest to separate the figure references "... by the short light pulsses (Fig EV1) ... closing kinetics (fig 1B)". It would also be easier to understand the sentence meaning if the charge transfer was quantified in fig EV1. Alternatively rewrite the sentence to clarify that charge transfer is illustrated by the normalized stationary current.

We thank the reviewer for the comment, which helped to improve clarity. We referred to the following relation. In transient currents evoked by brief light pulses that are shorter than ChR photocycle time, which is approximated by τ_{off} , the transferred charge (Q) is proportional to the current decay kinetics (ChR on-kinetics is usually beyond the time resolution of whole-cell patch-clamp experiments). For clarity, we added **Fig. EV 1 D**.

4-

Line 270 "oABRs mediated by f-ChR2 TC compared favorably to published results

obtained with ultrafast ChR variants: Chronos and f-Chrimson (Fig. 3 E-G, replotted from Keppeler et al. 2018 and Zerche et al. 2023)." The statement is a bit unfair, and lead to the notion that f-ChR2 TC outperform the other constructs. While true for Chronos, f-Chrimson seems on par whit f-ChR2 TC here. Adjust phrasing accordingly.

We thank the reviewer for the comment and adjusted the respective phrasing accordingly:

Lines 277- 282:

"We did not observe significant differences of blue-light activated f-ChR2 TC compared to red-light activated f-Chrimson, while f-ChR2 TC significantly outperformed Chronos for I) threshold: $p = 0.0002$; II) P1-N1 amplitude: $p = 0.001$ and III) P1-N1 latency: $p = 0.0002$ (unpaired Kruskal-Wallis test adjusted for multiplicity by Dunn's correction, **Fig. 3 E-G**, replotted from Keppeler et al. 2018 and Zerche et al. 2023)."

5- Major remark

My next point concerns the reported reduction in SGN density following injection. The authors should comment on the magnitude of this reduction, especially since the effect appears to vary between constructs. This variability weakens the argument that the observed SGN loss is solely due to injection pressure. To strengthen this hypothesis, it would be important to include a control group in which mice are injected with a saline solution or better yet an AAV with YFP alone. The authors could also discuss way to adapt the injection protocol to reduce this unwanted effect.

We understand the reviewers concern and appreciate the comment. We agree that the suggested inclusion of control groups such as injections of saline solutions and/or AAVs carrying EYFP alone would help better understand the mechanisms of the observed SGN reduction. Indeed, some of our previous studies have addressed these points. For example, based on a small sample size, we did not find significantly different SGN densities between postnatal injections of AAV-f-Chrimson or saline (analyzed at 3-months of age (Bali *et al*, 2022)). Moreover, as in the present MS, we have typically compared results in the AAV-injected ear to those of the non-injected ear (which typically also shows ChR expression due to CSF spread of AAV, (e.g. Mager *et al*, 2018; Keppeler *et al*, 2018; Bali *et al*, 2021, 2022). There, we typically find less pronounced SGN loss, suggesting an effect of injection and potentially of viral dose. We note that all findings are confounded by the substantial age-dependent SGN loss in C57 black 6 mice (Someya *et al*, 2009) which make it difficult to assess a treatment-related adverse effect (Bali *et al*, 2022).

Evidence for toxic effects of AAV as well as of fluorescent proteins or tagged ChR has been presented for other neurons/cells (Liu *et al*, 1999; Klein *et al*, 2006; Miyashita *et al*, 2013; Maimon *et al*, 2018).

In response to the reviewer's comment, we have performed additional experiments (**Appendix Fig. S5**). Given the report of immunological reactions to optogenetically rendered peripheral mammal neurons by microbial ChRs (Maimon et al. 2018) we probed sections of treated cochleae for signs of induced (CD3 positive cells) and innate (Iba-1 positive cells) immune response. In short, we did not encounter cells and so either the response proceeded before or was not prevalent. In addition, we performed a histopathological scoring based on hematoxylin and eosin (HE) stains of cochlear sections and found lower scores for neuron density and increased interstitial vacuolation compared with the contralateral, non-injected side.

Moreover, we have revised the **discussion** accordingly i) pointing out more translational AAV administration approaches such as via slow micropump-catheter systems with a pressure vent in the cochlea that are now also used in clinical *OTOF* gene therapy trials, and ii) discussing the need for further investigations to decipher the mechanism(s) of SGN reduction which we consider beyond the scope of this study.

Additionally, in Figure EV3, the rationale behind the left/right panel separation is unclear. If the right side corresponds to the injected cochlea and the left to the non-injected one, it is important to clarify what was injected in the control group, and why YFP-positive SGNs are visible on the right side in panel EV3-D.

We thank the reviewer for the remark and gladly provide clarification. As described above we compared the left, injected cochlea with the contralateral non-injected right cochlea from the same animal. In all of our studies with postnatally injected animals we found ChR expression in the contralateral non-injected cochlea likely due to AAV spread via the cochlea aqueduct(s) to/from the CSF space. This has now also been discussed more extensively in the manuscript. The black labelled control group is a non-injected littermate control of the injected animals, in which we did not detect any GFP-signal (zero eYFP+ SGNs/ $10^4\mu\text{m}^2$).

It is also important for a potential therapeutical development to assess if the reduction is due to the pressure injection, a toxic overexpression of the transgene, or a immune response. As the reduction in the number of cells is important, it should be possible to localise the dead cells, or identify the lost volume? It might be worth to relate the observation with what seems like a reduced PV expression in the f-ChR2 TC.

We thank the reviewer for this comment and refer to our above response. In brief, we performed additional experiments including labeling for immune cells and histopathology on cochleae treated with AAV-f-ChR2 TC and AAV-Chronos LC that we now included as **Appendix Figure S5**.

6-

I acknowledge the challenges involved in analyzing the subcellular localization of the optogene from cryosections, and the authors have made a commendable effort

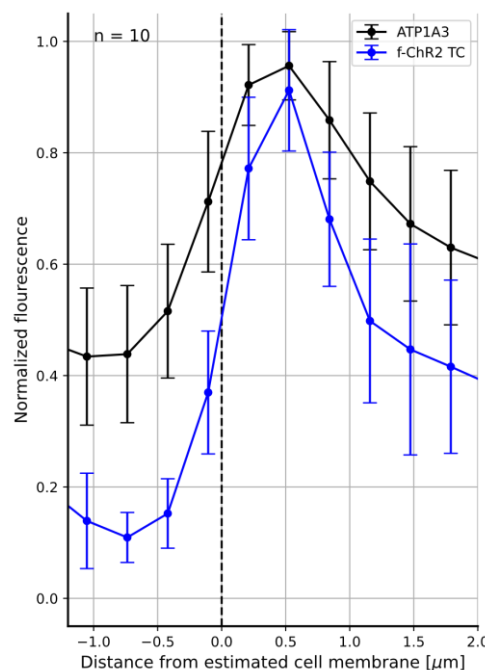
through careful quantification and appropriately cautious terminology in describing their findings. While the methods section outlines the procedure for the line analysis, I was unable to find a clear explanation of how the ratio presented in Figure 4-C was calculated. Furthermore, it would have been beneficial to perform membrane surface staining in at least a small subset of tissue to validate the quantification.

We thank the reviewer for appreciating our efforts. In response to the comment, we now present a clear explanation on how the fluorescence ratios based of the line profiles were calculated in the methods section:

Lines 1092-1094

“The ratio of membrane to intracellular fluorescence was calculated for each selected cell by dividing the mean fluorescence measured between 0.4 and 0.6 μm by the mean fluorescence measured between 1.4 and 1.6 μm .”

In addition, we further analyzed a small set of cryosections from a f-ChR2 TC transduced cochlea that we additionally immunolabeled for Na^+/K^+ ATPase (ATP1A3, which provides a nice plasma membrane staining) validating our quantification of the subcellular ChR distribution: the shapes of the line profiles of f-ChR2 TC and ATP1A3 (n = 10 cells) agree very well (Rebuttal Figure 1).



Rebuttal Figure 1

7-

line392 " injected (P5-P9) C57BL/6 wild-type mice after a standardized expression period of ~ 2 weeks" it is weird to have a standard being approximatively 2 weeks, please rephrase or clarify.

We thank the reviewer for the observation and rephrased accordingly:

Lines 401-403:

"These slices were obtained from postnatally AAV-injected (P5-P9) C57BL/6 wild-type mice after an expression period of 15 ± 0.4 days on average, as described before (Özçete and Moser 2021; Hain and Moser 2024)."

Line 466:

"... an expression period of 15 days, used for the slice measurements..."

8-

line 404 " Thus, by reducing the light irradiance, fewer inputs are recruited, which is reflected by smaller step sizes in the oeEPSC amplitude eventually arriving at monosynaptic input (single endbulb synapse)." From my understanding, this method is not used here. If it is indeed the case, please consider rephrasing, as it may confuse the reader.

We thank the reviewer for this observation and excuse the confusion. We revised the MS accordingly for clarity:

Lines 414-418:

"Given ChR expression differences and the variability of SGN membrane resistance, capacitance and spiking threshold, by reducing the light irradiance, fewer inputs are recruited, which is reflected by smaller oeEPSC amplitudes eventually arriving at monosynaptic input (single endbulb synapse)."

Line 409" Where positive," what is positive here? Typo?

We understand the potentially confusing phrasing and want to point out that not all the cells that are successfully patched have a light-triggered response. By using "where positive", we wanted to refer to the cells with light-driven EPSCs (optically-evoked EPSCs) at least at maximum irradiance. For clarity we decided to delete the statement "Where positive".

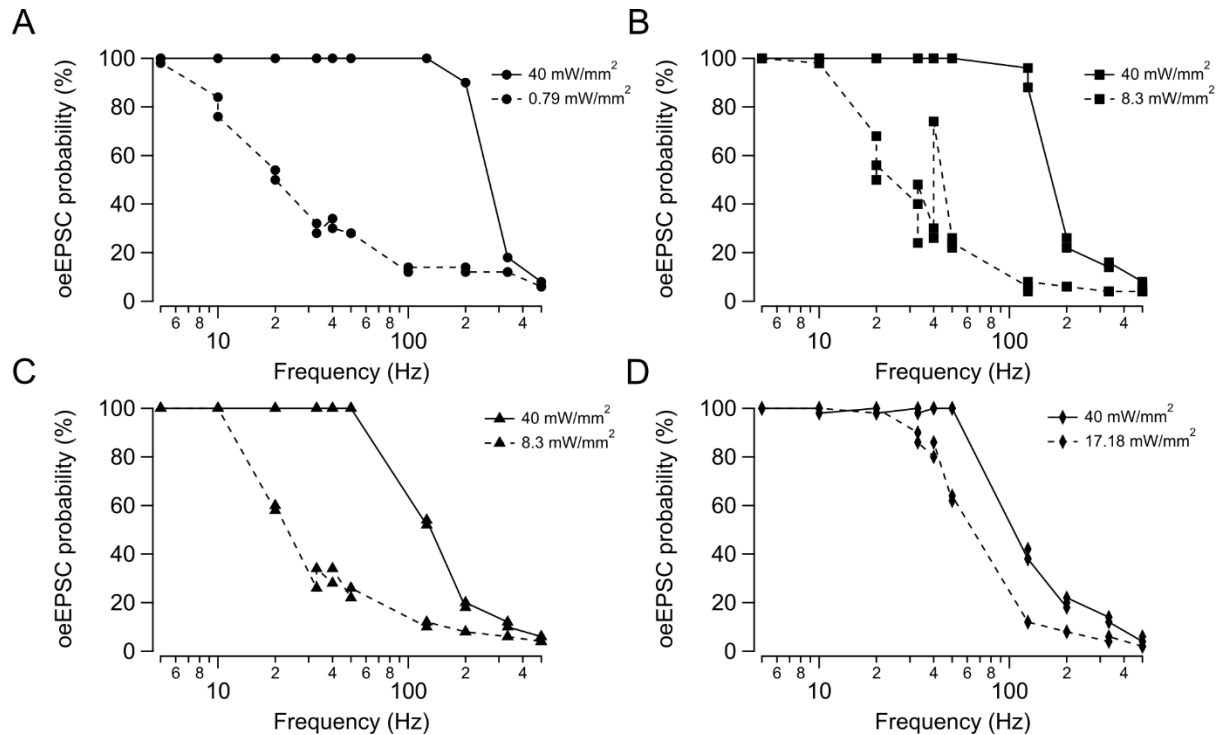
9- Major point

To measure the maximal frequency driving synaptic inputs the authors use variable frequency and saturating light (@ 40mW/mm²). While it help normalize the variability in light sensitivity, isn't it counterproductive in term of maximal frequency responses? I would assume that saturating light might increase the oeEPSC probability reduction observed in the train of stimuli.

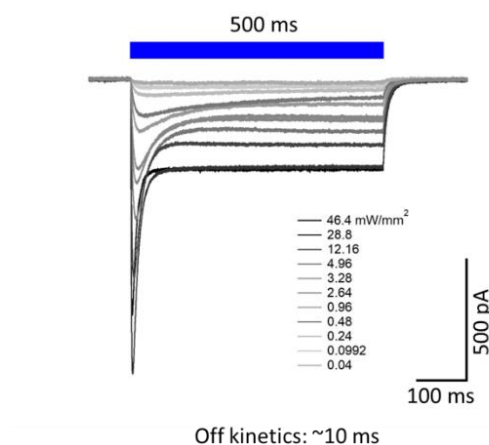
We thank the reviewer and accordingly provide further analyses of oeEPSC recordings following illumination with 50 stimuli of both saturating and subsaturating light at varying frequencies (see Rebuttal Fig. 2; recordings from four distinct bushy cells). As shown, reducing the light intensity consistently led to a decrease rather than an increase in oeEPSC probability within the high-frequency range in all cases.

We propose that this observation stems from the diminished photocurrent at lower light intensities, which resulted in longer latencies and fewer threshold-passing

events. Furthermore, we note that both the initial and sustained photocurrents increase not decrease with rising light intensities (see Rebuttal Fig. 3; f-ChR2 TC photocurrents at different light levels).



Rebuttal Figure 2



Rebuttal Figure 3

10- Major point
Line 416 "Quantification of oeEPSC probability versus irradiance and frequency of stimulation shows that bushy cells that are more light-sensitive in terms of higher oeEPSC probability and shorter synaptic delay also follow higher frequencies of stimulation (Fig. 5 E-G)."

This aspect is a crucial component of the synaptic findings and deserves a clearer representation. While the current data allow the reader to infer the conclusion by comparing individual cells, the current layout is suboptimal. Notably, the line colors in panels E/F and G are inconsistent. The authors should consider a more effective way to illustrate this relationship within a single panel. One possible solution would be to extract, for each cell, a single irradiance or frequency value corresponding to a 50% oEPSC probability, and plot these values against one another.

We apologize for the misunderstanding with the colors in panels E-G, we displayed the same cells with a dimmer tone of the same color to highlight the average in black. We now corrected and displayed all panels with the same color code to keep consistency and clarity.

11-

Line 421 "Given the fast-closing kinetics of f-ChR2 TC at physiological temperature ($\tau_{\text{off}} \sim 5$ ms, Table EV2), we reason that limited expression and photocurrent desensitization of f-ChR2 TC as well as a potential depolarization block hinder endbulb synapses to follow stimulation frequencies of more than 100 Hz."

The rationale behind this hypothesis is not fully elaborated. While the limited expression aspect is relatively straightforward, the desensitization observed with f-ChR2 TC should be more thoroughly discussed. Additionally, the role of depolarization block remains unclear. Since the authors refer to results from cells receiving "optogenetic inputs" rather than directly expressing the opsin, does this imply that the depolarization block occurs at the presynaptic terminals? Typically, a depolarization block would be expected in cells with higher-than-average opsin expression, which might be reflected by a shorter synaptic delay. Is such a correlation observed in this study?

We thank the reviewer for this important comment. As shown in Fig. 5, oeEPSC failures occur more frequently toward the end of the light stimulus train, with a lower oeEPSC probability in pulses 40–50 compared to pulses 1–10. We also observed an increase in oeEPSC latency over the course of the stimulus train (Appendix Fig. S8). Furthermore, pulsed light stimulation at high frequencies (>100 Hz) induces a level of desensitization comparable to that seen with continuous light stimulation (Fig. EV1 vs. Fig. 1E). Based on these observations, we reason that the reduced photocurrent amplitude at the end of the pulse train might be insufficient to trigger presynaptic terminal firing, thereby failing to induce postsynaptic oeEPSCs.

Regarding depolarization block, we agree with the reviewer that, intuitively, one might expect it to occur in cells with higher-than average expression levels. In such cases, light stimulation could induce a saturating photocurrent that impairs sustained neuronal firing and correlates with shorter synaptic delay. However, our dataset does not show that correlation (Appendix Fig. S8B). In contrast, high-performing endbulbs with higher oeEPSC probability at elevated stimulation frequencies also exhibit shorter synaptic latencies likely indicating high ChR expression. Given the absence of direct evidence for depolarization block in our data, we have revised the sentence in the results part accordingly.

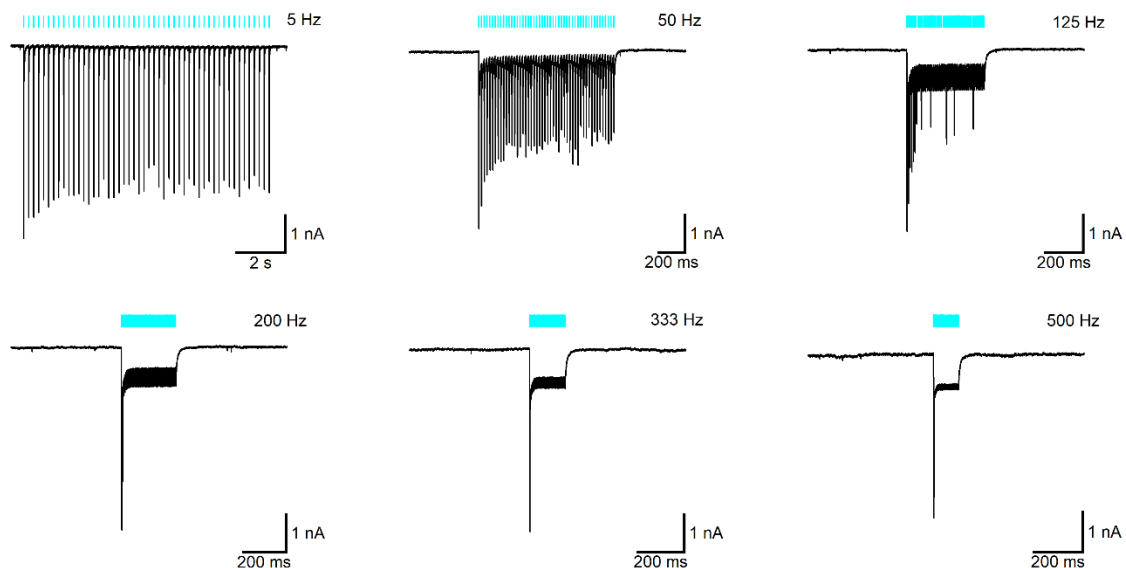
Lines 433-437:

„Given the fast-closing kinetics of f-ChR2 TC at physiological temperature ($\tau_{\text{off}} \sim 5$ ms, Table EV2), we reason that limited expression and photocurrent desensitization of f-ChR2 TC led to the observed heterogeneity of endbulb synaptic transmission to bushy cells and hindered most endbulb synapses to follow stimulation frequencies of more than 100 Hz.”

Our interpretation of the depolarization block was directed/referring to the fact that the decay of photocurrent is also limited by the closing kinetics of f-ChR2 TC at 33 – 34 °C = 4.1 ± 0.96 ms. We reasoned that a substantial remaining photocurrent could have induced a depolarization block in the endbulb of Held in some cases. We attach here an exemplary recording from a bushy cell that expressed f-ChR2 TC to illustrate the notion of the large stationary photocurrent (Rebuttal Fig. 4). Nonetheless, this statement remains somewhat speculative since we did not record directly from the endbulb. However, as we cannot rule out the occurrence of depolarization blocks, we added the following sentence to the discussion:

Lines 594-597:

“In addition, potential depolarization block resulting from prolonged spiral ganglion neuron (SGN) photodepolarization due to the limited closing kinetics of f-ChR2 may have contributed to suboptimal sustained photostimulation at high frequencies.”



Rebuttal Figure 4: Whole cell recording of bushy cell directly expressing f-ChR2 TC during pulsed stimulation (1 ms, 488 nm, ~ 40 mW/mm²) at different frequencies.

12-

Fig EV4: does the panels D and F relate to stellate cells and C and E to bushy cells? If so, it is not clearly indicated in the legend.

We thank the reviewer for pointing this out and accordingly adjusted the legend to provide more clarity.

13-

Line 437 "However, low oeEPSC probability was not always predicted by long latency, as shown exemplarily for the endbulb synapses depicted in Appendix Fig. S5 B-C, and compared to other synapses with similar oeEPSC probability."

I might misunderstand the point, but the example referred to in the legend of Fig S5 B-C seems to have a low probability AND a long latency (green dots). It may be the phrasing that is unclear as the legend indicate:

"Note that low oeEPSC probability was not always predicted by long latency, as shown exemplarily for the endbulb synapses depicted in green dots, and compared to other synapses with similar oeEPSC probability (displayed by purple hourglass icon () and red square). »
The current phrasing suggests that it is specifically for the green dots that the oeEPSC probability fails to predict long latency.

We agree with the reviewer that **Fig S5 B-C** were not clearly explained and did not convey the intended message. We prepared the new **Appendix Fig. S8** (renamed Fig S5). In Appendix **Fig. S8 B** the latency of the first oeEPSC is correlated to the oeEPSC probability determined for each single cell plotted in **Appendix Fig. S8 A**. Cells with similar latency can show very distinct oeEPSC probabilities upon light stimulation with a train of 50 stimuli (1 ms pulse length at 5 Hz). Nonetheless, we observe a general trend: endbulb synapses with shorter latencies tend to have higher oeEPSC probabilities.

We rephrased the text and figure legend accordingly:

Line 447-448:

"However, low oeEPSC probability was not always correlating to long latencies (**Appendix Fig. S8 B**)."

Line 1623-1626:

"Note that low oeEPSC probability was not always correlating to long latencies, as shown exemplarily for the endbulb synapses depicted in red square when compared to other synapses with similar latency (displayed by cyan rhombus, purple hourglass icon (⌵) and blue triangular star)."

14-

Fig S6 C legend: « Quantification of synaptic delay dependent on frequency of stimulation (~ 40 mW/mm², 1 ms; continuous line) and dependent on irradiance (10 Hz, 1 ms; dashed line). «

does not fit with the actual figure, i only have continuous line, and 1 X axis, rendering issue?

We are thankful to the reviewer and apologize for this mistake in the figure legend, which we corrected accordingly:

Lines 1635 - 1636:

"C, Quantification of synaptic delay dependent on frequency of stimulation (~ 40 mW/mm², 1 ms; continuous line)."

Discussion:

It would be valuable to further elaborate on strategies that could help minimize desensitization.

Since neurons can differ in their intrinsic properties-affecting, for instance, the maximal sustainable firing rate-the authors should also expand on how the results observed in hippocampal neurons relate to the later findings in SGNs or oABR recordings.

We thank the reviewer for the valuable comments. Regarding strategies to optimize ChR properties, such as minimizing photocurrent desensitization, we now provide Appendix Table S2 and refer to recently published work (Huet et al., 2024; Alekseev et al., 2025). We agree to the reviewer that neuronal subtype-specific differences, such as the maximal sustainable firing rate, need to be considered. In response to the reviewer's advice, we accordingly provide the following sections.

Lines 225-229:

"The investigation of high-rate neurostimulation is impeded by the limited and heterogeneous intrinsic maximal firing rate of rat hippocampal neurons which is typically 40 to 60 Hz (Gunaydin et al. 2010; Mager et al. 2018), well below that of fast spiking neurons such interneurons and SGNs (Mager et al. 2018). We accordingly turned to the investigation of optogenetic SGN stimulation in the auditory pathway of mice."

Lines 586-587:

"In order to investigate high-rate stimulation, we turned to experiments in SGNs, fast spiking auditory neurons.

Lines 560-562:

"ChR variants with balanced channel closing kinetics, robust plasma membrane targeted expression, and low photocurrent desensitization are desirable for efficient and safe neurostimulation at high rates (Huet et al. 2024).

Lines 643-650:

"A recent study has demonstrated the potential of the engineered green light-activated ChR variant ChReef, to mitigate this issue (Alekseev et al. 2025). ChReef

enables sustained control of excitable cell activity at low light intensities, owing to its good plasma membrane targeting, minimal photocurrent desensitization, and comparatively high unitary conductance. Yet, the temporal fidelity of SGN activation is limited by ChReef's slow channel closing kinetics. The generation and comprehensive characterization of both natural and engineered ChRs will further fuel the development of advanced, kinetically balanced variants that can be safely integrated into future blue- or green- μ LED-based oCIs."

Line 551 : "Limitations explaining failure or sustained high rate..." should be "...OF sustained high rate"

Although, that sentence remain unclear even with the proposed modification

We thank the reviewer for noticing this typo. We corrected the sentence and improved it for clarity.

Lines 589-593:

"Limitations explaining oeEPSC failure during sustained high-rate neurostimulation include limited expression levels of f-ChR2 TC in SGN terminals, as indicated by the correlation of oeEPSC probability and light sensitivity and the increased synaptic delay compared to electrically triggered EPSCs (Chanda and Xu-Friedman 2010)."

Methods:

Line 656: Toff is determined as a monoexponential fit of the current, but there is no quantification of the goodness of the fit? Could a double exponential lead to better fit?

It is then mentioned line 796 that 10/15 cells have a biexponential fit, without much explanation.

We want to thank the reviewer and provide more explanations. The quality of the fits was ensured by $R^2 > 0.98$ and residual analysis. The photocurrents were fitted monoexponentially. The EPSCs were fitted mono- or bi-exponentially, whereas either the time constants derived from the monoexponential fits or the faster time constants derived from the biexponential fits were used to classify principal cells as bushy cells ($T_{off} \leq 0.5$ ms) or stellate cells ($T_{off} > 0.5$ ms) as shown in previous studies (Isaacson and Walmsley 1995; Gardner et al. 1999; Cao and Oertel 2010; Chanda and Xu-Friedman 2010; Butola et al. 2021).

Line 663: the authors mention peak recovery of ChRs, where is that data presented? page 28 -line 745:

"The temperature dependence of the off-kinetics and peak recovery of ChRs and their mutants was investigated at temperatures closer to physiological conditions (33 to 34 °C)."

We thank the reviewer for the comment. The data showing the peak recovery is presented in the new **Appendix Fig. S2**.

Line 746-749:

“The peak current recovery and the closing kinetics of selected ChRs were investigated at temperatures closer to physiological conditions (33 to 34 °C). The peak current amplitude was fully recovered after a period of 30 s in the dark (**Appendix Fig. S2**).

Line 858 : the sentence is uncomplete (no verb).

We would like to thank the reviewer and have corrected the sentence accordingly.

11th Oct 2025

Dear Dr. Mager,

Thank you for the submission of your revised manuscript to EMBO Molecular Medicine. We have now received the enclosed report from the referees who were asked to re-assess it. As you will see, the referees are now supportive, and I am pleased to inform you that we will be able to accept your manuscript pending the following amendments:

1. The remaining minor issues raised by Ref #2.

On a more editorial level, please do the following:

1. Remove all figures from the main manuscript file. Keep the figure legends for both main and Extended View (EV) figures at the end of the manuscript file.

2. Delete the "Authors' Contributions" section from the manuscript file.

3. References: Use "et al." after listing the first 10 authors in each reference.

4. Funding

- Ensure the funding information in the submission system matches the manuscript text. The following funding details are currently missing in the submission system: Grant 1690, HORIZON TMA MSCA Postdoctoral Fellowship (OPTOCODE, grant 101107675), Scholarship from the Göttingen Promotionskolleg für Medizinstudierende, funded by the Jacob-Henle-Programm/Else-Kröner-Fresenius-Stiftung (Promotionskolleg für Epigenomik und Genomdynamik 2021_EKPK.04)
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- Remove all EV tables and their legends from the manuscript file.
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6. Appendix:

- Remove all appendix files, figures, and tables from the manuscript.
- Also delete individual appendix figures and tables from the submission system.
- Compile all appendix material into a single PDF file named "Appendix". The PDF must begin with a title page that includes a Table of Contents listing all items with their page numbers. Place the legend below each figure and table within the appendix.

7. Figures 6H-J are cited, but panel J is missing or not labeled in the figure. Please resolve this discrepancy.

8. During our routine image check, we noted the following:

- There is image reuse between Figure 1A and Appendix Figure S1D. This reuse should be explicitly stated in the figure legends.
- In Appendix Figures S5A and S5D, a side section appears to have been removed, which is visible under image filters(see attached). Please provide the original, unprocessed source data for these figures.

9. Data availability:

- Remove the subheadings "Data Availability Statement" and "Code Availability Statement" from the manuscript. Merge the relevant content into a single "Data Availability" section. Place the Data Availability section before the Acknowledgments section in the manuscript.
- The following statements must be removed: "The code used for analysis is available from the corresponding authors upon reasonable request." "The data that support the findings of this study is available from the corresponding authors upon reasonable request." Since the study does not generate large-scale datasets, formal data deposition is not required.
- However, the computer code used for analysis must be deposited in an appropriate public repository (e.g., GitHub, Zenodo). A direct access link to the repository must be provided in the Data Availability section.

10. Please address the following comments related to figure legends:

- Please note that the exact p values are not provided in the legends of figures 1F, 2E, 4B, C; EV2 C, E, F; EV3 A-C
- Please note that the box plots need to be defined in terms of minima, maxima, centre, bounds of box and whiskers, and percentile in the legends of figures 1B, F; 2B, E; 6C
- Please note that information related to n is missing in the legends of figures 3H, 5G, H
- Please note that the error bars are not defined in the legends of figures 1C, D; 3H, 5H, EV1 B, C

11. Please correct the order and headings of the manuscript sections to: Abstract / Keywords / The Paper Explained / Introduction / Results / Discussion / Methods / Data Availability / Acknowledgements / Disclosure and Competing Interests Statement / References / Main Figure Legends / Tables / Expanded View Figure Legends

We look forward to reading a new revised version of your manuscript as soon as possible.

Kind regards,
Jingyi

Jingyi Hou
Senior Editor
EMBO Molecular Medicine

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In the event of acceptance, this file will be published in conjunction with your paper and will include the anonymous referee reports, your point-by-point response and all pertinent correspondence relating to the manuscript. If you do NOT want this file to be published, please inform the editorial office at contact@embomolmed.org.

***** Reviewer's comments *****

Referee #1 (Comments on Novelty/Model System for Author):

The provided in vitro and in vivo data are consistent with the conclusions.

Referee #1 (Remarks for Author):

The authors have thoroughly revised the manuscript. All of my questions and concerns have been adequately addressed. I thank the authors for their efforts and congratulate them on their work.

Referee #2 (Comments on Novelty/Model System for Author):

The paper technical mastery is of the highest quality, experiment are well designed and executed and interpretations are thorough. it linked proteic mutation to physiological properties of channels and effect on auditory restoration in whole organism. The best candidate variant that arise from the present study englobe together all the necessary properties making it the new landmark for auditory restoration.

the potential for translation to clinical is straight forward. albeit the necessity for preclinical non human primate trials, which seem outside of the scope of this study.

The study englobe various model to test the gene candidate, with the gerbil model (establish previously as a model for audition restoration) demonstrating consistent results.

Referee #2 (Remarks for Author):

The authors have addressed all my comments thoroughly and with well-reasoned arguments.

The revisions made in response to my feedback, as well as that of the other reviewer, have greatly improved the manuscript's readability, making it more coherent and narrative-driven, with a stronger emphasis on the potential clinical implications.

I only noted an error in legend of figure 5(lines 498-498):

" Note the relationship between light-sensitivity and synaptic delay (e.g., synapses labeled in light blue horizontal hourglass icon () vs green cell vertical hourglass icon ())"

as far as i can tell, the icons mentioned are not on the figure.

The remaining minor issues raised by Ref #2.

I only noted an error in legend of figure 5(lines 498-498):

" Note the relationship between light-sensitivity and synaptic delay (e.g., synapses labeled in light blue horizontal hourglass icon () vs green cell vertical hourglass icon ())"

as far as i can tell, the icons mentioned are not on the figure.

We magnified the icons in order to improve the visibility and thank the reviewer for the comment, which helped to enhance clarity.

17th Nov 2025

Dear Dr. Mager,

Thank you for sending us your revised manuscript. We are pleased to inform you that your manuscript is accepted for publication and is now being sent to our publisher to be included in the next available issue of EMBO Molecular Medicine.

Your manuscript will be processed for publication by EMBO Press. It will be copy edited and you will receive page proofs prior to publication. Please note that you will be contacted by Springer Nature Author Services to complete licensing and payment information.

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Should you be planning a Press Release on your article, please get in contact with embo_production@springernature.com as early as possible in order to coordinate publication and release dates.

If you have any questions, please do not hesitate to contact the Editorial Office. Thank you for your contribution to EMBO Molecular Medicine.

Sincerely,
Jingyi

Jingyi Hou
Senior Editor
EMBO Molecular Medicine

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This checklist is adapted from Materials Design Analysis Reporting (MDAR) Checklist for Authors. MDAR establishes a minimum set of requirements in transparent reporting in the life sciences (see Statement of Task: [10.31222/osf.io/9sm4x](https://doi.org/10.31222/osf.io/9sm4x)). Please follow the journal's guidelines in preparing your

Please note that a copy of this checklist will be published alongside your article.

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The data shown in figures should satisfy the following conditions:

- the data were obtained and processed according to the field's best practice and are presented to reflect the results of the experiments in an accurate and unbiased manner.
- ideally, figure panels should include only measurements that are directly comparable to each other and obtained with the same assay.
- plots include clearly labeled error bars for independent experiments and sample sizes. Unless justified, error bars should not be shown for technical
- if $n < 5$, the individual data points from each experiment should be plotted. Any statistical test employed should be justified.
- Source Data should be included to report the data underlying figures according to the guidelines set out in the authorship guidelines on Data

2. Captions

Each figure caption should contain the following information, for each panel where they are relevant:

- a specification of the experimental system investigated (eg cell line, species name).
- the assay(s) and method(s) used to carry out the reported observations and measurements.
- an explicit mention of the biological and chemical entity(ies) that are being measured.
- an explicit mention of the biological and chemical entity(ies) that are altered/ varied/ perturbed in a controlled manner.
- the exact sample size (n) for each experimental group/condition, given as a number, not a range;
- a description of the sample collection allowing the reader to understand whether the samples represent technical or biological replicates (including how many animals, litters, cultures, etc.).
- a statement of how many times the experiment shown was independently replicated in the laboratory.
- definitions of statistical methods and measures:
 - common tests, such as t-test (please specify whether paired vs. unpaired), simple χ^2 tests, Wilcoxon and Mann-Whitney tests, can be unambiguously identified by name only, but more complex techniques should be described in the methods section;
 - are tests one-sided or two-sided?
 - are there adjustments for multiple comparisons?
 - exact statistical test results, e.g., P values = x but not P values < x;
 - definition of 'center values' as median or average;
 - definition of error bars as s.d. or s.e.m.

Please complete ALL of the questions below.
Select "Not Applicable" only when the requested information is not relevant for your study.

Materials

Newly Created Materials	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
New materials and reagents need to be available; do any restrictions apply?	Yes	New ChR variants are described in the Appendix Table S2; point mutations are explained in the main text and in Materials and methods.
Antibodies	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
For antibodies provide the following information: - Commercial antibodies: RRID (if possible) or supplier name, catalogue number and or/clone number - Non-commercial: RRID or citation	Yes	For antibodies the information can be found in the Materials and Methods and the Reagents and tools table.
DNA and RNA sequences	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
Short novel DNA or RNA including primers, probes: provide the sequences.	Yes	Primers used are listed in Appendix Table S3.
Cell materials	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
Cell lines: Provide species information, strain. Provide accession number in repository OR supplier name, catalog number, clone number, and OR RRID.	Yes	Materials and Methods and Reagents and tools table.
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Report if the cell lines were recently authenticated (e.g., by STR profiling) and tested for mycoplasma contamination.	Yes	Materials and Methods
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Human research participants	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
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Design

Study protocol	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
If study protocol has been pre-registered , provide DOI in the manuscript. For clinical trials, provide the trial registration number OR cite DOI.	Not Applicable	
Report the clinical trial registration number (at ClinicalTrials.gov or equivalent), where applicable.	Not Applicable	
Laboratory protocol	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
Provide DOI OR other citation details if external detailed step-by-step protocols are available.	Yes	Materials and Methods.
Experimental study design and statistics	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
Include a statement about sample size estimate even if no statistical methods were used.	Yes	Sample size for each experiment is stated in each Figure legend and in the Data availability section.
Were any steps taken to minimize the effects of subjective bias when allocating animals/samples to treatment (e.g. randomization procedure)? If yes, have they been described?	Not Applicable	
Include a statement about blinding even if no blinding was done.	Not Applicable	
Describe inclusion/exclusion criteria if samples or animals were excluded from the analysis. Were the criteria pre-established?	Yes	Inclusion/exclusion criteria are described in the Materials and methods where applicable.
If sample or data points were omitted from analysis, report if this was due to attrition or intentional exclusion and provide justification.		
For every figure, are statistical tests justified as appropriate? Do the data meet the assumptions of the tests (e.g., normal distribution)? Describe any methods used to assess it. Is there an estimate of variation within each group of data? Is the variance similar between the groups that are being statistically compared?	Yes	All statistical methods used are stated in each Figure legend and a specific section in Materials and methods describes all the tests performed.
Sample definition and in-laboratory replication	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
In the figure legends: state number of times the experiment was replicated in laboratory .	Yes	Figure legends and Data Availability section.
In the figure legends: define whether data describe technical or biological replicates .	Yes	Figure legends and Data Availability section.

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Ethics	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
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Studies involving human participants : Include a statement confirming that informed consent was obtained from all subjects and that the experiments conformed to the principles set out in the WMA Declaration of Helsinki and the Department of Health and Human Services Belmont Report.	Not Applicable	
Studies involving human participants : For publication of patient photos , include a statement confirming that consent to publish was obtained.	Not Applicable	
Studies involving experimental animals : State details of authority granting ethics approval (IRB or equivalent committee(s), provide reference number for approval. Include a statement of compliance with ethical regulations.	Yes	Materials and Methods.
Studies involving specimen and field samples : State if relevant permits obtained, provide details of authority approving study; if none were required, explain why.	Not Applicable	
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If you used a select agent, is the security level of the lab appropriate and reported in the manuscript?	Not Applicable	
If a study is subject to dual use research of concern regulations, is the name of the authority granting approval and reference number for the regulatory approval provided in the manuscript?	Not Applicable	

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The MDAR framework recommends adoption of discipline-specific guidelines, established and endorsed through community initiatives. Journals have their own policy about requiring specific guidelines and recommendations to complement MDAR.

Adherence to community standards	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
State if relevant guidelines or checklists (e.g., ICMJE, MIBBI, ARRIVE, PRISMA) have been followed or provided.	Not Applicable	
For tumor marker prognostic studies , we recommend that you follow the REMARK reporting guidelines (see link list at top right). See author guidelines, under 'Reporting Guidelines'. Please confirm you have followed these guidelines.	Not Applicable	
For phase II and III randomized controlled trials , please refer to the CONSORT flow diagram (see link list at top right) and submit the CONSORT checklist (see link list at top right) with your submission. See author guidelines, under 'Reporting Guidelines'. Please confirm you have submitted this list.	Not Applicable	

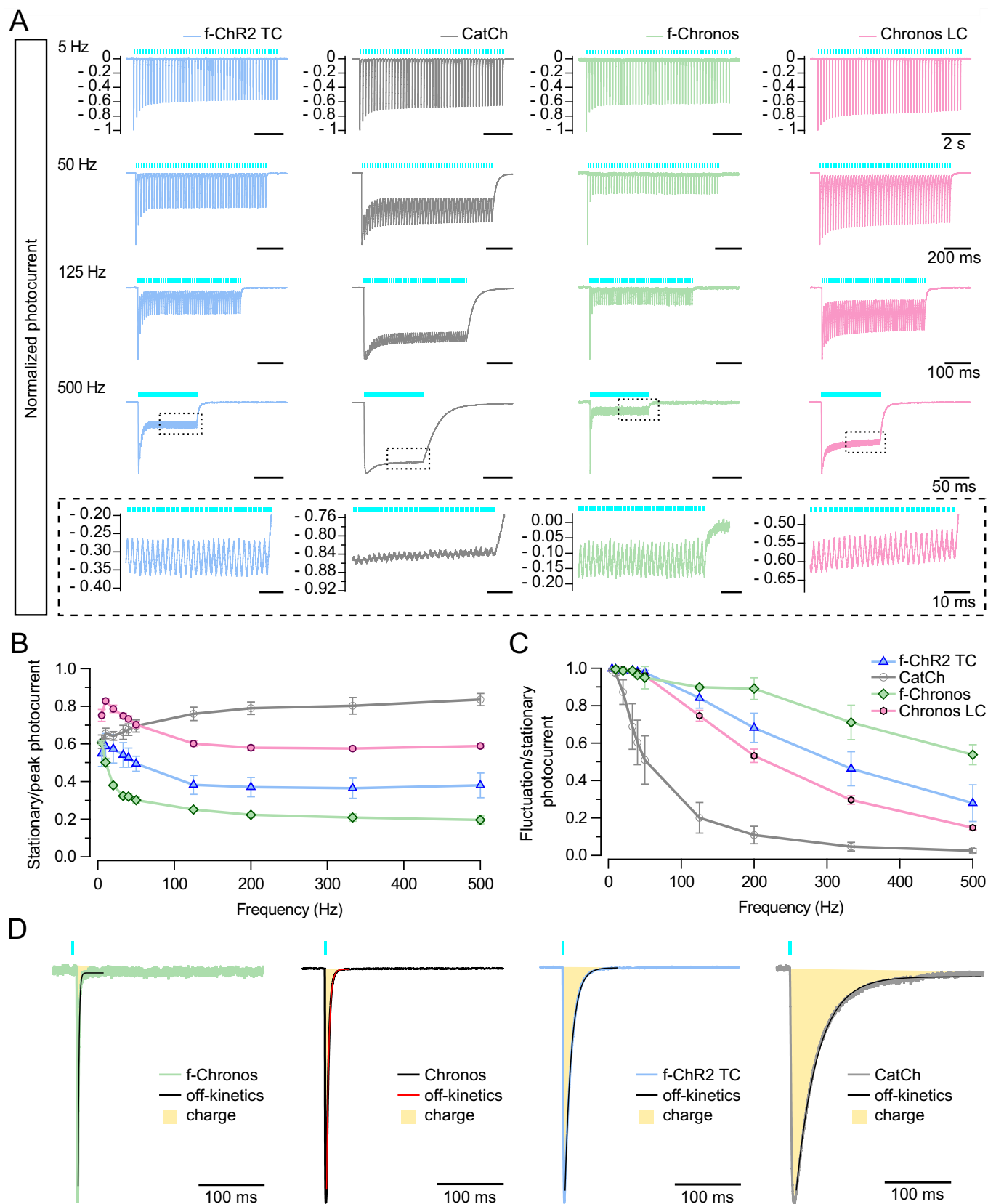
Data Availability

Data availability	Information included in the manuscript?	In which section is the information available? (Reagents and Tools Table, Materials and Methods, Figures, Data Availability Section)
Have primary datasets been deposited according to the journal's guidelines (see 'Data Deposition' section) and the respective accession numbers provided in the Data Availability Section?	Yes	Data Availability Section
Were human clinical and genomic datasets deposited in a public access-controlled repository in accordance to ethical obligations to the patients and to the applicable consent agreement?	Not Applicable	
Are computational models that are central and integral to a study available without restrictions in a machine-readable form? Were the relevant accession numbers or links provided?	Not Applicable	
If publicly available data were reused, provide the respective data citations in the reference list .	Not Applicable	

Expanded View Figures

Figure EV1. Photocurrent measurements of blue light-activated ChRs at physiological temperature.

(A) Exemplary peak-normalized photocurrents from whole-cell patch-clamp measurements at -34°C in NG108-15 cells. ChRs were activated by 50 light pulses of 1 ms at 488 nm ($\sim 40\text{ mW/mm}^2$) at different frequencies (5, 50, 125, and 500 Hz). The lower panels are magnifications of the 500 Hz traces showing photocurrent fluctuations at the stationary state. (B) Dependence of the stationary/peak ratio on light pulse frequency. Error bars depict SD. (C) Quantification of photocurrent fluctuations normalized to the stationary current amplitude. In panels (B, C): CatCh: $n = 5$, f-Chronos: $n = 3$, Chronos LC: $n = 3$, f-ChR2 TC: $n = 4$. Error bars depict SD. Statistical comparisons can be found in Table EV3. (D) Exemplary peak normalized photocurrents measured in NG cells expressing f-Chronos, Chronos, f-ChR2 TC, or CatCh showing the relation between photocurrent decay kinetics and the transferred charge (area under the curve shown in yellow).



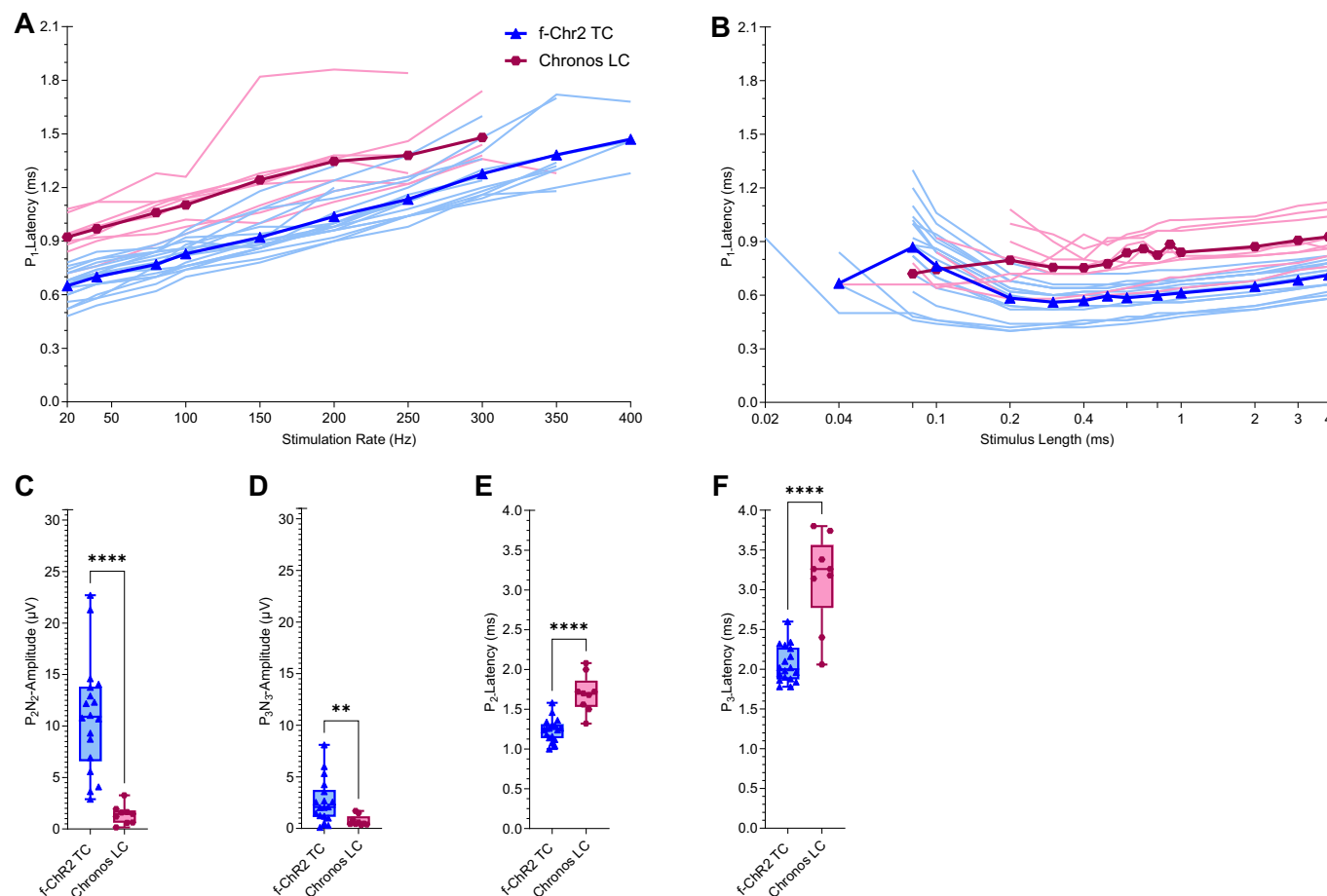


Figure EV2. Analysis of auditory pathway activity evoked by photostimulation of SGNs expressing optimized blue-light-sensitive ChRs.

(A) P1 latency of oABRs at varying repetition rate using 1 ms pulses at -38 to 45.6 mW (bold: mean; faint: all measurements), $n = 18$ mice for f-ChR2 TC, $n = 9$ mice for Chronos LC. (B) P1 latency of oABRs for varying pulse durations using -38 to 45.6 mW pulses at 10 Hz (bold: mean; faint: all measurements) for $n = 17$ mice for f-ChR2 TC, $n = 9$ mice for Chronos LC. (C-F) P₂-N₂, P₃-N₃ amplitudes and P₂, P₃ latencies of oABRs for $n = 18$ mice for f-ChR2 TC, $n = 9$ mice for Chronos LC depicting activation of the auditory pathway using -38 to 45.6 mW, 1 ms pulses at 10 Hz. Data were analyzed as mean \pm SD. Center lines represent median values. Boxes show the 25th and 75th percentile and error bars depict minima and maxima. **** $p = 8.53 \times 10^{-7}$ (C), $p = 1.09 \times 10^{-5}$ (E), $p = 2.62 \times 10^{-5}$ (F); ** $p = 0.0075$ by two-tailed Mann-Whitney test.

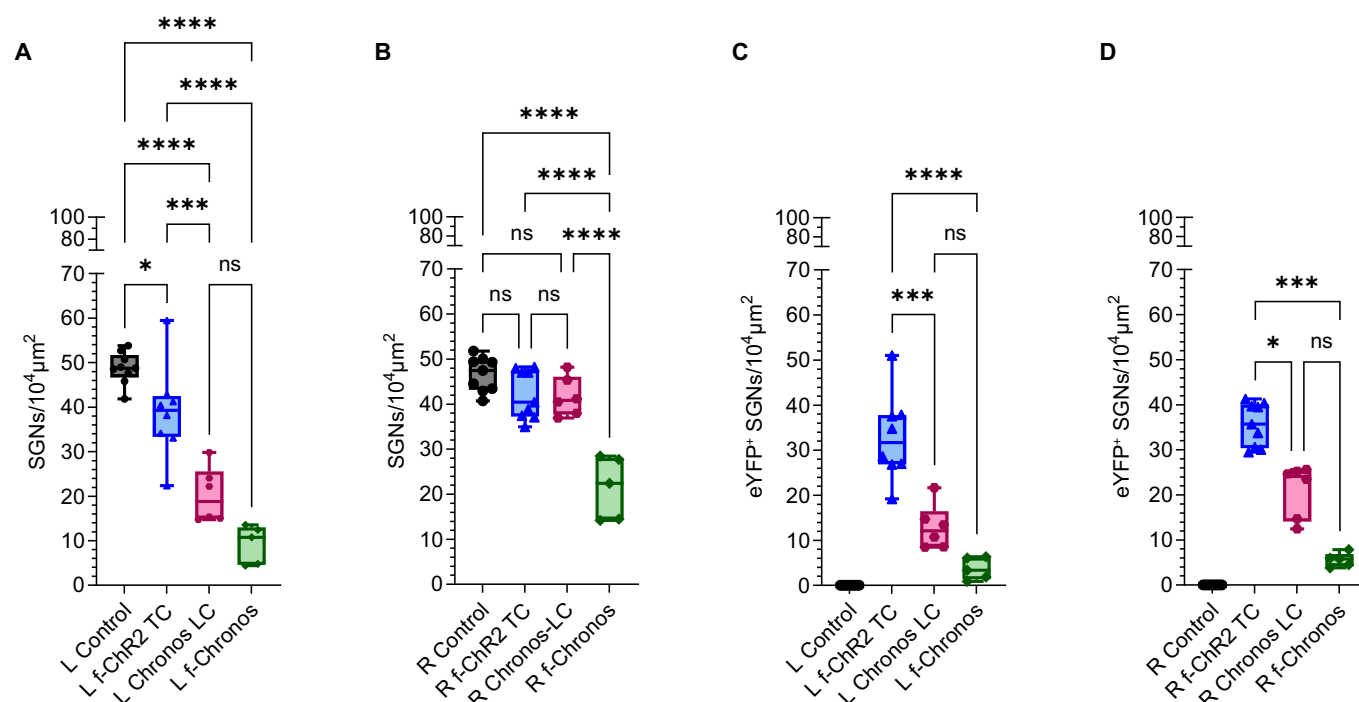


Figure EV3. Immunohistochemical quantification of SGNs expressing optimized blue-light sensitive ChRs.

(A–D) Box plots show statistics for the SGN density of left (injected, A) and right (non-injected, B) cochleae across all cochlear turns, as well as for the density of ChR-expressing (EYFP-positive) cells for the left (C) and the right (D) side. Quantification includes f-ChR2 TC (blue; $n = 8$ for the left and $n = 9$ for the right cochleae), Chronos LC (violet; $n = 6$ for left and right cochlea), f-Chronos (green; $n = 5$ for left and right cochlea), and non-treated wild-type cochleae (black; $n = 9$ cochlea for both sides each). Center lines represent median values. Boxes show the 25th and 75th percentile and error bars depict minima and maxima. **** $p = 2.68 \times 10^{-7}$ (A: L Control vs L Chronos LC), $p = 1.74 \times 10^{-9}$ (A: L Control vs. L f-Chronos), $p = 5.00 \times 10^{-7}$ (A: f-ChR2 TC vs. L f-Chronos), $p = 1.68 \times 10^{-8}$ (B: R Control vs. R f-Chronos), $p = 6.14 \times 10^{-7}$ (B: R f-ChR2 TC vs. R f-Chronos), $p = 3.53 \times 10^{-6}$ (B: R Chronos LC vs. R f-Chronos), $p = 6.06 \times 10^{-6}$ (C: L f-ChR2 TC vs. L f-Chronos); *** $p = 0.0002$ (A), $p = 0.0003$ (C), $p = 0.0002$ (D); * $p = 0.0458$ (A), $p = 0.0485$ (D) by ordinary one-way ANOVA corrected with Bonferroni's (A–C) and Kruskal-Wallis test corrected for multiple comparison with Dunn's (D).

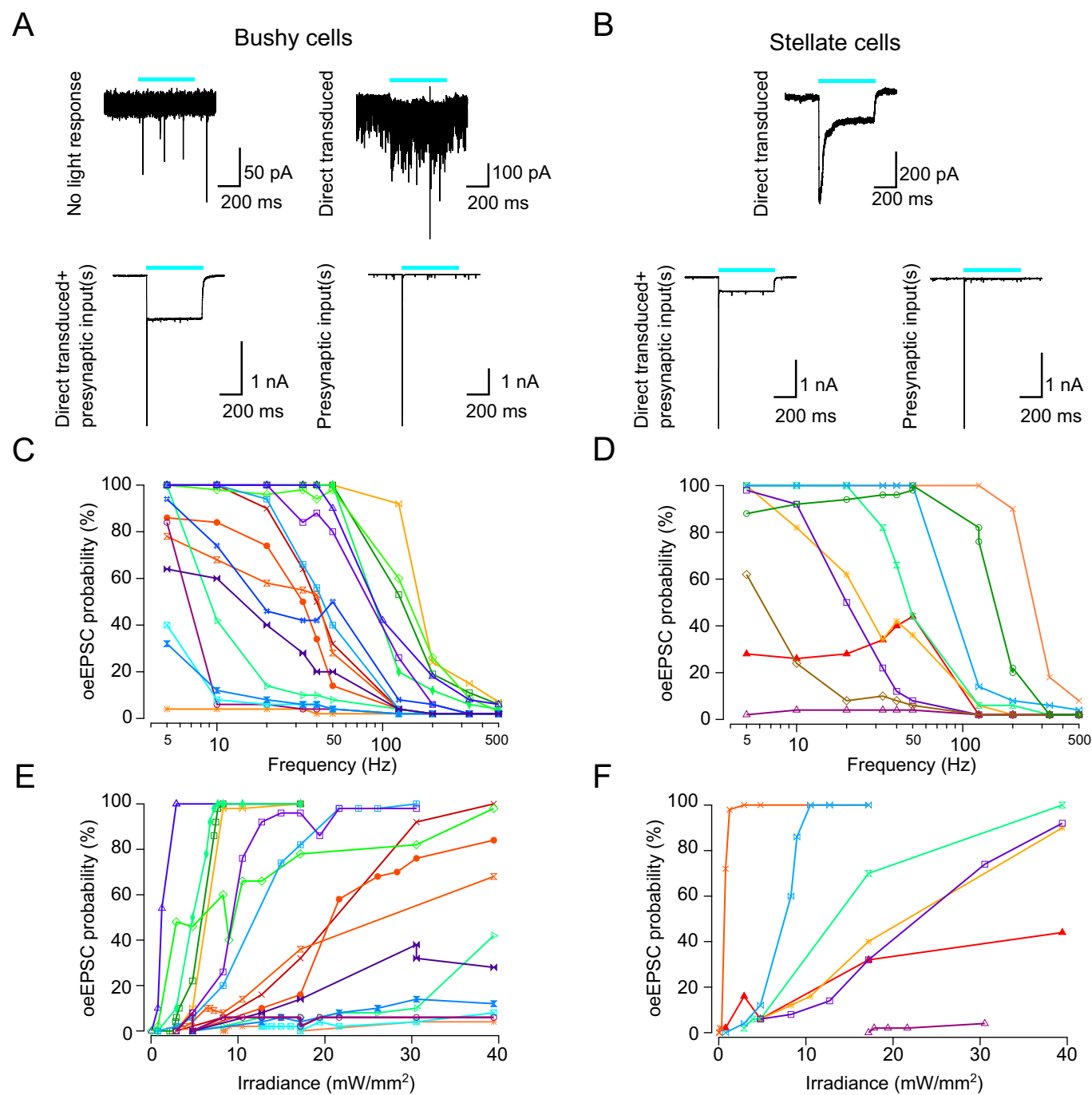


Figure EV4. Variability of photoresponses in principal cells of the AVCN.

(A, B), Recordings upon long light stimulation (500 ms, 488 nm, ~40 mW/mm²) of either bushy (A) or stellate (B) cells, indicating transduced principal cells (directly transduced) and non-transduced SGNs, a combination of directly transduced principal cells + transduced SGN presynaptic inputs), or none of them (no light response). Dependence of oeEPSC probability on the stimulation frequency ((C), for bushy cells; (D), for stellate cells) and the irradiance ((E), for bushy cells; (F), for stellate cells) in principal cells only receiving presynaptic input(s).