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1 **Olfactory rod cells: a rare cell type in the larval zebrafish olfactory epithelium**
2 **with an actin-rich apical projection**

3

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27 **Key words:** olfactory rod cell, olfactory placode, olfactory epithelium, actin, actin-rich
28 projection, Lifeact, zebrafish

29

30 **Abstract**

31 We report the presence of a rare cell type, the olfactory rod cell, in the developing
32 zebrafish olfactory epithelium. These cells each bear a single actin-rich rod-like
33 apical projection extending about 10 μm from the epithelial surface. Live imaging
34 with a ubiquitous Lifeact-RFP label indicates that the rods can oscillate. Olfactory
35 rods arise within a few hours of the olfactory pit opening, increase in numbers and
36 size during larval stages, and can develop in the absence of olfactory cilia. Olfactory
37 rod cells differ in morphology from the known classes of olfactory sensory neuron,
38 but express reporters driven by neuronal promoters. The cells also differ from
39 secondary sensory cells such as hair cells of the inner ear or lateral line, or sensory
40 cells in the taste bud, as they are not associated with established synaptic terminals.
41 A sub-population of olfactory rod cells expresses a Lifeact-mRFP_{ruby} transgene
42 driven by the *sox10* promoter. Mosaic expression of this transgene reveals that
43 olfactory rod cells have rounded cell bodies located apically in the olfactory
44 epithelium.

45

46 **Introduction**

47 The vertebrate olfactory epithelium (OE) enables the detection of chemical cues,
48 giving rise to the sense of smell (reviewed in [Axel, 1995]). The function of this
49 epithelium, which derives from paired cranial neurogenic placodes, is mediated by a
50 diverse set of cells that includes neuronal receptors and non-sensory cells. Olfactory
51 sensory neurons (OSNs) are bipolar neurons that extend a dendrite to the apical
52 surface of the OE, and an axon to the olfactory bulb (OB). In mammals, two broad
53 classes of sensory receptors — ciliated and microvillous OSNs — have been
54 identified on the basis of morphology, receptor expression and OB target.
55 Mammalian OSNs can act as both chemosensors and mechanosensors (Grosmaître
56 et al., 2007; Iwata et al., 2017). The OE of other vertebrates also contains ciliated
57 and microvillous neurons. In fish, additional classes of OSNs have been identified.
58 Each occupies a stereotyped position within the pseudostratified OE, with the
59 dendrite bearing a distinct and characteristic specialisation projecting into the
60 environment (Hansen & Zeiske, 1998; Hansen & Zielinski, 2005; Sato et al., 2005;
61 reviewed in [Maier et al., 2014]).

62

63 In zebrafish, ciliated neurons, which express olfactory marker protein (OMP) and
64 odorant receptor (OR) genes, have a cell body that lies deep within the OE, an axon
65 that projects to dorsal and medial regions of the OB, and a slender dendrite
66 extending to the surface of the olfactory pit. Here, the dendritic knob bears a cluster
67 of primary cilia that project into the olfactory cavity (Hansen & Zeiske, 1998; Hansen
68 & Zielinski, 2005; Sato et al., 2005). Microvillous neurons, characterised by the
69 expression of TrpC2 and vomeronasal (VR)-type pheromone receptors, have cell
70 bodies that lie in the intermediary layer of the OE, an axon that projects to the lateral
71 part of the OB, and a dendrite bearing a tuft of short, actin-rich microvilli (Hansen &
72 Zeiske, 1998; Hansen & Zielinski, 2005; Sato et al., 2005). Crypt neurons, less
73 abundant than ciliated or microvillous neurons, have rounded cell bodies that sit
74 apically in the OE, with both cilia and microvilli extending from a crypt within the cell
75 body (Hansen & Zeiske, 1998; Hansen & Zielinski, 2005; Parisi et al., 2014; Biechl et
76 al., 2016; Bettini et al., 2017). Kappe neurons lie in the superficial layers of the adult
77 zebrafish OE and are named for their apical actin-rich cap, presumed to be microvilli
78 (Ahuja et al., 2014). Pear-shaped neurons are also positioned superficially in the

79 adult OE and have short apical dendrites, but express some markers in common
80 with ciliated neurons (Wakisaka et al., 2017).

81

82 OSNs are surrounded and separated by a network of non-neuronal cell types. These
83 include sustentacular (support) cells, basal cells that replenish the OSNs, and goblet
84 cells that produce mucus (Hansen & Zeiske, 1993; Hansen & Zeiske, 1998; reviewed
85 in [Olivares & Schmachtenberg, 2019]; Demirler et al., 2019). In fish, multiciliated
86 cells, located around the rim of the olfactory pit, each bear multiple long motile cilia.
87 These have a characteristic 9+2 axoneme and beat at around 24 Hz, resulting in an
88 asymmetric flow that draws water and odorants into the olfactory cavity and flushes
89 them out again (Reiten et al., 2017).

90

91 In addition to common cell types, tissues may also contain rare or sparsely-
92 distributed cell types, which are difficult to detect by conventional histological
93 methods. These include stem cells, immune cells, or other rare cell types, which can
94 have critical functions (see, for example, [Montoro et al., 2018; Sui et al., 2018]).

95 Characterisation of the identity and lineage of every cell type of an organ system is a
96 goal of many contemporary single-cell and single-nucleus RNA-seq studies (Junker
97 et al., 2014; Satija et al., 2015; Hernández et al., 2018; Raj et al., 2018; Farnsworth
98 et al., 2020; Wattrus & Zon, 2020). Transgenic or other fluorescent markers, coupled
99 with high-resolution imaging of the whole embryo, can also help to identify cell types
100 that may previously have been overlooked (see, for example, [Kawakami et al.,
101 2010; Galanternik et al., 2017; van Lessen et al., 2017]).

102

103 We report here the existence of a rare cell type, the olfactory rod cell, in the
104 developing zebrafish OE. Olfactory rod cells are characterised by a single actin-rich
105 apical projection. The morphology of the rod matches brief descriptions of similar
106 structures in the OE of several other fish species, many of which were previously
107 dismissed either as senescent forms of OSNs or as fixation artefacts. Using a variety
108 of imaging techniques and transgenic lines, including live imaging, we show that
109 zebrafish olfactory rod cells are present in living fish and can be detected from early
110 stages of larval development.

111

112 **Materials and Methods**

113 **Zebrafish husbandry**

114 Zebrafish strains used in this study were wild type (AB strain – ZFIN), *ift88^{tz288b}*
 115 (TsujiKawa & Malicki, 2004), *sox10^{m618}* (Dutton et al., 2001), *Tg(actb2:Lifeact-*
 116 *RFP)^{e115}* (Behrndt et al., 2012), *Tg(actb2:Lifeact-GFP)^{e114}* (Behrndt et al., 2012),
 117 *Tg(Xla.Tubb:jGCaMP7f)^{sq214}* (Chia et al., 2019), *Tg(elavl3:GCaMP6f)^{jf1}* (Dunn et al.,
 118 2016), *Tg(elavl3:H2B-GCaMP6s)^{jf5}* (Dunn et al., 2016), *Tg(pou4f3:GAP-GFP)^{s356t}*
 119 (Xiao et al., 2005) and *Tg(sox10:Lifeact-mRFPruby)^{sh630}* (this study). Homozygous
 120 *sox10^{-/-}* mutant larvae were identified by their lack of body pigmentation at 5 days
 121 post-fertilisation (dpf). Adult zebrafish were kept in a 10 hours dark/14 hours light
 122 cycle at 28.5°C and spawned by pair-mating or marbling (Aleström et al., 2019).
 123 Eggs were collected and staged according to standard protocols (Kimmel et al.,
 124 1995; Nüsslein-Volhard & Dahm, 2002), and raised in E3 medium (5 mM NaCl, 0.17
 125 mM KCl, 0.33 mM CaCl₂, 0.33 mM MgSO₄, with 0.0001% methylene blue at early
 126 stages) at 28.5°C. For controlling the developmental rate to obtain embryos at
 127 stages 34-46 hours post-fertilisation (hpf), embryos were incubated at 25°C or 34°C
 128 in accordance with Kimmel's formula, $H_T = h \div (0.055T - 0.57)$ (Kimmel et al.,
 129 1995). For live imaging, zebrafish were anaesthetised with 0.5 mM tricaine mesylate
 130 in E3.

131

132 **Generation of the *Tg(sox10:Lifeact-mRFPruby)* transgenic line**

133 The *-4725sox10:Lifeact-mRFPruby* construct was generated using the Gateway Tol2
 134 kit (Kawakami, 2007; Kwan et al., 2007). The p5E *-4725sox10* promoter (Dutton et
 135 al., 2008; Rodrigues et al., 2012), pME-*Lifeact-mRFPruby* (Riedl et al., 2008), and
 136 p3E polyA sequences were cloned into pDestTol2pA3 through an LR Clonase
 137 reaction. The 12.1 kb final plasmid was sequenced and injected into the AB strain.
 138 Injected embryos were grown to adulthood and crossed to AB. Transgenic progeny
 139 from one founder male were selected based on mRFPruby expression in the inner
 140 ear and grown to adulthood to generate a stable line. Embryos with bright
 141 fluorescence, presumed to be homozygous for the transgene, were chosen for
 142 imaging.

143

144 **Immunohistochemistry and phalloidin staining**

145 Zebrafish embryos and larvae were fixed in 4% paraformaldehyde (PFA) in
146 phosphate-buffered saline (PBS) for two hours at room temperature or overnight at
147 4°C. Zebrafish were washed three or more times with PBS, and permeabilised by
148 incubation in PBS-Triton X-100 (0.2% Triton for 32-48 hpf embryos, 1% Triton for
149 later stages) for several hours at 4°C until staining.

150

151 To visualise F-actin, zebrafish were stained with either Alexa Fluor 488 phalloidin
152 (Cell Signaling Technology; 1:150), Alexa Fluor 568 (Invitrogen ThermoFisher; 1:50),
153 or Alexa Fluor 647 phalloidin (Invitrogen ThermoFisher; 1:50) in PBS overnight at
154 4°C. After staining, zebrafish were washed four times in PBS over two or more hours
155 before imaging.

156

157 For antibody staining, after fixing and washing, zebrafish were washed a further
158 three times in PBS-0.2% Triton and incubated in blocking solution (10% sheep
159 serum in PBS-0.2% Triton for acetylated α -tubulin staining; 1% bovine serum
160 albumin (BSA) in PBS-0.1% Triton for SV2 staining) for 60 minutes at room
161 temperature. Primary antibodies were mouse IgG1 anti-acetylated α -tubulin antibody
162 (Sigma-Aldrich; 1:100) and mouse IgG1 anti-SV2 antibody (deposited in
163 Developmental Studies Hybridoma Bank by K. M. Buckley; 1:100). Staining was
164 carried out in blocking solution containing 1% dimethyl sulfoxide (DMSO; Sigma-
165 Aldrich) overnight at 4°C. Zebrafish were washed three times in PBS-0.2% Triton,
166 and a further four times over two or more hours. The secondary antibody was Alexa
167 647-conjugated goat anti-mouse IgG1 (Invitrogen ThermoFisher; 1:200). For double
168 stains with phalloidin, Alexa Fluor 488 phalloidin (1:150) and DMSO (1%) were
169 added together with the secondary antibody in blocking solution overnight at 4°C.
170 Zebrafish were then washed four times in PBS-0.2% Triton and stored at 4°C until
171 imaging. Controls with no primary antibody yielded no staining (not shown).

172

173 **Ototoxin treatment**

174 For neomycin treatment, a concentration of 500 μ M was chosen, as it was an
175 effective concentration used by Harris et al. (2003) for minimum lateral line hair cell
176 survival, as measured by DASPEI staining. A 5 mM solution was made by adding
177 neomycin trisulfate salt hydrate (Sigma-Aldrich) to MilliQ water and used at a 1:10
178 dilution in E3 fish medium. *Tg(pou4f3:GFP)* transgenic zebrafish were treated for 60

179 minutes at 28.5°C. An equivalent volume of MilliQ water in E3 was used for the
180 control group. Zebrafish were washed three times in fresh E3 and left at 28.5°C for
181 two hours. GFP signal was screened using widefield fluorescence microscopy to
182 analyse hair cell damage. Zebrafish were fixed and stained with Alexa Fluor 647
183 phalloidin as above.

184

185 **Fluorescence imaging**

186 For confocal imaging, fixed zebrafish embryos and larvae were mounted in 1.5% low
187 melting point (LMP) agarose in PBS, and live zebrafish were mounted in 1.5% LMP
188 agarose in E3 in WillCo glass-bottomed dishes (mounted in frontal view for 32-48
189 hpf, dorsal view for later stages). Zebrafish were imaged on a Zeiss LSM880
190 Airyscan confocal microscope equipped with a Plan-Apochromat 20×/0.8 M27 air
191 objective, LD LCI Plan-Apochromat 40×/1.2 Imm Korr DIC M27 water immersion
192 objective, or Plan-Apochromat 63×/1.4 oil DIC M27 objective. Images were acquired
193 in Airyscan SR mode, Airyscan Fast scan mode with SR sampling, or Airyscan Fast
194 scan mode with Opt sampling. Zebrafish were also imaged on a Zeiss LSM 800
195 attached to an upright microscope with a W Plan-Apochromat 40×/1.0 DIC M27 or
196 63×/1.0 M27 water dipping objective. The laser lines used were 488, 561, and 633
197 nm. Widefield imaging was performed on a Zeiss Axio Zoom.V16 fluorescence
198 stereo zoom microscope equipped with a Zeiss 60N-C 1" 1.0× C-mount and
199 AxioCam MRm camera. For fast-capture time series imaging, live zebrafish larvae
200 were mounted in 0.9% LMP agarose in E3 and imaged on a Zeiss Z1 Light-sheet
201 microscope, with 4% tricaine in E3 in the sample chamber. Imaging was performed
202 with a W Plan-Apochromat 20× objective using brightfield illumination and the 561
203 nm laser line. Images were acquired at a rate of 50.07 frames per second (fps).

204

205 **Scanning electron microscopy**

206 For scanning electron microscopy, *ift88* homozygous mutant and phenotypically wild-
207 type sibling larvae at 4 dpf were fixed overnight in 2.5% glutaraldehyde/0.1M sodium
208 cacodylate buffer. Samples were washed in buffer, post-fixed in 2% aqueous
209 osmium tetroxide for 1 hour, washed in buffer again and then dehydrated through a
210 graded ethanol series (50%, 75%, 95%, 100%) before being dried in a mixture of
211 50% hexamethyldisilazane (HMDS) in 100% ethanol. Final drying was in 100%

212 HMDS. After removal of the final HMDS wash, samples were left to dry in a fume
 213 hood overnight. Samples were mounted onto a pin stub using a Leit-C sticky tab and
 214 Leit-C mounting putty, gold-coated using an Edwards S150B sputter coater, and
 215 examined in a Tescan Vega3 LMU Scanning Electron Microscope at an operating
 216 voltage of 15 kV and imaged using a secondary electron detector.

217

218 ***Image processing, quantification, and statistical analyses***

219 Zeiss LSM880 Airyscan confocal images were subjected to Airyscan processing on
 220 Zen Black 2.3 software (Zeiss) using “Auto” Airyscan processing parameters. Further
 221 processing was performed on Fiji (Schindelin et al., 2012). 3D rendering was
 222 performed using the 3D Viewer plugin (Schmid et al., 2010) on Fiji. Rod projection
 223 lengths were measured in 3D from confocal images using Fiji, and calculated in
 224 Microsoft Excel using the PyT method (based on the Pythagorean theorem) from
 225 Dummer et al. (2016). All quantifications were exported into GraphPad Prism 8,
 226 which was then used for performing statistical analyses and making graphs.

227

228 Statistical analyses were carried out in GraphPad Prism 8. Datasets were
 229 considered normally distributed if they passed at least one of four normality tests
 230 (Anderson-Darling, D’Agostino & Pearson, Shapiro-Wilk, and Kolmogorov-Smirnov
 231 tests). Statistical tests used are stated in the figure legends. Bars on graphs indicate
 232 mean \pm standard error of the mean (S.E.M.), unless stated otherwise. *P* values are
 233 indicated as follows: *P* > 0.05 (not significant, ns), *P* < 0.05 (*), *P* < 0.01 (**), *P* <
 234 0.001 (***), *P* < 0.0001 (****).

235

236 For mapping spatial distributions of rod cells within the olfactory pit, 2D maximum
 237 intensity projection images were imported into the Desmos Graphing Calculator
 238 (desmos.com). The positions and sizes of the images were adjusted to align the rims
 239 of olfactory pits with an ellipse to fit the shape of the rim, defined by $\frac{(x-35)^2}{5} +$
 240 $\frac{(y-33)^2}{10} = 7.6^2$. The positions of the base of each rod, relative to the ellipse, were
 241 plotted as coordinates onto the graph. The resulting graphs were exported as .png
 242 image files.

243

244 Figures were prepared using Adobe Photoshop.

245 **Results**246 ***Actin-rich rod-like apical projections, distinct from microvilli and cilia, are***
247 ***present in the olfactory epithelium of larval and juvenile zebrafish***

248 Staining of the wild-type larval and juvenile zebrafish OE with fluorescently-
249 conjugated phalloidin, which binds to F-actin, reveals the presence of several actin-
250 rich rod-like projections ('rods') in each olfactory pit (Figure 1A-B'). These projections
251 differ in number, distribution, size and morphology from any of the described apical
252 projections of zebrafish OSNs. The projections extend from below the apical surface
253 of the OE and project about 10 μm above it, tapering to a point. This is an order of
254 magnitude longer than OSN microvilli, which are typically 0.5-0.8 μm in length
255 (Hansen & Zeiske, 1998). Olfactory rods are shorter than the surrounding phalloidin-
256 negative olfactory cilia (Fig. 1C-D'), and do not label with an anti-acetylated α -tubulin
257 antibody (Figure 1C-C'''). Rods are not evenly distributed across the OE, but are
258 mostly clustered posterolaterally in each olfactory pit, although there is variation
259 between individuals (Figure 1E). At low magnification, the olfactory rods appear
260 similar to the actin-rich stereociliary bundle of mechanosensory hair cells of the inner
261 ear and lateral line. However, higher magnification images reveal that the olfactory
262 rod is not oligovillous, but appears to be a single structure (Figure 1B', C''', D'). This
263 contrasts with the stepped array of multiple stereocilia present on the apical surface
264 of mechanosensory hair cells (Figure 1F).

265

266 To characterise the timing of appearance and development of the olfactory rods
267 during embryonic and larval stages, we stained fixed samples from 36 hpf, just after
268 formation of the olfactory pits (Hansen & Zeiske, 1993), to 5 dpf. Occasional rods
269 were present in olfactory pits at 36 hpf, but were only consistently present beyond 46
270 hpf (Figure 2A, B). Although the number of rods per olfactory pit varied at each
271 stage, the average number increased over time. By 5 dpf, each olfactory pit
272 contained 10.7 ± 2.9 (mean \pm standard deviation, s.d.) rods (Figure 2B). After
273 measuring the rods in 3D, we found an increase in projection length (from the base
274 of the phalloidin-positive projection to the tip) from 36 hpf to 5 dpf, with the most
275 significant increase occurring by 48 hpf, despite a relatively large range in length at
276 each stage. At 5 dpf in fixed samples, the mean projection length was 10.4 ± 2.2
277 (s.d.) μm , with the largest measuring 17.5 μm (Figure 2C).

278

279 Olfactory rod cell projections can develop in the absence of olfactory cilia

280 As described above, olfactory rods differ from olfactory cilia in terms of size, shape,
281 cytoskeletal composition, and distribution in the OE. We therefore hypothesised that
282 olfactory rod cell projections would not be affected by mutations that disrupt the
283 formation of cilia. To test this, we examined fish mutant for *ift88*, which codes for a
284 component of the intraflagellar transport machinery necessary for the normal
285 formation and maintenance of cilia (Tsujikawa & Malicki, 2004). A phalloidin stain
286 revealed that olfactory rods were present in the OE of *ift88*^{-/-} mutants at 5 dpf (Figure
287 3A, B).

288

289 The absence of cilia in *ift88*^{-/-} mutants allowed us to examine morphology of the rods
290 using scanning electron microscopy (SEM). In the phenotypically wild-type sibling
291 OE, the rods were almost completely obscured by olfactory cilia, with only the
292 occasional tip of a projection visible (Figure 3C-E”). However, SEM images of the
293 olfactory pit of *ift88*^{-/-} mutants at 4 dpf, which lack cilia, revealed the presence of rod-
294 like projections with a similar size, number, smoothly tapering morphology, and
295 spatial distribution to the actin-rich projections described above (Figure 3F-I). At their
296 base, olfactory rods are wider in diameter (about 0.6 µm) than the olfactory cilia in
297 wild-type larvae (0.2 µm in diameter, as is typical for many cilia). We conclude that
298 olfactory rods can develop in the absence of cilia.

299

300 Olfactory rods can be labelled in the live larva

301 To visualise olfactory rods in live larvae, we imaged the *Tg(actb2:Lifeact-RFP)*
302 transgenic line at 4 and 6 dpf, and *Tg(actb2:Lifeact-GFP)* at 5 dpf (Behrndt et al.,
303 2012). We found fluorescent apical projections in the olfactory pits of live larvae in all
304 cases (*N* of fish = 4) (Figure 4A-C, Supplementary Movie 1). These matched the
305 size, shape, and posterolateral distribution of rod cells present in fixed samples
306 (Figure 4D, E). Despite potential shrinkage due to fixation, there was no overall
307 difference in the lengths of projections between live and fixed samples (Figure 4E).
308 The zig-zag pattern exhibited by RFP-positive olfactory rods in raster-scanned
309 images of live larvae suggested that rods were moving during image capture (Figure
310 4B). Fast-capture time series imaging of the *Tg(actb2:Lifeact-RFP)* transgenic line
311 allowed us to observe that the projection oscillates (Supplementary Movie 2),
312 possibly as a result of ciliary beating.

313

314 Neuronal promoters drive reporter expression in olfactory rod cells

315 To test whether rod cells are neuronal, we imaged two transgenic lines that have
316 broad neuronal expression of cytoplasmic fluorescent reporters –
317 *Tg(Xla.tubb:jGCaMP7f)* (Chia et al., 2019) (*N* of olfactory pits = 4) and
318 *Tg(elavl3:GCaMP6f)* (Dunn et al., 2016) (*N* = 5). Dendrites and dendritic knobs of
319 OSNs were clearly labelled by both lines. In some examples, we observed faintly-
320 labelled projections extending from the surface of the olfactory epithelium, with a
321 similar length and morphology to olfactory rods (Figure 5A-B). Imaging of double-
322 transgenic *Tg(elavl3:GCaMP6f);Tg(actb2:Lifeact-RFP)* larvae at 5 dpf indicated that
323 rods were GCaMP6f-positive (*N* of fish = 3; Figure 5C-C’). These observations
324 suggest that olfactory rod cells may be neurons.

325

326 Olfactory rod cells are not hair-cell-like cells

327 Given the superficial similarity in appearance of the olfactory rod to hair-cell
328 stereocilia in phalloidin stains, and a report of a rare cell type bearing stereocilia-like
329 microvilli in the rat OE (Menco & Jackson, 1997), we tested whether there is any
330 similarity between olfactory rod cells and mechanosensory hair cells of the inner ear
331 and lateral line. As shown in Figures 1 and 3, the zebrafish olfactory rod appears to
332 be a single structure rather than a collection of microvilli or stereocilia. To test
333 whether olfactory rod cells express sensory hair cell markers, we performed an
334 Alexa-phalloidin co-stain on the *Tg(pou4f3:GFP)* transgenic line, a known marker for
335 hair cells (Xiao et al., 2005). At 5 dpf, the stereociliary bundle of lateral line
336 neuromast hair cells was clearly marked by both GFP and phalloidin, which acted as
337 our positive control (Figure 6A-A’). However, the GFP did not co-localise with the
338 phalloidin signal in the olfactory rods, or in the cell body beneath a phalloidin-positive
339 rod (Figure 6B-B’). Additionally, confocal images of an antibody stain against
340 synaptic vesicle protein 2 (SV2), a marker of synaptic terminals of afferent neurons
341 contacting secondary sensory cells such as mechanosensory hair cells and taste
342 receptors (Buckley & Kelly, 1985; Portela-Gomes et al., 2000; Zachar & Jonz, 2012),
343 showed that SV2-positive synapses are not present in the olfactory epithelia at 4 dpf
344 (Figure 6C, D). This contrasted with strong staining of synaptic terminals in cranial
345 lateral line neuromasts in the vicinity of the olfactory pits (Fig. 6C, D).

346

347 We next investigated whether treatment with neomycin, an aminoglycoside antibiotic
348 and well-described ototoxin, has the same damaging effect on olfactory rod cells as
349 on lateral line hair cells (Harris et al., 2003). Following neomycin treatment at 500
350 μ M for 60 minutes on 3 dpf *Tg(pou4f3:GFP)* larvae, lateral line hair cells were lost or
351 severely damaged, as determined by a decrease in the number of GFP-positive cells
352 in both cranial and trunk neuromasts and a change in morphology of any remaining
353 cells (Figure 6E-H). By contrast, olfactory rods appeared unaffected (Figure 6I, J),
354 with no significant change in the number of rods present in each olfactory pit (Figure
355 6K). Taken together, the smooth appearance of the olfactory rods, lack of hair cell
356 and synaptic vesicle marker expression, and resistance to neomycin indicate that
357 olfactory rod cells are not closely related to hair cells.

358

359 ***A sub-population of olfactory rod cells expresses a Lifeact transgene driven***
360 ***by the sox10 promoter***

361 Sox10 is a known marker of both neural crest and otic epithelium (Dutton et al.,
362 2001). Robust transgene expression driven by the *sox10* promoter has been
363 reported in the OE and other tissues in the zebrafish (Mongera et al., 2013; Saxena
364 et al., 2013). We have generated a *Tg(sox10:Lifeact-mRFPruby)* transgenic line to
365 visualise actin localisation and dynamics in the live embryo in *sox10*-expressing
366 tissues. As reported for the *Tg(sox10:eGFP)* transgene (Saxena et al., 2013), we
367 observed OSNs expressing *Tg(sox10:Lifeact-mRFPruby)* in the OE at 4 and 5 dpf;
368 based on morphology, most of these cells were microvillous neurons. However,
369 staining with Alexa-phalloidin on fixed samples revealed the co-expression of Lifeact-
370 mRFPruby in a sub-population of phalloidin-positive olfactory rod cell projections
371 (Figure 7A-B''). Not all olfactory rod cells expressed the transgene; an average of
372 64.4% of rod cells marked by phalloidin (*N* of olfactory pits = 5, *n* of olfactory rods =
373 59) also expressed Lifeact-mRFPruby (Figure 7C). As for the rods labelled with
374 Lifeact-RFP, rods labelled with Lifeact-mRFPruby oscillated (Supplementary Movie
375 3).

376

377 The sparse expression of the *Tg(sox10:Lifeact-mRFPruby)* transgene allowed us to
378 visualise the morphology of the cell body of olfactory rod cells and ask whether they
379 have an axon. Lifeact-mRFPruby-expressing cell bodies were positioned apically in
380 the OE and were rounded in shape (Figure 7B-B'', E). They were morphologically

381 distinct from the well-described microvillous neurons (Figure 7D, E) as well as
382 ciliated and crypt OSNs. The axons of microvillous OSNs were visible in those cells
383 labelled by the transgene (Figure 7D). However, with this marker, we were unable to
384 observe an axon extending from the cell body of olfactory rod cells (N of olfactory
385 pits = 5, n of cells = 9; Figure 7E).

386

387 To test whether the development of olfactory rod cells is dependent on *sox10*
388 function, we stained *sox10*^{-/-} homozygous mutants (Dutton et al., 2001) with Alexa-
389 phalloidin. Olfactory rods were present in *sox10*^{-/-} mutants at 5 dpf, but variable in
390 number (N of olfactory pits = 8, n of olfactory rods = 53; Figure 8). Taken together,
391 the data from *Tg(sox10:Lifeact-mRFP_{ruby})* transgenic and *sox10*^{-/-} mutant larvae
392 indicate that *sox10* function is not essential for the formation of olfactory rods.

393

394 Discussion

395 Through the use of phalloidin staining, immunohistochemistry, transgenic zebrafish
396 lines, SEM and high-resolution fluorescence confocal imaging, we have identified a
397 rare cell type, the olfactory rod cell, in the zebrafish larval and juvenile OE. Olfactory
398 rod cells, which have not previously been described in zebrafish to our knowledge,
399 are morphologically distinct from the well-characterised OSNs and other known cell
400 types in terms of their apical projections, cell shape, and distribution and positioning
401 within the OE.

402

403 *The olfactory rod: an actin-rich apical projection*

404 The spectacular actin-rich projection of the olfactory rod cell adds to the rich
405 repertoire of known F-actin-based cellular specialisations, which include microvilli,
406 stereocilia, lamellipodia, filopodia, cytonemes and microridges (reviewed in [Heath &
407 Holifield, 1991; Theriot & Mitchison, 1991]; Ramírez-Weber & Kornberg, 1999; Pinto
408 et al., 2019; Inaba et al., 2020). Many classes of sensory cell, in both fish and
409 mammals, bear actin-rich mechano- or chemosensory microvillous projections,
410 including the stereocilia of sensory hair cells (Tilney et al., 1980; reviewed in
411 [Gillespie & Müller, 2009; Barr-Gillespie, 2015]), and the microvilli of olfactory and
412 vomeronasal microvillous neurons, solitary chemosensory cells (SCCs) of the skin
413 and barbel (Kotrschal et al., 1997; Finger et al., 2003), taste bud cells (Hansen et al.,
414 2002; Zachar & Jonz, 2012), spinal cerebrospinal fluid-contacting neurons (CSF-
415 cNs; Djenoune et al., 2014; Desban et al., 2019), Merkel cells, retinal Müller glia
416 (Sekerková et al., 2004), and the brush and tuft cells of mammalian respiratory and
417 intestinal epithelia, respectively (reviewed in [Reid et al., 2005; Schneider et al.,
418 2019]). As a single structure with a smoothly tapering morphology, the zebrafish
419 olfactory rod differs from these oligovillous structures. Adult zebrafish SCCs, found
420 distributed over the entire body surface (Kotrschal et al., 1997), and mature light
421 cells of the zebrafish taste bud (Hansen et al., 2002) each bear a single microvillus,
422 but at 1-3 μm in length, these are much shorter than the olfactory rods we describe.

423

424 A note on terminology: olfactory rod cells are distinct from rodlet cells, which have
425 been reported in many different epithelial tissues (including OE) of marine and
426 freshwater fish, including zebrafish, and contain several intracellular electron-dense
427 rodlets within a thick cuticular-like wall (Bannister, 1966; reviewed in [Morrison &

428 Odense, 1978]; Hansen & Zeiske, 1998 Dezfuli et al., 2007; DePasquale, 2020).
429 Recently, phalloidin staining has demonstrated that the rodlets, which can be
430 extruded from the cell, are not composed of F-actin (DePasquale, 2020). Thus,
431 zebrafish olfactory rod cells, which are unique to the OE at the larval stages we have
432 described, are not related to rodlet cells.

433

434 **Olfactory rod cells in other teleost species**

435 Previous studies have provided brief descriptions of similar cell types in other teleost
436 species, including the common minnow (Bannister, 1965), several eel species
437 (Schulte, 1972; Yamamoto & Ueda, 1978), goldfish (Breipohl et al., 1973; Ichikawa &
438 Ueda, 1977), rainbow trout (Rhein et al., 1981), common bleak (Hernádi, 1993),
439 catfish (Datta & Bandopadhyay, 1997), and several cave fish and cave loach species
440 (Waryani et al., 2013; Waryani et al., 2015; Zhang et al., 2018).

441

442 Using transmission electron microscopy (TEM), Bannister (1965) reported sparsely-
443 populated rod-shaped protrusions, approximately 4 μm in length and shorter than
444 surrounding sensory and non-sensory olfactory cilia, in the OE of adult (3.7 cm)
445 common minnow (*Phoxinus phoxinus*). Here, the rod-like projection consisted of
446 several bundles of fibres, consistent with the appearance of F-actin, extending from
447 deep within the cell (Bannister, 1965). Similarly, using TEM and SEM respectively,
448 Schulte (1972) and Yamamoto and Ueda (1978) reported the presence of olfactory
449 rod cells in the OE of several adult eel species: European eel (*Anguilla anguilla*),
450 Japanese eel (*A. japonica*), white-spotted conger (*Conger myriaster*), buffoon snake
451 eel (*Microdonophis erabo*), and brutal moray (*Gymnothorax kikado*). In European
452 eels, the cells were described as a receptor with a single rod-shaped appendage,
453 measuring 0.8 μm in diameter and extending 4 μm above the apical surface of the
454 epithelium (Schulte, 1972). Olfactory rods in the other four species measured 1 μm
455 in diameter and 10 μm in length. Rods were either found to exist solitarily or in a
456 group; interestingly, it was noted that olfactory cilia were sparse in areas where rods
457 occurred in a group (Yamamoto & Ueda, 1978).

458

459 More recent reports include comparisons of the surface structures of olfactory
460 epithelia in different adult cave fish and loaches. SEMs in *Sinocyclocheilus jii* and *S.*
461 *furcodorsalis* cave fish, and in *Oreonectes polystigmus* and *O. guananensis* cave

462 loaches revealed that olfactory rods were clustered in different regions of olfactory
463 rosette lamellae (Waryani et al., 2013; Waryani et al., 2015). Another SEM study on
464 the variations in olfactory systems of adult cave fish species of different habitats
465 reported not just one, but three different cell types all classified as 'rod cilia' in the
466 olfactory epithelia of *S. anshuiensis* and *S. tianlinensis*. The first cell type had a long
467 base with an oval apex, the second contained an oval base with a thin apex, while
468 the third was rod-shaped and thin from base to tip, measuring 2.01-3.08 μm in length
469 (Zhang et al., 2018). Despite the shorter length, this third type appeared
470 morphologically consistent with zebrafish olfactory rod cells. Unlike other teleosts,
471 olfactory rod cells were reported as the dominant cell type over ciliated and
472 microvillous neurons in the OE of *S. jii* (Waryani et al., 2013). This may be an
473 example of the known compensatory enhancement of the olfactory system in blind
474 morphs of cave fish (Bibliowicz et al., 2013; reviewed in [Krishnan & Rohner, 2017]).

475

476 Although there appear to be variations in the numbers and sizes of olfactory rod cells
477 reported in these other teleost species, some of these cells may be homologous to
478 the olfactory rod cells we describe in zebrafish larvae. However, all of these previous
479 studies were limited to fixed adult samples by means of TEM and SEM, and none
480 have tested or confirmed the molecular composition of the rod.

481

482 ***Olfactory rod cells differ from known olfactory sensory neurons or secondary*** 483 ***receptor cells***

484 We have detected weak expression of cytoplasmic fluorescent markers driven by
485 neuronal promoters in olfactory rod cells, suggesting they are a type of OSN.
486 However, we were unable to detect an axon in nine individual olfactory rod cells
487 imaged with a Lifeact-mRFP_{ruby} transgene at 4-5 dpf. Of note, Ichikawa and Ueda
488 (1977) performed olfactory nerve bundle transection in adult goldfish to determine
489 which cell types are olfactory receptors. As expected, transection caused retrograde
490 degeneration of both ciliated and microvillous OSNs. Olfactory rod cells, however,
491 were still identifiable by SEM in the OE 10 days after nerve transection. The authors
492 concluded that adult goldfish olfactory rod cells are not sensory receptor cells;
493 however, the result could also indicate differences in regeneration kinetics between
494 different neuronal cell types.

495

496 The shape, position and actin-rich projection of zebrafish olfactory rod cells,
497 however, do have some similarities to those of kappe OSNs, identified to date only in
498 adult zebrafish OE (Ahuja et al., 2014). The actin-positive apical specialisation of
499 kappe cells differs in morphology from the rods we describe, and is thought to
500 consist of microvilli, although this has not been confirmed at an ultrastructural level.
501 Kappe neurons were reported to have an axon, and were thus interpreted as a class
502 of OSN (Ahuja et al., 2014). It will be important to determine whether olfactory rod
503 cells are present in the adult zebrafish OE and how their morphology relates to
504 kappe cells.

505

506 Olfactory rod cells do not appear to share characteristics with sensory hair cells. As
507 a monovillous structure, the olfactory rod is quite unlike the stereociliary bundle of a
508 hair cell; moreover, the cells do not express known hair cell markers, and are
509 insensitive to the aminoglycoside antibiotic neomycin, a potent ototoxin. In addition,
510 the entire olfactory pit was negative for the synaptic vesicle and neuroendocrine cell
511 marker SV2 (Buckley & Kelly, 1985; Portela-Gomes et al., 2000), indicating that
512 olfactory rod cells are unlikely to be secondary sensory cells innervated by afferent
513 neurons.

514

515 ***Olfactory rod cells as artefact***

516 Since the first report of olfactory rod cells, several studies have proposed that they
517 may represent senescent forms of OSNs or fixation artefacts (Muller & Marc, 1984;
518 Moran et al., 1992; reviewed in [Hansen & Zielinski, 2005]). A study in the goldfish
519 (*Carassius auratus*) and channel catfish (*Ictalurus punctatus*), using TEM, SEM and
520 filling with horseradish peroxidase, concluded that rods are most likely a result of
521 fusion of olfactory cilia or microvilli – an indicator of ageing OSNs (Muller & Marc,
522 1984). A later study on the ultrastructure of olfactory mucosa in brown trout (*Salmo*
523 *trutta*) also classified rods as products of the fusion of olfactory cilia during fixation
524 (Moran et al., 1992). Indeed, TEM images in this study showed multiple ciliary
525 axonemes surrounded by a single membrane (Moran et al., 1992). The presence of
526 such fixation artefacts has led to frequent dismissal of olfactory rod cells in the
527 literature, for example in juvenile and adult European eels (Sola et al., 1993). In the
528 zebrafish, however, the olfactory rods we describe are clearly not a fixation artefact,
529 as they are present in the live larva. Moreover, they are not formed by fusion of cilia,

530 as the olfactory rods are F-actin-positive, do not stain with an anti-acetylated α -
531 tubulin antibody, and are present in *ift88*^{-/-} mutants, which lack cilia.

532

533 **Possible functions of olfactory rod cells**

534 Actin-rich projections on sensory cells are known to have mechanosensory
535 (reviewed in [Gillespie & Müller, 2009]), chemosensory (Höfer & Drenckhahn, 1999;
536 Hansen et al., 2002; Zachar & Jonz, 2012), or multimodal functions (for example in
537 CSF-cNs in zebrafish; Djenoune et al., 2014; Desban et al., 2019). A
538 mechanosensory role for zebrafish olfactory rod cells, for example in detecting ciliary
539 movement or ciliary-driven fluid flow, or a chemosensory role in detecting odorants,
540 could aid olfactory perception in the larva. The olfactory rod cell has previously been
541 referred to as a sensory receptor cell in studies of other teleost species (Bannister,
542 1965; Schulte, 1972; Breipohl et al., 1973; Waryani et al., 2013; Waryani et al.,
543 2015). Although functions were not explicitly investigated, Bannister (1965)
544 speculated that based on their internal structure, minnow olfactory rod cells perform
545 specific chemosensory roles. Another possibility is that they could correspond to
546 brush or tuft cells in air-breathing mammals, which have important roles in immunity
547 (Andres, 1975; reviewed in [Reid et al., 2005]; Howitt et al., 2016; reviewed in
548 [Schneider et al., 2019]). These ideas remain to be tested.

549

550 **Possible origins of olfactory rod cells**

551 Our work does not address the developmental origin of olfactory rod cells, but it is of
552 interest that they express a *sox10*-driven transgene, albeit in a mosaic fashion.
553 *Sox10* mRNA is frequently described as a neural crest marker, but is also expressed
554 strongly in otic epithelium (Dutton et al., 2001), a placodally-derived tissue. The use
555 of *sox10*-driven transgenic lines to identify neural crest derivatives remains
556 controversial. Expression of a *sox10:eGFP* transgene together with photoconversion
557 studies has led to the conclusion that a subpopulation of microvillous neurons in the
558 OE is derived from neural crest (Saxena et al., 2013), and use of an inducible
559 *sox10:ER^{T2}-Cre* transgenic line has identified previously 'contested' neural crest
560 derivatives, including cells in the sensory barbels (Mongera et al., 2013). However,
561 using lineage reconstruction through backtracking and photoconversion experiments,
562 Aguillon et al. (2018) have argued that all olfactory neurons, including OSNs and
563 gonadotropin-releasing hormone 3 (GnRH3) cells, are derived entirely from

564 preplacodal progenitors. We think it likely that olfactory rod cells are placodally-
565 derived, but this remains to be confirmed.

566

567 The *Tg(sox10:Lifeact-mFRPruby)* line is expressed in a subset of both olfactory rod
568 cells and of microvillous neurons, with variation in the proportion of expressing cells
569 between individuals. This could reflect true heterogeneity in the olfactory rod cell and
570 microvillous neuron populations, or it could be a result of mosaic or leaky expression
571 of the transgene. Mosaic expression is typical for many transgenes (Mosimann et al.,
572 2013), while leaky expression, which can be explained through the lack of
573 appropriate silencer elements (Jessen et al., 1999), is suspected for the *sox10*
574 promoter fragment used in our transgenic construct (reviewed in [Tang & Bronner,
575 2020]). Nevertheless, the *Tg(sox10:Lifeact-mFRPruby)* line has proved a fortuitous
576 tool for visualising olfactory rod cells in the live larva.

577

578 **Concluding remarks**

579 As a key model organism for the study of the olfactory system (reviewed in [Kermen
580 et al., 2013; Calvo-Ochoa & Byrd-Jacobs, 2019]), a complete inventory of the cell
581 types present in the zebrafish OE will be an important resource and reference point
582 for further study. Olfactory dysfunction can signify underlying cellular disorders and
583 can also be implicated in neurodegenerative diseases (reviewed in [Whitlock, 2015];
584 Bergboer et al., 2018). OSNs project directly to the OB, and thus provide an entry
585 route for pathogens to the brain (reviewed in [Dando et al., 2014]). Cells in the OE
586 can themselves be damaged by viral infection, leading to a reduction, change, or
587 loss of sense of smell, a phenomenon that has attracted much recent attention due
588 to the damaging action of SARS-CoV-2 on the human olfactory system (Brann et al.,
589 2020; Gupta et al., 2020). In the zebrafish, new functions of the OE, such as the
590 detection of sodium and chloride ions (Herrera et al., 2020), continue to be
591 uncovered. Addressing the functions of the olfactory rod cell will be an important next
592 step.

593

594 **Ethics**

595 All zebrafish work in Sheffield was undertaken under licence from the UK Home
596 Office and according to recommended standard husbandry conditions (Aleström et
597 al., 2019). All experiments in Singapore were performed under guidelines approved
598 by the Institutional Animal Care and Use Committee of Biopolis.

599

600 **Conflict of Interest**

601 The authors declare no competing interests.

602

603 **Author Contributions**

604 Designed the research: KYC, TTW, SJJ. Conducted the experiments: KYC, SJJ,
605 TTW, SB, NJvH, MM, CJH. Data analysis: KYC, SJJ, TTW. Writing (original draft):
606 KYC, TTW; writing (review and editing): KYC, TTW, SJJ, with additional
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608

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615 sheet microscopy. Work in the SJ lab was funded by a start-up grant from the Lee
616 Kong Chian School of Medicine.

617

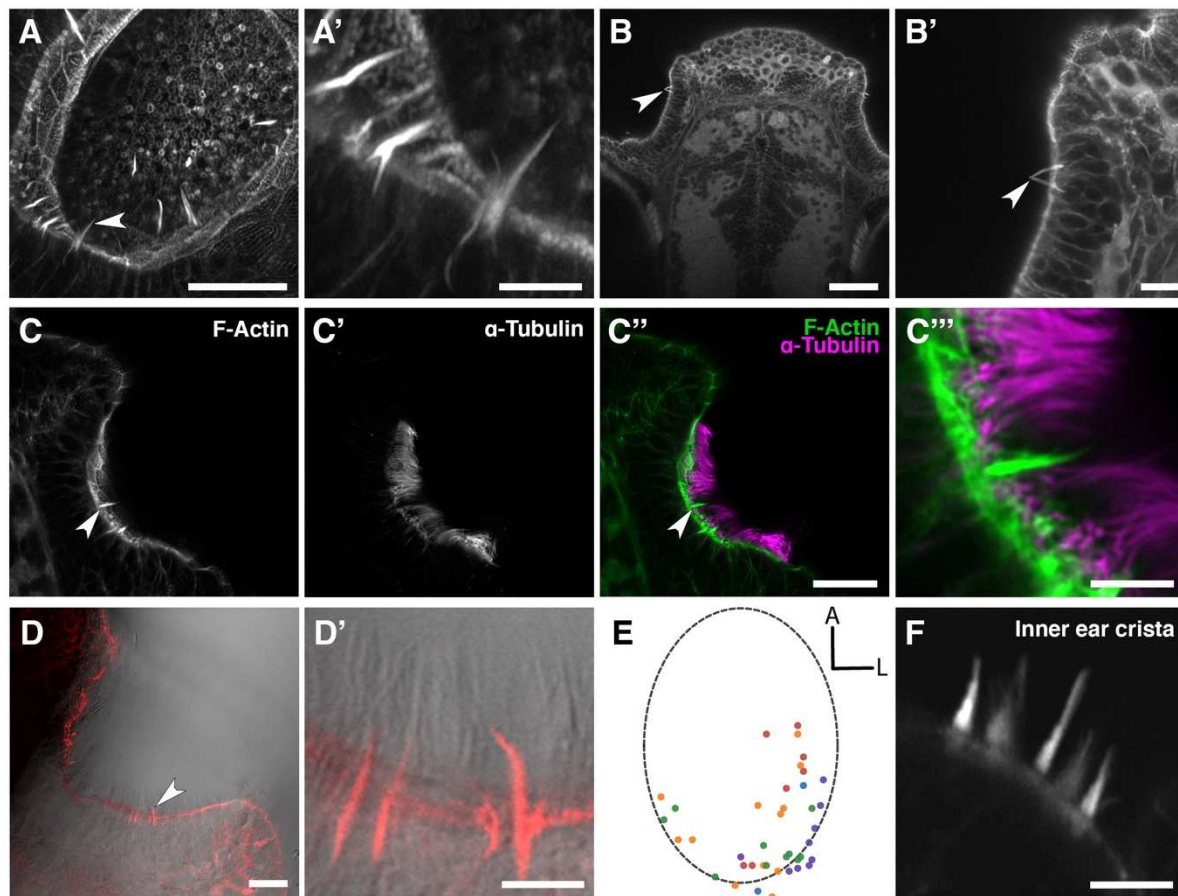
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627

628 **Figures**

629



630

631

632 **Figure 1. Phalloidin staining reveals the presence of actin-rich rod-like projections, distinct**
 633 **from microvilli and cilia, in the zebrafish larval and juvenile olfactory epithelium.**

634 (A) Maximum intensity projection of an Airyscan confocal image of phalloidin stain in an olfactory pit of
 635 a 5 dpf wild-type larva; anterior to the top right, lateral to the bottom right. Arrowhead marks one
 636 example olfactory rod. Scale bar = 20 μm . (A') Enlargement of olfactory rods in A. Scale bar = 5 μm .

637 (B) Dorsal view low power image of phalloidin stain in the head of an 18 dpf (5 mm) wild-type juvenile
 638 zebrafish; anterior to the top. Arrowhead marks the position of two olfactory rods in an olfactory pit.

639 Scale bar = 50 μm . (B') Enlargement of OE in B. Arrowhead marks two olfactory rods. Scale bar = 10

640 μm . (C-C'') Airyscan confocal image of Alexa-phalloidin signal (C), acetylated α -tubulin
 641 immunohistochemistry signal (C'), and merged signals (C'') in an olfactory pit of a 4 dpf wild-type

642 larva; anterior to the top, lateral to the right. Arrowhead marks one example olfactory rod. Scale bar =
 643 20 μm . (C''') Enlargement of olfactory rod in C''. Scale bar = 5 μm .

644 (D) Differential interference contrast (DIC) image and phalloidin stain (red) in an olfactory pit of a 5 dpf wild-type larva;
 645 anterior to the top, lateral to the right. Arrowhead marks one example olfactory rod. Scale bar = 20 μm .

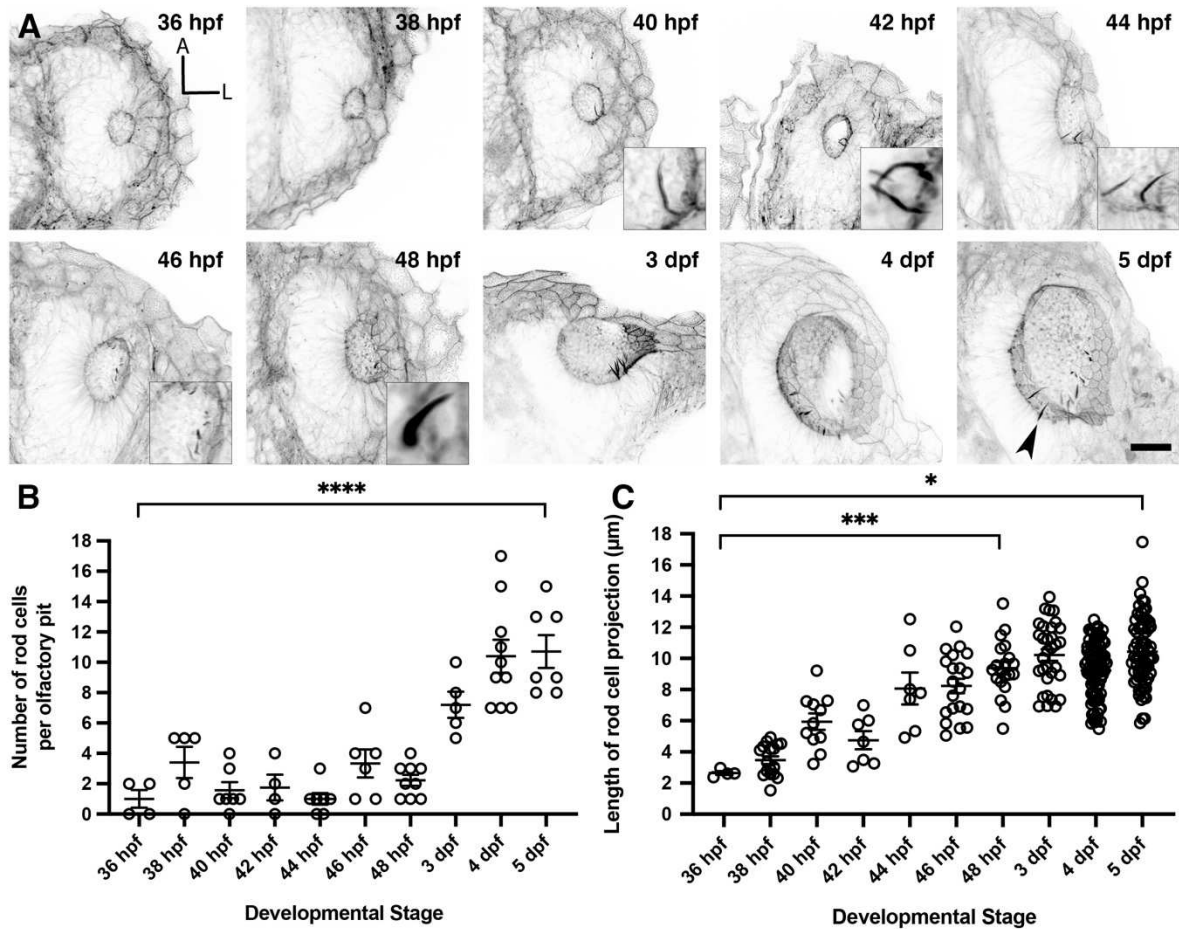
646 (D') Enlargement of olfactory rods in D. Surrounding olfactory cilia are visible and unlabelled by Alexa-
 647 phalloidin. Scale bar = 5 μm .

648 (E) A map of the positions of olfactory rod cell projection bases in
 649 olfactory pits of 4 dpf wild-type larvae (N of olfactory pits = 5), based on 2D maximum intensity
 650 projections of confocal images of phalloidin stains; anterior 'A' to the top, lateral 'L' to the right. One

651 dot represents one olfactory rod. Different coloured dots represent rods from different larvae. (F)

652 Airyscan confocal image of phalloidin stain in an inner ear crista of a 5 dpf wild-type larva. Hair cell
 653 stereocilia are labelled with Alexa-phalloidin, and are arranged in a stepped array. In the stereociliary
 654 bundle on the extreme left, four different stereociliary lengths are visible (compare with A'). Scale bar

655



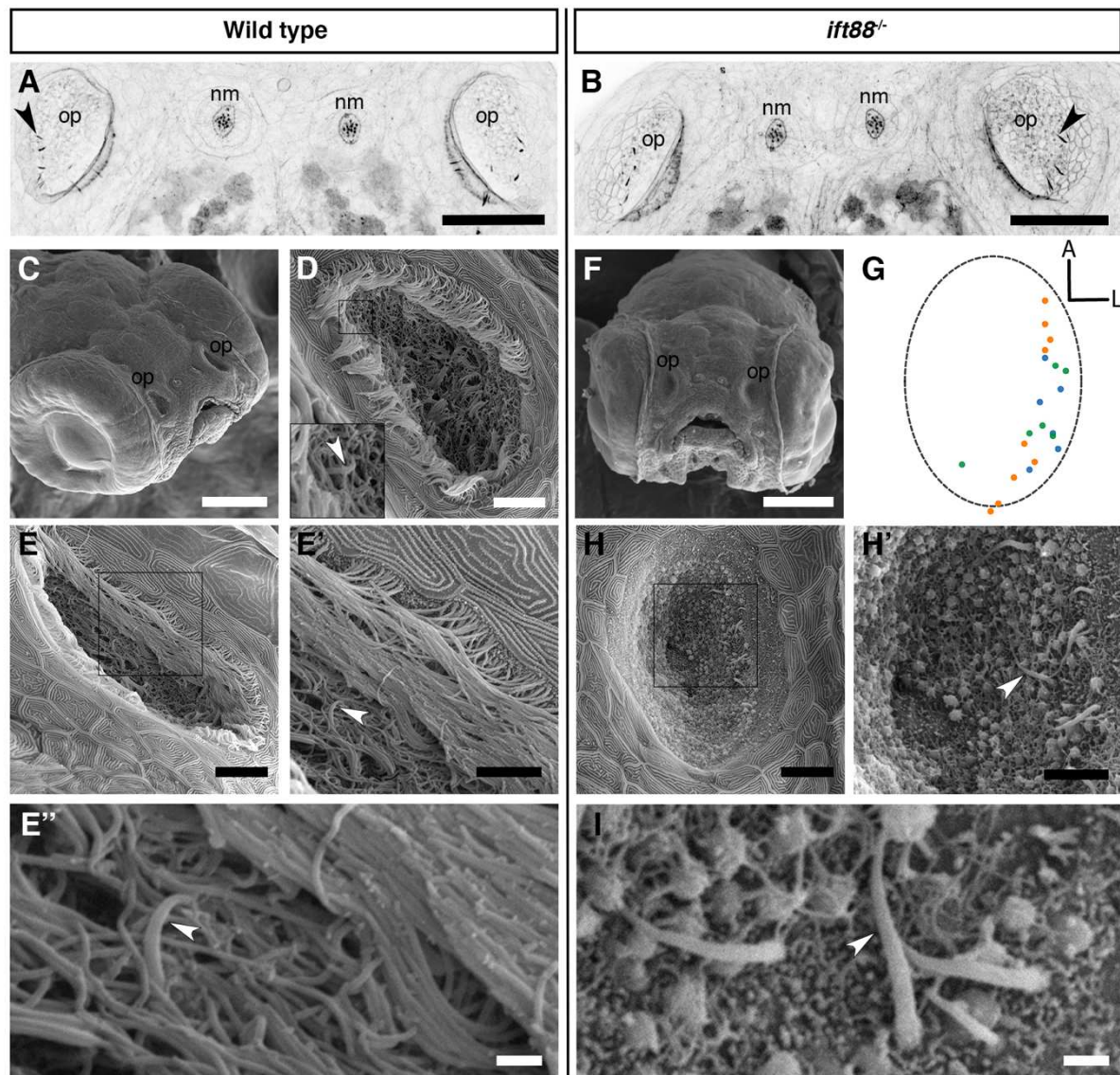
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657

658 **Figure 2. Olfactory rod cells arise early during zebrafish olfactory pit development.**

659 (A) Maximum intensity projections of Airyscan confocal images showing the wild-type development of
 660 olfactory pit and olfactory rod cells at various embryonic and larval stages, using Alexa-phalloidin as a
 661 marker; anterior 'A' to the top, lateral 'L' to the right. Grayscale values from the original fluorescence
 662 image have been inverted. Arrowhead marks one example olfactory rod. Scale bar = 20 μm. Selected
 663 inserts show olfactory rods at higher magnification. (B) The change in number of olfactory rod cells
 664 per olfactory pit during embryonic development – 36 hpf (N of olfactory pits = 4), 38 hpf (N = 5), 40 hpf
 665 (N = 7), 42 hpf (N = 4), 44 hpf (N = 7), 46 hpf (N = 6), 48 hpf (N = 9), 3 dpf (N = 5), 4 dpf (N = 10), and
 666 5 dpf (N = 7). Bars indicate mean \pm S.E.M. for each stage. Linear regression analysis; **** indicates P
 667 < 0.0001 . (C) The change in lengths of olfactory rod cell projections during embryonic development –
 668 36 hpf (N of olfactory pits = 2, n of olfactory rods = 4), 38 hpf (N = 4, n = 17), 40 hpf (N = 6, n = 11),
 669 42 hpf (N = 3, n = 7), 44 hpf (N = 5, n = 7), 46 hpf (N = 6, n = 20), 48 hpf (N = 9, n = 20), 3 dpf (N = 5,
 670 n = 32), 4 dpf (N = 10, n = 82), and 5 dpf (N = 8, 71). Bars indicate mean \pm S.E.M. for each stage.
 671 Linear regression analysis; * indicates $P = 0.0251$, *** indicates $P = 0.0009$.

672



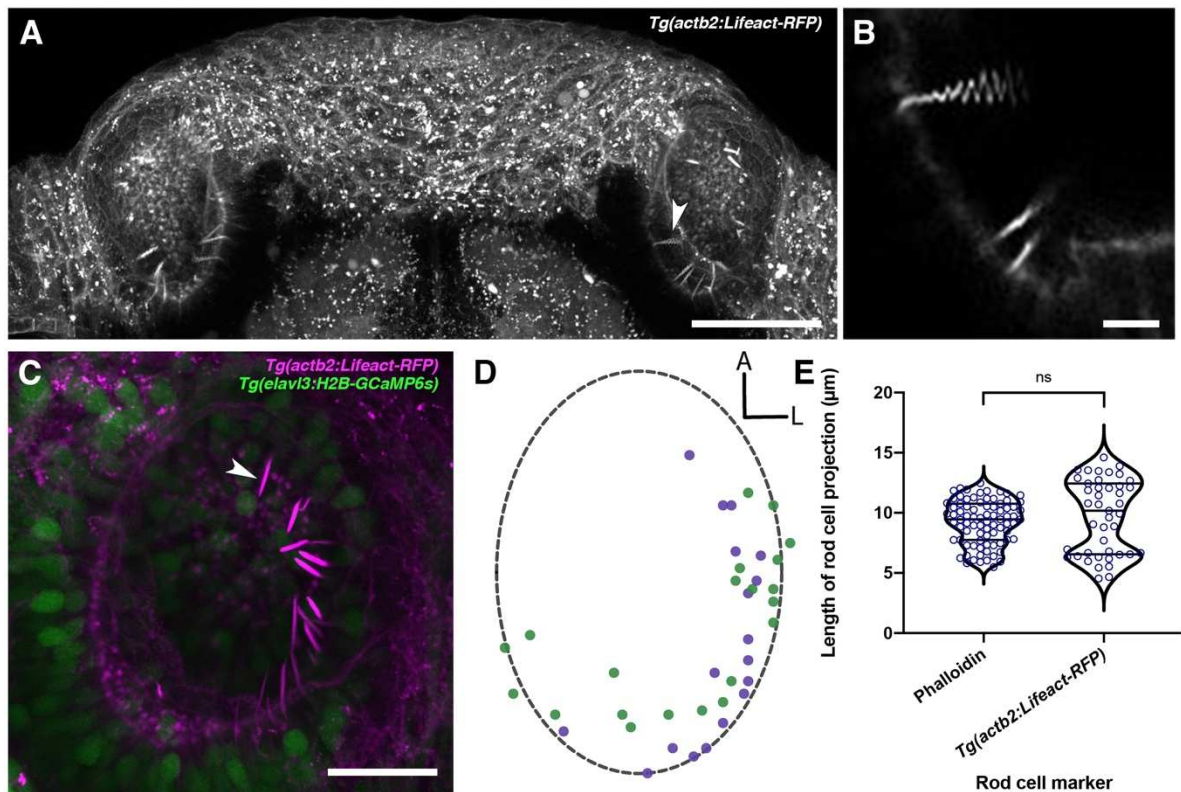
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674

675 **Figure 3. Olfactory rod cells are present in the olfactory epithelia of *ift88*^{-/-} zebrafish mutants,**
 676 **which lack cilia.**

677 (A, B) Maximum intensity projections of Airyscan confocal images of phalloidin stains of a 5 dpf wild-
 678 type (A) and *ift88*^{-/-} mutant (B) larva; dorsal views, anterior to the top. Grayscale values from the
 679 original fluorescence image have been inverted. Abbreviations: nm, cranial neuromast; op, olfactory
 680 pit. Several olfactory rods (arrowheads mark examples) are visible in each olfactory pit. Scale bar =
 681 50 μ m. (C) SEM of the head of a 4 dpf wild-type larva. Scale bar = 100 μ m. (D-E) SEM of 4 dpf larval
 682 wild-type olfactory pits (enlarged from panel C). Scale bars = 10 μ m. Insert in D shows enlarged view
 683 of boxed area in D. Arrowhead marks the tip of a rod cell apical projection surrounded by olfactory
 684 cilia. (E') Enlarged view of boxed area in E. Arrowhead marks one olfactory rod. Scale bar = 5 μ m.
 685 (E'') Enlargement of olfactory rod in E' (arrowhead). Scale bar = 1 μ m. (F) Frontal view SEM of the
 686 head of a 4 dpf *ift88*^{-/-} mutant larva. Scale bar = 100 μ m. (G) A map of the positions of olfactory rod
 687 cell projection emergence through the OE in *ift88*^{-/-} mutant larvae (N of olfactory pits = 3), based on
 688 SEM images at 4 dpf; anterior 'A' to the top, lateral 'L' to the right. One dot represents one olfactory
 689 rod. Different coloured dots represent rods from different larvae. (Compare with Figure 1E.) (H) SEM
 690 of 4 dpf larval *ift88*^{-/-} mutant olfactory pit (enlarged from panel F). Scale bar = 10 μ m (H') Enlarged
 691 view of boxed area in H. Arrowhead marks one example olfactory rod cell projection present despite
 692 the loss of cilia. Scale bar = 5 μ m. (I) Enlarged SEM of olfactory rods (arrowhead marks example) in 4
 693 dpf larval *ift88*^{-/-} mutant olfactory pit (from a different individual). Scale bar = 1 μ m.

694



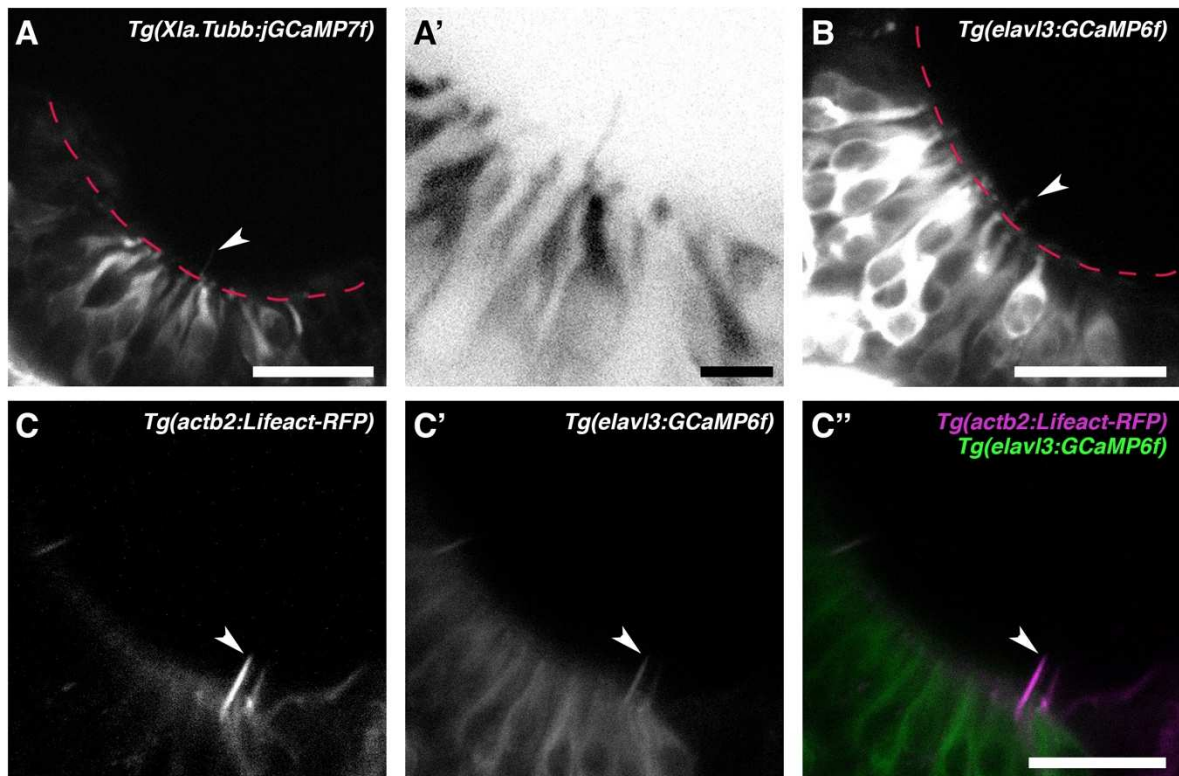
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697 **Figure 4. Olfactory rods are labelled in the olfactory epithelia of live zebrafish larvae by the**
 698 ***Tg(actb2:Lifect-RFP)* transgene.**

699 (A) Maximum intensity projection of dorsal view image of the olfactory pits of a live 6 dpf
 700 *Tg(actb2:Lifect-RFP)* transgenic larva; anterior to the top. Arrowhead marks one example olfactory
 701 rod positive for the Lifect-RFP transgene. Scale bar = 50 μ m. (B) Enlargement of olfactory rods in A
 702 (arrowhead in A) oscillating during raster-scanned image capture. (Raster scanning was performed
 703 from top to bottom in the image, as it has been rotated 90° clockwise.) (See Supplementary Movie 2.)
 704 Scale bar = 5 μ m. (C) Maximum intensity projection image of a live 4 dpf *Tg(actb2:Lifect-*
 705 *RFP);Tg(elavl3:H2B-GCaMP6s)* double-transgenic larval olfactory pit; anterior to the top, lateral to the
 706 right. Arrowhead marks one example olfactory rod positive for the Lifect-RFP transgene (magenta).
 707 Neuronal nuclei are labelled in green. Larvae were fully mounted in agarose, so rods were not
 708 moving. Scale bar = 20 μ m. (See Supplementary Movie 1.) (D) A map of the positions of olfactory rod
 709 cell projection bases in olfactory pits of 4 dpf *Tg(actb2:Lifect-RFP);Tg(elavl3:H2B-GCaMP6s)*
 710 double-transgenic larvae (N of olfactory pits = 2), based on 2D maximum intensity projections of
 711 confocal images; anterior 'A' to the top, lateral 'L' to the right. One dot represents one olfactory rod.
 712 Different coloured dots represent rods from different larvae, with purple corresponding to panel C.
 713 (Compare with Figure 1E.) (E) A quantitative comparison of the lengths of olfactory rod cell
 714 projections in fixed larvae, using Alexa-phalloidin as a marker ($N = 10$, n of olfactory rods = 82) versus
 715 live larvae, using Lifect-RFP as a marker ($N = 2$, $n = 43$). Violin plot; bars indicate the median and
 716 lower and upper quartiles for each group. Mann-Whitney U test; ns, not significant ($P = 0.232$).

717



718

719

720 **Figure 5. Olfactory rod cells are labelled by the cytoplasmic neuronal markers**721 ***Tg(Xla.Tubb:jGCaMP7f)* and *Tg(elavl3:GCaMP6f)*.**722 (A) Olfactory pit of a 4 dpf *Tg(Xla.Tubb:jGCaMP7f)* larva; anterior to the top, lateral to the right. Red

723 dotted line outlines the apical surface of the OE; arrowhead marks one olfactory rod, albeit faintly

724 labelled. Scale bar = 20 μm. (A') Enlargement of olfactory rod marked by arrowhead in A (grayscale

725 values inverted). Scale bar = 5 μm. (B) Olfactory pit of a 5 dpf *Tg(elavl3:GCaMP6f)* larva; anterior to

726 the top, lateral to the right. Red dotted line outlines the apical surface of the OE; arrowhead marks

727 one example olfactory rod, albeit faintly labelled. Scale bar = 20 μm. (C-C'') Lifect-RFP signal (C),

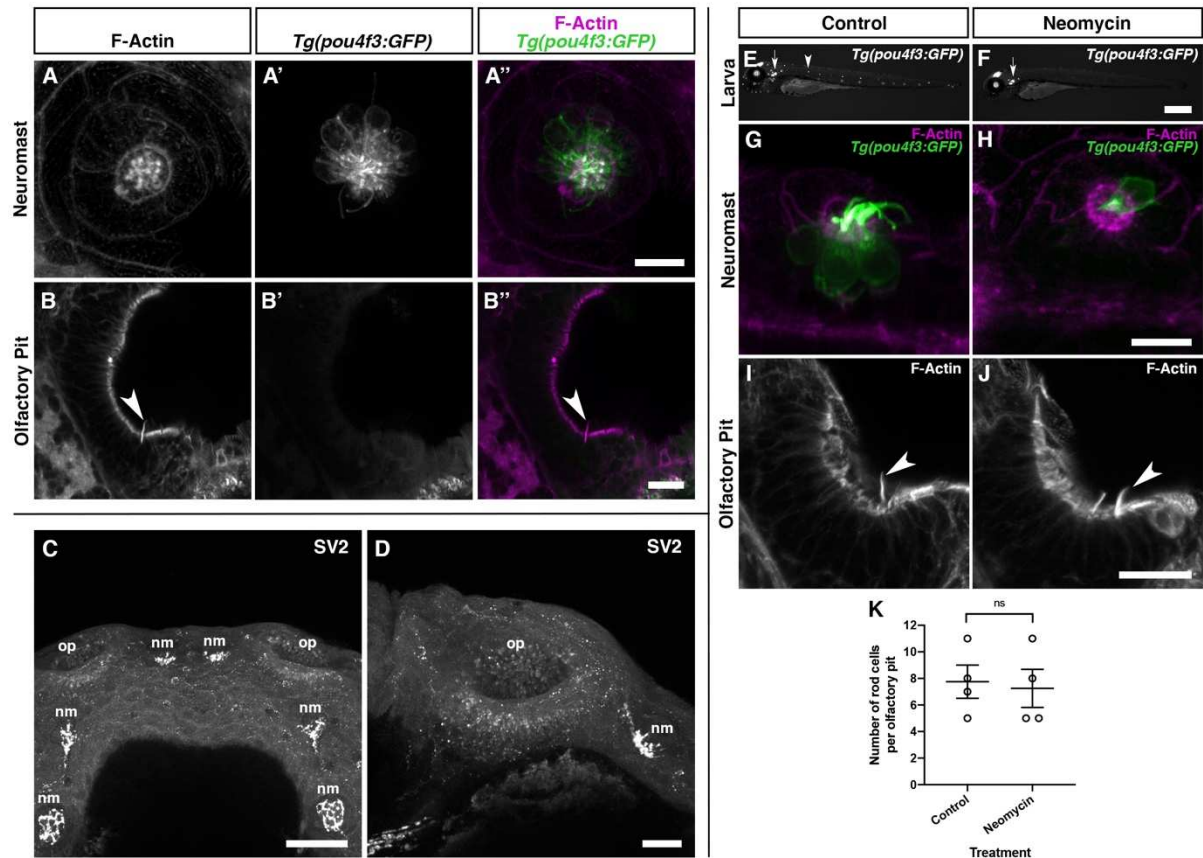
728 GCaMP6f signal (C'), and merged signals (C'') in an olfactory pit of a 5 dpf

729 *Tg(elavl3:GCaMP6f); Tg(actb2:Lifect-RFP)* double-transgenic larva; anterior to the top, lateral to the

730 right. Arrowhead marks one example olfactory rod, positive for both Lifect-RFP and GCaMP6f. Scale

731 bar = 20 μm.

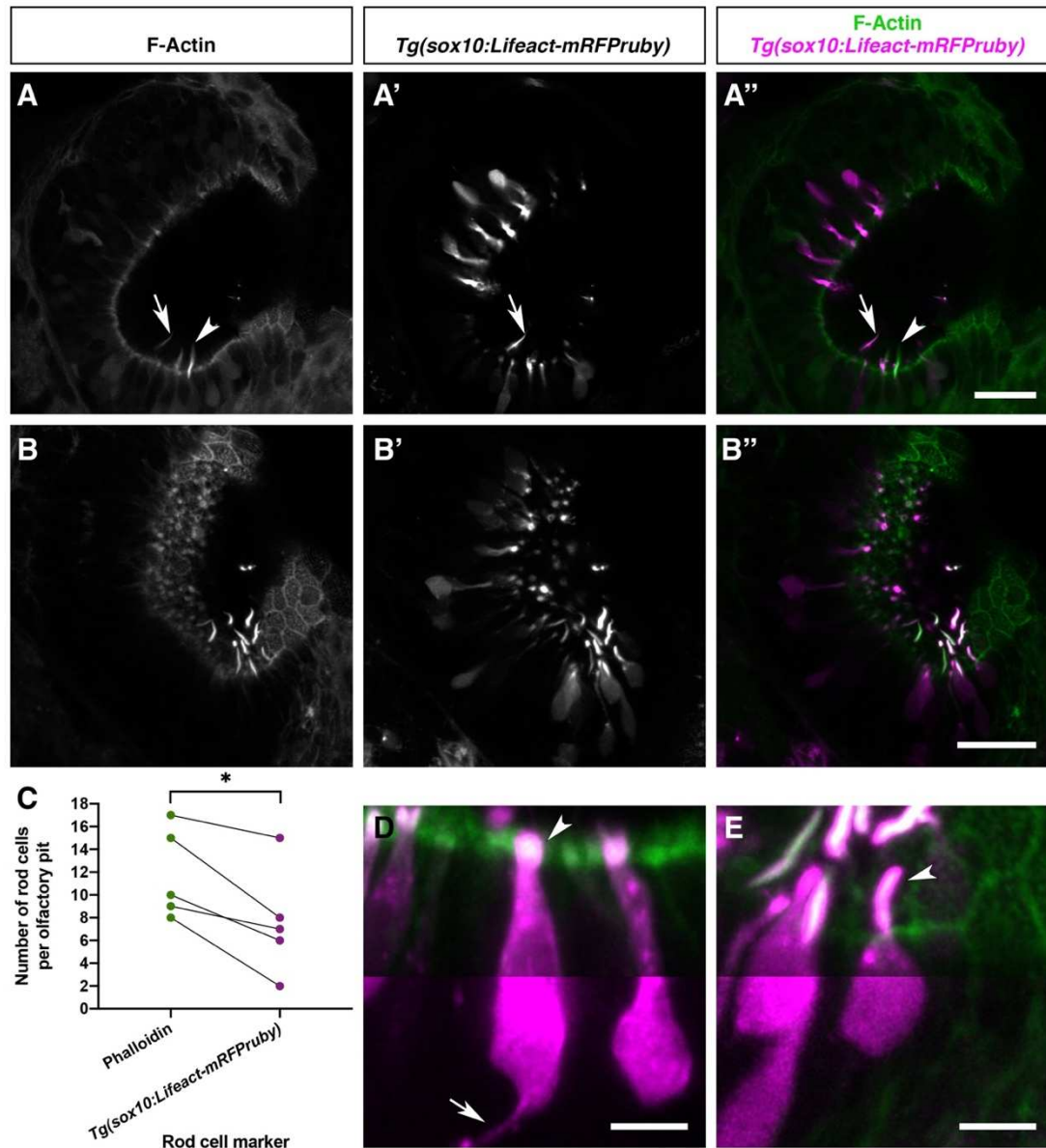
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733

734

735 **Figure 6. Olfactory rod cells in the zebrafish olfactory epithelium are not hair-cell-like.**
 736 (A-A'') Maximum intensity projection of Airyscan confocal image of Alexa-phalloidin signal (A),
 737 *Tg(pou4f3:GFP)* signal (A'), and merged signals (A'') in a cranial neuromast of a 5 dpf larva. Scale bar
 738 = 10 μ m. (B-B'') Airyscan confocal image of Alexa-phalloidin signal (B), *Tg(pou4f3:GFP)* signal (B'),
 739 and merged signals (B'') in an olfactory pit of a 5 dpf larva; anterior to the top, lateral to the right.
 740 Arrowhead marks one olfactory rod. Scale bar = 20 μ m. (C-D) Maximum intensity projections of
 741 confocal images of synaptic vesicle 2 (SV2) protein immunohistochemistry signal in the heads of 4 dpf
 742 wild-type larvae. Abbreviations: nm, cranial neuromast; op, olfactory pit. (C) Anterior to the top. Scale
 743 bar = 50 μ m. (D) Anterior to the top, lateral to the right. Scale bar = 20 μ m. (E-F) Widefield imaging of
 744 3 dpf *Tg(pou4f3:GFP)* larvae showing the damaging effects of 500 μ M neomycin treatment for 60
 745 minutes on lateral line neuromast hair cells. Fluorescence is lost or greatly reduced in both trunk
 746 (arrowhead) and cranial neuromasts, whereas fluorescence in hair cells of the inner ear maculae and
 747 cristae (arrow) is unaffected. Scale bar = 500 μ m. (G-H) Maximum intensity projections of Airyscan
 748 confocal images showing the damaging effects of 500 μ M neomycin treatment for 60 minutes on hair
 749 cells in a cranial neuromast of a 3 dpf larva, using *Tg(pou4f3:GFP)* (green) and Alexa-phalloidin
 750 (magenta) as markers. Scale bar = 10 μ m. (I-J) Maximum intensity projections of Airyscan confocal
 751 images showing no effect of 500 μ M neomycin treatment for 60 minutes on olfactory rods, using
 752 Alexa-phalloidin as a marker; anterior to the top, lateral to the right. Arrowheads mark olfactory rods.
 753 Scale bar = 20 μ m. (K) The number of olfactory rod cell projections per olfactory pit of 3 dpf
 754 *Tg(pou4f3:GFP)* larvae after 500 μ M neomycin treatment for 60 minutes (N of olfactory pits = 4),
 755 compared with an untreated group (N = 4). Welch's unpaired two-tailed t -test; ns, not significant (P =
 756 0.8018).
 757

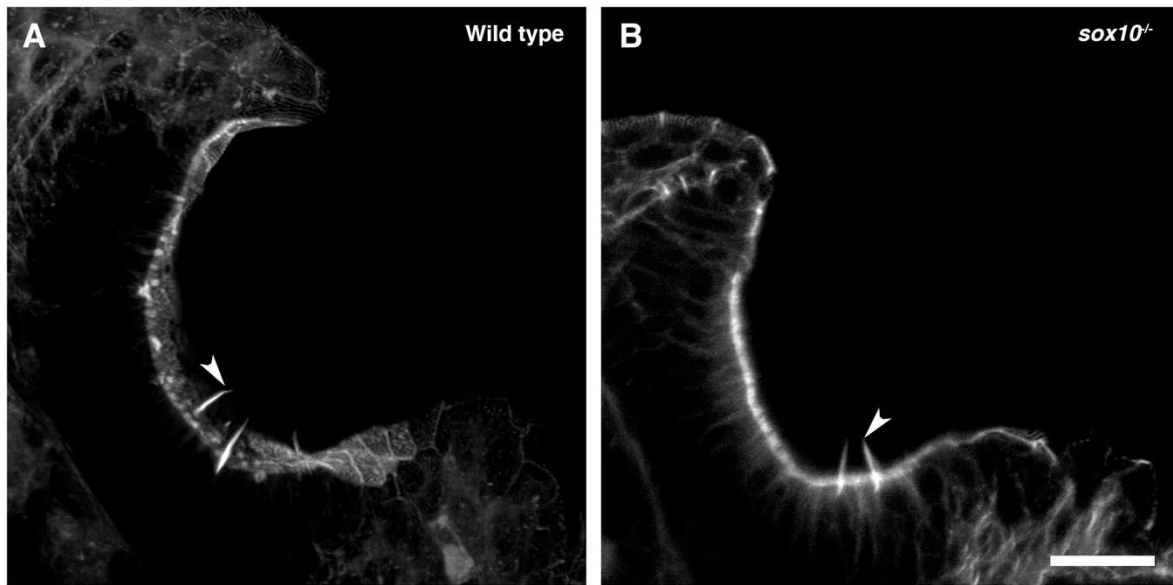


758

759

760 **Figure 7. Olfactory rod cells are apically located in the zebrafish olfactory epithelium, with a**
 761 **rounded cell body and no detectable axon.**
 762 (A-B'') Airyscan confocal image of Alexa-phalloidin signal (A, B), *Tg(sox10:Lifeact-mRFP*) signal
 763 (A', B'), and merged signals (A'', B'') in olfactory pits of 4-5 dpf larvae; anterior to the top, lateral to the
 764 right. Arrowhead marks one olfactory rod negative for Lifeact-mRFP. Arrow marks one olfactory
 765 rod positive for Lifeact-mRFP. Scale bars = 20 μ m. (C) Number of olfactory rod cells positively
 766 marked by Alexa-phalloidin (n of olfactory rods = 59), compared with the number of those also
 767 marked by *Tg(sox10:Lifeact-mRFP)* (n = 38), in olfactory pits of 4-5 dpf larvae (N of olfactory pits
 768 = 5). Connecting lines indicate rods from the same olfactory pit. Paired two-tailed t -test; * indicates P
 769 = 0.0146. (D) Enlargement of two microvillous OSNs, expressing Lifeact-mRFP, in the OE of a 4
 770 dpf larva; Alexa-phalloidin signal (green), *Tg(sox10:Lifeact-mRFP)* signal (magenta). Arrowhead
 771 marks the microvillous apical projections. The gamma value for the magenta channel in the bottom
 772 half of the panel has been adjusted to show the axon from one of the cells (arrow). Scale bar = 5 μ m.
 773 (E) Enlargement of olfactory rod cells (of which both the apical actin projections and cell bodies are
 774 labelled by the *Tg(sox10:Lifeact-mRFP)* transgene) in the OE of a 4 dpf larva; Alexa-phalloidin
 775 signal (green), *Tg(sox10:Lifeact-mRFP)* signal (magenta). Arrowhead marks a rod cell apical
 776 projection, positive for both markers. The gamma value for the bottom half of the panel has been
 777 adjusted as in D; no axon is visible. Scale bar = 5 μ m. See also Supplementary Movie 3.

778



779

780

781 **Figure 8. Olfactory rod cells are present in the olfactory epithelia of *sox10^{-/-}* zebrafish mutants.**

782 (A) Maximum intensity projection of Airyscan confocal image of phalloidin stain in a 5 dpf larval wild-

783 type olfactory pit; anterior to the top, lateral to the right. Arrowhead marks one example olfactory rod.

784 Scale bar = 20 μm . (B) Airyscan confocal image of phalloidin stain in a 5 dpf larval *sox10^{-/-}* mutant

785 olfactory pit; anterior to the top, lateral to the right. Arrowhead marks one example olfactory rod. Scale

786 bar = 20 μm .

787

788 **Supplementary Material**

789

790 **Supplementary Movie 1. Olfactory rods are labelled in the olfactory epithelia of**
791 **live zebrafish by the *Tg(actb2:Lifeact-RFP)* transgene.**

792 3D rendering of a confocal image of a 4 dpf *Tg(actb2:Lifeact-RFP);Tg(elavl3:H2B-*
793 *GCaMPs)* double-transgenic larval olfactory pit; anterior to the top. Olfactory rods are
794 labelled in magenta; neuronal nuclei are labelled in green.

795

796 **Supplementary Movie 2. Olfactory rods labelled with Lifeact-RFP in the**
797 **olfactory epithelia of live zebrafish larvae oscillate.**

798 Fast-capture time series confocal imaging (4.35 fps) of olfactory rods in a 6 dpf
799 *Tg(actb2:Lifeact-RFP)* larva; anterior to the top, lateral to the right. Playback speed
800 of the movie is 4 fps. Scale bar = 20 µm.

801

802 **Supplementary Movie 3. Olfactory rods labelled with Lifeact-mRFP_{ruby} in the**
803 **olfactory epithelia of live zebrafish larvae oscillate.**

804 Fast-capture time series light-sheet imaging (50.04 fps) of a 5 dpf *Tg(sox10:Lifeact-*
805 *mRFP_{ruby})* larval olfactory pit; anterior to the top left, lateral to the top right. Beating
806 olfactory cilia are visible in brightfield (grayscale), and oscillating olfactory rods are
807 labelled by Lifeact-mRFP_{ruby} (magenta). Playback speed of the movie is 7 fps.
808 Scale bar = 20 µm.

809

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811

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