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Glial-Specific Deletion of Med12 Results in Rapid Hearing Loss via Degradation of the Stria Vascularis

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47 **Abstract**

48 Mediator protein complex subunit 12 (Med12) is a core component of the basal
49 transcriptional apparatus and plays a critical role in the development of many tissues.
50 Mutations in Med12 are associated with X-linked intellectual disability syndromes and
51 hearing loss; however, its role in nervous system function remains undefined. Here, we
52 show that temporal conditional deletion of Med12 in astrocytes in the adult central
53 nervous system results in region specific alterations in astrocyte morphology.
54 Surprisingly, behavioral studies revealed rapid hearing loss after adult deletion of
55 Med12 that was confirmed by a complete abrogation of auditory brainstem responses.
56 Cellular analysis of the cochlea revealed degeneration of the stria vascularis, in
57 conjunction with disorganization of basal cells adjacent to the spiral ligament and
58 downregulation of key cell adhesion proteins. Physiological analysis revealed early
59 changes in endocochlear potential, consistent with strial-specific defects. Together, our
60 studies reveal that Med12 regulates auditory function in the adult by preserving the
61 structural integrity of the stria vascularis.

62 **Significance Statement**

63 Mutations in Mediator protein complex subunit 12 (Med12) are associated with X-linked
64 intellectual disability syndromes and hearing loss. Using temporal-conditional genetic
65 approaches in CNS glia, we found that loss of Med12 results in severe hearing loss in
66 adult animals through rapid degeneration of the stria vascularis. Our study describes the
67 first animal model that recapitulates hearing loss identified in Med12-related disorders
68 and provides a new system in which to examine the underlying cellular and molecular
69 mechanisms of Med12 function in the adult nervous system.

70 **Introduction**

71 Glial cell support of neuronal function is a hallmark feature of nervous systems across
72 evolution, with both glial abundance and specialization correlated with increasing
73 neuronal- and organism-complexity (Freeman & Rowitch 2013). Accordingly,
74 mammalian systems contain a host of specialized glial cells that subserve neuronal
75 function, with astrocytes serving as an archetype for glial support (Allen & Lyons 2018,
76 Nagai et al 2020). Among the key functions performed by astrocytes include buffering
77 neurotransmitters, maintenance of ion gradients, and providing metabolic support, all of
78 which directly influence the function of neurons and associated circuits (Khakh &
79 Deneen 2019).

80 In the mammalian auditory system hair cells use mechanosensation to convert
81 energy from soundwaves into neurophysiological signals that are transmitted to the
82 brainstem (Brownell et al 1985, Chan & Hudspeth 2005, Muller 2008) Hair cell function
83 relies on the endocochlear potential, which is an ion gradient established in the
84 endolymph and maintained by the stria vascularis (Gulley & Reese 1976, Patuzzi 2011).
85 The resting potential of the endolymph is regulated by the transport of K^+ ions across
86 the stria vascularis into the cochlear duct (Boettger et al 2002, Grunder et al 2001, Mittal
87 et al 2017). This function is mediated by potassium channels, including Kir4.1 which is
88 the predominant channel used by astrocytes to buffer neuronal K^+ (Chen & Zhao 2014,
89 Rozengurt et al 2003). Critically, disruption of these electrochemical gradients in the
90 endolymph can lead to altered hair cell function and survival, culminating in a loss of
91 hearing (Mittal et al 2017, Nin et al 2016). While much progress has been made in

92 understanding how the stria vascularis regulates endocochlear potential, the
93 transcriptional mechanisms that control stria vascularis function remain poorly defined.

94 Glial development offers a venue in which to identify transcription factors that
95 regulate mature glial cell function. For example, Sox10 and NFIA are key regulators of
96 oligodendrocyte and astrocyte development, respectively, and the physiological
97 functions of these cells in the adult CNS (Huang et al 2020, Stolt et al 2002). These
98 parallels also apply to auditory support cells, where Sox2, Sox9, and Sox10 are
99 expressed in developing and mature CNS glia and support cells in the cochlea
100 (Loponen et al 2011, Oesterle et al 2008, Watanabe et al 2000). Drawing on this parallel,
101 we identified Mediator Complex Subunit 12 (Med12) as a developmental transcription
102 factor that is enriched in subsets of astrocytes in the adult brain (John Lin et al 2017).
103 Studies in development have shown that Med12 cooperates with Sox10 to regulate
104 oligodendrocyte differentiation (Vogl et al 2013), however its role in mature astrocytes is
105 undefined. Furthermore, mutations in Med12 are linked to intellectual disability
106 syndromes and coupled with hearing loss (Donnio et al 2017, Lesca et al 2013,
107 Rubinato et al 2020, Schwartz et al 2007, Vulto-van Silfhout et al 2013). These
108 observations implicate Med12 as a critical transcription factor contributing to a host of
109 nervous systems functions related to neuronal circuits, myelination, and auditory
110 function. Despite its widespread expression in neurons and glia, the precise role of
111 Med12 in the adult CNS remains unknown.

112 To dissect the contributions of Med12 towards brain function we generated
113 temporal-conditional mouse knockout models that enabled us to delete Med12 in
114 astrocytes. Analysis across brain regions revealed reduced astrocyte complexity in the

115 hippocampus, while behavioral studies revealed that mice lacking astrocytic-Med12 are
116 deaf and have no auditory brainstem response. Cellular analysis of the cochlea
117 revealed that Med12 is required to maintain the integrity of the stria vascularis, which
118 eventually results eventual loss of hair cells. Together, these studies identify Med12 in
119 the stria vascularis as a key regulator of auditory function.

120

121

122 **Materials and Methods**

123 *Animals*

124 All experimental animals were treated in compliance with the US Department of Health
125 and Human Services, the NIH guidelines, and Baylor College of Medicine IACUC
126 guidelines. All mice were housed with food and water available ad libitum in a 12-hour
127 light/dark environment. Both female and male mice were used for all experiments, and
128 littermates of the same sex were randomly allocated to experimental groups. For
129 histological analyses of brains and behavior assays, adult mice aged 4-month were
130 used unless otherwise described. For auditory brainstem response (ABR) test and
131 histological analyses of inner ears, examinations or tissue collections were performed at
132 either 1, 2, 3, or 5 weeks after the tamoxifen treatment as described. All mice used in
133 this study were maintained on the C57BL/6J background. Med12 conditional knockout
134 mice were generated by crossing Med12 fl/fl or fl/y conditional mutant mice (Rocha et
135 al., 2010) with Aldh111-CreER (The Jackson Laboratory; RRID:IMSR_JAX:029655),
136 resulting in Med12fl/fl; Aldh111-CreER or Med12fl/y; Aldh111-CreER (A12-KO) and
137 Med12fl/fl or Med12fl/y (A12-Con) littermate controls. For histological analysis, the

138 Aldh111-GFP mouse was crossed with A12-KO, resulting in Med12fl/fl (or Med12fl/y);
139 Aldh111-CreER; Aldh111-GFP (AG12-KO) and Med12fl/fl (or Med12fl/y); Aldh111-GFP
140 (AG12-Con) mice. To mark the Cre-targeted cells, Rosa-CAG-LSL-tdTomato-WPRE
141 mice was crossed with A12-KO, resulting in Med12fl/fl (or Med12fl/y); Aldh111-CreER;
142 Rosa-CAG-LSL-tdTomato-WPRE (AT12-KO) and Med12+/+ (or Med12+/y); Aldh111-
143 CreER; Rosa-CAG-LSL-tdTomato-WPRE (AT12-Con) mice. To induce deletion of
144 Med12 in the adult, four-week-old mice were gavage-fed with 150 mg/kg body weight of
145 Tamoxifen (Sigma-Aldrich, cat no. T5648) dissolved in corn oil twice per day for 5 days.
146 Above experiments were approved by Baylor College of Medicine IACUC.

147

148 *Immunofluorescence on frozen tissue sections*

149 Mice were anesthetized under isoflurane inhalation and perfused transcardially with 1X
150 PBS pH 7.4 followed by 4% paraformaldehyde (PFA). Brains were removed, post-fixed
151 in 4% PFA for 8 hours, and placed in 20% sucrose for 16 hours before embedded in
152 OCT. Inner ears were dissected from temporal bones and fixed in 4% PFA overnight,
153 decalcified in 0.2 M EDTA for 24 hours. Decalcified inner ears were incubated in 7.5%
154 gelatin solution (1X PB/10% sucrose) at 37°C for 1 hour before embedded in OCT.
155 Brain sections of 30 μ m and cochlea sections of 12 μ m were made on a cryostat. For
156 immunostaining, tissue sections were washed with 1X PBS 5 min X3, blocked with 10%
157 goat serum in PBS with 0.3% Tween20, and then incubated with primary antibodies in
158 blocking solution overnight. On the next day, sections were incubated with secondary
159 antibodies in PBS with 0.1% Tween20 for 1 h RT, followed by incubation with DAPI for 5
160 min, and mounted with VECTASHIELD Antifade Mounting Media (Vector Laboratories,

161 H-1000). The following primary antibodies were used: Chicken anti-GFP (1:1000;
162 abcam, ab13970), rabbit anti-Med12 (1:1000; Novus Biologicals, NB100-2357), chicken
163 anti-GFAP (1:1000; abcam, ab4674), mouse anti-GFAP (1:1000, EMD millipore,
164 MAB360), mouse anti-NeuN (1:500, millipore, MAB377), rabbit anti-Myosin-VIIa (1:1000,
165 Proteus BioSciences, 25-6790), mouse anti-Myosin-VIIa (1:500, DSHB, MYO7A), rabbit
166 anti-Sox2 (1:1000, Millipore, AB5603), rat anti-Sox2 (1:1000, Invitrogen, 14-9811-82),
167 rabbit anti-GLUT1 (1:500, abcam, ab115730), mouse anti-ZO-1 (1:500, Invitrogen, 33-
168 9100), mouse anti- E-Cadherin (1:200, BD Biosciences, 610181), and rabbit anti-
169 connexin 31 (1:250, Proteintech Group, 12880-1-AP). The following secondary
170 antibodies were used (1:500): Alexa Fluor 488 goat anti-chicken (Thermo Fisher
171 Scientific, A11039), Alexa Fluor 488 goat anti-mouse (Thermo Fisher Scientific,
172 A32723), Alexa Fluor 488 goat anti-rabbit (Thermo Fisher Scientific, A32731), Alexa
173 Fluor 568 goat anti-mouse (Thermo Fisher Scientific, A-11004), Alexa Fluor 568 goat
174 anti-rabbit (Thermo Fisher Scientific, A11036), Alexa Fluor 647 goat anti-mouse
175 (Thermo Fisher Scientific, A32728), and Alexa Fluor 647 goat anti-rabbit (Thermo
176 Fisher Scientific, A32733).

177

178 *Confocal imaging and image analysis*

179 To measure astrocyte morphology, fluorescent images were acquired using a Zeiss
180 LSM 880 laser scanning confocal microscope with 63X oil immersion objective with
181 frame size at 1024 x 1024 and bit depth at 12. Serial images at z axis were taken at an
182 optical step of 1 μ m, with overall z axis range encompassing the whole section. Images
183 were imported to Imaris Bitplane software, and only astrocytes with their soma between

184 the z axis range were chosen for further analysis (Lanjakornsiripan et al., 2018). We
185 performed 3D surface rendering using the Imaris Surface module, and color-coded the
186 reconstructed surface images based on the surface area of each astrocyte.
187 Morphological analysis was performed using the Imaris Filament module. Astrocyte
188 branches and processes were outlined by Autopath with starting point set at 10 mm and
189 seed point set at 0.5 mm, and statistical outputs including “filament number Sholl
190 intersections” were extracted and plotted with Prism software. Data were generated
191 from 3 brain sections per region per mouse with 3 mice per genotype. To analyze
192 number of astrocytes and Med12 knockout efficiency, fluorescent images were acquired
193 using a Zeiss LSM 880 laser scanning confocal microscope with 20X objective. Cell
194 numbers were quantified by the QuPath software Cell Detection function (Bankhead et
195 al., 2017). To examine the cochlea, fluorescent images were acquired using a Zeiss
196 LSM 880 laser scanning confocal microscope with 20X objective.

197

198 *Behavioral tests*

199 *Fear conditioning*

200 Fear conditioning was performed as previously described (Samaco et al., 2008). Mice
201 were tested at 4-month-old in a chamber that contains a grid floor that can deliver an
202 electric shock (Actimetrics chamber system, Med Associates, St. Albans, VT, USA). On
203 Day 1 of the test, mice were placed in the chamber and left undisturbed for 2 min
204 followed by a 30 s white noise sound pulse (‘cue’). At the end of the cue, the mouse
205 was shocked (2 s, 0.4 mA). 2 min later, a second pairing of sound cue followed by

206 shock was delivered. Thirty seconds after the final shock, the animal was removed and
207 replaced to the home cage. The following day, the animals were replaced to the same
208 chamber ('context test') and freezing behavior was recorded for 6 min. Freezing
209 behavior was recorded automatically by the instrument. One hour after the context test,
210 the animals were placed into a chamber which had been cleaned with an unfamiliar
211 agent (ethanol) and the wall color, the chamber shape and the odor (artificial vanilla)
212 had been changed to remove the contextual cues of the chamber. The animals were
213 then monitored for 3 min. After 3 min, the white noise cue was started and lasted 3 min.
214 The amount of freezing was recorded separately for the first 3 min and for the last 3 min
215 (cue test). The number of freezing intervals was converted to a percentage of freezing
216 for both the context test and the cue test, and the data were analyzed using a one-way
217 ANOVA.

218

219 *Acoustic startle response*

220 Acoustic startle response and prepulse inhibition was performed as previously
221 described (Samaco et al., 2008). Mice at 4-month old were subjected to acoustic
222 prepulse inhibition test. The acoustic prepulse inhibition task consists of presenting the
223 animal with two closely paired sound pulses: a prepulse at +0 dB, +4 dB (74 dB), +8 dB
224 (78 dB), +12 dB (82 dB) and over background followed 100 ms later by a pulse of 120
225 dB. The amount of startle the pulse induces in the animal is recorded using a startle
226 chamber for mice (SR-Lab, San Diego Instruments, San Diego, CA, USA) which
227 records activity for 65 ms after the pulse. The maximum amplitude recorded over the 65

228 ms is recorded and compared using an ANOVA. No prepulse inhibition result is
229 presented because the Med12-deficient mice showed no acoustic startle response.

230

231 *Auditory brainstem response recording*

232 Auditory brainstem response recording was performed as previously described (Manalo
233 et al., 2020). Mice were intraperitoneally injected with ketamine-xylazine (100: 10
234 mg/kg). Depth of anesthesia was verified by the absence of toe-pinch reflex. Testing
235 was performed in a soundproof faraday cage booth while mice were placed on heating
236 pad to maintain normal body temperature throughout the procedure. Pure tone bursts
237 (0.1 ms rise/fall, 2 ms duration, 21 presentations/s) from 4 to 48 kHz were generated
238 using System 3 digital signal processing hardware and software (Tucker Davis
239 Technologies). The intensity of the tone stimuli was calibrated using a type 4938 one-
240 quarter inch pressure-field calibration microphone (Bruël & Kjær). EC1 ultrasonic, low-
241 distortion electrostatic speakers were coupled to the ear canal to deliver stimuli within 3
242 mm of the tympanic membrane. Response signals were recorded with subcutaneous
243 needle electrodes inserted at the vertex of the scalp (channel 1), the postauricular bulla
244 region (reference), and the back leg (ground), and averaged over 500 presentations of
245 the tone bursts. Electrode-recorded activity was filtered (high pass, 300 Hz; low pass, 3
246 kHz; notch, 60 Hz) before averaging to minimize background noise. Auditory thresholds
247 were determined by decreasing the sound intensity of each stimulus to 10 dB from 90
248 dB in 5 dB steps until the lowest sound intensity with reproducible and recognizable
249 waveforms was detected. Thresholds were determined to within 5 dB for each
250 frequency by two raters to ensure reliability. SD (dB SPL) were plotted as a function of

251 stimulus frequency (kilohertz) and analyzed for group differences by a two-way ANOVA,
252 followed by a multiple-comparison test, to reveal overall trends.

253

254 *Distortion Product Otoacoustic Emission*

255 DPOAEs (2f1-f2) were measured from the left ears in sessions separate from
256 ABR recording. Twelve A12-KO mice (3 female) and 9 A12-Con mice (1 female) were
257 prepared as for ABR measures. Input/output curves were obtained for primary stimulus
258 frequencies f1 and f2, where $f1=f2/1.2$ at fixed level differences L1 and L2, where
259 $L2=L1-10$ dB. DPOAEs were measured for f2=12, 18, and 24 kHz, and L2 levels
260 ranging from 0-80 dB SPL in 5 dB steps. Stimuli were presented using 2 TDT EC1
261 speakers configured in a closed acoustic system along with a Knowles probe
262 microphone to record speaker and cochlear output. Primary and DPOAE levels were
263 recorded using EMAN software in conjunction with TDT and custom hardware. DPOAE
264 thresholds were defined as a 2f1-f2 response of at least -15 dB SPL. This level was
265 chosen as it represented the lowest criterion response that was reliably above
266 background noise levels and was generally within 10 dB of ABR thresholds.

267

268 *Endocochlear Potential recording*

269 EP recordings were obtained from the left ear of 10 A12-KO mice (2 female) and
270 9 A12-Con mice (1 female). To record the EP, animals were anesthetized with 80
271 mg/kg ketamine, 15 mg/kg xylazine, i.p., and positioned ventral-side-up in a custom
272 head holder. Core temperature was maintained at 37.5 ± 1.0 °C using a DC electric

273 heating pad in conjunction with a rectal probe (FHC). After blunt dissection of the jaw
274 musculature to reveal the left auditory bulla and opening of the posterior bulla with
275 forceps, a hole was made in the cochlear capsule directly over scala media of the lower
276 basal turn using a fine drill. Glass capillary pipettes (20-30 M Ω) filled with 0.15 M KCl
277 were mounted on a hydraulic microdrive (Frederick Haer) and advanced until a stable
278 positive potential was observed that did not change with increased electrode depth.
279 The signal from the recording electrode was led to an AM Systems Model 1600
280 intracellular amplifier.

281

282 *QUANTIFICATION AND STATISTICAL ANALYSIS*

283 Sample sizes and statistical tests can be found in accompanying Figure legends. Offline
284 analysis was carried out using SPSS 20 and Excel software. We assessed the
285 significance of data for comparison by Student's two-tailed unpaired/paired t test or two-
286 tailed unpaired t test. For multiple comparisons, we used the one-way ANOVA with
287 Tukey's test and two-way ANOVA with Sidak's test. In general, we assumed data were
288 normally distributed but this was not formally tested. Data are presented as mean \pm
289 SEM (standard error of the mean). Levels of statistical significance are indicated as
290 follows: * ($p < 0.05$), ** ($p < 0.01$), *** ($p < 0.001$).

291

292

293

294 **Results**

295 **Deletion of Med12 Selectively Reduces Morphological Complexity of Astrocytes**

296 To decipher whether Med12 plays a role in astrocyte function in the adult brain, we
297 evaluated its expression in astrocytes by co-immunostaining with GFP and Med12 in
298 16-week-old *Aldh111-GFP* reporter mice. This analysis revealed that approximately 70%
299 of *Aldh111-GFP* astrocytes express Med12 in the olfactory bulb (73.1±2.3%) and around
300 50% across other examined brain regions (**Figures 1A-1D, 1R**, cortex: 52.0±2.3%,
301 hippocampus: 48.5±8.8%, brainstem: 57.9±5.2%); Med12 also exhibited expression in
302 NeuN positive neurons in these same brain regions.

303 Med12 plays an important role in numerous developing tissues, rendering the
304 germline knockout embryonic lethal (Rocha et al 2010a, Rocha et al 2010b) and
305 requiring the generation of temporal and lineage-specific knockout alleles to study its
306 role in mature astrocytes. To specifically knock out *Med12* in astrocytes in the adult
307 brain, we acquired mice containing a floxed *Med12* allele (*Med12^{fl/fl}*) and intercrossed it
308 with the *Aldh111-CreER* line and the *Aldh111-GFP* reporter, producing the *Aldh111-
309 CreER; Aldh111-GFP; Med12^{fl/fl}* and *Aldh111-CreER; Aldh111-GFP; Med12^{fl/y}* (AG12)
310 mouse line. Tamoxifen (TM) was administered at post-natal day 30 (P30), which
311 allowed us to bypass possible developmental roles for Med12. Mice were harvested two
312 months after TM treatment (P80-P90) and stained for Med12. We observed loss of
313 Med12 in more than 50% of *Aldh111-GFP* positive astrocytes in the olfactory bulb,
314 cortex, hippocampus, and brainstem two months after treatment (**Figures 1A-1H, 1R**;
315 *Aldh111-GFP* astrocytes that express Med12 in AG12-Con: olfactory bulb: 73.1±2.3%,
316 cortex: 52.0±5.1%, hippocampus: 48.5±8.8%, brainstem: 57.9±5.2%; in AG12-KO:
317 olfactory bulb: 28.0±5.6%, cortex: 31.7±4.5%, hippocampus: 9.4±4.7%, brainstem:

318 28±12%). We were unable to achieve higher KO efficiency because prolonging TM
319 treatment led to severe gastrointestinal illness, which ultimately resulted in death.

320 To determine whether deletion of Med12 impacts astrocyte homeostasis, we
321 assessed their morphology using confocal imaging of the Aldh1l1-GFP reporter in
322 AG12-Con and AG12-KO brains. Three-dimensional reconstructions of these images
323 were subjected to Sholl Analysis to evaluate the morphological complexity of astrocytes.
324 This analysis revealed that astrocytes from AG12-KO mice demonstrate reduced
325 complexity in the prefrontal cortex and CA1 of the hippocampus, whereas astrocytes
326 from the olfactory bulb and brainstem were unaffected (**Figures 1J-1Q; 1S-1V**).

327 Quantification of the gross numbers of Sox9-expressing astrocytes did not reveal any
328 differences between AG12-Con and AG12-KO mice (Aldh1l1-GFP⁺/DAPI ratio in AG12-
329 Con: olfactory bulb: 15.5±2.8%, cortex: 13.6±2.2%, hippocampus: 25.7±6.8%,
330 brainstem: 10.3±2.3%; in AG12-KO: olfactory bulb: 14.9±1.6%, cortex: 15.6±1.9%,
331 hippocampus: 26.1±2.7%, brainstem: 9.6±2.2%), indicating that loss of Med12 does not
332 lead to cell death (**Figure 1W**). Together, these studies indicate that Med12 is
333 selectively required to maintain the morphological complexity of astrocytes in the
334 hippocampus and cortex, but is not required in the olfactory bulb or brainstem.

335

336 **Elimination of Med12 Leads to Rapid Hearing Loss**

337 The foregoing observation that loss of Med12 results in decreased morphological
338 complexity of astrocytes in the hippocampus and prefrontal cortex, led us to examine
339 whether these cellular phenotypes lead to behavioral deficits associated with these
340 brain regions. Since the hippocampus demonstrated the most efficient knockout of

341 Med12, we conducted fear conditioning assays, which are associated with hippocampal
342 function. As shown in Figure 2A, AG12-Con mice showed significantly increased
343 freezing behaviors after the auditory cue (AG12-Con mice: no cue: $25.3 \pm 1.5\%$, cued:
344 $75.2 \pm 3.2\%$); however, AG12-KO mice demonstrated no change in freezing before and
345 during the auditory cue (**Figure 2A**, AG12-KO mice: no cue: $6.7 \pm 1.7\%$, cued:
346 $17.1 \pm 5.4\%$). Further analysis revealed that the AG12-KO mice did not respond to
347 acoustic stimulation (**Figure 2B**, AG12-Con mice: 2495.2 ± 492.1 a.u., AG12-KO mice:
348 58.0 ± 11.9 a.u.), indicating that the apparent deficits in fear conditioning are likely the
349 result of impaired hearing. Consistent with this interpretation, we did not observe any
350 changes in short term potentiation (STP) or long term potentiation (LTP) in the
351 hippocampus of AG12-KO compared to AG12-Con (**Figure 2C-E**, LTP: AG12-Con mice:
352 $120.69 \pm 14.43\%$, AG12-KO mice: $124.8 \pm 25.67\%$).

353 The loss of hearing in the AG12-KO mice prompted us to further examine
354 auditory function by measuring auditory brainstem responses (ABRs). We first
355 established a baseline ABR for AG12-Con and AG12-KO mice prior to TM-induced
356 deletion of Med12, finding no differences in ABR thresholds between two groups
357 (**Figure 3B**, 4 to 32 KHz). Next, we evaluated ABR in weekly intervals post-TM induced
358 deletion (**Figure 3A**), finding dramatic elevation of auditory thresholds at one-week
359 post-TM treatment (at 4 KHz, AG12-Con mice: 63.3 ± 1.7 dB SPL, AG12-KO mice:
360 77.9 ± 4.9 dB SPL. at 16 KHz, AG12-Con mice: 49.2 ± 2.4 dB SPL, AG12-KO mice:
361 65.7 ± 6.3 dB SPL), culminating in a complete loss of auditory response to all stimulation
362 by three weeks post TM-treatment (**Figure 3C-3D**, 4 to 32 KHz, AG12-KO mice with no
363 response). To identify the possible cause of hearing loss we further examined the

364 recording of auditory brainstem responses from each group at three weeks after TM
365 induction. ABR responses at 8kHz in AG12-Con mice revealed five clearly defined
366 maxima which represent major ABR peaks I to V (**Figure 3F**). In contrast, we did not
367 observe any of the major ABR peaks in AG12-KO mice three weeks after TM (**Figure**
368 **3G**). ABR wave I corresponds to compound auditory nerve conduction, therefore its loss
369 in the AG12-KO mice indicates that Med12 deletion results in dysfunction of the inner
370 ear, which inhibits initiation of the auditory response.

371

372 **Med12 is Required to Maintain the Structural Integrity of the Stria Vascularis**

373 Having established that the loss of hearing in AG12-KO mice is likely due to defects in
374 the inner ear, we next examined the cellular integrity of the cochlea. Starting with the
375 organ of Corti, analysis of AG12-KO mice at three weeks post-TM induction did not
376 reveal any overt structural changes in supporting cells (Sox2+) or hair cell (Myo7a+)
377 populations (**Figure 4A-H**, 3 animals in each group were analyzed). Moreover, analysis
378 at five weeks post-TM induction also did not show any changes in the cellular integrity
379 of the organ of Corti (**Figure 4I-P**, 3 animals in each group were analyzed). However,
380 analysis at 18 weeks post-TM induction did reveal an eventual loss of Myo7a-
381 expressing hair cells, coupled with disorganization of Sox2-expressing support cells and
382 a general breakdown in organ of Corti integrity (**Figure 4Q-X**, 3 animals in each group
383 were analyzed). Nevertheless, the latent loss of hair cells at 18 weeks after Med12
384 deletion does not coincide with the observed loss of ABR at 2-3 weeks post-deletion
385 (**Figure 3**), indicating that Med12-dependent loss of ABR is likely mediated through
386 other cellular mechanisms in the adult cochlea.

387 Next, we examined additional structures in the cochlea, finding that Aldh111-GFP
388 is also co-localized to the spiral ligament and basal cells of the stria vascularis (**Figure**
389 **5A-D**). Strikingly, at three weeks post-TM induction the stria vascularis of AG12-KO
390 mice exhibited drastic alterations in its integrity, highlighted by alterations in the
391 structure of ATP1A1-expressing intermediate cells at the apical turn (**Figure 5C-D v**
392 **Figure 5G-H; yellow arrows**). Analysis at five weeks post-TM revealed that ATP1A1-
393 expressing intermediate cells are completely lost from middle to apical turns in AG12-
394 KO mice, which further disrupts the architecture of these structures (**Figure 5I-P**). Given
395 the essential role of the stria vascularis in maintaining the ionic composition of
396 endolymph, these findings suggest that progressive loss of the stria vascularis results in
397 ionic imbalance of the endolymph, which compromises hair cell function, leading to
398 hearing loss in AG12-KO mice.

399 To decipher which cell populations in the cochlea are dependent on Med12
400 expression, we used immunostaining to determine its pattern of expression. Staining for
401 Med12 was observed in the stria vascularis in AG12-con mice three weeks post-TM
402 induction (**Figure 5Q-R**). Nuclear localization of Med12 was observed in basal cells and
403 fibrocytes in the spiral ligament (**Figure 5R, yellow arrowheads**). Analysis of Med12
404 expression in AG12-KO mice, revealed that the nuclear localized expression in basal
405 cells was significantly decreased (**Figure 5R v 5T**). The percentage of Med12 positive
406 basal cells reduced from $91.2 \pm 5.9\%$ (AG12-con mice, n=3) to $25.1 \pm 1.2\%$ (AG12-KO
407 mice, n=3) (**Figure 5W**). These observations suggest that Med12 deletion is occurring
408 in the basal cells of the stria vascularis in the AG12-KO mouse line.

409

410 **Med12 Deletion Disrupts Basal Cell Organization**

411 To track which populations in the cochlea undergo TM-induced Cre-recombination in
412 the *Aldh111*-CreER line we replaced the *Aldh111*-GFP reporter with a Cre-inducible
413 reporter, generating the *Aldh111-CreER; Rosa-CAG-LSL-tdTomato; Med12^{fl/fl}* mouse line
414 (i.e. AT12 line). We treated AT12-Con mice with TM and harvested the cochlea two
415 weeks after induction, finding that the tdTomato reporter labeled the Glut1-expressing
416 basal cells of the stria vascularis and some fibrocytes in the spiral ligament (**Figure 6A-**
417 **D; yellow arrowheads**), but not spiral ganglion cells (not shown). Consistent with a
418 requirement for Med12 expression in basal cells, examination of AT12-KO mice two
419 weeks after TM induction revealed a loss of basal cell organization, where Glut1- and
420 tdTomato- expressing basal cells appear diffuse and lose cell-cell contact (**Figure 6E-H;**
421 **unfilled arrowheads**, 3 animals in each group analyzed, phenotype observed in all
422 AT12-KO mice). Together, these finding indicate that Med12 is required to maintain the
423 organization and integrity of basal cells in the stria vacularis.

424 Adhesion between cell types within the stria vascularis is required to generate
425 and preserve ion gradients in the endolymph, which are essential for maintaining
426 endocochlear potential and auditory function (Ciuman 2009, Wan et al 2013). Our
427 observed cellular phenotypes in basal cells of the AT12-KO line suggests that cell
428 adhesion within the stria vascularis is disrupted. To test this, we evaluated the
429 expression of key junction proteins associated with hearing loss in this region, ZO1, E-
430 cadherin (Ecad), and Connexin 31 (Cx31). In AT12-Con animals, ZO1 is concentrated
431 next to the cell membrane of basal cells, however, ZO1 expression level was
432 dramatically decreased in AT12-KO mice (**Figure 6I-L; white arrowheads**, the intensity

433 ratio reduced to 0.696 ± 0.026 compared to AT12-Con, $n=3$ in each group). Cx31 protein
434 was highly concentrated between basal cells and fibrocytes in spiral ligament in AT12-
435 Con mice and demonstrated disorganized localization in AT12-KO mice (**Figure 6M-P**,
436 **white arrowheads**; **Figure 6U** the intensity ratio was reduced to 0.689 ± 0.103
437 compared to AT12-Con, $n=3$ in each group). Ecad expression was mainly observed in
438 both intermediate cells and basal cells (**Figure 6Q-R**). Although the distribution of Ecad
439 was altered in AT12-KO mice (**Figure 6S-T**), the intensity was not significantly different
440 compared to AT12-Con mice (**Figure 6Q-T, U**, intensity ratio: 1.036 ± 0.047 compared to
441 AT12-Con, $n=3$ in each group). These finding suggested that Med12 is necessary to
442 maintain the expression of these critical junction and cell adhesion proteins in the stria
443 vascularis and spiral ligament.

444

445 Reduced Endocochlear Potential in Med12 Mutant Mice

446 The forging studies indicate that loss of Med12 disrupts the organization of the basal
447 cells as early as two weeks after TM-induced deletion, suggesting that the cascade of
448 cellular events that lead to hearing loss originates in the stria vascularis. Therefore, we
449 next measured the endocochlear potential (EP) in AG12-Con and AG12-KO mice 10-14
450 days after TM-induced deletion. Consistent with our prior results (**Figure 3**), AG12-KO
451 mice showed ABR thresholds that were 15-25 dB higher than those in AA2-Con mice.
452 These differences were significant overall ($p < .001$), with no interactions by frequency
453 (**Figure 7A**). The overall relative 'flat' threshold shift in the AG12-KO mice compared to
454 AG12-Con supports a non-frequency specific influence, consistent with isolated strial
455 dysfunction (Schuknecht and Gacek, 1993). EP recordings from the same cochleas

456 **(Figure 7B)** revealed no overlap of EP values by group ($p < .001$), with AG12-Con
457 showing typical values for C57BL/6 mice (>100 mV) (Ohlemiller, et al. 2009) and AG12-
458 KO mice averaging ~ 30 mV lower (76.8 versus 106.7 mV).

459 To confirm that these mice, at this timepoint, have stria-specific defects we next
460 performed distortion product otoacoustic emission (DPOAE) tests on these same mice
461 (Ueberfuhr, et al 2016). In the presence of an isolated strial defect, maximum DPOAE
462 values should be similar for normal and stria-impaired mice, and DPOAE thresholds
463 should be spared relative to ABR thresholds at the same frequencies (Mills, 2003).
464 Since the DPOAE input/output results did not vary with f_2 frequency, data from $f_2 = 12,$
465 $12, 24$ kHz were all included from each animal. This analysis revealed that the
466 maximum DPOAE values did not differ significantly between AG12-Con and AG12-KO
467 mice, although control mice gave slightly higher values (30.5 versus 36.2 dB SPL)
468 **(Figure 7C)**. Finally, a stria-specific defect is also predicted to yield DPOAE thresholds
469 that are lower relative to ABR thresholds, meaning that ABR-DPOAE threshold
470 differences should be higher when stria is compromised. Indeed, this metric both
471 averaged (0.83 versus -5.04) and skewed higher in the AG12- KO **(Figure 7D)**,
472 supporting the existence of isolated strial dysfunction in the majority of the AG12-KO
473 mice. In sum, ABR, EP, and DPOAE data supported a central tendency toward isolated
474 strial dysfunction in the AG12-KO mice as early as 10-14 days after TM induced
475 deletion.

476

477

478 **Discussion**

479 Mutations in MED12 are linked to a broad spectrum of genetic disorders with X-linked
480 intellectual disability (Donnio et al 2017, Lesca et al 2013, Rubinato et al 2020,
481 Schwartz et al 2007, Vulto-van Silfhout et al 2013), however embryonic lethality in
482 Med12-null and heterozygous mice have hindered the study of these Med12-associated
483 phenotypes (Rocha et al 2010a, Rocha et al 2010b) Here, we applied temporal and
484 tissue-specific approaches to study Med12 function in adult astrocytes in the CNS.
485 Critically, the lethality rate dramatically increased with prolonged Tamoxifen treatment,
486 which likely prevented us achieving a high KO efficiency in astrocytes in the CNS (see
487 figure 1R). This technical limitation may have masked some CNS related phenotypes
488 associated with Med12 function. Surprisingly, we found that Aldh111-CreER mediated
489 Med12 deletion in the adult led to rapid hearing loss. Although hearing loss has been
490 reported in cases of Med12-related disorders (Rubinato et al 2020) the underlying
491 mechanisms remained undefined. We found that Med12 is necessary for maintaining
492 the integrity of the stria vascularis in the adult, likely by regulating the expression of ZO-
493 1, E-cadherin, and Cx31. These changes in gene expression alter cell adhesion
494 between basal cells and the spiral ligament, which would likely impede the generation of
495 normal endolymph and culminates in loss of hair cell mechanotransduction.

496 Our studies revealed that basal cells and the spiral ligament of the adult cochlea
497 are transcriptionally dependent on Med12 to maintain their overall structural integrity.
498 We recently identified similar transcriptional dependencies in the adult brain, where the
499 transcription factor NFIA is required to maintain the morphological complexity of
500 hippocampal astrocytes (Huang et al 2020). Together, these studies highlight adult- and
501 region- specific roles for developmental transcription factors in maintaining key aspects

502 of cell structure in mature populations. Med12 functions, in part, to maintain the
503 expression of key proteins that maintain tight junctions (TJs) and gap junctions (GJs),
504 which are critical for normal cochlear function (Wan et al 2013, Wangemann 2006).
505 Prior studies indicate that genetic deletion of either E-cadherin (Trowe et al 2011), or
506 Cx31 (Lopez-Bigas et al 2001) during development results in hearing loss, suggesting
507 that their regulation by Med12 contributes to the observed hearing loss in the adult.
508 While our studies suggest that downregulation of these structural proteins is responsible
509 for the effects of Med12-loss in the cochlea, it is possible that Med12 regulates the
510 expression of other genes that are also important for these effects. Further studies on
511 Med12 target gene networks in basal cells will be critical for deciphering these
512 regulatory networks and associated mechanisms.

513 Our findings demonstrate that loss of Med12 in basal cells and the spiral
514 ligament leads to defects of the stria vascularis. Progressive degeneration of the stria
515 vascularis underlies a form of age-related hearing loss known as strial or metabolic
516 presbycusis (Schuknecht & Gacek 1993). However, we cannot exclude the possibility
517 that loss of Med12 in other tissues contributes to hearing loss in humans. Med12
518 expression is also observed in supporting cells in cochlea and in major cell types in
519 nervous system, raising the possibility that it has related functions in these systems.
520 Nevertheless, our study describes the first animal model that recapitulates hearing loss
521 identified in Med12-related disorders and provides a new system in which to examine
522 the underlying cellular and molecular mechanisms. Moreover, the TJ and GJ proteins
523 that are downregulated in the Med12-deficient cochlea also play important roles in other
524 systems; for example, ZO1 is highly expressed in circumventricular organs of the CNS

525 and is required for the integrity of blood brain barrier (Petrov et al 1994), and a Cx31
526 mutation leads to peripheral neuropathy in addition to sensorineural hearing impairment
527 in patients (Lopez-Bigas et al 2001). Therefore, our studies lay the foundation for further
528 examination of Med12 function in other systems linked to Med12-disorders.

529 To ascertain whether the initial defects in the AG12-KO cochlea are due to strial
530 dysfunction we measured a series of physiological parameters associated with cochlear
531 function at timepoints where basal cell organization is altered and the organ of corti is
532 intact. The most essential evidence for strial pathology in the AG12-KO mouse derives
533 from direct EP recordings, where AG12-KO and AG12-Con mice showed no overlap of
534 EP values, with the AG12-KO averaging ~30 mV decrease in EP. Thus, all of the AG12-
535 KO mice exhibited the single functional requirement for strial pathology, although this
536 does not directly address whether other defects are present at these relatively early
537 timepoints (i.e. 10-14 days post-TM). Other evidence for strial-specific defects at these
538 early timepoints are based on no changes in the DPOAE input/output data between the
539 AG12-KO and AG12-Con, coupled with a relative sparing of DPOAE thresholds versus
540 ABR (DPOAE-ABR) thresholds at the same frequency (Mills, et al. 2003) which yielded
541 higher positive values (ABR threshold higher) in the AG12-KO line. In sum, four
542 functional metrics supported an initial, delimited, strial pathology in AA12-KO mice. EP
543 reduction was the requisite criterion, but since EP recording is a terminal procedure it
544 cannot be used clinically. While we did not first conceive of the threshold and DPOAE-
545 based metrics we used to evaluate strial pathology in our AG12-KO mice, we believe
546 we are the first to apply all three criteria. The principal limitation to such an approach at
547 present is simply the lack of suitable models. Thus, the Med12 KO model may

548 represent a useful system for additional studies aimed at detecting and characterizing
549 isolated chronic EP reduction.

550

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656

657 **Figure Legends**

658

659 **Figure 1: Deletion of Med12 in Mature Astrocytes Selectively Reduces**

660 **Morphological Complexity**

661 **(A–H)** Co-expression of Med12 and Aldh111-GFP in AG12-Con and AG12-KO mice.
 662 Filled arrowheads show co-expression of Aldh111-GFP and Med12; unfilled show
 663 Aldh111-GFP cells that do not express Med12. **(I)** Tamoxifen treatment paradigm. **(J–Q)**
 664 High-magnification confocal images of Aldh111-GFP from AG12-Con or AG12-KO mice
 665 showing reduced complexity in the cortex and hippocampus. **(R)** Quantification of

666 Med12/Aldh111-GFP co-expression in AG12-Con and AG12-KO mice. Data are derived
667 from 3 mice per genotype, 3 slides per region, per mouse: at least 500 cells per region,
668 per mouse. One-way ANOVA. **(S–V)** Scholl analysis of astrocyte complexity in the
669 cortex, olfactory bulb, brainstem, and hippocampus. Data are derived from 3 mice per
670 genotype, 3 slides per region, per mouse, with at least 30 cells per region, per genotype;
671 Two-way repeated-measures ANOVA. **(W)** Quantification of Aldh111-GFP/DAPI co-
672 expression in AG12-Con and AG12-KO mice. Data are derived from 3 mice per
673 genotype, 3 slides per region, per mouse: at least 500 cells per region, per mouse. One-
674 way ANOVA.

675

676 **Figure 2. Deletion of Med12 Results in Hearing Loss and No Changes in Synaptic**
677 **Plasticity in the Hippocampus**

678 **(A)** Quantification of freezing in cued fear conditioning and **(B)** startle responses in
679 acoustic startle. In A, 4 AG12-Con and 5 AG12-KO mice were used; in B, 3 AG12-Con
680 and 4 AG12-KO mice were used. Two-tailed unpaired t test. AG12-Con denotes
681 Med12fl/fl; Aldh111-GFP and Med12fl/y; Aldh111-GFP. AG12-KO denotes Med12fl/fl;
682 Aldh111-CreER; Aldh111-GFP and Med12fl/y; Aldh111-CreER; Aldh111-GFP. *p < 0.05;
683 **p < 0.01. **(C)** Schematic of LTP recording experimental setting. **(D)** LTP traces from
684 AG12-Con and AG12-KO hippocampal slices. **(E)** Quantification of LTP. All
685 electrophysiological experiments are derived from 3 mice from each genotype, ranging
686 from 7 to 13 cells total for each experiment. LTP: long-term potentiation; STP: short
687 term potentiation; NS: no significance; *p < 0.05; **p < 0.001; Student's two-tailed
688 paired **(L)** t test. AG12-Con denotes Med12fl/fl; Aldh111-GFP and Med12fl/y; Aldh111-

689 GFP. AG12-KO denotes Med12fl/fl; Aldh111-CreER; Aldh111-GFP and Med12fl/y;
690 Aldh111-CreER; Aldh111-GFP.

691

692 **Figure 3. Med12 Deletion Causes Rapid Peripheral Hearing Loss in Adult Mice**

693 **(A)** Tamoxifen treatment paradigm for weekly auditory brainstem response (ABR) tests.

694 **(B-E)** ABR thresholds at different frequencies before **(B)**, post-1-week **(C)**, post-2-

695 weeks **(D)**, and post-3-weeks **(E)** of the tamoxifen treatment. Dashed lines mark the

696 maximum pressure, 90dB, used in the tests. In **(B)**, 9 AG12-Con and 8 AG12-KO mice

697 were recorded; in **(C)**, 6 AG12-Con and 7 AG12-KO mice were recorded; in **(D)**, 4

698 AG12-Con and 6 AG12-KO mice were recorded; in **(E)**, 9 AG12-Con and 10 AG12-KO

699 mice were recorded. **(F-G)** ABR to stimuli of 60-90 dB sound pressure level (8 kHz) in

700 AG12-Con **(F)** and AG12-KO **(G)** mice. AG12-Con denotes Med12fl/fl; Aldh111-GFP and

701 Med12fl/y; Aldh111-GFP. AG12-KO denotes Med12fl/fl; Aldh111-CreER; Aldh111-GFP

702 and Med12fl/y; Aldh111-CreER; Aldh111-GFP. Two-way repeated-measures ANOVA. *p

703 < 0.05; **p < 0.01.

704

705 **Figure 4: Deletion of Med12 has Latent Effects on the Structure of Organ of Corti**

706 **(A-X)** Detection of the markers of supporting cells, Sox2, and hair cells, Myo7a, by

707 immunofluorescence on midmodiolar sections of cochleae of AG12-Con and AG12-KO

708 mice. Figures show the organ of corti in the basal turn. **(A-H)** A Comparison of

709 midmodiolar sections of cochleae of AG12-Con and AG12-KO mice 3 weeks post-TM,

710 **(I-P)** 5 weeks post-TM. **(Q-X)** 18 weeks post-TM. AG12-Con denotes Med12fl/fl;

711 Aldh111-GFP and Med12fl/y; Aldh111-GFP. AG12-KO denotes Med12fl/fl; Aldh111-
712 CreER; Aldh111-GFP and Med12fl/y; Aldh111-CreER; Aldh111-GFP. TM: Tamoxifen
713 treatment. Scale bar in D is 20 μ m.

714

715 **Figure 5: Med12 Expression in Basal Cells is Required to Maintain the Stria**
716 **Vascularis**

717 **(A-P)** Detection of the stria vascularis marker, ATP1A1, by immunofluorescence on
718 midmodiolar sections of cochleae of AG12-Con and AG12-KO mice. Figures show the
719 stria vascularis in the apical turn. Yellow arrowheads mark the regions of stria vascularis.

720 **(A-H)** A Comparison of midmodiolar sections of cochleae of AG12-Con and AG12-KO
721 mice 3 weeks post-TM, **(I-P)** 5 weeks post-TM. **(Q-T)** Immunostaining of Med12 in stria
722 vascularis of midmodiolar sections of cochleae of AG12-Con and AG12-KO mice at 2
723 weeks post-TM. Figures show the stria vascularis in the middle turn. **(R)** and **(T)** are the
724 regions from dashed squares in **(Q)** and **(S)**. Unfilled arrowheads mark the side of
725 intermediate cells; yellow arrowheads mark the side of basal cells and the spiral
726 ligament. AG12-Con denotes Med12fl/fl; Aldh111-GFP and Med12fl/y; Aldh111-GFP.
727 AG12-KO denotes Med12fl/fl; Aldh111-CreER; Aldh111-GFP and Med12fl/y; Aldh111-
728 CreER; Aldh111-GFP. TM: Tamoxifen treatment. **U** Quantification of Med12 expression
729 in Aldh111-GFP basal cells; *p<0.05. Scale bar in **D** is 50 μ m and **Q** is 20 μ m.

730

731 **Figure 6: Med12 is Required to Maintain Expression of Cell Adhesion Proteins**

732 **(A-H)** Detection of Glut1, endothelial and basal cell marker, by immunofluorescence on
733 midmodiolar sections of cochleae of AT12-Con and AT12-KO mice at 2 weeks post-TM.
734 Figures show the stria vascularis in the middle turn. **(B-C)** Yellow arrowheads denote
735 Glut1-positive basal cells, labeled by AT12-Con tdTomato reporter; **(F-G)** Unfilled
736 arrowheads denote disorganized Glut-1/tdTomato basal cells in AT12-KO. **(I-T)**
737 Detection of ZO1, E-cadherin, and Connexin-31 proteins by immunofluorescence on
738 midmodiolar sections of cochleae of AG12-Con and AG12-KO mice at 2 weeks post-TM.
739 **(I,M,Q)** Filled arrowheads denote AG12-Con expression of makers. **(K,O,S)** Unfilled
740 arrows denote altered expression of markers in AG12-KO. **U** Quantification of
741 immunostaining from I-T; * $p < 0.05$. Scale bars in **D** and **L** are 50 μm .

742

743 **Figure 7: Loss of Med12 Results in Stria Vascularis-Specific Defects**

744 **(A)** ABR thresholds were significantly higher in KOs versus controls (2-way ANOVA,
745 $F=46.188$, $DF=6$, $p < 0.001$) with no significant interactions. **(B)** EPs were significantly
746 lower in KOs (t-test for unequal variance, $DF=11$, $t=-7.23$, $p < 0.001$). **(C)** $2f_1-f_2$
747 maximum values were not significantly different by group (t-test for unequal variance,
748 $DF=60$, $t=-1.84$, $p < 0.07$). **(D)** Difference in ABR versus DPOAE threshold (ABR-
749 DPOAE for -15 dB SPL criterion) was significantly higher in KOs (t-test for unequal
750 variance, $DF=59$, $t=2.32$, $p < 0.02$). DPOAE metrics were derived from input/output
751 curves for $f_2=12$, 18, and 24 kHz (See horizontal bar in A). Since these curves did not
752 differ by f_2 (not shown), data from all three frequencies were included in C and D

Figure 1

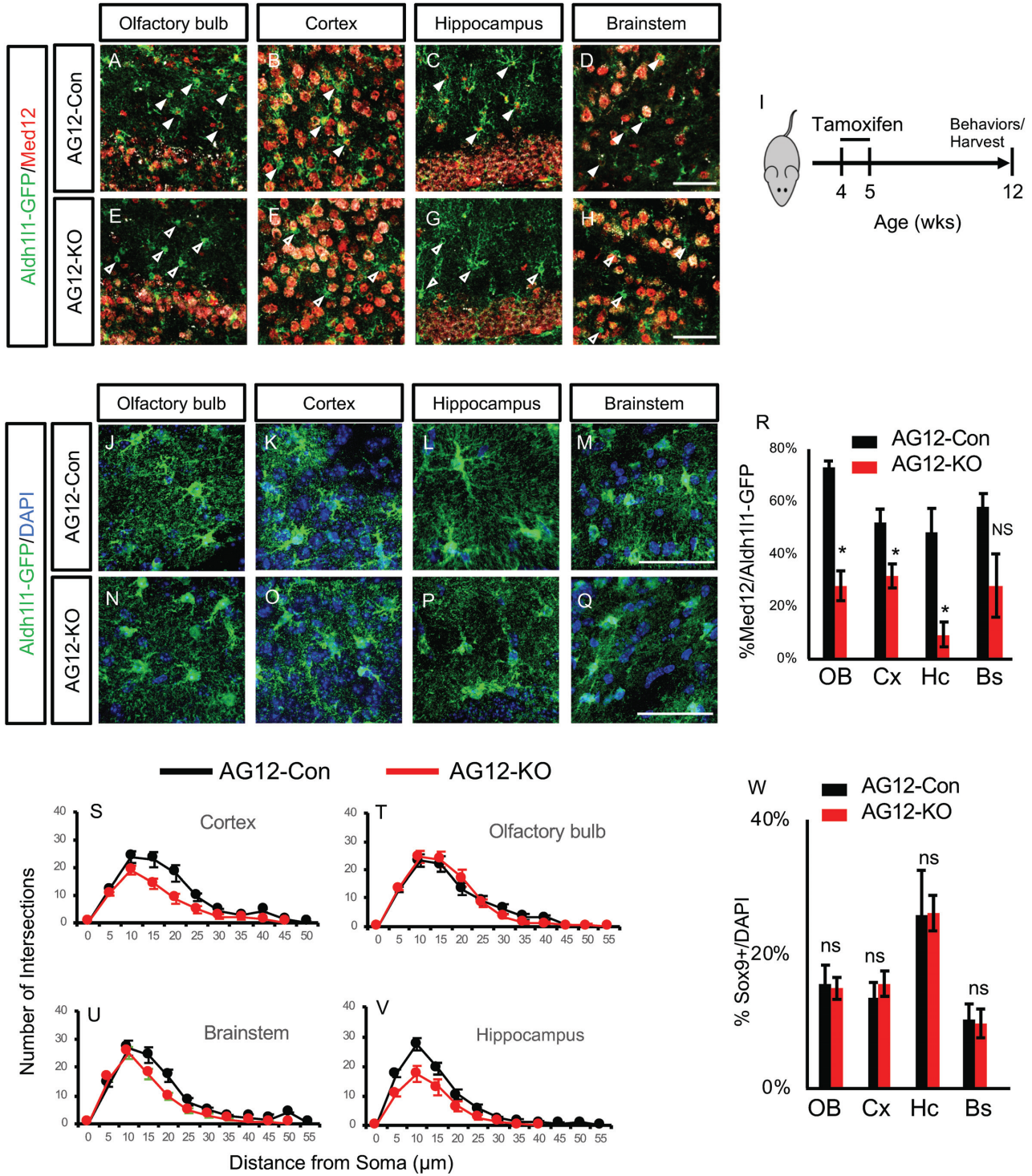


Figure 2

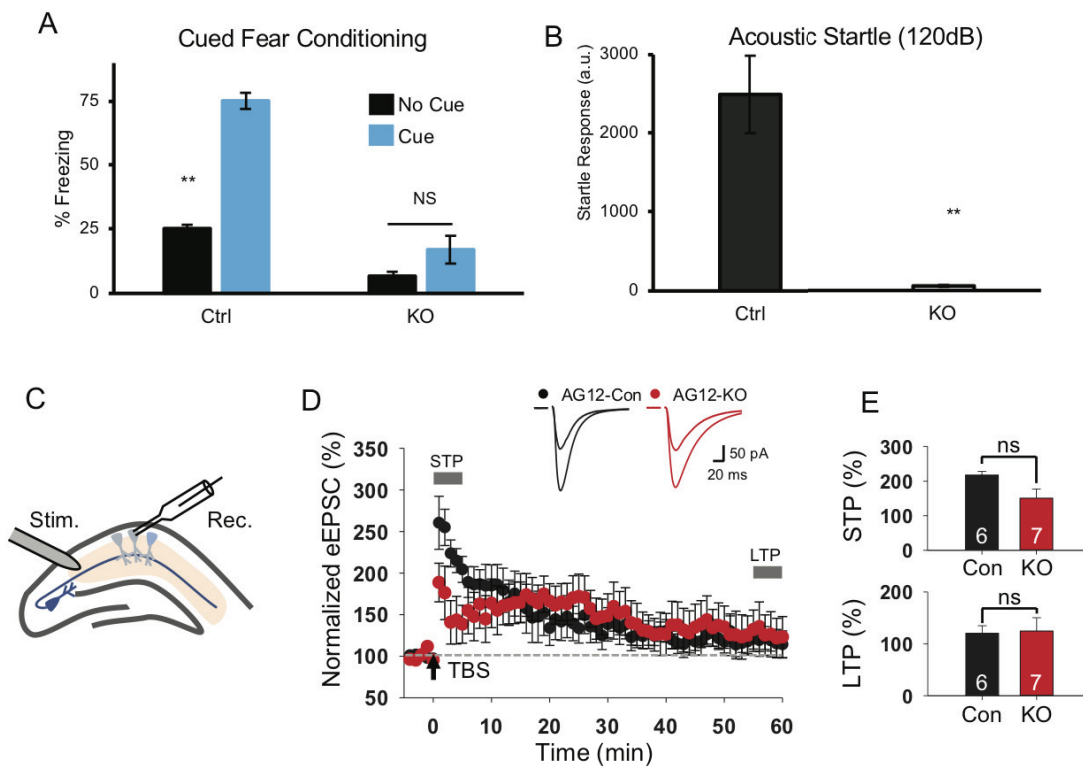


Figure 3

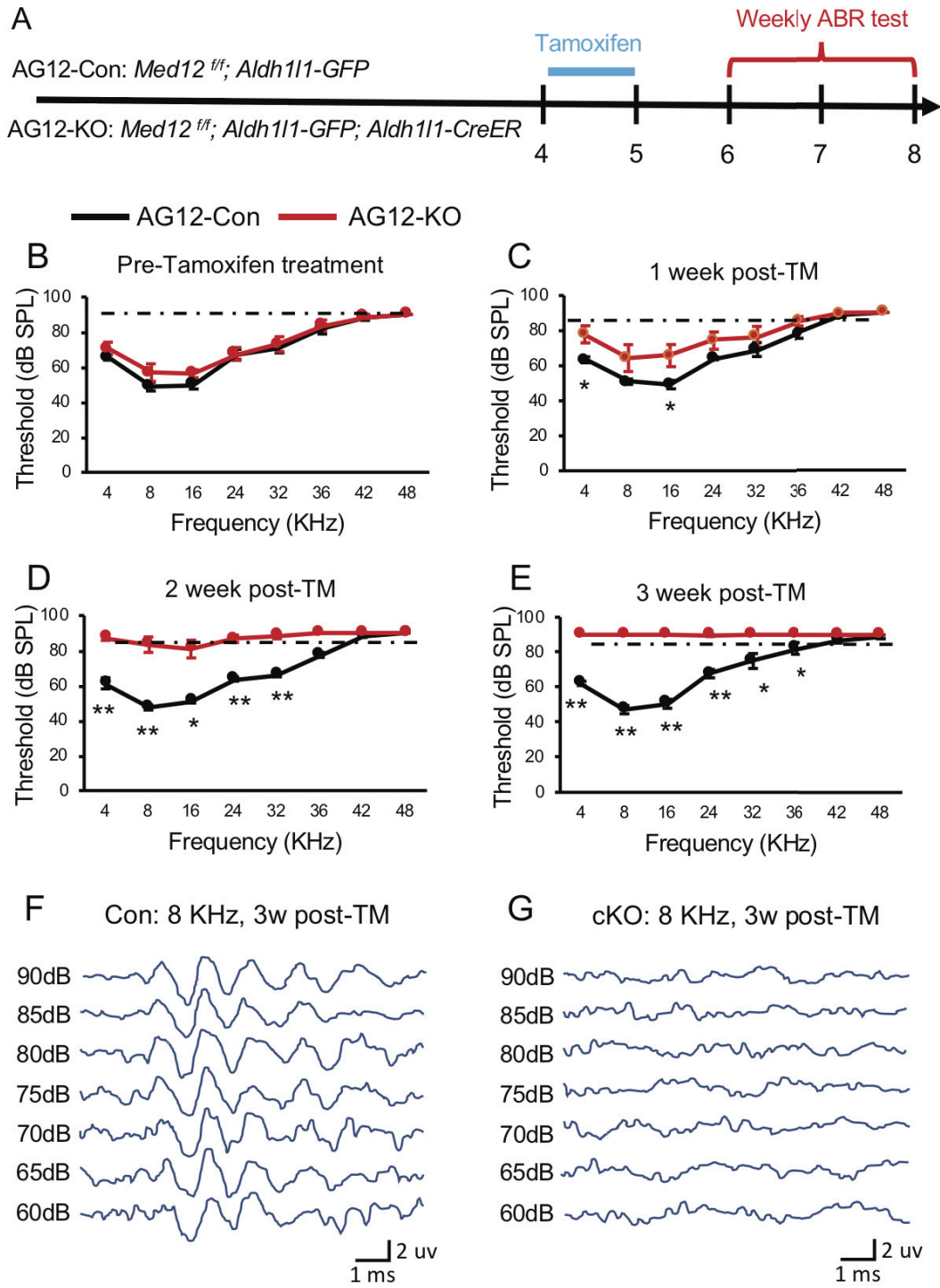


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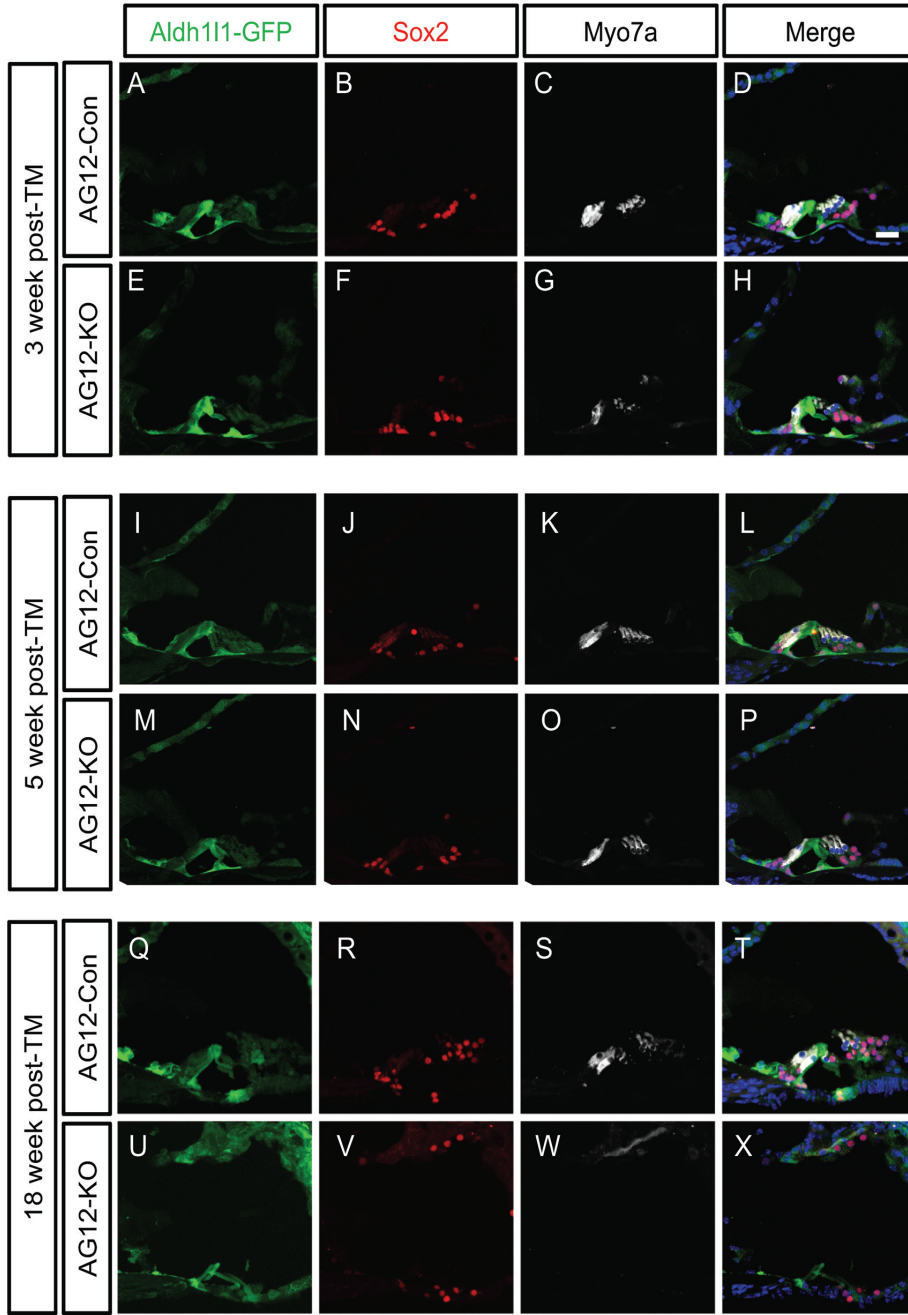


Figure 5

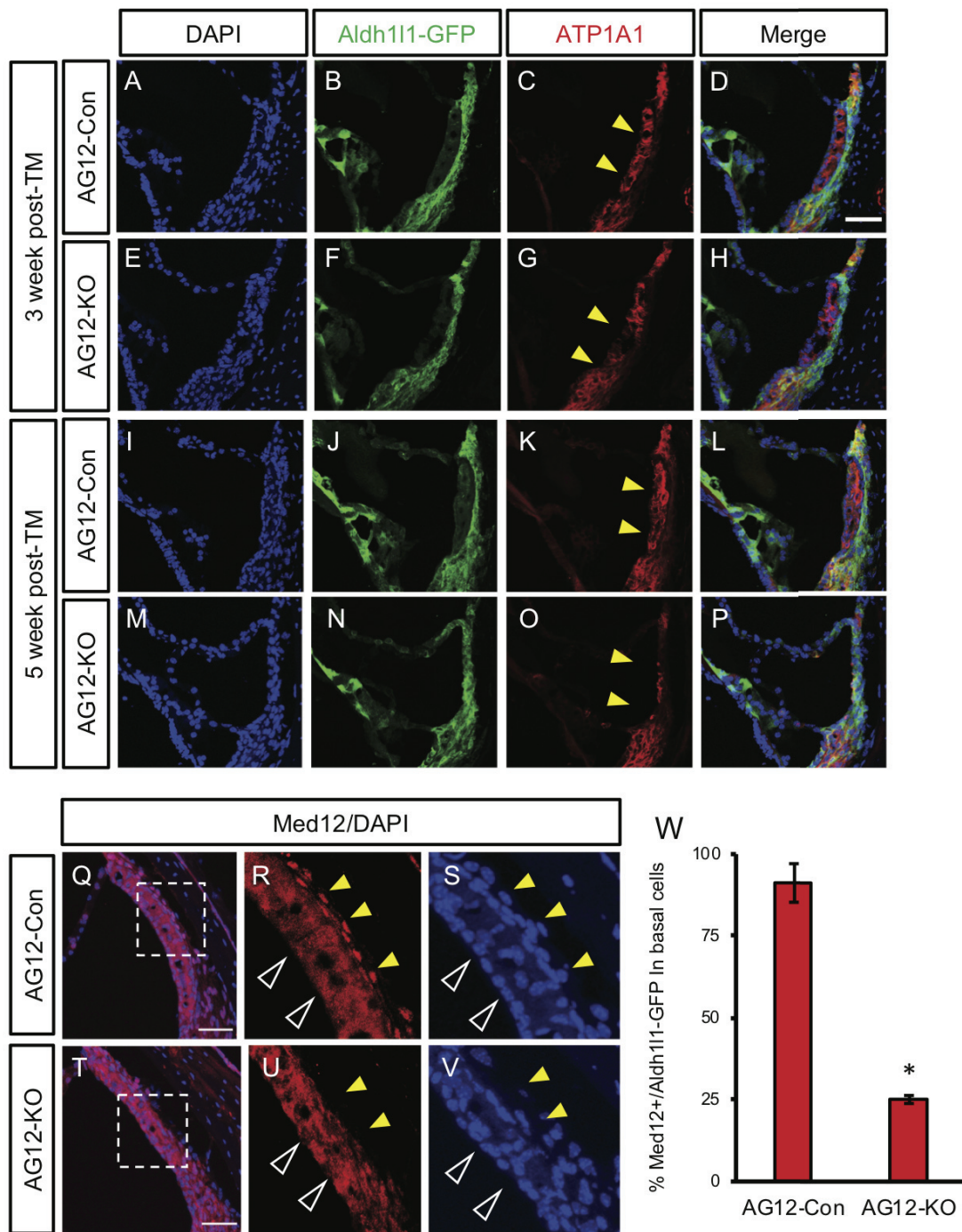


Figure 6

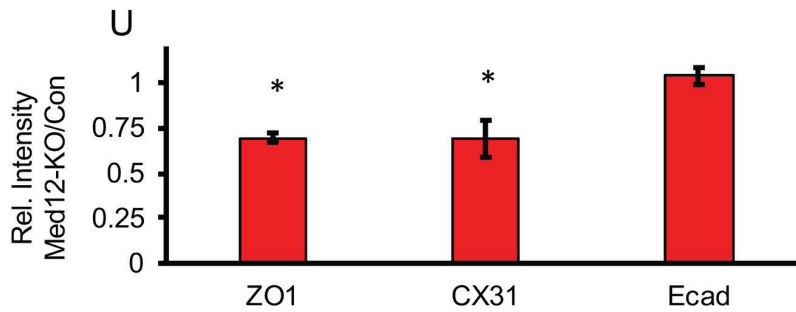
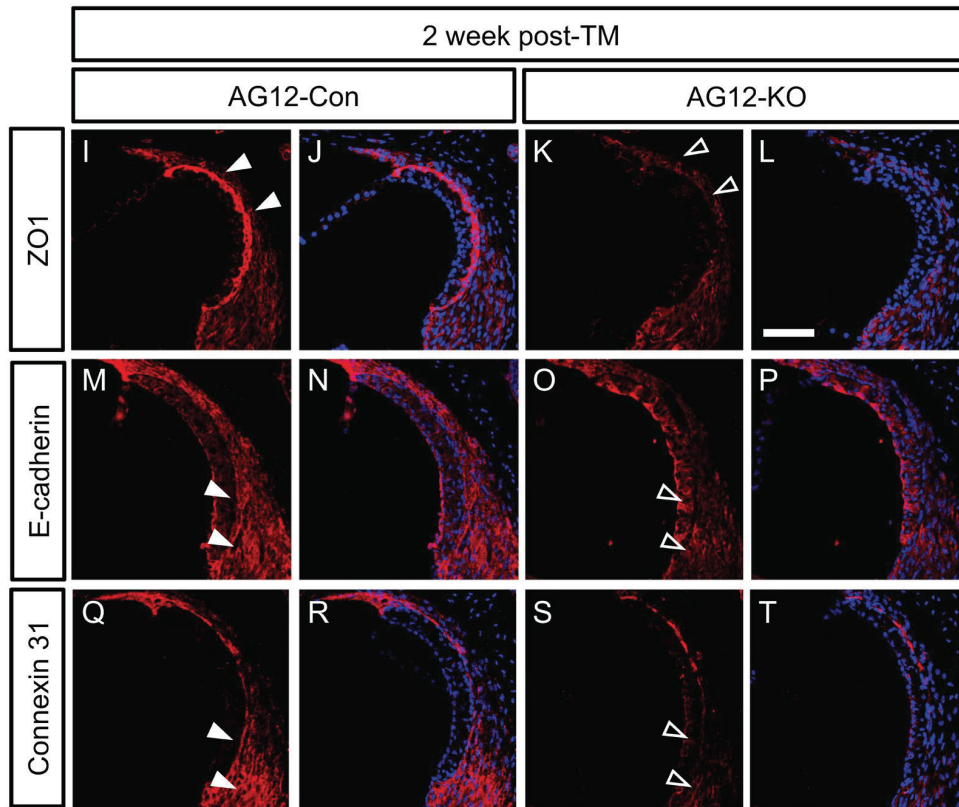
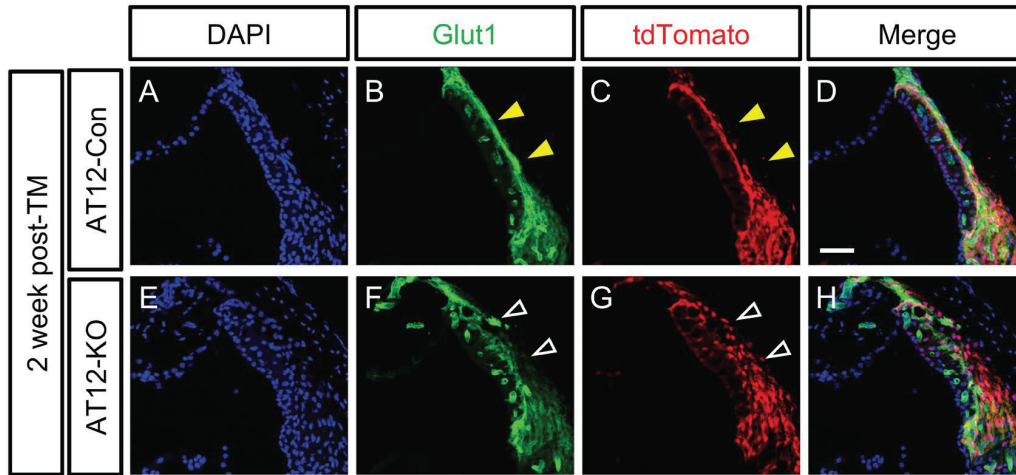


Figure 7

