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Age-related hearing loss is dominated by damage to inner ear sensory cells, not the cellular battery that powers them.

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Abstract

12 Age-related hearing loss arises from irreversible damage in the inner ear, where sound is
13 transduced into electrical signals. Prior human studies suggested that sensory-cell loss is rarely the
14 cause; correspondingly, animal work has implicated the stria vascularis, the cellular “battery” driving the
15 amplification of sound by hair cell “motors”. Here, quantitative microscopic analysis of hair cells,
16 auditory nerve fibers and stria tissues in 120 human inner ears obtained at autopsy, most of whom had
17 recent audiograms in their medical records, shows that the degree of hearing loss is well predicted from
18 the amount of hair cell loss and that inclusion of stria damage does not improve the prediction.
19 Although many aging ears showed significant stria degeneration throughout the cochlea, our statistical
20 models suggest that, by the time stria tissues are lost, hair cell death is so extensive that the loss of
21 battery is no longer important to pure-tone thresholds and that audiogram slope is not diagnostic for
22 stria degeneration. These data comprise the first quantitative survey of hair cell death in normal-aging
23 human cochleas, and reveal unexpectedly severe hair cell loss in low-frequency cochlear regions, and
24 dramatically greater loss in high-frequency regions than seen in any aging animal model. Comparison
25 of normal-aging ears to an age-matched group with acoustic-overexposure history suggests that a
26 lifetime of acoustic overexposure is to blame.

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Significance Statement

29 This report upends dogma about the causes of age-related hearing loss. Our analysis of over 120
30 autopsy specimens shows that inner-ear sensory cell loss can largely explain the audiometric patterns
31 in aging, with minimal contribution from the stria vascularis, the “battery” that powers the inner ear,
32 previously viewed as the major locus of age-related hearing dysfunction. Predicting inner ear damage
33 from the audiogram is critical, now that clinical trials of therapeutics designed to regrow hair cells are
34 underway. Our data also show that hair cell degeneration in aging humans is dramatically worse than
35 that in aging animals, suggesting that the high-frequency hearing losses that define human presbycusis
36 reflect avoidable contributions of chronic ear abuse to which aging animals are not exposed.

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Introduction

41 Presbycusis is age-related hearing loss that cannot be explained by recognized otologic diseases,
42 known ototoxins or hereditary factors. Hearing loss decreases the quality of life and is a major
43 contributor to social isolation and cognitive decline in the elderly (Goman and Lin, 2018). Roughly 50%
44 of 85 yr olds have hearing loss, thus the problem will grow as the population ages. The classic
45 audiometric pattern in presbycusis shows a monotonic decrease of hearing sensitivity with increasing
46 pitch of the test-frequency tones (Fig. 1D).

47 Presbycusis arises in the inner ear (Fischer et al., 2019; Keithley, 2019; Tawfik et al., 2019), where
48 sound is transduced, by inner hair cells (IHCs), into electrical signals carried by auditory nerve fibers to
49 the brain (Fig. 1A,C). The stria vascularis (Fig. 1B) is specialized for ion transport (Patuzzi, 2011), and
50 maintains the electric potential that, like a battery, drives current into hair cells when their transduction
51 channels are opened by sound. The outer hair cells (OHCs) also transduce sound into electrical
52 signals, but also reverse-transduce those electrical signals into mechanical motion, which powers a
53 cochlear amplifier that gives the inner ear its sensitivity (Dallos, 2008). This sensory epithelium is
54 mechanically tuned to high frequencies at the basal end of the spiraling cochlear duct, and to low
55 frequencies at the apex.

56 Because the inner ear lies deep within the temporal bone, it cannot be biopsied, and its sensory
57 structures cannot be resolved with clinical imaging. Thus, inner-ear pathology has been studied in
58 histological sections of tissues extracted after death (Schuknecht, 1974). In classic studies
59 (Schuknecht, 1955, 1964; Schuknecht, 1974; Schuknecht et al., 1974; Pauler et al., 1988; Ramadan
60 and Schuknecht, 1989; Schuknecht and Gacek, 1993), Schuknecht and colleagues concluded that
61 presbycusis was rarely explained by hair cell loss. Rather, it was associated with stria atrophy and/or
62 loss of ANFs and often was uncorrelated with any cochlear histopathology (Ramadan and Schuknecht,
63 1989; Bhatt et al., 2001; Merchant and Nadol, 2010). Correspondingly, studies in aging gerbils also
64 concluded that age-related high-frequency hearing loss is due to stria atrophy, rather than hair cell loss
65 (Tarnowski et al., 1991; Schulte and Schmiedt, 1992; Schmiedt et al., 2002; Dubno et al., 2013).

66 Understanding the functionally significant structural changes in human presbycusis is critical as
67 therapies for rebuilding a damaged inner ear move to clinical trials (Kujawa and Liberman, 2019), and
68 inclusion criteria are based on audiometric data. Clearly, a therapy to elicit hair cell regeneration would
69 be ineffective in presbycusis if the predominate pathology is stria atrophy. Here, using high-resolution
70 light microscopy (Wu et al., 2019), we provide the first quantitative view of inner ear damage in human
71 presbycusis. By applying rigorous statistical modeling, rather than the case-based approaches of the
72 past, we show that hearing loss is well predicted by hair cell loss, that stria atrophy or auditory-nerve
73 loss contribute little or nothing to the audiometric pattern and that audiogram slope is not diagnostic for
74 the stria degeneration. (Pauler et al., 1988; Dubno et al., 2013). We also show that patterns of hair cell
75 loss in aging humans are fundamentally different from those in animal models of aging, and suggest
76 that these differences arise because of the routine ear abuse of life in industrialized societies.

77

78

Methods

79 **Subjects and groups:** Archival slide sets of serially sectioned, celloidin-embedded temporal
80 bones from the Massachusetts Eye and Ear collection were analyzed in the present study. A survey of
81 the entire collection yielded 120 ears that met the following inclusion criteria for this study of “normal
82 aging”: 1) adequate preservation for reliable microscopic analysis, 2) no genetically determined or
83 developmental defects, 3) no history of infections, neoplastic growth, vascular disorders, or bone
84 malformation in the ear, 4) no history of otologic surgery, 5) no history of vertigo, dizziness, or sudden
85 sensorineural hearing loss, 7) no autoimmune disease, and 8) no history of exposure to ototoxic drugs.
86 The selected individuals ranged in age from birth to 104 yrs, and included 75 males and 45 females, all
87 have white, non-Hispanic origins. Of these 120 cases, 77 had audiograms taken within 5.5 yrs of death,
88 and 54 also had word-recognition scores (expressed as a percentage of monosyllabic, phonetically
89 balanced words correctly identified when presented in quiet at an audible level). All procedures and

90 protocols for the study of human tissues included informed consent and were approved by the
 91 Institutional Review Board of the Massachusetts Eye and Ear. The custom computer code as well as
 92 the datasets generated and/or analyzed during the current study are available from the corresponding
 93 author upon reasonable request.

94 **Histological Analysis:** For all histological analysis, observers were blinded to medical history and
 95 audiometric data. For each case, the set of 40-50 archival slides contains ~ 100 cuts through the
 96 spiraling cochlear duct. These slide sets were used for quantification of hair cell loss and stria atrophy.
 97 The cochlear spiral in each case was graphically reconstructed (Merchant and Nadol, 2010) to allow
 98 the percent distance along the spiral of each section through the cochlear duct to be computed and
 99 converted into frequency (kHz) using a cochlear frequency map for human (Greenwood, 1990), as
 100 modified to produce apical-most and basal-most best frequencies of 0.1 and 20 kHz respectively.

101 **Tissue Processing.** When originally archived, each temporal bone was aldehyde fixed, decalcified,
 102 embedded in celloidin, and serially sectioned at 20 μm in the horizontal plane. Every 10th section
 103 was stained with hematoxylin/eosin and slide-mounted (Merchant and Nadol, 2010), while
 104 intervening (unstained) sections were stored in 80% alcohol. For the present study, selected
 105 unstained sections were retrieved, de-celloidinized (O'Malley et al., 2009) and stained with a
 106 fluorescent membrane dye (CellMask[®] Orange, ThermoFisher #C10045) at 1:1000 for 5 min to label
 107 myelin sheaths. The sections were then coverslipped with Vectashield.

108 **Hair Cell counts.** Fractional survival of inner and outer hair cells (IHCs and OHCs) was quantified in
 109 all archival sections from each case using a 100x oil-immersion objective (N.A. = 1.3) and
 110 differential interference optics on a Nikon E800 microscope. In each section through the cochlear
 111 duct, the number of remaining cells in each hair cell row was counted, and the number of missing
 112 cells was estimated as described in more detail elsewhere (Wu et al., 2019). Dividing the two
 113 values yields fractional survival, which needs no further normalization. Hair bundles and cuticular
 114 plates were the main criterion for surviving hair cells, rather than cell nuclei, as done in all prior
 115 work. The hair cell analysis in each case is based on observations at ~100 positions along the
 116 cochlear spiral.

117 **Peripheral axon counts.** Neural degeneration was assessed by counting peripheral axons of
 118 auditory nerve fibers (ANFs; Fig. 1A), rather than their cell bodies, because cell bodies can survive
 119 for years after peripheral axon loss (Chen et al., 2006; Kujawa and Liberman, 2009; Liu et al.,
 120 2015). ANFs were quantified at five locations along the cochlear spiral, one at the tangential section
 121 through each half turn, where the axons are cut in cross section and can be precisely counted (Wu
 122 et al., 2018a). Previously unstained sections were selected from the archive for each half turn,
 123 seeking the section near the junction between the limbus and Tectorial membrane (Fig. 1). After
 124 Cellmask staining (see above), confocal z-stacks were acquired with 0.33 μm z-spacing on a Leica
 125 SP8 using a 63X glycerol objective (N.A. = 1.3). Image stacks spanning the top 5 μm of the section
 126 were acquired. The brightest focal plane was selected, and axons were counted within a 250 μm
 127 mask. Raw data were normalized to average counts from the appropriate cochlear region in six
 128 young normal cases, aged 0 - 5.4 yrs (Wu et al., 2018a).

129 **Strial analysis:** Strial atrophy was quantified in the archival slide sets by measuring the cross-
 130 sectional area of the stria at 13 cochlear locations, equally spaced at 1/8 of each cochlear turn,
 131 avoiding regions where the stria is cut tangentially. At each locus, healthy strial tissue was outlined
 132 and measured using computer-aided anatomy software (NeuroLucida[®]). The sections were
 133 displayed through a video camera connected to a Nikon E600 using a 20x objective (NA = 0.75).
 134 One focal plane, near the middle of the section, was chosen for tracing. Raw data were normalized
 135 to average data from the appropriate cochlear region in young normals (n=5), aged 0 - 5.4 yrs.

136 **Statistical Modeling:** To select the functionally most important histological structures and to fit a
 137 model, linear regression was performed via the LASSO (Least Absolute Shrinkage and Selection
 138 Operator) method using the *glmnet* package in R 3.6.1. Before fitting in the LASSO linear regression,
 139 the data were pre-processed: 1) histopathological measurements were averaged across audiometric
 140 frequency to produce a mean survival and treated as continuous variables; 2) continuous variables (i.e.

141 histopathological measurements, age and hearing level) were normalized (z-score). The best model
 142 was selected via nested 5x10-fold cross-validation. The statistical significance of between-group
 143 differences, when analyzed as a function of cochlear distance/frequency, was assessed using profile
 144 analysis, with the *profileR* package in R 3.6.1.

145

146

Results

147 The present study is based on quantitative histopathological analysis of 120 cases, aged 1 – 104
 148 yrs, from the archival temporal bone collection at the Massachusetts Eye and Ear (Merchant et al.,
 149 2008). In each case, the fractional survival of IHCs and OHCs, ANFs and the stria vascularis was
 150 computed at numerous locations along the cochlear spiral. Using published maps relating cochlear
 151 location to best frequency (Greenwood, 1990), the histopathological data were correlated with the pre-
 152 mortem audiometric results from the same individuals, which were available for 77 of the cases, tested
 153 within 5.5 yrs of death. The example cases in Figure 2 illustrate the type of data obtained in each case,
 154 as well as a number of general trends in the histopathology. Although there is age-related hair cell
 155 death throughout the cochlea, the IHC loss tends to be greater in the basal (high-frequency) half of the
 156 cochlea than the apical half, whereas the apex-base gradient of OHC death is more variable. ANF
 157 degeneration is seen throughout the cochlea, but is usually worse in the cochlear base. In contrast,
 158 stria degeneration is typically worse in the cochlear apex.

159 To consider these apical-basal gradients more quantitatively, and to probe the interactions among
 160 the histopathological metrics, we performed K-means clustering on each measure, across all cases >
 161 50 yrs old (Figure 3). The cluster analysis of stria degeneration profiles identified a “good-stria” group
 162 and a “bad-stria” group of roughly equal size (Fig. 3A₁). The group means suggest that, when present,
 163 stria degeneration tends to be worse in the cochlear apex. In contrast, the mean data derived from
 164 ANF or IHC clustering show clear apex-to-base gradients of degeneration (Figs. 3B₂, 3D₃), while OHCs
 165 show both apical and basal foci of cell death (Fig. 3C₃).

166 Contrary to prevailing views (Pauler et al., 1988; Merchant and Nadol, 2010), the two stria clusters
 167 did not differ significantly with respect to hearing loss (Fig. 3A₄; $p = 0.764$ by profile analysis), and the
 168 audiometric slope in the “bad-stria” group was not flat, as has been suggested (Pauler et al., 1988).
 169 Indeed, the cluster analysis suggests that stria condition is uncorrelated with any of the other
 170 measures: i.e. with IHC or OHC loss (Fig. 3A₃), neural loss (Fig. 3A₂), word-recognition scores or age
 171 (Fig. 3A₅).

172 In contrast, the analyses of ANFs, IHCs and OHCs show significant inter-cluster differences in
 173 audiometric thresholds (Figs. 3B₄, C₄, D₄), as well as age (Figs. 3B₅, C₅, D₅). The largest audiometric
 174 differences are seen when clustering is based on OHCs (Fig. 3C₄), as expected given the role of these
 175 cells in powering the cochlear amplifier needed for hearing sensitivity (Dallos, 2008). The threshold
 176 differences between the ANF clusters (Fig. 3B₄), especially at low frequencies, is surprising since
 177 neural degeneration, *per se*, should not strongly affect audiometric thresholds (Schuknecht and
 178 Woellner, 1955). This association likely reflects the strong correlation between ANF degeneration and
 179 OHC loss (see below). The only significant differences in word scores are seen when clustering for
 180 ANF degeneration (Fig. 3B₅), which fits with the hypothesized importance of neural density in
 181 discrimination of complex acoustic signals, rather than in signal detection in quiet (Wu et al., 2018b).

182 To more directly identify the functionally important structural changes underlying age-related
 183 hearing loss, we averaged each histopathological measure, in each case, in the region surrounding
 184 each audiometric measure. Plotting each of the four histopathological predictors vs. hearing level for all
 185 cases (Fig. 4A,B,C,D) shows that, when only one predictor is used in the regression, hearing level is
 186 best predicted by OHC survival in both the high- and low-frequency halves of the hearing range (see r^2
 187 values in Fig. 4E). However, in the high-frequency half of the cochlea, the single-variable predictions for
 188 IHCs and ANFs are very close to those for OHCs. The slope of the relation between OHCs and
 189 hearing level (Fig. 4B) is steeper at high frequencies, as expected based on the larger contribution of
 190 the OHC-based cochlear amplifier at high frequencies (Liberman et al., 2002; Schmiiedt et al., 2002). As

191 expected based on the cluster analysis, the contributions of strial atrophy to hearing level are
192 insignificant in both the high- and low-frequency regions (Figs. 4D,E).

193 Parsing the relative contributions of ANF, IHC and OHC survival to hearing loss is complicated by
194 the collinearity among the various histopathological measures (Pearson's r values for pairwise
195 correlations are represented graphically in Fig. 4F; scatterplots are shown in Fig. 5). To deal with these
196 collinearities, we adjusted the joint effect of histopathological predictors, plus age, into a multivariable
197 linear regression model using the LASSO method (Yang and Zou, 2015). Recognizing the important
198 differences between the apical and basal halves of the cochlea, e.g. with respect to the contribution of
199 OHCs to cochlear amplification (Schmiedt et al., 2002), we divided the six audiometric frequencies into
200 low-frequency and high-frequency halves, and fit separate regression models for each (Fig. 6). All the
201 predictors were normalized to the same scale, thus the absolute values of the LASSO coefficients (Fig.
202 6B) indicate the relative importance of each predictor to its respective model. In all, 65.3% of the
203 variance in hearing level is explained by the model (Fig. 6A). In both high- and low-frequency regions,
204 OHC and IHC survival are the most important histological predictors (Fig. 6B). ANF survival contributes
205 to the prediction of hearing loss only in the basal half of the cochlea, and strial condition does not
206 contribute in either cochlear region. The fact that age improves the predictive power of both models
207 indicates there are important age-related effects that have not been captured by the present
208 histopathological analysis of the cochlea. These uncaptured effects could be subtler pathologies in
209 surviving structures (e.g. damage to the hair cell stereocilia (Liberman and Dodds, 1984)) or changes in
210 the central auditory pathways outside the scope of the present investigation.

211

212

Discussion

213

A. Differences from prior studies of human presbycusis

214

215 Most forms of acquired hearing loss, including presbycusis, arise in the inner ear, but understanding
216 the underlying damage is limited by the inability to biopsy the inner ear or to reveal cellular detail non-
217 invasively. Prior studies of human otopathology (for review (Merchant and Nadol, 2010)) have been
218 case-based, and the relations between histopathology and hearing loss have rarely been quantitatively
219 examined. A rigorous understanding of these relationships is increasingly important, as biological
220 therapeutics for inner ear disorders are on the horizon, and subject-inclusion criteria depend on
221 predicting the degree and pattern of hair cell or neural loss from non-invasive tests of hearing function.

221

222 Almost all human presbycusis studies are based on light-microscopic evaluation of semi-serial
223 temporal-bone sections. However, prior studies have "binarized" the hair cell analysis, rating cell
224 survival in each section as either 0 or 100% (Ramadan and Schuknecht, 1989; Schuknecht and Gacek,
225 1993; Nelson and Hinojosa, 2006; Merchant and Nadol, 2010; Landegger et al., 2016; Linthicum et al.,
226 2017). Using high-power DIC objectives to optically section the slides, we assessed fractional survival
227 of hair cells (Wu et al., 2019), and conclusions from the two approaches dramatically differ (Fig. 7B,C).
228 In a binarized analysis, only complete hair cell loss is noted (Fig. 7E,F). Thus, Schuknecht's influential
229 studies of presbycusis (Schuknecht, 1955, 1964; Schuknecht, 1974; Pauler et al., 1988; Ramadan and
230 Schuknecht, 1989; Schuknecht and Gacek, 1993) concluded that hair cell loss rarely explains the
231 audiometric pattern, and that strial degeneration or neuronal loss more often responsible.

231

232 Cases analyzed here included all 35 exemplars from Schuknecht's presbycusis studies
233 (Schuknecht, 1955, 1964; Schuknecht, 1974; Schuknecht et al., 1974; Pauler et al., 1988; Ramadan
234 and Schuknecht, 1989; Schuknecht and Gacek, 1993; Merchant and Nadol, 2010): the original slide
235 sets are still available from the Massachusetts Eye and Ear collection. In our multivariable regression
236 analysis of audiogram prediction, coding the data according to Schuknecht's presbycusis classification,
237 i.e. sensory, strial, neural or indeterminate (Schuknecht and Gacek, 1993; Merchant and Nadol, 2010),
238 suggests that that all four presbycusis "types" are equally well "explained" by the same weighting of
239 OHC, IHC and neural losses (Fig. 6A), and thus that this classic, and widely cited, typology is not
240 informative of the underlying causes of age-related hearing loss.

240

241 **B. Inner ear pathology and audiometric slopes: metabolic vs sensory presbycusis**

242 The gold-standard hearing test for humans is the audiogram. Animal studies show that thresholds
 243 provide a sensitive measure of OHC function, because OHCs are biological amplifiers, enhancing the
 244 sound-induced vibrations of the sensory epithelium (Dallos, 2008). Consistent with this, OHC survival
 245 was the most important histopathological predictor of thresholds in the present analysis (Fig. 6B). IHCs
 246 are also critical for hearing, because myelinated ANFs contact only IHCs (Spoendlin, 1972; Liberman,
 247 1982). However, animal studies suggest that IHC loss, *per se*, does not affect thresholds until the loss
 248 exceeds 80% (Lobarinas et al., 2013), because thresholds are tested at octave-frequency intervals, and
 249 the normal IHC density of 500 cells per octave provides massive redundancy (Wu et al., 2018b).
 250 Consistent with this, IHC survival was less important than OHC survival as a threshold predictor (Fig.
 251 6B). Although we need ANFs to hear, thinning the ANF population, without hair cell loss, does not
 252 elevate thresholds (Schuknecht and Woellner, 1955). Therefore, a minor role of ANF survival in
 253 hearing-level prediction (Fig. 6B) was expected. As the neural loss increases, it degrades the
 254 representation of complex stimuli in central pathways, thus, only ANF survival was significantly
 255 correlated with word-recognition scores (Fig. 3B₅).

256 Strial atrophy was common in the aging ears we studied: in atrophic cases, the strial epithelium was
 257 reduced by almost 50% across the entire cochlea (Fig. 3A₁). The poor correlation between strial
 258 damage and either hair cell or neural survival (Fig. 4F, Fig. 5) suggests that a chronic decrease in
 259 endolymphatic polarization does not jeopardize either hair cell or neural health. Nevertheless, the
 260 prevalence of strial atrophy presents a challenge for therapeutics designed to restore hearing:
 261 regenerating hair cells will not restore normal thresholds to ears with strial atrophy, and present results
 262 suggest no clearcut way to diagnose the degree of strial damage.

263 A decrease in endolymphatic potential should raise thresholds, as demonstrated in aging gerbils
 264 (Schmiedt et al., 2002). Surprisingly, strial pathology was uncorrelated with thresholds in our study (Fig.
 265 3A₂, Fig 4D) and was not useful in hearing-level prediction (Fig. 6). Apparently, by the time there is
 266 significant strial atrophy, there is already so much OHC loss (Fig. 3A₃) that the cochlear amplifier is
 267 dysfunctional. Without the amplifier, a reduction of endolymphatic potential from, for example, +100 to
 268 +50 mV, reduces the driving force for IHC transduction currents from 160 mV to 110 mV (IHC
 269 intracellular potential is -60 mV (Fettiplace, 2017)). In a linear regime (without cochlear amplification),
 270 this decreases receptor currents by only 50/160 (i.e. 3.25 dB), which is audiologically insignificant.

271 Prior human and animal studies suggested that two types of presbycusis, i.e. “metabolic” (due to
 272 strial atrophy) vs. “sensory” (due to hair cell loss), can be diagnosed by audiogram slope: i.e. shallow
 273 down-sloping (metabolic) vs. steeply down-sloping (sensory) (Pauler et al., 1988; Schmiedt et al., 2002;
 274 Dubno et al., 2013). Our analysis showed no effect of strial atrophy on slope (Fig. 3A₄) rather that
 275 audiogram slope is determined by apical-basal gradients of IHC and OHC loss (Fig. 2, Fig. 6A). The
 276 idea that sensory cell losses lead to steeply sloping audiograms (Schuknecht and Gacek, 1993) arose
 277 from binarization of hair cell counts: if only complete hair cell loss is noted, the associated profound
 278 hearing loss will necessarily produce a precipitous drop in the audiogram (e.g. Fig. 2C). Pure strial
 279 atrophy will likely result in a gently down-sloping audiogram, and pure strial atrophy may be the
 280 dominant phenotype of the aging gerbil (Schmiedt et al., 2002), however, a pure strial phenotype is
 281 rarely seen in human presbycusis.

282

283 **C. Human presbycusis vs. animal models of age-related hearing loss**

284 Age-related hearing impairment in humans is extensively documented (e.g. (Gordon-Salant, 2005)),
 285 and average audiograms for our cases (Fig. 8B₁) are within 10 dB of age-matched norms (Rosen et al.,
 286 1962). Many laboratories have studied cochlear dysfunction and histopathology in aging animals
 287 (Ulehlova, 1975; Bhattacharyya and Dayal, 1985; Tarnowski et al., 1991; Chen et al., 2009; Ohlemiller
 288 et al., 2010; Cai et al., 2018); however, without quantitative histopathological norms for aging humans,
 289 it has been impossible to judge which aging animal is most relevant. Several mammals (rats, mice and
 290 gerbils) show an apical focus of OHC loss (Fig. 8A₃), similar to that in humans (Fig. 8B₃). This aspect of

291 human otopathology has been overlooked, because the loss is fractional, and fractional loss has been
292 ignored. Furthermore, apical hair cell loss is not reflected in the audiogram, because 1) the apex is
293 tuned to lower frequencies than those tested (Greenwood, 1990), and 2) OHCs contribute less cochlear
294 amplification at low frequencies (Liberman et al., 2002). Nevertheless, the resultant cochlear
295 dysfunction likely contributes to impairment of speech perception, for which low-frequency information
296 is critical.

297 Humans have more mid- and high-frequency threshold shift (Fig. 8B₁) than animals (Fig. 8A₁) and,
298 correspondingly, show more hair cell loss (Fig. 8B_{2,3} vs. Fig. 8A_{2,3}). The contrast is most dramatic for
299 best-studied model, the aging gerbil (Tarnowski et al., 1991) (Fig. 8A₃). In gerbil, apical hair cell loss
300 may explain the low-frequency threshold shifts, but high-frequency losses (Fig. 8A₁) have been
301 convincingly correlated with striae degeneration (Gratton et al., 1996; Schmiedt et al., 2002): the
302 associated reduction in endolymphatic potential has been measured, and the threshold elevations are
303 quantitatively appropriate (Schmiedt et al., 2002). This comprehensive body of animal work has
304 reinforced the idea that presbycusis is not *sensory*, but *metabolic* (Dubno et al., 2013). Although these
305 ideas fit with Schuknecht's typology, the gerbil patterns are not representative of aging humans (Fig. 8).
306 Indeed, none of the animal models mimics the massive IHC or OHC loss in the basal half of the
307 cochlea, documented here for the first time in humans.

308

309 **D. Age-related hearing loss vs. noise-induced hearing loss**

310 Animal studies show that the first signs of permanent noise-induced hearing loss are typically at
311 high frequencies, regardless of the frequency spectrum of the noise exposure (Liberman and Kiang,
312 1978). For decades, researchers have wondered if age-related high-frequency hearing loss is an
313 avoidable result of daily ear abuse from the ubiquitous noise of industrialized society. Classic
314 audiometric studies of aged individuals from isolated societal groups supported this idea, e.g. by
315 showing 60 dB less high-frequency hearing loss in aging Sudanese tribesmen than in age-matched
316 Americans (Rosen et al., 1962). Other studies have reached similar conclusions with experimental
317 designs that controlled for genetic factors (Goycoolea et al., 1986).

318 One obvious difference between animal and human studies is that laboratory animals are born and
319 raised in highly controlled acoustic environments. To gain insight into the role of acoustic overexposure
320 in age-related cochlear histopathology in humans, we separated our normal-aging cohort into those
321 with vs. without a noise-exposure history, i.e. occupational exposures in a noisy workplace or in the
322 military. Although the noise metric was relatively crude, our noise-history group showed significantly
323 more hair cell loss, especially in the basal half of the cochlea, where the differences with animal data
324 are also greatest (Fig. 8B_{2,3}). This additional loss in the noise group was most striking for OHCs, and
325 especially among the 50 – 75 yr old group. This, along with the animal data, is consistent with the idea
326 that the apical focus of hair cell damage is inherently age-related, whereas the larger, and more
327 functionally significant, basal loss in humans is largely noise-induced. If true, the bad news is that we
328 are all abusing our ears, to our significant functional detriment, as we age. The good news is that raised
329 consciousness about the dangers of noisy environments could lead to an improved hearing prognosis
330 for the aging population.

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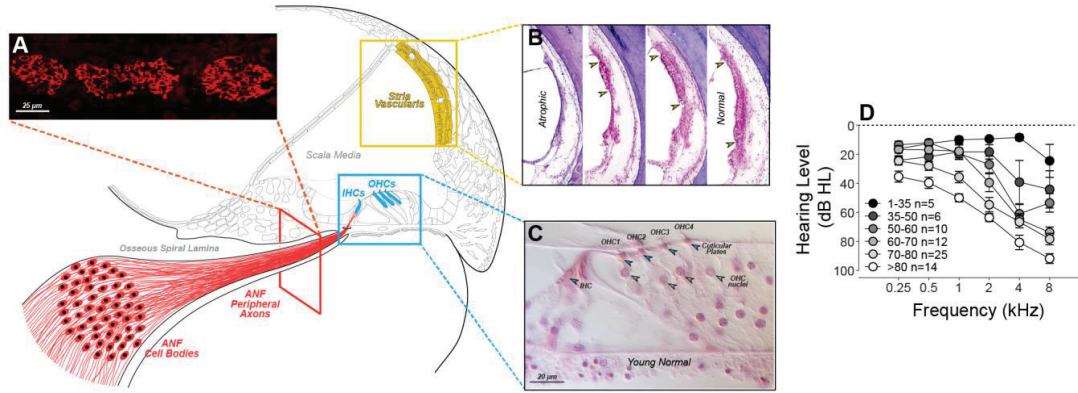
334 Research supported by a grant from the NIDCD (P50 DC 015857) and a grant from the Lauer
335 Tinnitus Center at the Massachusetts Eye and Ear.

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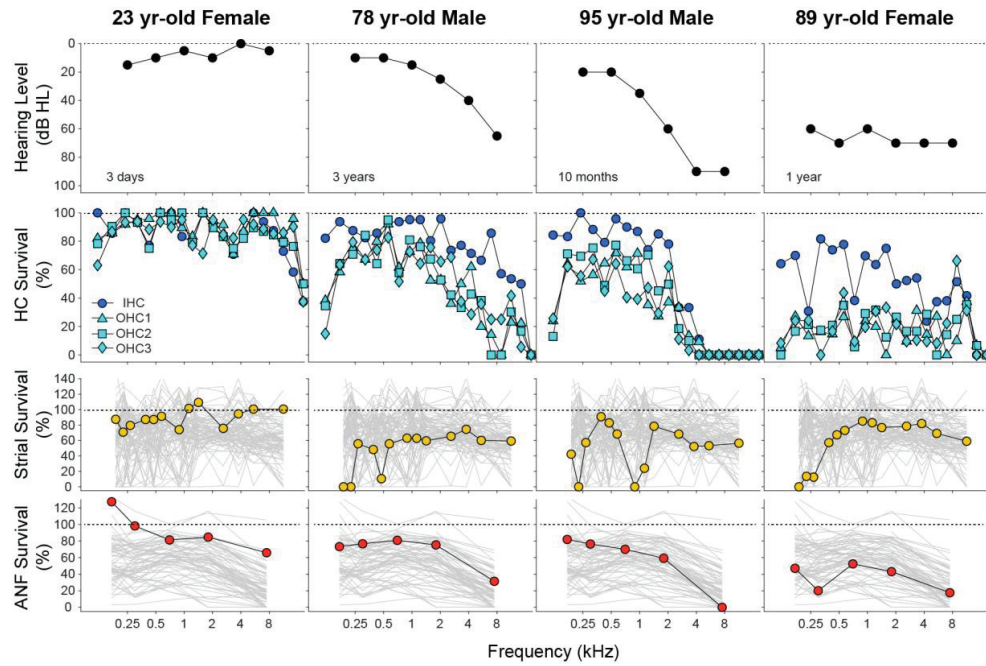
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Figures with Captions



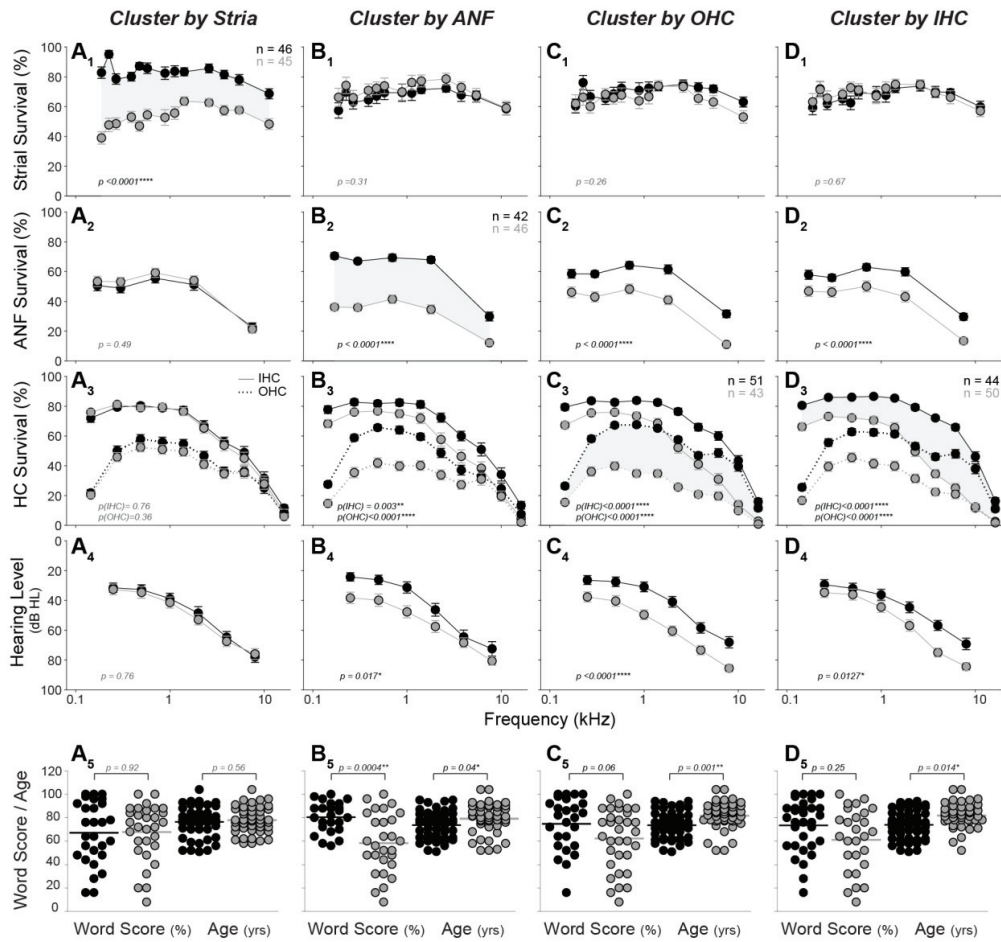
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Figure 1: Three histological features (**A,B,C**) were quantified as histological predictors of hearing level (**D**). Peripheral axons of auditory nerve fibers (ANFs) were counted in cross-sections (**A**). The stria vascularis (**B**), shown from four cases (Merchant and Nadol, 2010), was quantified by measuring surviving epithelial area (regions between the arrowheads). IHCs and OHCs (**C**) were counted using stereocilia and cuticular plates (open arrowheads) as markers of survival (Wu et al., 2019). Audiograms (**D**) show group mean hearing loss (\pm SEMs) for all cases analyzed, except 5 that were excluded (from this plot only) because they were chosen for prior study based solely on their unusual audiometric pattern (Pauler et al., 1988). Cases are separated into six age groups as shown in the key. The male (M) - female (F) mix for each group were: age 1-35, 1 M - 4 F; age 35-50, 5 M - 1 F; age 50-60, 10 M - 0 F; age 60-70, 11 M - 1 F; age 70-80, 17 M - 8 F, and age >80, 8 M - 6 F.



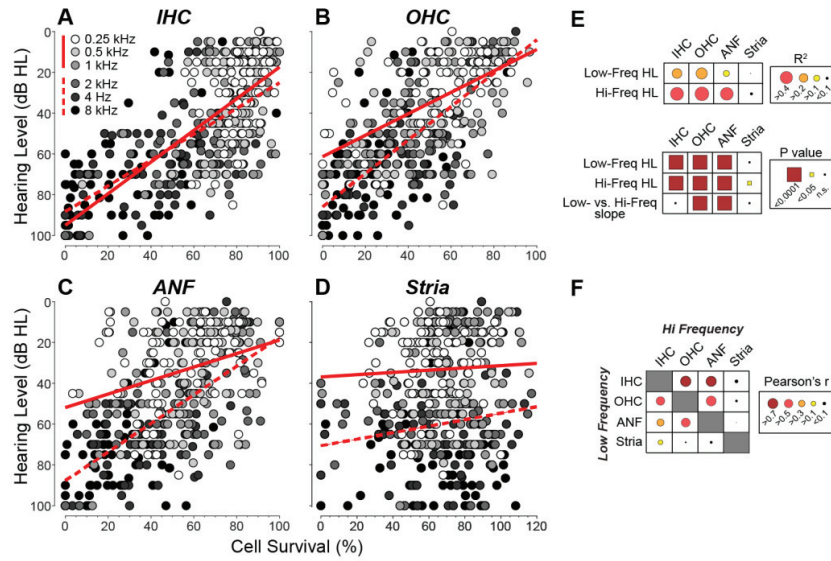
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Figure 2: Histopathological analyses from four of the normal-aging cases in the present study. For each case, the most recent audiogram is shown in the top row, and the histopathological data for hair cells, stria vascularis and auditory nerve fibers are summarized in the 2nd, 3rd and 4th rows, respectively. Hair cell data are binned into 5% increments of cochlear length, and the key at the left applies to all columns. Strial and neuronal data are unbinned and are plotted with the profiles from all cases studied (gray). For the histopathological measures, cochlear location has been converted into frequency according to the cochlea frequency map for human (Greenwood, 1990).



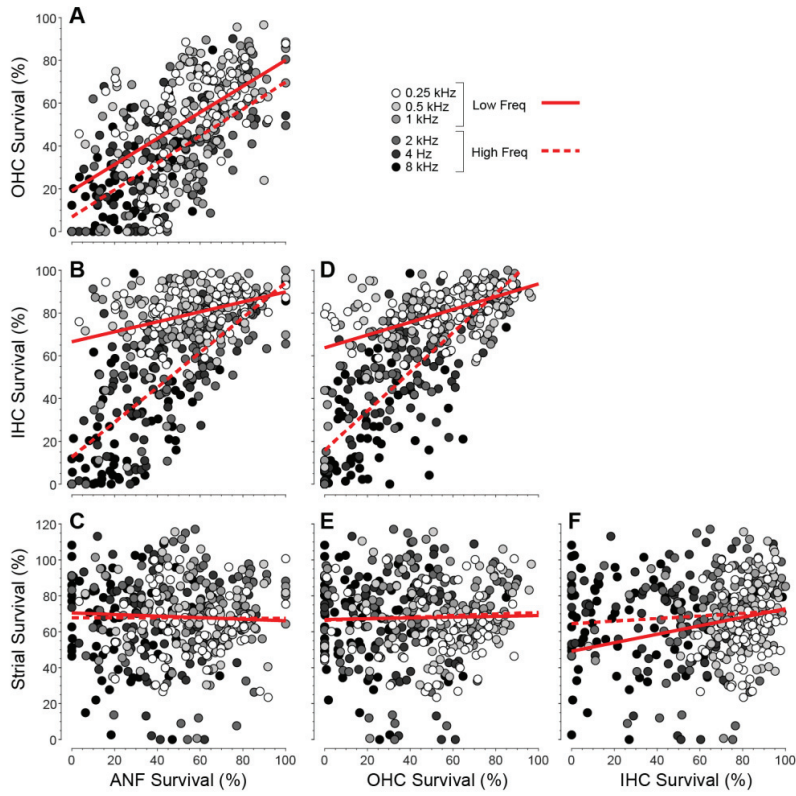
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Figure 3: Cluster analysis shows interactions among predictors and the contribution of each to hearing level and word scores. In each column, K-means clustering was performed based on fractional survival of a different histological metric, as indicated in the header: **A**₁₋₅, stria; **B**₁₋₅, ANF, **C**₁₋₅, OHCs and **D**₁₋₅, IHCs. Clustering was adjusted to define two groups of roughly equal size, as shown by the paired functions separated by grey shading (n's are group sizes). For the top three rows, mean values (\pm SEMs) are shown for the histological metric indicated on the y axis at the left. The bottom two rows show the outcomes for each cluster: means (\pm SEMs) for audiograms (**A**₄, **B**₄, **C**₄, **D**₄) and dot plots for word scores (**A**₅, **B**₅, **C**₅, **D**₅). Ages are shown in the bottom row, along with statistics for the pairwise comparisons: students t-test for the bottom row; profile analysis for the top three rows. The data set for hair cell counts includes all cases older than 50 yrs (n=94: 58 M - 36 F); strial measurements were impossible in 3 cases due to gross port-mortem artifact (n=91: 57 M - 34 F) and ANF counts were impossible in 6 cases due to faint staining (n=86: 56 M - 30 F).



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Figure 4: Regression analyses of the four histological predictors of hearing level (**A-E**), and the pairwise correlations among them (**F**), show that strial atrophy is relatively independent of the other three metrics and is poorly predictive of threshold. Each point in each scatterplot represents the mean of the relevant metric, binned over cochlear loci appropriate to the audiometric frequency, as coded in grayscale (key in **A** also applies to **B**, **C** and **D**). The regression was assessed separately for low (0.25, 0.5 and 1.0) vs. high (2.0, 4.0 and 8.0) frequencies, and the best-fit straight lines are shown by solid and dashed lines, respectively. The r^2 values and p values for the least-squares regression are indicated graphically in **E**, along with p values for the significance of the difference of slope between low- and high-frequency regressions. The Pearson's r values for the pairwise correlations between each of the histological predictors are indicated graphically in **F**: the raw scatterplots are in Fig. 5. Hair cell measures were completed in 77 cases (43 M – 34 F), ANF measure in 72 cases (39 M – 33 F) and strial measures in 76 cases (43 M – 33 F).



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Figure 5: Pairwise correlations among the histological predictors. As in Figure 4, the histological data in each case are binned and averaged over the cochlear locations appropriate to each audiometric frequency: thus each case produces six points on each graph. Data were derived from the 71 cases (39 M – 32 F) with recent audiograms and complete histological data. Least-squares best-fit straight lines are separately computed for low- vs high-frequency regions (solid vs. dashed lines).

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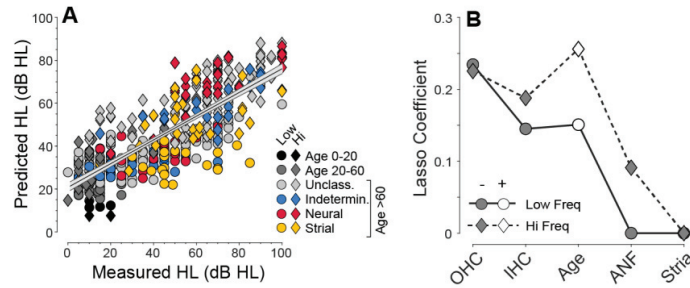
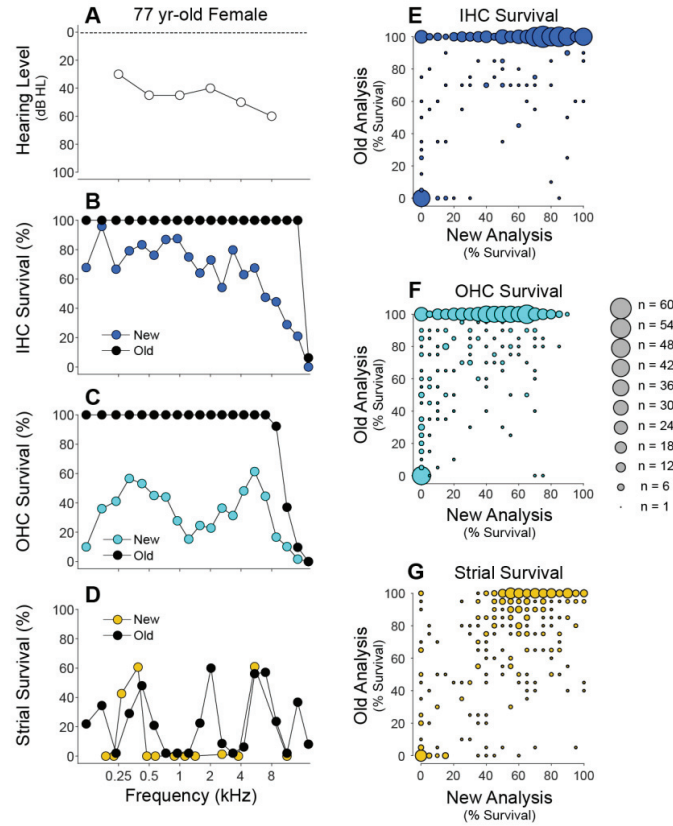


Figure 6: Multivariable LASSO regression was used to derive the best weightings (coefficients) of the histological metrics (plus age; **B**) should be used to predict hearing level (**A**). As for Figure 4, the regression analysis was separately conducted for low- vs. high-frequency regions (circles vs. diamonds in **A** and **B**). The regression line, plus 95% confidence interval, is shown for both frequency regions combined. The differences in coefficient polarity (**B**) arise, because hearing level is expressed as a loss and is positively correlated with age, while histological metrics are expressed as survival and are negatively correlated with hearing level. The data is derived from the 71 cases (39 M – 32 F) with recent audiograms (<5.5 yrs) and complete histological data. The younger ears (<60 yrs) are grouped as shown in key (age 0 - 20, 0 M – 1 F; age 20 – 60, 6 M – 4 F, Unclassified, 25 M – 11 F; Indetermin, 5 M - 3 F; Neural, 1 M – 7 F; strial, 2 M – 6 F). The older ears (≥ 60 yrs) comprise all the exemplars of the four presbycusis types previously studied ($n= 24/35$) (Schuknecht and Gacek, 1993; Merchant and Nadol, 2010), except 11 that were excluded (from this plot only) because ANF counts were impossible or recent audiograms (<5.5 yrs) were unavailable. The Unclassified group comprises additional normal-aging individuals from the Mass. Eye and Ear collection not previously studied.

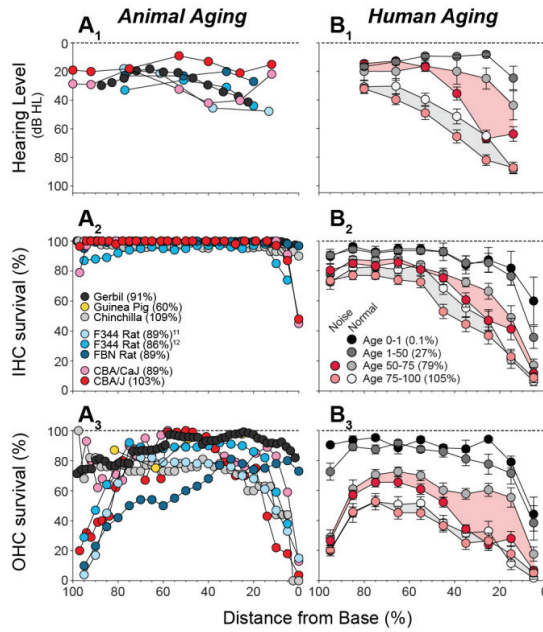
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Figure 7: Comparison of hair cell and strial survival metrics from the same slide sets: published data, i.e. “Old analysis” (Schuknecht and Gacek, 1993) vs. the “New Analysis” in the present study. **A,B,C,D:** One example case, including the audiogram (**A**) and a comparison of results for IHCs (**B**), OHCs (**C**) and stria (**D**). **E,F,G:** Summary comparisons from all the exemplar cases (n=35) from the presbycusis studies of Schuknecht and colleagues (Schuknecht and Gacek, 1993; Merchant and Nadol, 2010). To create the summary plots in **E,F,G**, we scanned the published graphs, digitized and extracted the histogram values, and converted the data into mean survival for each cell type, within each 5% cochlear-length bin. For the same cases, we computed the mean survival for each cell type over the same 5% length bin as assessed in the present study. To display the results, we rounded each mean value to the nearest 5% survival bin, and created these scatterplots in which the area of the symbol is proportional to the number of observations falling within each bin of the matrix.

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Figure 8: Histological and audiometric measures of age-related hearing loss in animals (**A_{1,2,3}**) vs. humans (**B_{1,2,3}**). **A₁**: Cochlear threshold shifts from animal studies. **B₁**: Mean audiograms from the present study, grouped by age as indicated (panel **B₂**), and separated into those with or without a noise-exposure history (age 0 - 1, 3 M - 1 F; age 1 - 50, 5 M - 5 F, age 50 - 75 with noise history, 19 M - 0 F; age 50 - 75 without noise history, 3 M - 9 F; age 75 - 100 with noise history, 21 M - 3 F; age 75 - 100 without noise history, 7 M - 9 F). **A_{2,3}**, **B_{2,3}**: Mean values of IHC or OHC survival as indicated. Data in **A_{1,2,3}** were from the following studies: mouse (Ohlemiller et al., 2010), gerbil (Tarnowski et al., 1991), rat (Turner and Caspary, 2005; Bielefeld et al., 2008), chinchilla (Bohne et al., 1990) and guinea pig (Ulehlova, 1975). Shading in **B_{1,2,3}** aid visual pairing of normal and noise-history groups within each age range. Parenthetical values in keys (**A₂**, **B₁**) indicate mean age, expressed as percentage of median lifespan: gerbil 39 months (Arrington et al., 1973), Chinchilla 15 yrs (Nowal, 2018), Guinea pig 5 yrs (Nowal, 2018), F344 rat 28 months (Solleveld et al., 1984), FBN rat 36 months (Lipman et al., 1996), CBA/CaJ mouse 27 months (Yuan et al., 2012), and CBA/J mouse 23 months (Fox et al., 1997).

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