TGFβ2 -induced senescence during early inner ear development

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Supplementary Information Figure S1. Senescence during the early development of the chicken embryo. Senescence was visualised in specific regions of chicken embryos by SA β G staining. A) Overview of SA β G staining in the HH20 embryo; B) Close up the otic vesicle from an HH20 embryo showing senescence in the otic epithelium particularly intense at the endolymphatic duct; C) SA β G staining in the optic cup, in the lens vesicle and in the olfactory placode at HH17; D) SA β G staining in the pineal gland of an HH17 stage embryo; E) Overview of the head of an HH17 embryo showing senescence in the diencephalon, midbrain, pineal gland; F) SA β G staining in the dorsal region of the pharyngeal clefts; G) Higher magnification of the epiphysis and diencephalon in E showing SA β G staining. Abbreviations: De, diencephalon; Ed, endolymphatic duct; LV, lens vesicle; LVp, lens vesicle pore; Me, midbrain; OIP, olfactory placode; OpC, optic cup; PC, pharyngeal cleft; PG, pineal gland; OV, otic vesicle. Scale bars: 500 μ m (A, E); 250 μ m (C, D, F, G); 125 μ m (B). Representative microphotographs are shown of at least n=2-3 stained embryos.

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Supplementary Information Figure S2. Senescence and apoptosis in explanted chicken otic vesicles. Otic vesicles were isolated from HH18 chicken embryos and incubated ex vivo for 20 h in serum-free culture medium without additives (0S, A-B,F), in the presence of the pan-caspase inhibitor BOC (200 μ M; C-D) or in the presence of the senolytic navitoclax (1 μ M; G). SA β G staining was increased by 1.9-fold in BOC-treated otic vesicles (A,C, quantification in E), whilst apoptotic cell death visualized by TUNEL staining (green) was reduced by 0.9-fold (B,D, quantification in E). Elimination of senescent cells by navitoclax reduced SA β G staining (Figure 2 A,E, quantification in M) and increased apoptosis (F,G, quantification in H). SA β G staining is represented with respect to the total otic vesicle area. TUNEL-positive cells were measured from compiled confocal microscopy projections and normalized to the 0S condition. At least n=4 otic vesicles per condition were studied. Data are shown as mean \pm SEM.*p<0.05, ***p<0.001 vs 0S. of otic vesicles. Orientation: A, anterior; D, dorsal. Scale bars, 150 μ m.

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Supplementary Information Figure S3. *Neurog1* as a biomarker of early ventral region. *Neurog1* mRNA expression levels were measured by RT-qPCR in non-cultured HH18 and HH19 otic vesicles and in dissected dorsal and ventral regions HH19 otic vesicles. Statistical significance was determined with the Student's t-test: ***P<0.005, versus HH18; ###P<0.005, versus dorsal region.

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Supplementary Information S4. Full length blots. To improve the clarity of the information, we grouped cropped blots in Figure 4 western blottings in which we excluded conditions that were not objective of these manuscript (shown in the images as "-").

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Supplementary Information S5 Full length blots. To improve the clarity of the information, we grouped cropped blots in Figure 5western blottings in which we excluded conditions that were not objective of these manuscript (shown in the images as "-"). The antibodies used (Supplementary Information Table S1) have been previously characterized upon the same conditions in otic vesicles cultures (references 17 and 19).

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Antibody	Type ¹	Source/Cat #	Technique
Anti-Akt1/2	GP	Santa Cruz/sc-1619	1:1000 (WB)
Anti-pAkt ^{Ser473}	RbP	Cell Signaling/9271	1:1000 (WB)
Anti-pSMAD2	RbP	Cell Signaling/3101	1:1000 (WB), 1:200 (IHF)
Anti-ERK MAPK	RbP	Cell Signaling/9102	1:1000 (WB)
Anti-pERK MAPK	RbP	Cell Signaling/ 9101	1:1000 (WB)
Anti-phistone 3	Rbp	Upstate/06-570	1:200 (IHF)

Supplementary Information Table S1. Antibodies used

Summary of antibodies used for immunohistofluorescence (IHF) and western blotting (WB). ¹Antibody type: RbP, rabbit polyclonal; GP, goat polyclonal.