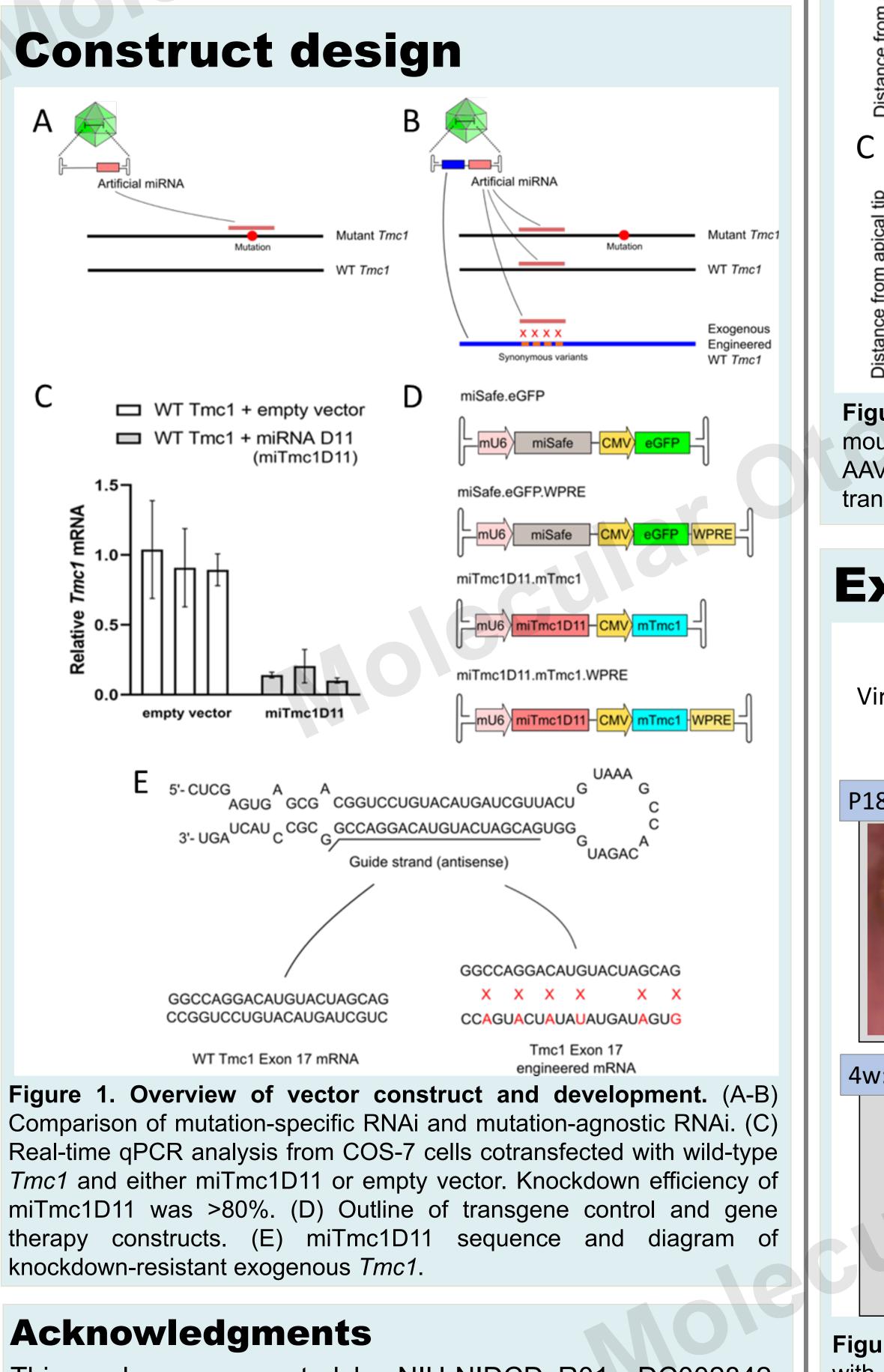


Mutation Agnostic RNA Interference with Concomitant Engineered Gene **Replacement Rescues Hearing in a Mature Murine Model of DFNA36**

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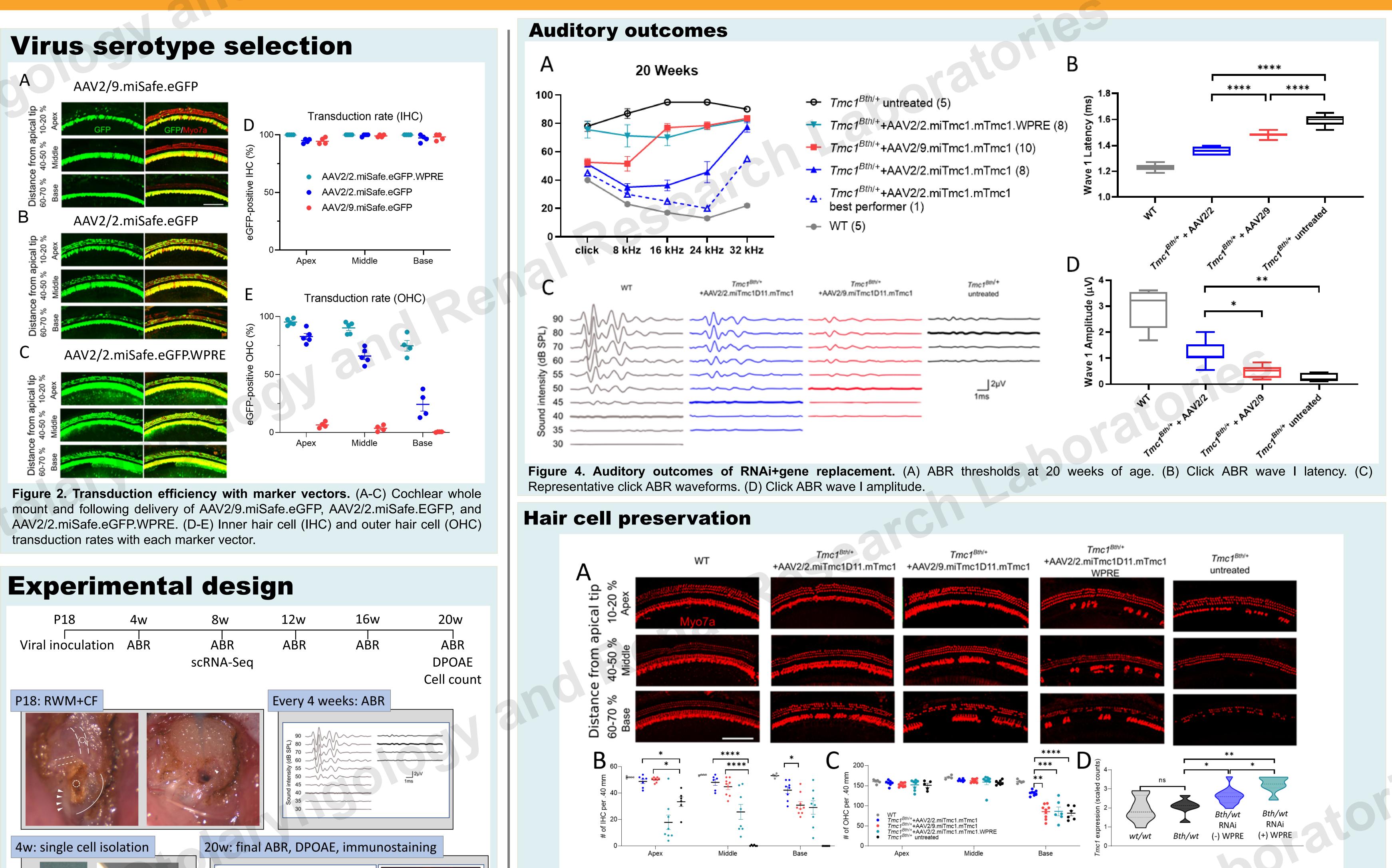
Introduction

TMC1 encodes the pore-forming subunit of the hair cell mechanotransduction channel. Mutations in TMC1 cause dominant (DFNA36) and recessive hearing loss (DFNB7/11) which comprise ~2% of genetic hearing loss. Murine models of DFNA36 have been successfully treated with mutation-targeted RNA interference (RNAi), but these strategies require individually validated constructs for each causative variant. We present a mutation-agnostic approach for RNAi with concomitant gene replacement for preservation of hearing in a murine model of DFNA36.



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P18	4w	8w	12w	16w	20w
iral inoculation	ABR	ABR scRNA-Seq	ABR	ABR	ABR DPOAE
		SCHNA-SCQ			Cell count
8: RWM+CF			Every 4 w	eeks: ABR	
			Sound intensity (dB SPL) 80 40 40 55 00 45 00 45 00 00 00 00 00 00 00 00 00 0		2μV Ims
v: single cell isolation 20w: final ABR, DPOAE, immunostaining					
			v ^t -treated Th	mc1 ^{Bth/wt}	

Figure 3. Summary of experimental timeline. Round window membrane injection with canal fenestration was performed at P16-18. Auditory brainstem response (ABR) was performed every 4 weeks, up to 20 weeks. Manual micropipetting was used to isolate outer hair cells (OHCs) for single-cell RNA sequencing for differential expression analysis. Cell counting was used to characterize hair cell preservation.



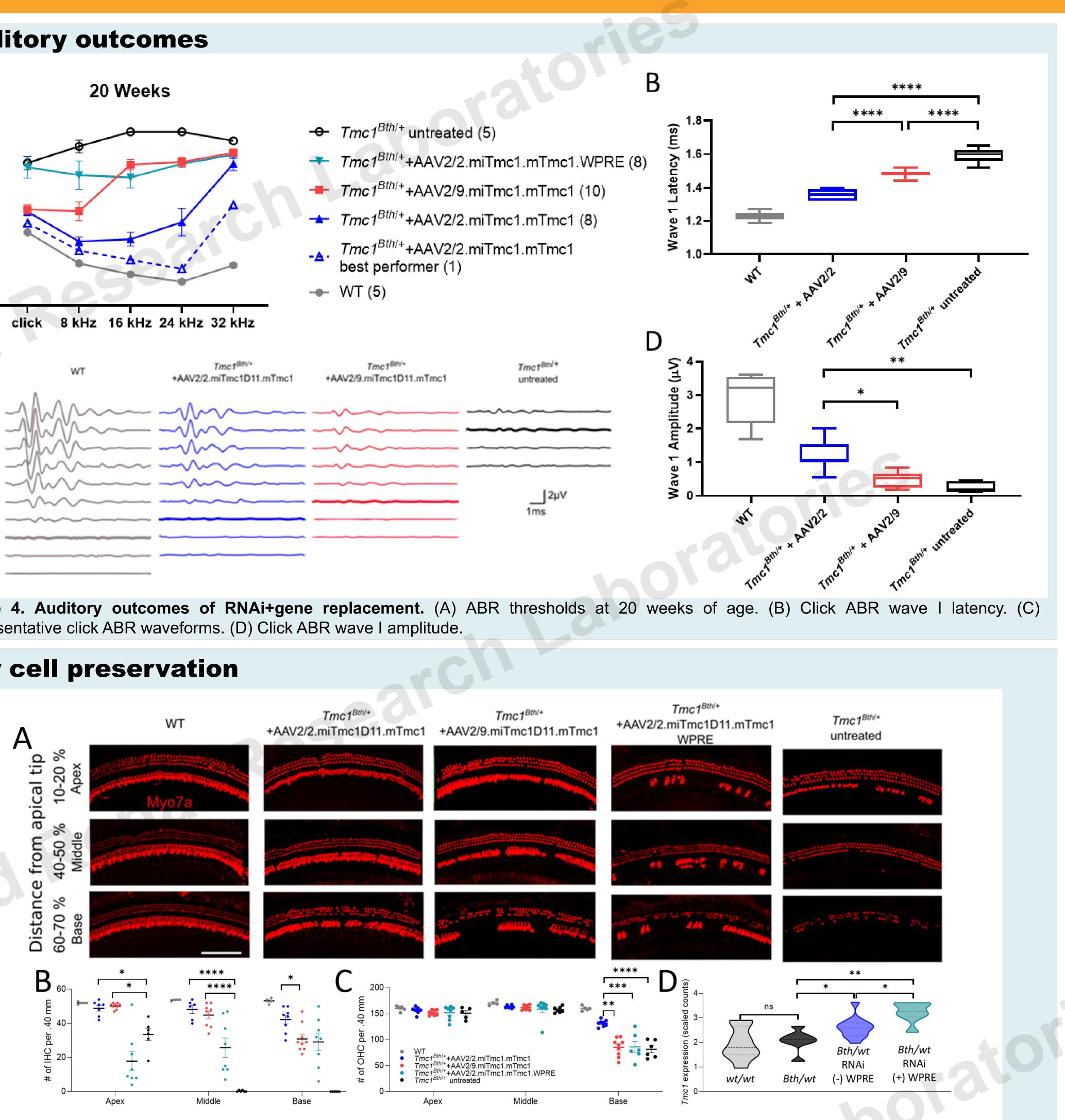


Figure 5. Hair cell preservation in animals which received gene therapy. (A) Cochlear whole mount of treated animals and wild-type and heterozygous controls. (B) Inner hair cell (IHC) counts in 0.4 mm segments of the apical, middle basal turns of the cochlea. (C) Outer hair cell (OHC) counts in apical, middle basal turns of the cochlea. (D) RNAi+gene replacement vectors resulted in increased Tmc1 expression relative to untreated animals; expression was significantly higher in mice which received WPRE-carrying vector.

Conclusions

Mutation-agnostic RNAi with engineered, exogenous replacement achieved preservation of cochlear hair cells and auditory function in the treatment of Tmc1-related hearing loss. This strategy is likely to be broadly applicable in autosomal dominant nonsyndromic hearing loss. The inferior performance of WPRE-carrying vectors suggests that optimization of transgene dosage is critical for optimal auditory outcomes in gene therapy for TMC1-related hearing loss.

