Physiological Profiling of Sensorineural Hearing Loss for Predicting Speech-in-Noise Outcomes



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Introduction

Although the audiogram is the cornerstone of clinical hearing assessment, individuals with similar hearing thresholds often differ in their performance on suprathreshold listening tasks. This variation is likely related to underlying differences in cochlear pathologies that affect neural coding of sound but are hidden from the audiogram, such as cochlear synaptopathy [1], auditory nerve damage [2], and inner-hair-cell dysfunction [3]. Physiological biomarkers have been suggested to identify these hidden pathologies in listeners with normal hearing thresholds, but diagnosis becomes more complicated when outer-hair-cell loss is co-occurring. Given the multi-factorial nature of sensorineural hearing loss, we aim to identify subtypes of *Complex SNHL* based on a battery of biomarkers. Improved diagnostic precision is critical for personalizing hearing loss interventions, including emerging pharmaceutical treatments.

In this study, we assessed whether our battery of physiological biomarkers predicted individual speech-in-noise outcomes. These data, combined with our coordinated experiments in chinchillas with known cochlear pathologies, test the hypothesis that variations in the health of the periphery contribute to individual variation in suprathreshold listening.

Framework Physiological Through coordinated cross-species experiments in chinchillas with Biomarkers Animal Models of SNHL experimentally-induced cochlear Compare biomarker profiles in chinchillas with known pathologies and humans with a SNHL subtypes range of sensorineural hearing loss, we gain mechanistic insight Cochlear into differences in cochlear Pathology pathologies and correlate Profile Identification _abel SNHL subtype clusters diagnostics with outcomes. in humans with predicted pathology Methods **Participants**

- 36 participants
- 18 female, 17 male, 1 non-binary
- 15 with hearing loss (thresholds > 25 dB HL) in the standard clinical range (250-8000 Hz),
- Mean age of participant: 41 years old
- Age range: 19-68 years

Speech in Noise

Speech was presented at 65 dB SPL. A hearing aid simulator fit to DSL v5.0 was used to increase audibility for participants with hearing loss.

	Hea	10
Modified Rhyme Test (MRT)	,	
Inharmonic tone complex masker [4] Maximizes energetic maskir	ng	

-20			
0			
20			\mathbf{X}
40			
60			
80	PTA (.5,	1, 2, 4 kł	-lz):
100	0.25	0.5	1
			Freque
	Ç	uickSl	N

Common clinical measure of speech-in-noise ability



Measure	Sensitive to	Chinchilla	
Hearing Thresholds	OHC function, limited sensitivity to IHC, neural function	Estimated via ABR thresholds at 500, 1000, 2000, 4000, 8000 Hz	Beh bon
Swept DPOAE	OHC function	Swept 500-16000 Hz at 7	
Swept SFOAE	OHC Function, cochlear tuning	Swept 500-16000 Hz using suppressor pa level	
Wideband Middle Ear Muscle Reflex (WB-MEMR)	Cochlear synaptopathy [6]	Wideband probe with broadband elicitors from 45 dB to 105 dB FPL	Wide elicito
Envelope Following Response to RAM- stimuli (RAM-EFR)	Cochlear synaptopathy [7], IHC dysfunction	4 kHz tone with rectangular amplitude modulation at 223 Hz, vertical montage of 3 subdermal electrodes	
Click Auditory Brainstem Response (ABR)	EHF OHC function, Cochlear synaptopathy [6]	Click at 90 dB SPL, vertical montage of 3 subdermal needle electrodes	Hig pe

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Human

navioral audiometry (air and ne conduction) .250-16 kHz

75/65 dB FPL

aradigm at 40 dB FPL probe

band probe with broadband tors from 46 dB to 88 dB SPL

kHz tone with rectangular litude modulation at 223 Hz, 32 channel EEG cap

sh-pass-filtered click at 115 eSPL, 32 channel EEG cap

Cross-Species Biomarkers

Chinchillas were exposed to either noise or an ototoxic drug to create specific cochlear pathologies. Two exposures (CA, TTS) result in little to no change in hearing thresholds, while the other exposures elevate thresholds (PTS, GE).







Predicting Speech-in-Noise

Biomarkers in Human Participants

- PTA, High frequency DPOAE amplitudes, and RAM EFR magnitude are all individually correlated with speech-in-noise performance.
- Though some biomarkers are correlated with each other (like the audiogram and DPOAEs), other measures may provide unique information.



Cluster Analysis

- One metric was selected or calculated from each biomarker:
 - Pure tone average (PTA), 3-8 kHz,
 - Average DPOAE amplitude, 3-8 kHz,
 - Average SFOAE amplitude 3-8 kHz,
 - WB-MEMR threshold,
 - RAM-EFR amplitude, • ABR Wave I/V ratio
- After scaling the metrics, k-means analysis (k=2) was used to identify clusters in the data.
- Clusters from the biomarkers and PTA show differences in MRT thresholds
- In addition to PTA and DPOAE, RAM-EFR and WB-MEMR contribute to PC1

Noise induced temporary threshold shift (TTS) Cochlear synaptopathy (n = 8)cell loss and stereocilia

Complex sensorineural dysfunction (n = 11)<u>Gentamicin (GE)</u>

Outer and inner hair cell loss (n = 5)

Our biomarkers show group-specific effects that are not predicted by the audiogram • For example, WB-MEMR shows differences across the two "hidden" hearing loss groups



Correlations between many variables show a triangular shape like the figure to the left. • Participants with poor DPOAEs have poor RAM EFRs, but participants with good DPOAEs show a range of EFR magnitudes.



(LEFT) The first two principal components accounted for 46.06% and 15.25% of variance. (RIGHT) The clusters significantly predicted MRT threshold (p = 0.03).





- **OHC** function (OAEs).

Additional data collection and analyses

- **Predicting pathology**

References: [1] Kujawa & Liberman, J Neuro 2009, [2] Wu, O'Malley, de Gruttola, & Liberman, J Neuro 2020 [3] Lobarinas, Salvi, Ding, Hear Res 2013 [4] Stone & Moore, JASA 2014 [5] Kidd, Mason, Swaminanthan, Roverud, Clayton, & Best, JASA 2016 [6] Bharadwaj, Hustedt-Mai, Ginsberg, Dougherty, Muthaiah, Hagedorn, Simpson & Heinz, Comm Bio 2022 [7] Vasilkov, Garrett, Mauerman, & Verhulst Hear Res 2021

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Developing Biomarker Profiles

Conclusions

• The classical view of SNHL is that broadened tuning related to OHC dysfunction is the major contributor to speech-in-noise deficits, but differences between individuals with similar audiograms are highlighted by measures other than those that directly reflect

• Expanding the clinical test battery has potential to improve clinical decision making and prepares clinicians to individualize treatment and identify candidates for new therapeutics that target specific mechanisms of hearing loss.

Future Directions

• Our statistical analyses perform better with more data, so data collection is ongoing. These biomarkers used in this study show promise and warrant future analysis to optimize useful individual and combinations of metrics.

• Cochlear histology on exposed chinchillas will provide additional insight to assist with prediction of cochlear pathologies in humans.

<u>Correlations with hearing loss etiology and self-report of hearing ability</u>

• Human participants each completed questionnaires regarding noise exposure history, hearing health history, and subjective report of hearing ability (e.g. SSQ)

References