

# A MULTILINEAR (TENSOR) FRAMEWORK FOR HRTF ANALYSIS AND SYNTHESIS

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## ABSTRACT

This paper introduces a multilinear (tensor) framework for the analysis and synthesis of the head-related transfer function (HRTF). The HRTF is the result of the confluence of two factors, sound location and person (anatomy). Our multilinear modeling technique employs a tensor extension of the conventional matrix singular value decomposition (SVD), known as the  $N$ -mode SVD [1] which explicitly represents the HRTF in terms of its constituent factors. Anatomical data is mapped to our multilinear HRTF model using regression. This mapping defines a data-driven model capable of producing different personalized HRTFs from easily obtained anatomical measurements. We show that our approach yields objectively superior results to those of a mapping based on principle components analysis (PCA).

**Index Terms**— Audio systems, HRTF, Tensors, Multilinear algebra

## 1. INTRODUCTION

The head related transfer function (HRTF) and its corresponding impulse response, the head related impulse response (HRIR), are essential components of many approaches to binaurally-based spatial audio synthesis. The HRTF describes, for a particular person and sound source direction, the frequency response that results from a complex interaction of anatomically-based reflection and diffraction effects. Along with interaural time differences and interaural level differences, these filtering effects are believed to be the primary cues used to discern sound direction [2]. Thus by filtering a sound sample with the appropriate HRTF, convincing virtual auditory environments can be constructed.

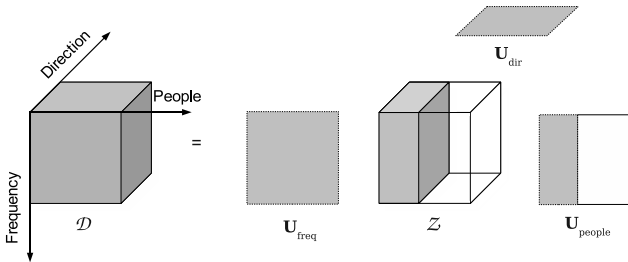
The HRTF is a function of two variables, sound source direction and the external morphology (head, torso, and outer ear) unique to each person. In general, a set of HRTFs recorded from one person will yield poor results when used to spatialize audio for someone else. Use of a mismatched set of HRTFs often results in front-back confusion as well as difficulty perceiving source elevation [3]. Unfortunately, it is difficult and time-consuming to measure HRTFs, making empirically-based customization unrealistic. This has given

rise to significant interest in model-based approaches. Several recent papers describe approaches to HRTF customization using anthropometric data [4, 5]. Systems based on anatomical measurements are a particularly attractive approach as the data is easy to gather and computer vision systems promise to make this process even easier [6]. Other prior modeling work [7, 8] has used PCA to analyze HRTF data. However, PCA works best with functions of one variable as it can only capture the total variation present in a dataset. This inability to distinguish between variations due to sound source direction and variations due to morphology can result in PCA disposing of information that is important for spatial localization.

In this paper, we introduce a nonlinear, multifactor model of HRTFs that generalizes conventional PCA. Whereas PCA employs linear (matrix) algebra, our approach exploits multilinear (tensor) algebra. Multilinear algebra, the algebra of higher order tensors, is able to learn the interactions of the multiple factors inherent to HRTFs and separately encode each of the modes. We exploit the dataset’s natural factoring by using the  $N$ -mode SVD, a tensor decomposition which allows us to explicitly represent each of the factors in the dataset (frequency, people, and direction) as orthogonal vector spaces. Each of these factor spaces (location and person) encodes a representation, statistically invariant of the other, that describes how the HRTF data is influenced by that particular factor.

Next, we define a data-driven model capable of producing different personalized HRTFs from easily obtained anatomical measurements by mapping anthropometric data to our HRTF data model. Anthropometric features are mapped to the people factor space produced by the  $N$ -mode SVD through regression. Then, given anatomical measurements of a person not in the database, we use the regression model to solve for that person’s representation in people space. This representation is then combined with the results of the  $N$ -mode SVD to generate a custom set of HRTFs. This approach is similar in spirit to that of Inoue et al. [9] who use a multiple regression model to map anthropometric features to the coefficients representing each HRTF after applying PCA.

## 2. TENSOR ALGEBRA



**Fig. 1.**  $N$ -mode SVD decomposition illustrated for a 3-mode tensor,  $\mathcal{D}$ . The result is core tensor  $\mathcal{Z}$  and mode matrices  $\mathbf{U}_{freq}$ ,  $\mathbf{U}_{people}$ , and  $\mathbf{U}_{dir}$ . Dimensionality reduction, shown for mode  $\mathbf{U}_{people}$ , is accomplished by truncating the rows of  $\mathbf{U}_{people}$  as well as the corresponding portions of  $\mathcal{Z}$ .

A tensor is a higher order generalizations of a scalars (0<sup>th</sup>-order tensor), a vectors (1<sup>st</sup>-order tensor), and matrices (2<sup>nd</sup>-order tensor).<sup>1</sup> A tensor  $\mathcal{X} \in \mathbb{R}^{I_1 \times I_2 \times \dots \times I_N}$  is said to be of *order*  $N$ . The *mode- $n$  product* of a tensor,  $\mathcal{X} \in \mathbb{R}^{I_1 \times I_2 \times \dots \times I_n \times \dots \times I_N}$ , by a matrix,  $\mathbf{M} \in \mathbb{R}^{J_n \times I_n}$ , is denoted as  $\mathcal{X} \times_n \mathbf{M}$ . This product yields a tensor  $\mathcal{Y} \in \mathbb{R}^{I_1 \times \dots \times I_{n-1} \times J_n \times I_{n+1} \times \dots \times I_N}$  whose entries are given by  $y_{i_1, \dots, i_{n-1}, j_n, i_{n+1}, \dots, i_N} = \sum_{i_n} x_{i_1, \dots, i_{n-1}, i_n, i_{n+1}, \dots, i_N} m_{j_n i_n}$ .

The singular value decomposition (SVD), which forms the basis of principle components analysis (PCA), factors a matrix  $\mathbf{D} = \mathbf{U}_1 \mathbf{S} \mathbf{U}_2^T$ , where  $\mathbf{U}_1$  is a set of orthonormal basis vectors spanning column space,  $\mathbf{S}$  is the diagonal singular value matrix, and  $\mathbf{U}_2$  is a set of orthonormal basis vectors spanning row space. Using the mode- $n$  product, this decomposition can be written as mode- $n$  product of the two orthonormal vector spaces  $\mathbf{D} = \mathbf{S} \times_1 \mathbf{U}_1 \times_2 \mathbf{U}_2$ . The more general  $N$ -mode SVD decomposes a tensor of order  $N$  into its  $N$  constituent vector spaces:

$$\mathcal{D} = \mathcal{Z} \times_1 \mathbf{U}_1 \times_2 \mathbf{U}_2 \dots \times_n \mathbf{U}_n \dots \times_N \mathbf{U}_N \quad (1)$$

The core tensor  $\mathcal{Z}$  plays a role analogous to the eigenvalue matrix  $\mathbf{S}$  in the SVD (although note that its non-diagonal entries are not necessarily zero) and modulates the interaction between modes. The columns of each mode matrix,  $\mathbf{U}_i$ , span the space of that factor while the rows encode the particular instances of that factor that are present in  $\mathcal{D}$ . For a more detailed discussion of the  $N$ -mode SVD, see Vasilescu and Terzopoulos [1].

Perhaps one of the most useful aspects of the  $N$ -mode SVD is the ability to perform *targeted* dimensionality reduc-

<sup>1</sup>We use uppercase calligraphic letters to denote tensors, uppercase bold letters to denote matrices, and lowercase bold letters to denote vectors.

tion. In contrast to traditional PCA where dimensionality reduction affects all aspects of the data, we can truncate the basis vectors of each mode matrix separately, affording a much finer degree of control. Dimensionality reduction through truncation has a bounded error that is equal to the sum of the eigenvalues of the truncated vectors [10]. Figure 1 illustrates the  $N$ -mode SVD as well as dimensionality reduction of one mode.

## 3. DATA

We use the publicly available CIPIC database [11] which contains head related impulse responses (HRIRs) recorded for both ears of 45 subjects over 1250 directions (25 azimuths  $\theta \in [-80..80]$  and 50 elevations  $\phi \in [-45..230.625]$  specified in interaural polar coordinates) spaced roughly uniformly over the head sphere. Each HRIR is 200 samples long (approximately 4.5 ms) and was recorded at a 44.1 kHz sampling rate in 16-bit resolution. Each HRIR was transformed into an HRTF by a 512-point FFT and then filtered to contain only frequencies between 500 Hz and 16 kHz, leaving 181 frequencies in each HRTF. We then combined left and right ear data for each direction and person by concatenating each pair of HRTFs into a single vector of length 362.

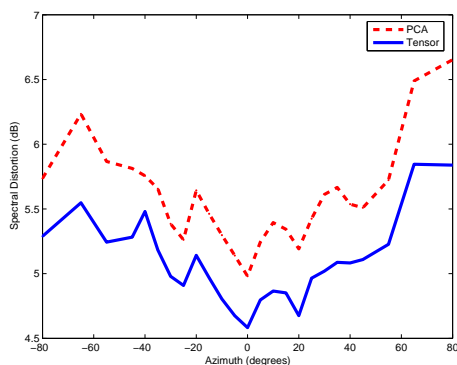
The database also contains anthropometric data for each of the subjects, although eight subjects are missing one or more morphological measurements and therefore were not used in our experiments. For the remaining 37 subjects, we used the following morphological features: head width, cavum concha height, cymba concha height, cavum concha width, fossa height, pinna height, pinna width, intertragal incisure width, cavum concha depth, pinna rotation angle, and pinna flare angle. The measurements for both ears of each person were concatenated together and PCA was performed on the resulting vectors. The first 10 principle components were retained as they explained over 90% of the data's variance. A column of 37 ones was then prepended to the matrix of anthropometric coefficients to provide a constant term for the regression model. We refer to the resulting dataset as  $\mathbf{A}$ .

## 4. HRTF CUSTOMIZATION

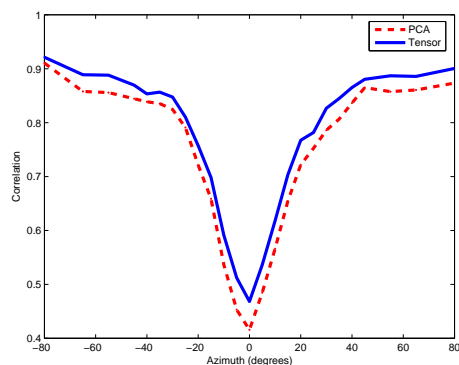
Using the HRTF data described above, we define a data tensor,  $\mathcal{D} \in \mathbb{R}^{F \times P \times D}$  where  $F$  is the number of frequencies (362),  $P$  is the number of people (37), and  $D$  is the number of directions (1250). Using the  $N$ -mode SVD, we decompose the tensor as follows (see Figure 1):

$$\mathcal{D} = \mathcal{Z} \times_1 \mathbf{U}_F \times_2 \mathbf{U}_P \times_3 \mathbf{U}_D \quad (2)$$

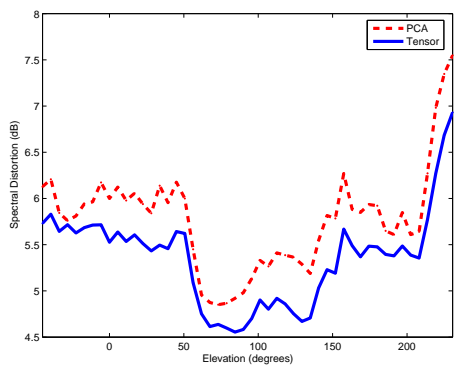
In our experiments, we reduced the dimensionality of the people mode,  $\mathbf{U}_P$ , from 37 to 6, giving  $\hat{\mathbf{U}}_P$ . The direction model,  $\mathbf{U}_D$ , was also reduced from 1250 down to 1041, yielding  $\hat{\mathbf{U}}_D$ . We now wish to find a mapping  $\mathbf{A} \mapsto \hat{\mathbf{U}}_P$ . This can be done with the following regression model,



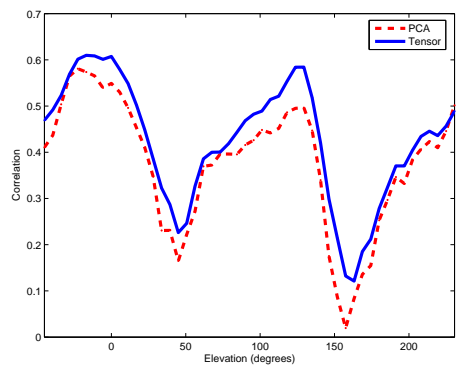
(a) Spectral distortion values for  $\phi = 0, \theta = [-80..80]$



(b) Correlation values for  $\phi = 0, \theta = [-80..80]$



(c) Spectral distortion values for  $\theta = 0, \phi = [-45..230.625]$



(d) Correlation values for  $\theta = 0, \phi = [-45..230.625]$

**Fig. 2.** Comparison of representative tensor and PCA modeling results for one fixed elevation and one fixed azimuth. Both spectral distortion scores as well as correlation values between synthesized and real HRTFs are shown.

$$\hat{\mathbf{U}}_P^T = \mathbf{B}\mathbf{A}^T \quad (3)$$

where  $\mathbf{B}$  is a matrix of coefficients. Using  $\mathbf{A}^+$  to denote the pseudoinverse of  $\mathbf{A}$ , we can easily solve for  $\mathbf{B}$  as:

$$\mathbf{B}^T = \mathbf{A}^+ \hat{\mathbf{U}}_P \quad (4)$$

Given a vector of anthropometric measurements,  $\mathbf{a}_{\text{new}}$  for a person not in the database, we solve for the vector of coefficients,  $\hat{\mathbf{u}}_{\text{p}_{\text{new}}}$ , that correspond to the new person in people space.

$$\hat{\mathbf{u}}_{\text{p}_{\text{new}}}^T = [\mathbf{1} \ \mathbf{a}_{\text{new}}^T] \mathbf{B}^T \quad (5)$$

We can now solve for the complete set of HRTFs,  $\mathcal{D}_{\text{new}}$ , using  $\hat{\mathbf{u}}_{\text{p}_{\text{new}}}$ :

$$\mathcal{D}_{\text{new}} = \mathcal{Z} \times_1 \mathbf{U}_F \times_2 \hat{\mathbf{u}}_{\text{p}_{\text{new}}}^T \times_3 \mathbf{U}_D \quad (6)$$

## 5. EXPERIMENTS

To test the effectiveness of our approach, we conducted a suite of experiments comparing our multilinear approach to PCA. For the PCA approach, we considered the HRTFs across subjects for each direction separately. For each direction,  $d$ , we performed PCA on the corresponding HRTFs of all subjects, reducing their dimensionalities down to 5. This yielded a matrix,  $\hat{\mathbf{C}}_d$ , containing the coefficient vectors for direction,  $d$ , for each subject. As with the tensor framework, regression was then applied to map the anthropometric data to  $\hat{\mathbf{C}}_d$ . Note, that the PCA model retained a total of  $5 \times 1250 = 6250$  basis vectors while the tensor model retained  $6 \times 1041 = 6246$  basis vectors. Also, note that the PCA approach requires a separate regression model for each direction (1250 in total), while the tensor model requires only one.

Performance was evaluated using a cross-validation ap-

proach. For each of the 37 subjects we constructed both a tensor as well as a PCA-based mapping using the other 36 subjects' data. These models were then used to synthesize the held-out subject's HRTFs from their anthropometric data. We then computed the reconstruction error and correlation between each of the held-out subjects' synthesized and real HRTFs. These correlation and error values were then averaged across subjects. The spectral distortion was used as an error metric between the real and synthesized HRTF data:

$$SD(H, \hat{H}) = \sqrt{\frac{1}{N} \sum_{i=1}^N \left( 20 \log \frac{|H_i|}{|\hat{H}_i|} \right)^2} \quad (7)$$

where  $|H_i|$  is the magnitude of the  $i^{th}$  frequency of the true HRTF,  $|\hat{H}_i|$  is the magnitude of the  $i^{th}$  frequency of the synthesized approximation, and  $N$  is the number of frequencies.

Figure 2 shows a pair of representative experimental results. Spectral distortion and correlation values are shown for all azimuths with a fixed elevation (Figure 2a and Figure 2b) and for all elevations with a fixed azimuth (Figure 2c and Figure 2d). In both cases 0 was arbitrarily chosen for the fixed value.

As can be seen in Figure 2, the tensor approach resulted in lower spectral distortion and higher correlation values in a variety of experimental conditions. In fact, in all 1250 directional cases, the spectral distortion was lower and the correlation was higher for the tensor model than for the PCA model.

## 6. CONCLUSIONS

We have presented a novel tensor framework for modeling person-specific HRTFs. Our generative model is fully data-driven and capable of producing individualized sets of HRTFs from easily obtained anatomical measurements. We have demonstrated that our approach compares favorably to a PCA-based framework, achieving consistently lower spectral distortion scores and consistently higher correlation values across all directional conditions. Furthermore, our system maps anthropometrics to HRTFs using a single regression model while the PCA approach requires a separate regression model for each spatial direction.

We are currently planning listening experiments to provide a subjective measure of our system's abilities. In addition, we are investigating how the tensor framework might be used in other areas of HRTF research, such as spatial interpolation and source position estimation [12].

## 7. REFERENCES

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