

AUDIO SYNCHRONISATION WITH A TUNNEL MATRIX FOR TIME SERIES AND DYNAMIC PROGRAMMING

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ABSTRACT

Precise multimodal studies require precise synchronisation between audio and video signals. However, raw audio and video from video recordings can be out of sync for several reasons. In order to re-synchronise them, a dynamic programming (DP) approach is presented here. Traditionally, DP is performed on the rectangular distance matrix comparing each value in signal *A* with each value in signal *B*. Previous work limited the search space using for example the Sakoe Chiba Band (Sakoe and Chiba, 1978). However, the overall space of the distance matrix remains identical. Here, a tunnel matrix and its according DP-algorithm are presented. The matrix contains merely the computed distance of two signals to a pre-specified bandwidth and the computational cost is equally reduced. An example implementation demonstrates the functionality on artificial data and on data from real audio and video recordings.

Index Terms— Audio-video Synchronisation, Image-loss Compensation, Tunnel Matrix, Tunnel DP-algorithm, Storage Requirements

1. INTRODUCTION

The number of studies that analyse audio and video signals has increased with the attempt to account for the multimodality of social interaction involving amongst others speech and gestures [1, 2] and investigations into the natural habitat of speech: face-to-face interaction. Although conversation analysts often have to restrain themselves to low quality audio and video recordings in order to preserve the naturalness of talk-in-interaction [3], studies in interactional phonetics [4] that combine qualitative and quantitative methods by applying computation techniques on automatically extracted features, e.g. Kurtic et al. [5] Gorisch et al. [6] and Bertrand et al. [7], require high quality audio recordings from close-speaking microphones. The audio has to be synchronised

with the video recordings, if the multimodality of talk should be analysed for the same data. Usually, precautions are taken to ensure high standards for audio-video synchronisation, such as by Edlund et al. [8] who enhanced their recording setup and included a rotating turntable with an LP-disk that was scratched at a certain place and a visible marker. Audio and video signals were then synchronised in subsequent steps in order to render the data analysable. The analysis of desynchronised audio and video signals, however, is almost impossible. One source of error can be a clocking difference of two recording devices that capture the same scene. Another source of error can be the hardware equipment itself. For example, video recordings or capturing systems that imply MiniDV technology regularly introduce errors, i.e. a loss or surplus of image frames. Both problems were encountered with recordings of a multimodal speech corpus [9]. Previously, such problems have been tackled by inserting or deleting frames at specific places by hand, as was performed for the CID corpus [10]. A more efficient and re-usable solution that does not have to involve additional equipment, such as turntables, was envisaged for the alignment of the signals, as e.g. of the audiovisual MapTask corpus [9].

2. APPROACH

The advantage of video recordings is that the camera usually includes an audio channel that captures the same scene as the separate audio recordings do. It means that two similar signals of the same quality, i.e. acoustics, exist for the computation of their alignment. Thus, the aim for synchronising audio and video is to insert or delete images from the video according to the alignment of the audio signals that stem from two different devices.

Dynamic Programming (DP) and Dynamic Time Warping (DTW) are methods that can be applied here, as they are used to align time series in an optimal way. They allow to determine where one signal has to be stretched in time or shrunk in time in order to match the other signal [11]. These methods had their revival in speech processing in recent years, e.g. with algorithms that search for re-occurring acoustic patterns for modelling infant speech acquisition [12] and algorithms that evaluate the similarity in prosodic patterns of conversa-

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tional participants performing basic social actions [13]. Developments have been performed by Salvador and Chan [14] who increased the time efficiency of DTW algorithms. They solved a similar problem by matching two acoustic signals in order to synchronise them with a video signal. However, a problem of limited space and computational cost remains unsolved.

2.1. Space and computational cost problem of DP

Many DP problems today take large time series as underlying data. For example, an attempt to align audio recordings of the length of half an hour (1,800s), even using a relatively sparse sampling rate of 25 samples/s (as standard video frames), results in a distance- or similarity matrix of $(25 \times 1,800)^2$, that are 2,025,000,000 value pairs. Additional to the space requirements of this matrix comes the computational cost of dynamic programming to evaluate the best alignment path of the two underlying vectors. The Sakoe Chiba Band [15] or the Itakura Parallelogram [16] are techniques that can help to reduce that computation by limiting the search space to a band around the diagonal of that matrix from the start point to the end point of the underlying vectors.

One problem of current matrix processing is the limited storage space that is available on e.g. personal computers. Sometimes, it is physically not possible to create the distance matrix for two time series. Fortunately, this is even not necessary because the majority of values are not required in the case for the audio-out-of-sync problem. If the DP algorithm that is supposed to be applied to such data is not required to look at the edges of the distance matrix, that space is used without need. A sparse matrix, as introduced by Gilbert et al. (1992) [17], could be used in order to implement such a band. However, DP has not been demonstrated on sparse matrices, yet. In order to save space and computation power to solve the desynchronisation problem at hand, a new structure for the values of the distance matrix is suggested in section 3.

2.2. Conceptual and aesthetic problem

A more conceptual and probably aesthetic problem of a rectangular distance matrix (1) is the distribution of the axes.

a_1b_1	a_1b_2	a_1b_3	a_1b_4	a_1b_5	a_1b_6	a_1b_7	a_1b_8	a_1b_9
a_2b_1	a_2b_2	a_2b_3	a_2b_4	a_2b_5	a_2b_6	a_2b_7	a_2b_8	a_2b_9
a_3b_1	a_3b_2	a_3b_3	a_3b_4	a_3b_5	a_3b_6	a_3b_7	a_3b_8	a_3b_9
a_4b_1	a_4b_2	a_4b_3	a_4b_4	a_4b_5	a_4b_6	a_4b_7	a_4b_8	a_4b_9
a_5b_1	a_5b_2	a_5b_3	a_5b_4	a_5b_5	a_5b_6	a_5b_7	a_5b_8	a_5b_9
a_6b_1	a_6b_2	a_6b_3	a_6b_4	a_6b_5	a_6b_6	a_6b_7	a_6b_8	a_6b_9
a_7b_1	a_7b_2	a_7b_3	a_7b_4	a_7b_5	a_7b_6	a_7b_7	a_7b_8	a_7b_9

(1)

While one time series evolves along the x-axis and the other time series evolves along the y-axis, the overall time evolves along the diagonal. This is traditionally visualised with two vectors, e.g. $\vec{a} = a_1a_2 \dots a_7$ and $\vec{b} = b_1b_2 \dots b_9$ that combine

in a distance matrix as shown in (1) with time elapsing for $\vec{a}$ top-down, for $\vec{b}$ left-right and for overall time from top-left to bottom-right. From a conceptual point of view where time elapses from left to right, or from top to bottom, the traditional matrices, as they are produced by Matlab® or R, are not optimal in this sense. Both, the reduction of the search space and the orientation of the time axis can be realised by introducing a tunnel matrix that represents the Sakoe Chiba Band (shaded area in (1)) rotated by 45 degrees to the left for horizontal or to the right for vertical propagation of time. This paper proposes an implementation of such a distance matrix in tunnel form at the initialisation stage, i.e. already before the calculation of the alignment path.

3. IMPLEMENTATION OF A TUNNEL MATRIX

The aim is to rotate the Sakoe Chiba Band from a traditional distance matrix around the diagonal by -45 degrees in order to let time elapse from left to right. As can be seen in (1), the diagonal contains the isochronous pairs $a_1b_1, a_2b_2, \dots, a_7b_7$. This should be the mid row of the new matrix. It can also be seen that the values of $\vec{a}$ are constant for each column of and increase from row to row. For $\vec{b}$ it is the inverse: the values are constant for each row and increase from column to column. In order to illustrate the new tunnel arrangement, two intermediate matrices are prepared. For all values of $\vec{a}$ and $\vec{b}$ that fall into the bandwidth of the Sakoe Chiba Band, a ‘tunnel A’ (2) and a ‘tunnel B’ (3) is created.

NA	NA	a_1	a_2	a_3	a_4	a_5	a_6	a_7	
NA	a_1	a_2	a_3	a_4	a_5	a_6	a_7	NA	
a_1	a_2	a_3	a_4	a_5	a_6	a_7	NA	NA	
a_1	a_2	a_3	a_4	a_5	a_6	a_7	NA	NA	
a_1	a_2	a_3	a_4	a_5	a_6	a_7	NA	NA	

(2)

b_1	b_2	b_3	b_4	b_5	b_6	b_7	b_8	b_9	NA	NA
b_1	b_2	b_3	b_4	b_5	b_6	b_7	b_8	b_9	NA	NA
b_1	b_2	b_3	b_4	b_5	b_6	b_7	b_8	b_9	NA	NA
NA	b_1	b_2	b_3	b_4	b_5	b_6	b_7	b_8	b_9	NA
NA	NA	b_1	b_2	b_3	b_4	b_5	b_6	b_7	b_8	b_9

(3)

For illustration purposes, a width of three values in each dimension was chosen. Both intermediate matrices are then merged in order to create the final AB Tunnel Matrix by cross-multiplying ‘tunnel A’ and ‘tunnel B’, as shown in (4). Note that these values represent here similarity instead of distance.

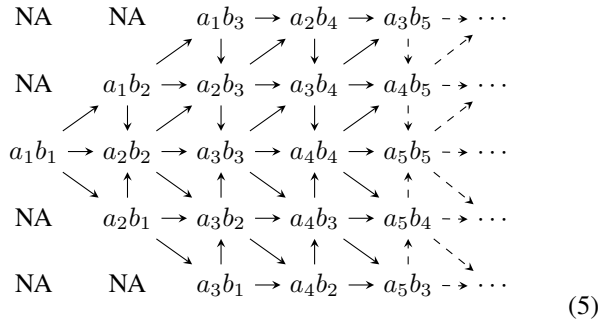
NA	NA	a_1b_3	a_2b_4	a_3b_5	a_4b_6	a_5b_7	a_6b_8	a_7b_9	
NA	a_1b_2	a_2b_3	a_3b_4	a_4b_5	a_5b_6	a_6b_7	a_7b_8	NA	
a_1b_1	a_2b_2	a_3b_3	a_4b_4	a_5b_5	a_6b_6	a_7b_7	NA	NA	
NA	a_2b_1	a_3b_2	a_4b_3	a_5b_4	a_6b_5	a_7b_6	NA	NA	
NA	NA	a_3b_1	a_4b_2	a_5b_3	a_6b_4	a_7b_5	NA	NA	

(4)

4. THE TUNNEL DP-ALGORITHM

For the newly created similarity matrix, the traditional DP algorithms are inappropriate. The start point (a_1b_1) and the end point (a_7b_9) are no more at the same location. And therefore, the possible directions that an alignment path might take are also different. For a traditional similarity matrix, the most common step-pattern is from a_1b_1 to a_2b_2 without insertion or deletion, to a_1b_2 by insertion in $\vec{a}$ or to a_2b_1 by insertion in $\vec{b}$ (or deletion from $\vec{a}$).

This translates into the new tunnel matrix, however it has to be adapted depending where in the matrix the algorithm is currently working. The principle is the same: it is merely possible to continue in both vectors or to stay at the same value in one vector while proceeding in the other. In the mid row of the tunnel matrix, imagine to start at a_1b_1 in (5), the steps indicated by arrows are possible. In the top half of the matrix (imagine the prior step went from a_1b_1 to a_1b_2), one arrow points into a different direction. That means the possible steps are different. In the bottom half of the matrix (imagine the prior step went from a_1b_1 to a_2b_1), again, arrows point in different directions.



In order to find the path through the whole Similarity Matrix with the highest quality (inverse of cost), DP accumulates for each point of the matrix the maximum quality path up to that point in a Quality Matrix. Therefore, at each point of that matrix, the previously accumulated quality is added to the current similarity. The direction from where the lowest cost path came from is memorised in order to find the way back, once the whole matrix has been passed. The algorithm for filling the Quality Matrix according to the similarity matrix is again different from the traditional procedure. It is not possible to proceed row wise and column wise from one corner of the matrix (formerly the start point) to the opposite corner (formerly the end point), even if the step-patterns are clear for each point. Because the DP algorithm always looks backwards from the current point in order to evaluate the highest quality path and accumulates in the current point the new quality-to-this-point, all possible points from where the path can come from need already be evaluated. If they are not, the new evaluation would be based on both, points that include the accumulated similarities (the quality) and points that have stored the mere individual similarity. This would bias the de-

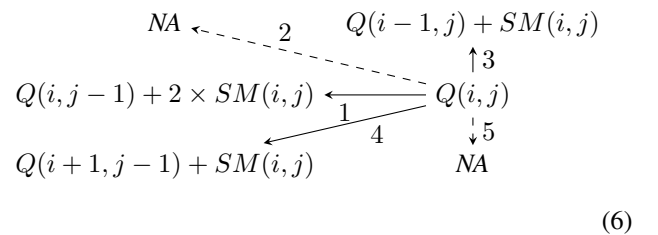
cision in favour of the latter. A solution is suggested in the following section.

4.1. Quality Tunnel Matrix and Backtrack Matrix

A Quality Matrix Q stores the accumulated similarities. A Backtrack Matrix ϕ stores at each point the direction from where the path with the highest quality came from. In a first step, Q is filled with the values of the similarity matrix SM and surrounded on top, bottom and left column, by a line of NAs (Not Applicable). This border is necessary in order to have initial values to be compared with. Once the DP algorithm has finished, the end point of that matrix contains the overall quality for the alignment of the two vectors $\vec{a}$ and $\vec{b}$. The algorithm iterates over the columns via j and over the rows via i . $Q(i, j)$ is the point in the Quality Matrix for which the accumulated quality is to be calculated, i.e. the maximum quality that accumulates until that point. As mentioned above, it is important that each time a potentially proceeding path is evaluated for the maximum quality, the values are either part of the border, or already filled. Therefore, a specific evaluation pattern is established.

An overall trend from left to right is used in order to reach from the start in the first column to the end in the last column. The rows are divided into three parts. The first part is the top half of the matrix, the second part is the bottom part of the matrix and the third part is the mid row. The rows of the top and bottom part are evaluated from the outside to the middle. For the top part (6), it means that i increases from 2 to the *middle* - 1 (row above the mid). For the bottom part (7), it means that i decreases from the *end* - 1 (previous last row) to the *middle* + 1 (row below the mid). At the last iteration per column, the value of the mid row is evaluated (8).

The following comparisons are performed for the three parts at each point $Q(i, j)$. NAs (Not Applicable) are inserted in the diagrams in order to keep the indexing function for the backtrack path constant: The horizontal step is always 1, the vertical steps are always 3 (top) and 5 (down), and the diagonal steps are always 2 and 4. The lowest value is added to the current point $Q(i, j)$. The direction from where this highest quality path came from is added to the Backtrack Matrix at point $\phi(i, j)$. Arrows with solid lines indicate valid steps; arrows with dashed lines indicate invalid steps. **The top half of the matrix** (row 2 to *mid* - 1) evaluates $Q(i, j) = \max(Q(i, j-1) + 2 \times SM(i, j), NA, Q(i-1, j) + SM(i, j), Q(i+1, j-1) + SM(i, j), NA)$.



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