

Nonlinear Analysis of VR Synchronization

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1 Abstract

There exists a literature dealing with time series analysis in the nonlinear sciences in the case of synchronization, or the adjustment of rhythms of oscillating objects to coincide with each other due to weak interaction. We attempt many of these methods in analyzing an instance of the phenomenon of interpersonal synchrony in a virtual environment. We also propose a mapping from time series to networks with a deterministic inverse, useful for exploratory analysis.

2 Introduction

Before defining what we mean by synchronization in this context, we note that there exists a long history of the study of synchronization outside of social science. Synchronization was first reported in physical systems by Huygens in 1665. He saw that two pendulum clocks in the boat that he was testing the clocks in were mysteriously ticking in unison, and found that the cause was a support beam which connected the clocks, which moved imperceptibly and therefore maintained the synchronization[1].

Centuries later, the physical and biological systems which were later found to synchronize include mechanical rotators, lasers, fireflies emitting synchronous patterns of light, syn-

chronous contraction of heart cells, synchronization of human circadian rhythm to a solar light cycle, and synchronous interaction of human beings[2], [3].

Now, we should define the terms. Interpersonal synchrony is defined in the social science literature as individuals' temporal coordination during social interaction[4]. In the physical sciences, synchronization is defined as an adjustment of rhythms of oscillating objects due to weak interaction, with generalizations possible for chaotic systems, which depend upon a phase of the chaotic system existing[3].

Although social synchrony is one of many other synchronies, many of which do not necessarily serve a function of any kind, it is observed that social synchrony serves a function in human social groups. There exists evidence that synchronization acts as a cooperation-inducing mechanism[5], that it acts to induce rapport[6], and that it in actuality, independent of its effect on rapport, enhances the ability to pursue joint goals in tandem for those who are synchronized[7]. This suggests immediately that it must be measured in a systematic way, to study phenomena of that kind.

In the beginning of the social synchronization literature, most psychologists did not use the then-developing automated signals processing techniques for the detection and measurement of synchronization, instead using manual methods to detect and rate the presence or absence of synchronization, with trained raters and validated measuring systems[11]. Although these measurements have been validated[12], they depend upon human raters and therefore are less replicable and less convenient than automated systems.

Given that the signals created by an individual during social interaction have a phase, and social interaction can be construed as a weak interaction between the two individual systems, it should be clear that the definition of interpersonal synchrony given in the social psychological literature is a subset of the definition given in the physics literature. This has often been noted[4], and has therefore spawned a cross-disciplinary field wherein signals processing techniques are used to measure interpersonal synchrony.

As a specific instance of a domain where signals processing techniques are used, a large

problem in synchronization is the definition of the signal itself and its extraction from observations of social interaction. To this end, many methods have been used, including extraction of coordination of movement features and speech features, movement of single and dyadic body parts, image processing techniques and video tracking techniques.

An important analogy exists between the collection of time series data about the physiology of the body in order to assess synchrony and the collection of time series data about the body in order to assess health. Indeed, there exists a literature on the synchronization properties and the time series analysis of heart rate variability which we have derived much inspiration from, not necessarily in the usage of correlational metrics, for which there is a literature dealing in social synchronization, but more sophisticated metrics for which there is not such a literature[16][17]. The essential premise is that we treat social synchronization as an instance of an abstract problem of physical synchronization, which justifies using measures used by other physiological measurements.

Measuring synchronization from virtual reality body tracking has many benefits in comparison to other measures, as argued in Blascovich et al. To wit, much of the typical tradeoff between experimental control and ecological validity is avoided because the ecological validity can be increased by improved graphics and worlds, but total experimental control is always held. Replicability is much better because the hardware and software can be of standard make. And data can be collected without video markers because the tracking apparatus can collect data by itself. [13]

2.1 Justification for Nonlinear Measures

This project will attempt to use some of the already existing tools for the analysis of time series data on VR time series data, as well as apply some tools which have not previously been used to analyze synchronization in VR time series data of social synchronization in a virtual world. — summary of what exists, talk only about the fact that they exist, as a preliminary summary of the essay. we will go deeper and actually formal with this one.

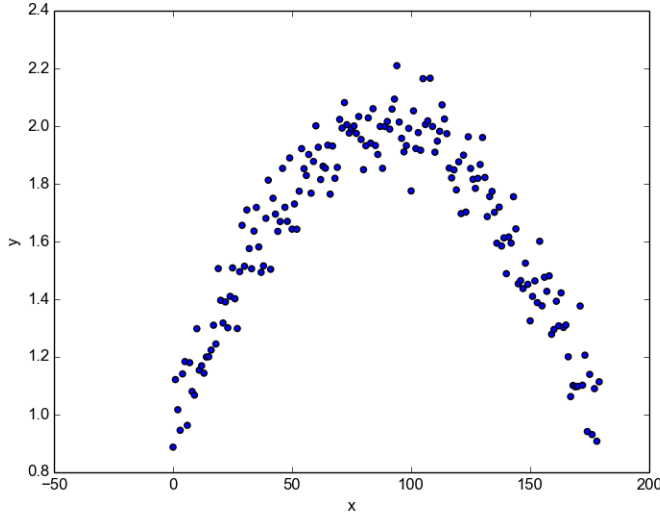


Figure 1: Example of simple nonlinear relation between two variables

correlation, MI in physics, spectral methods, analytic signal, graph analysis

One might wonder why the existing methods of measurement of synchronization do not already serve for the purpose of measuring synchronization of body movements in VR. They do not serve for the same reasons that they do not serve in physical time series. That is, they are fundamentally *linear* measures of the relation that they have to each other, meaning that measuring only correlation implicitly puts in the assumption that an affine transform relates the two signals to each other. It is possible to learn about higher order dependences if correlations of higher moments are used, but this becomes rapidly time-consuming very quickly[14].

One example which should help develop a theoretical intuition about nonlinearity and synchronization is the sine wave relation example. Consider the figure 2.1. According to a linear correlation measure, taking the whole domain of x into account at once, the two variables x and y cannot disprove the null hypothesis of independence ($p = 0.45$), but by inspection, one can say that actually $y = \sin(x) + \epsilon$, where ϵ is a small amount of Gaussian noise. But nonlinear measures *will* take this nonlinear relation into account. For example, mutual information will say that the mutual information between the two variables x and y

is 7.5 bits, which indicates that the variables determine each other to a great degree.

This aspect of the analysis, the question as to the linearity of the measures, is mostly at a lower level than the vast literature on the role of synchronization in psychological processes. However, there is a distinction which might be of interest, which is the distinction between phase and amplitude synchronizations.

Nonlinear analyses have been considered for modelling in the sensorimotor literature, and often rejected because linear models often are a good approximant. However, in the *measurement* of degrees of synchronization, or of the real experimental distinguishing of synchronization state, it does not necessarily follow that avoiding the assumption of linearity is as intractible as fitting a nonlinear *model* would be. Importantly, there do exist nonlinear models for phase-entrainment dynamics of physiological movement.[19]

In other fields of physiological measurement, the use of nonlinear methods is commonplace.

2.2 Data Description

— This is the place where we will describe the data we got, the ad hoc analyses made previously to fold in the data into small number of dimensions. Discuss the data incompleteness problem.

— Why the dimensionality reduction? we suffer from curse of dimensionality, might as well do it ad hoc. Note that it's a terribly naive method of dimensionality reduction, but it shouldn't matter because it's consistent between data

3 Time Domain Analysis

3.1 Correlation Analysis

One analysis commonly used with relaxation oscillators like ECG data is the Poincare map, not to be confused with the Poincare plot. However, because there are not clear demarcations

as to where each oscillation is, the Poincare map is less suited to this dataset and will not be used[15]

3.2 Mutual Information Analysis

In order to introduce mutual information, we must introduce Shannon information and Shannon entropy.

Shannon information is defined on a measurement drawn from some set (let the set be discrete, $X \in 1...n$, for ease of notation: there exist continuous analogues). Given that measurement from some set, the Shannon information in bits is defined as:

$$-\log_2 P_X(i)$$

where $P_X(i)$ is the probability that a measurement will find the system in state i . That is, a lower probability answer is more surprising, and therefore more information is found in it.

Shannon entropy, denoted $H(X)$ is defined on a data stream (a measurement stream) X . It is the average amount of information contained in every message received, and so is also measured in bits. An easy intuition about entropy can be gained by noting that a *fair* coin flipping process is the most entropic possible coin flip, because we are most surprised by it. Formally, entropy is:

$$H(X) = - \sum_i P_X(i) \log_2 P_X(i)$$

Information and entropy are also defined on joint probability distributions in an analogous way. The information of a joint probability distribution on variables X and Y is:

$$-\log_2 P_{XY}(i, j)$$

And the entropy, therefore, is:

$$H(X, Y) = - \sum_i \sum_j P_{XY}(i, j) \log_2 P_{XY}(i, j)$$

Now, if we sample two random variables simultaneously, the relationship has a quantity defined upon it which is called the *mutual information*. Because it is defined on the relation, it has an auto-mutual information variant and a cross-mutual information variant, like correlation does. However, mutual information measures something different, which is the *information which one of the random variables gives me about the other random variable*. It is defined as:

$$I(X, Y) = H(X) + H(Y) - H(X, Y)$$

And one sees that this quantity is symmetric with respect to X and Y .

3.2.1 AMI and CMI

Auto-mutual information (AMI) and cross mutual information (CMI) are the mutual information analogues to autocorrelation and cross-correlation. They have the same advantages over the correlation variants as mutual information itself has to correlation. Therefore, they are better measurements for synchronization.

—— actual content of Mutual information analysis goes here.

4 Frequency Domain Analysis

There is more than one way to represent time series data, even given only the data itself and not any underlying function that generates it. The most important alternate representation is the frequency domain representation, which is taken in finite data with the discrete Fourier transform. Later, we will show the use of the frequency representation in re-representing our data so that it shows a numerical current state of the phase, using the related Hilbert

transform.

The frequency domain represents data in terms of the linear combination of simple sine and cosine waves: the analogy often used is that of a chord being hit on a piano, which can be represented with its time domain signal but can also be represented as the amplitude of the individual notes within the chord, which presents a better decomposition of the sound in and of itself. Likewise, we can attempt to decompose our time domain signals into the frequencies of the sinusoids which make it up.

The discrete fourier transform is an essential tool for the further analyses below. It converts a finite discrete sample from the time to the frequency domain in a computable and fast way.

— add back in the math of DFT — add back in the coherence measures

4.1 Hilbert Transform and γ

A signal is called an *analytic signal* if it does not have any negative-frequency components. The Hilbert transform is a way to get the analytic *representation* for a signal in the time domain, which throws away the superfluous information which would be in a spectrum of that signal. This allows the expression of the analytic signal in terms of its time-variant magnitude and phase, allowing a study of the phase in an *unwrapped form*[18].

Formally speaking, given that the original signal is $s(t)$, the analytic signal ζ is:

$$\zeta(t) = s(t) + js_H(t) = A(t)e^{j\phi(t)}$$

$$s_H(t) = \frac{1}{\pi}PV \int_{-\infty}^{\infty} \frac{s(\tau)}{t - \tau} d\tau$$

s_H is called the Hilbert transform; *PV* indicates that the integral is a Cauchy principal value integral. The Hilbert transform is thus formally equivalent to convolution of $s(t)$ with $\frac{1}{\pi t}$, so it can easily be calculated from the fast Fourier transform.

— finish the Hilbert transform, add the justification (you can take out the amplitude entirely as a factor, whereas coherence and that sort of thing ends up not doing this). — add pretty pictures of unwrapped hilbert transform, and the description of the gammadifference measures — -then- add the gamma results and the hilbert transform results themselves

References

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[19] Schmidt, R. C., & Turvey, M. T. (1994). Phase-entrainment dynamics of visually coupled rhythmic movements. *Biological cybernetics*, 70(4), 369-376.