PyIF: A Fast and Light Weight Implementation to Estimate Bivariate Transfer Entropy for Big Data

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Abstract—Transfer entropy is an information measure that quantifies information flow between processes evolving in time. Transfer entropy has a plethora of potential applications in financial markets, canonical systems, neuroscience, and social media. We offer a fast open source Python implementation called PyIF that estimates Transfer Entropy with Kraskov's method. PyIF utilizes KD-Trees, multiple processes by parallelizing queries on said KD-Trees, and can be used with CUDA compatible GPUs to significantly reduce the wall time for estimating transfer entropy. We find from our analyses that PyIF's GPU implementation is up to 1072 times faster (and it's CPU implementation is up 181 times faster) than existing implementations to estimate transfer entropy on large data and scales better than existing implementations.

Index Terms—Transfer Entropy, Parallel Processing

I. INTRODUCTION

Information theory provides a framework for studying the quantification, storage, and communication of information [1]. This theory defines entropy as the amount of uncertainty or disorder in a random process. Mutual Information is another measure in this theory which quantifies the amount of information shared across random variables. While similar to mutual information, transfer entropy (TE) also considers the dynamics of information and how these dynamics evolves in time. [2]. Put simply, TE quantifies the reduction in uncertainty in one random process from knowing past realizations of another random process. This is a particularly useful property of TE as many real-world phenomena, from stock market prices to neural signals, are dynamic processes evolving in time. TE is also an asymmetric measure of information transfer. Ergo, TE computed from process A to process B may yield a different result than TE computed from B to A. The information theoretic framework and these measures have led to a variety of applications in different research areas [1], [5].

A. Applications of Transfer Entropy

TE is particularly useful for detecting information transfer in financial markets [5]. Marschinski and Kantz used TE to perform index-to-index analysis with the Dow Jones Share Market (DJIA) and the Frankfurt Stock index (DAX) [4] and document the extent to which one index drives the behavior of the other. Following Marschinski and Kantz, other researchers apply TE to examine related research questions about financial markets. These include measuring TE from market indexes, such as S&P 500 or the DJIA, to individual equities as well as between individual equities [5].

Network Inference is another application area of TE. An objective of network inference is to infer a network by indentifying relationships between individual processes in the data. Computational neuroscience, financial market analysis, gene regulatory networks, social media, and multi-agent systems are areas where TE has been used to model networks. Early approaches that used TE for network inference either measure pairwise TE between all pairs of variables in a network or threshold the TE values to select connections between nodes in a network [12], [13], and [14]. Recent approaches have used statistical significance tests of pairwise TE to determine whether links exist [15] and [16]. [5] offers more examples of TE applications.

B. Outline

In the next section we formally define TE. The following section discusses TE estimation methods. We then discuss our our proposed implementation called PyIF to estimate bivariate TE. Next, we describe a comparative analysis between PyIF and existing implementations that estimate bivariate TE. Lastly, we conclude the paper with a discussion and future work.

II. DEFINITION OF TRANSFER ENTROPY

In 2000, Schreiber [2] discovered TE and coined the name "transfer entropy," although Milian Palus [3] also independently discovered the concept as well. Let the function I represent mutual information between two probability distributions. Lagged mutual information $I(X_t:Y_{t-k})$ can be used as a time-asymmetric measure of information transfer from

Y to X where X and Y are both random processes, k is a lag period, and t is the current time period. However, lagged mutual information is unsatisfactory as it does not account for a shared history between the processes X and Y [6].

TE considers the shared history between two processes via conditional mutual information. Specifically, TE conditions on the past of X_t to remove any redundant or shared information between X_t and its past. This also removes any information in the process Y about X at time t that is in the past of X [7]. Transfer entropy T (where the transfer of information occurs from Y to X) can be defined as:

$$T_{Y \to X}(t) \equiv I(X_t : Y_{t-k} | X_{t-k}).$$
 (1)

Kraskov [8] shows that transfer entropy can be expressed as the difference between two conditional mutual information computations:

$$T_{Y \to X}(t) = I(X_t | X_{t-k}, Y_{t-k}) - I(X_t | X_{t-k})$$
 (2)

The intuition of this definition is that TE measures the amount of information in Y_{t-k} about X_t after considering the information in X_{t-k} about X_t . Put differently, TE quantifies the reduction in uncertainty about X_t from knowing Y_{t-k} after considering the reduction in uncertainty about X_t from knowing X_{t-k} .

III. ESTIMATING TRANSFER ENTROPY

There are many techniques for estimating mutual information. Khan et al. explored the utility of different methods for mutual information estimation [10] and many of the methods they considered are applicable to estimate TE.

A. Kernel Density Estimator

Kernel Density Estimators can be used to estimate TE [11]. For a bivariate dataset of size n with variables X and Y, Mutual Information can be estimated as:

$$\hat{I}(X,Y) = \frac{1}{n} \sum_{i=1}^{n} \ln \frac{\hat{p}_{XY}(x_i, y_j)}{\hat{p}_{X}(x_i)\hat{p}_{Y}(y_i)}$$
(3)

where $\hat{p}_X(x_i)$ and $\hat{p}_Y(y_i)$ are the estimated marginal probability density functions and $\hat{p}_{XY}(x_i,y_j)$ is the joint estimated probability density function. For a multivariate dataset containing: $x_1, x_2, ..., x_n$ where each x is in a d-dimensional space, the multivariate kernel density estimator with kernel K is defined by:

$$\hat{p}(x) = \frac{1}{nh^d} \sum_{i} = 1^n K(\frac{x - x_i}{h})$$
 (4)

where h is the smoothing parameter, and in this case, K is a standard multivariate normal kernel defined by $K(x) = (2\pi)^{-d/2}e^{\frac{x^Tx}{2}}$. Moon et al. outlined a procedure to estimate Mutual Information using marginal and joint probabilities with Kernel Density Estimators [11].

B. Kraskov Estimator

Transfer Entropy can be estimated using k-nearest neighbors [8]. Note that entropy can be estimated with:

$$\hat{H}(X) = -\frac{1}{n} \sum_{i=1}^{n} ln\hat{p}(x_i)$$
 (5)

Kraskov et al. expanded this definition to estimate entropy to:

$$\hat{H}(X) = -\frac{1}{n} \sum_{i=1}^{n} \psi(n_x(i)) - \frac{1}{k} + \psi(n) + \ln(c_{d_x}) + \frac{d_x}{n} \sum_{i=1}^{n} \ln(\epsilon(i))$$
 (6)

where n are the number of data points, k are the nearest neighbors, d_x is the dimension of x, and c_{d_x} is the volume of the d_x -dimensional unit ball. For two random variables X and Y, let $\frac{\epsilon(i)}{2}$ be the distance between (x_i,y_i) and it's kth neighbor be denoted by (kx_i,ky_i) . Let $\frac{\epsilon_x(i)}{2}$ and $\frac{\epsilon_y(i)}{2}$ be defined as $||x_i-kx_i||$ and $||y_i-y_i||$ respectively. $n_x(i)$ is the number of points x_j such that $||x_i-x_j|| \leq \epsilon_x(i)/2$, $\psi(x)$ is the digamma function where

$$\psi(x) = \Gamma(x)^{-1} d\Gamma(x) / dx \tag{7}$$

and $\Gamma(x)$ is the ordinary gamma function. Lastly $\psi(1) = -C$ where C = 0.5772156649 and is the Euler-Mascheroni constant. To estimate the entropy for the random variable Y, Y can be substituted into $\hat{H}(X)$.

Joint entropy between X and Y can then be estimated as:

$$\hat{H}(X,Y) = -\psi(k) - \frac{1}{k} + \psi(n) + \ln(c_{d_x}c_{d_y}) + \frac{d_x + d_y}{n} \sum_{i}^{n} \ln(\epsilon(i))$$
(8)

where d_y is the dimension of y, and c_{d_y} is the column of the d_y -dimensional unit ball. Using $\hat{H}(X), \hat{H}(Y)$, and $H(\hat{X}, Y)$ mutual information can be estimated as:

$$\hat{I}(X,Y) = \psi(k) - \frac{1}{k} - \frac{1}{n} \sum_{i=1}^{n} [\psi(n_x(i)) + \psi(n_y(i))] + \psi(n)$$
(9)

where $n_y(i)$ is the number of points y_j such that $||y_i - y_j|| \le \frac{\epsilon_y(i)}{2}$. This method has been referred to as the Kraskov estimator in literature.

C. Additional Estimators

Khan et al. also explored the utility of Edgeworth approximation of differential entropy to calculate Mutual Information and adaptive partitioning of the XY plane to estimate the joint probability density, which can be used to estimate mutual information. Ultimately Khan et al. found that a KDE estimator and Kraskov estimator outperform other methods with

respect to their ability to capture the dependence structure of random processes. Currently our software supports estimating bivariate TE using the Kraskov estimator with plans to add other estimators in the future.

IV. PYIF

Our proposed software implementation PyIF 1 is an open source implementation. PyIF currently only supports using the Kraskov estimator to estimate TE. PyIF utilizes recent advancements in hardware to parallelize & optimize operations across CPUs and Cuda compatible GPUs [9] and CPUs. In particular we focus our efforts on the parallelization & optimization across operations to obtain n_x & n_y in Eq. 9 faster.

PyIF is a python only implementation which utilizes 5 well-known and actively supported python libraries: SciPy [17], NumPy [18], scikit-learn [19], nose [20], and numba [21]. SciPy is an open source Python library used for a variety of STEM applications. NumPy is a part of SciPy's ecosystem and is an open source package that provides convenient ways to perform matrix manipulations and useful linear algebra capabilities. The library scikit-learn is a popular open source library for machine learning and nose is another open source library that is useful for testing code to ensure that it will produce the correct outcome. Lastly, numba is a python compiler that can compile Python code for execution on multicore CPUs and CUDA-capable GPUs.

PyIF's interface only requires you to supply X and Y, two numpy arrays with Nx1 dimensions. Optional arguments can be passed in such as k which controls the number of neighbors used in KD-tree queries, embedding which controls how many lagged periods are used to estimate transfer entropy and a boolean argument GPU can be used to specify if you want to use a CUDA compatible GPU. Lastly another boolean argument safetyCheck can be used to check for duplicates rows in your dataset. This boolean argument is there to help prevent a more subtle error that can occur when multiple data points in a bivariate dataset have identical coordinates. This essentially can lead to several points that have an identical distance to a query point which violates assumptions of the Kraskov estimator. A solution that is used in practice and that we recommend is to add a small amount of noise to your dataset to avoid this error.

V. COMPARATIVE ANALYSIS

We compare PyIF's ability to estimate Transfer Entropy against existing implementations with respect to computational performance. We present all of the data and code used to estimate TE for all implementations ². Each implementation in this comparative analysis estimates TE on four simulated bivariate datasets of different sizes. The estimated TE values are roughly the same for each implementation and we forgo

¹PyIF can freely be downloaded from: https://github.com/lcdm-uiuc/PyIF
²The data and code can freely be downloaded from: https://github.com/lcdm-uiuc/Publications/tree/master/2020_Ikegwu_Traguer_McMullin_Brunner

comparing the actual values since this is random simulated data. We make the assumption that there is relatively little to no information transfer between the random processes. We run each of the implementations (excluding Transfer Entropy Toolbox) on nano, a cluster of eight SuperMicro servers with Intel Haswell/Broadwell CPUs and NVIDIA Tesla P100/V100 GPUs hosted by the National Center of Super Computing Applications at the University of Illinois at Urbana-Champaign. We used one node which contains two E5-2620 v3 Intel Xeon CPU's and 2 NVIDIA P100 GPUs with 3584 cores. We refer to this analysis as Analysis 1.

We conduct the same analysis on different hardware to compare PyIF to Transfer Entropy toolbox because of MATLAB licensing issues with the National Center of Super Computing Applications. We use an Engineering Workstation with an Intel Xeon Processor E5-2680 v4 hosted by Engineering IT shared services at the University of Illinois at Urbana-Champaign. We use a single CPU core and up to 16GB of RAM to estimate TE with Transfer Entropy toolbox and PyIF. This workstation does not offer CUDA compatiable GPUs to use for either PyIF or Transfer Entropy Toolbox so we forgo comparing the GPU implementations. This workstation has a CPU time limit of 60 minutes meaning that if any process uses 100% of a CPU core for more than 60 minutes the process is terminated. We refer to this analysis as Analysis 2.

A. IDTxl

The first implementation is the Information Dynamics Toolkit xl (IDTxl). IDTxl is an open source Python toolbox for network inference [22]. Currently IDTxl relies on NumPy, SciPy, CFFI (which is another open source library that provides a C interface for Python code), H5py which is a Python package that is used to interface with HDF5 binary data format, JPype which is a Python module that provides a Java interface for Python code, and Java jdk which is a developer kit to develop Java applications and applets. IDTxl has additional functionaility besides estimating TE however we only use IDTxl's capability to estimate TE on a bivariate dataset.

B. TransEnt

TransEnt is a R package that estimates transfer entropy [23]. Currently TransEnt relies on Rcpp which acts as a interface to C++ from R. TransEnt also relies on a C++ library called Appromixate Nearest Neigbors (ANN) [24] which performs exact and approximate nearest neighbor searches. Currently the package has been removed from CRAN, however this software can be used and installed from [23]'s github repo

C. RTransferEntropy

RTransferEntropy is a R package that estimates transfer entropy between two time series [25]. Currently the RTransfer-Entropy package relies on Rcpp, and the future package which supports performing computations in parallel to decrease the

³ [23] Github Repo: https://github.com/Healthcast/TransEnt

wall time. We include both the parallel implementation of RTransferEntropy and the default implementation for completeness in the results.

D. Transfer Entropy Toolbox

Transfer Entropy Toolbox is an open source MATLAB toolbox for transfer entropy estimation [26]. This code's dependencies include: the Statistics & Machine Learning toolbox which provides functions to analyze and model data; the FieldTrip toolbox which is used for EEG, iEEG, MEG, and NIRS analysis; the parallel computing toolbox that performs parallel computations of multicore CPUs and GPUs; the signal processing toolbox that provides functions to analyze, preprocess, and extract features from sampled signals; the TSTOOL toolbox which is a toolbox for nonlinear time series analysis. TSTOOL no longer exists and cannot be download from it's official homepage 4. Nevertheless, the developers of Transfer Entropy toolbox include pre-compiled mex files of TSTOOL that will work with this implementation. At the time of writing this paper Transfer Entropy toolbox has not been updated since 2017.

E. Data

We create four bivariate datasets for this comparative analysis. Each dataset contains two time series with randomly generated values between 0 and 1. The first dataset contains 1000 observations, the second dataset contains 10,000 observations, the third dataset contains 100,000 observations, and the fourth dataset contains 1,000,000 observations. We used the seed 23 for the pseudo-random number generator for reproducibility. We will refer to the first dataset, second dataset, third dataset, and fourth dataset as the micro dataset, small dataset, medium dataset, and the large dataset respectively.

VI. RESULTS

We report the results for Analysis 1 in Table I. After estimating TE using all of the implementations outlined in the comparative analysis section we found that PyIF scales better on larger data. Excluding the TransEnt implementation, the CPU implementation of PyIF (or PyIF (CPU)) takes less time to estimate TransferEntropy than all other implementations. The R package TransEnt has a better performance in terms of speed than PyIF (CPU) for the micro dataset and the small dataset. However PyIF (CPU) is able to estimate transfer entropy in less time than all other implementations for the medium dataset and large dataset. PyIF (GPU) outperforms PyIF (CPU) for the small, medium and large datasets. Figure 1 visualizes this explanation. We suspect that the optimizations performed by Numba contribute to PyIF having a larger wall time than TransEnt on the micro and small datasets.

The results for Analysis 2 are in Table II. Although the Transfer Entropy Toolbox exceeds the CPU time limit for the large dataset, the results show that PyIF is able to scale better than Transfer Entropy Toolbox for the other three datasets. PyIF's wall times are less than Transfer Entropy toolbox's

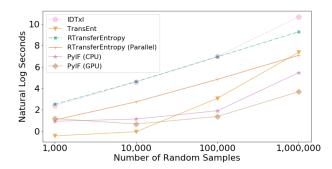


Fig. 1. This figure shows the natural log time (in seconds) to estimate Transfer Entropy for each implementation (excluding Transfer Entropy Toolbox) for each dataset used in this study.

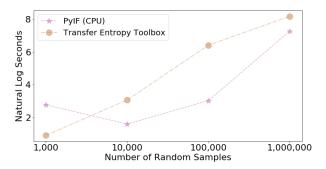


Fig. 2. This figure shows the natural log time(in seconds) to estimate Transfer Entropy between PyIF and Transfer Entropy Toolbox on an Engineering Workstation as described in the section Comparative Analysis. Transfer Entropy Toolbox exceeded the maximum allowable CPU runtime for the Large Dataset.

wall times excluding the Micro Dataset. Figure 2 visualizes this explanation.

VII. CONCLUSION

An important issue is addressed regarding Big Data with respect to estimating bi-variate Transfer Entropy. We introduce a fast solution to estimate Transfer Entropy with a small amount of dependencies. On large data our implementation PyIF is up to 1072 times faster utilizing GPUs and up to 181 times faster utilizing CPUs than existing implementations that estimate bi-variate TE. PyIF is also open sourced and publicly available on github for anyone to use. For future work we plan to improve the existing code base to increase the computation performance of PyIF even further. In addition to this we plan to implement additional estimators outlined in the section entitled "Estimating Transfer Entropy" to estimate bi-variate TE. This boost in computational performance will enable researchers to estimate bi-variate TE much faster for a variety of research applications.

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⁴http://www.dpi.physik.uni-goettingen.de/tstool/

TABLE I

THE WALL TIME TO ESTIMATE TRANSFER ENTROPY FOR A VARIETY OF IMPLEMENTATIONS ON THE DIFFERENT DATA SETS DESCRIBED IN THE SECTION COMPARATIVE ANALYSIS. THE HIGHER THE WALL TIME THE LONGER IT TOOK FOR THE IMPLEMENTATION TO ESTIMATE TRANSFER ENTROPY. THE NUMBER IN THE RELATIVE PERFORMANCE INDICATES HOW MANY TIMES FASTER (OR SLOWER) PYIF (CPU) IS TO A PARTICULAR IMPLEMENTATION.

Implementation	Wall	Relative	
	Time (in	Performance	
	seconds)	to PyIF (CPU)	
Micro Dataset Results (1000 Obs.)			
IDTxl	10.98	4.28	
TransEnt	0.656	0.25	
RTransferEntropy	12.492	4.87	
RTransferEntropy (Parallel)	2.876	1.12	
PyIF (CPU)	2.564	1.00	
PyIF (GPU)	3.282	1.28	
Small Dataset Results (10,000 Obs.)			
IDTxl	100.23	31.94	
TransEnt	0.968	.308	
RTransferEntropy	102.228	32.57	
RTransferEntropy (Parallel)	15.703	5.00	
PyIF (CPU)	3.138	1.00	
PyIF (GPU)	1.98	0.63	
Medium Dataset Results (100,000 Obs.)			
IDTxl	1070.749	152.89	
TransEnt	21.708	3.03	
RTransferEntropy	1036.661	152.00	
RTransferEntropy (Parallel)	127.281	18.66	
PyIF (CPU)	6.82	1.00	
PyIF (GPU)	3.996	0.58	
Large Dataset Results (1,000,000 Obs.)			
IDTxl	43150.129	181.97	
TransEnt	1585.942	6.68	
RTransferEntropy	10592.77	44.67	
RTransferEntropy (Parallel)	1188.636	5.01	
PyIF (CPU)	237.122	1.00	
PyIF (GPU)	40.231	0.16	

TABLE II

THE WALL TIME AND RELATIVE PERFORMANCE TO PYIF (CPU) TO ESTIMATE TRANSFER ENTROPY BETWEEN PYIF AND TRANSFER ENTROPY TOOLBOX ON AN ENGINEERING WORKSTATION MACHINE AS DESCRIBED IN THE SECTION COMPARATIVE ANALYSIS.

Implementation	Wall Time (in	Relative
_	seconds)	Performance
		to PyIF (CPU)
Micro Dataset Results (1000 Obs.)		
PyIF (CPU)	16.049	1.00
Transfer Entropy Toolbox	2.5012	0.15
Small Dataset Results (10,000 Obs.)		
PyIF (CPU)	4.989	1.00
Transfer Entropy Toolbox	21.6880	4.347
Medium Dataset Results (100,000 Obs.)		
PyIF (CPU)	20.915	1.00
Transfer Entropy Toolbox	616.8712	29.49
Large Dataset Results (1,00,000 Obs.)		
PyIF (CPU)	1455.725	1.00
Transfer Entropy Toolbox	> 3600	> 2.47

utilizes resources provided by the Innovative Systems Laboratory at the National Center for Supercomputing Applications at the University of Illinois at Urbana-Champaign. Lastly, we would like to thank Alice Perng for helpful work in Analysis 2.

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