

# On the Need for Time Series Data Mining Benchmarks: A Survey and Empirical Demonstration

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**Abstract.** In the last decade there has been an explosion of interest in mining time series data. Literally hundreds of papers have introduced new algorithms to index, classify, cluster and segment time series. In this work we make the following claim. Much of this work has very little utility because the contribution made (speed in the case of indexing, accuracy in the case of classification and clustering, model accuracy in the case of segmentation) offer an amount of “improvement” that would have been completely dwarfed by the variance that would have been observed by testing on many real world datasets, or the variance that would have been observed by changing minor (unstated) implementation details.

To illustrate our point, we have undertaken the most exhaustive set of time series experiments ever attempted, re-implementing the contribution of more than two dozen papers, and testing them on 50 real world, highly diverse datasets. Our empirical results strongly support our assertion, and suggest the need for a set of time series benchmarks and more careful empirical evaluation in the data mining community.

**Keywords:** time series, data mining, experimental evaluation

## 1. Introduction

In the last decade there has been an explosion of interest in mining time series data. Literally hundreds of papers have introduced new algorithms to index, classify, cluster and segment time series. In this work we make the following claim. Much of the work in the literature suffers from two types of experimental flaws, implementation bias and data bias (defined in detail below). Because of these flaws, much of the work has very little generalizability to real world problems.

In particular, we claim that many of the contributions made (speed in the case of indexing, accuracy in the case of classification and clustering, model accuracy in the case of segmentation) offer an amount of “improvement” that would have been completely dwarfed by the variance that would have been observed by testing on many real world datasets, or the variance that would have been observed by changing minor (unstated) implementation details.

In order to support our claim we have conducted the most exhaustive set of time series experiments ever attempted, re-implementing the contribution of more than 25 papers and testing them on 50 real word datasets. Our results strongly support our contention.

We are anxious that this work should not be taken as being critical of the data mining community. We note that several papers by the current first author are among the worst offenders in terms of weak experimental evaluation. While preparing the survey we read more than 360 data mining papers and we were struck by the originality and diversity of approaches that researchers have used to attack very difficult problems. Our goal is simply to demonstrate that empirical evaluations in the past have often been inadequate, and we hope this work will encourage more extensive experimental evaluations in the future.

For concreteness we begin by defining the various tasks that occupy the attention of most time series data mining research.

- **Indexing (Query by Content):** Given a query time series  $Q$ , and some similarity/dissimilarity measure  $D(Q,C)$ , find the nearest matching time series in database DB.
- **Clustering:** Find natural groupings of the time series in database DB under some similarity/dissimilarity measure  $D(Q,C)$ .
- **Classification:** Given an unlabeled time series  $Q$ , assign it to one of two or more predefined classes.
- **Segmentation:** Given a time series  $Q$  containing  $n$  datapoints, construct a model  $\bar{Q}$ , from  $K$  piecewise segments ( $K \ll n$ ) such that  $\bar{Q}$  closely approximates  $Q$ .

Note that segmentation has two major uses. It may be performed in order to determine when the underlying model that created the time series has changed (Gavrilov et al., 2000, Ge and Smyth, 2000), or segmentation may simply be performed to create a high level representation of the time series that supports indexing, clustering and classification (Ge and Smyth, 2000, Keogh and Pazzani, 1998, Keogh and Smyth, 1997, Lavrenko et al., 2000, Li et al., 1998, Park et al., 2001, Polly and Wong, 2001, Pratt and Fink, 2002, Qu et al., 1998, Shatkay and Zdonik, 1996, Wang and Wang, 2000b).

As mentioned above, our experiments were conducted on 50 real world, highly diverse datasets. Space limitations prevent us from describing all 50 datasets in detail, so we simply note the following. The data represents the many areas in which time series data miners have investigated, including finance, medicine, biometrics, chemistry, astronomy, robotics, networking and industry. We also note that all data and code used in this paper is available for free by emailing the first author.

The rest of this paper is organized as follows. In Section 2 we survey the literature on time series data mining, and summarize some statistics about the empirical evaluations. In Section 3, we consider the indexing problem, and demonstrate with extensive experiments that many of the published results do not generalize to real world problems. Section 4 considers the problem of evaluating time series classification and clustering algorithms. In Section 5 we show that similar problems occur for evaluation of segmentation algorithms. Finally in Section 6 we summarize our findings and offer concrete suggestions to improve the quality of evaluation of time series data mining algorithms.

## 2. Survey

In order to assess the quality of empirical evaluation in the time series data mining community we begin by surveying the literature. Although we reviewed more than 360 papers, we only included the subset of 57 papers actually referenced in this work when assessing statistics about the number of datasets etc. The subset was chosen based on the following (somewhat subjective) criteria.

- Was the paper ever referenced? Self-citations were not counted. The rule was relaxed for paper published in the last year because of publishing delays. We used ResearchIndex (<http://citeseer.nj.nec.com/cs>) to make this determination.
- Was the paper published in a conference or journal likely to be read by a data miner? For example, several interesting time series data mining papers have appeared in medical and signal processing conferences, but are unlikely to come to the attention of the data mining community.

The survey is very comprehensive, but was not intended to be exhaustive. Such a goal would in any case be subjective (should a paper which introduces a new clustering algorithm, and mentions that it could be used for time series be included?). In general the papers come from high quality conferences and journals, including (SIG)KDD (11), ICDE (11), VLDB (5), SIGMOD/PODS (5), and CIKM (6).

Having obtained the 57 papers, we extracted various statistics (discussed below) from them about their empirical evaluation. In most cases this was easy, but occasionally a paper was a little ambiguous in explaining some feature of its empirical evaluation. In such cases we made an attempt to contact the author for clarification, and failing that, used our best judgment.

In presenting the results of the survey, we echo the caution of Prechelt, that “*while high numbers resulting from such counting cannot prove that the evaluation has high quality, low numbers (suggest) that the quality is low*” (1995).

### 2.1. Size of test datasets

We recorded the size of the test dataset for each paper. Where two or more datasets are used, we considered only the size of the largest.

The results are quite surprising; the median size of the test database was only 10,000 objects. Approximately 89% of the test databases could comfortably fit on a 1.44 Mb floppy disk.

### 2.2. Number of rival methods

Another surprising finding of the survey is the relative paucity of rival methods to which the contribution of the paper is compared. The median number is 1 (The average is 0.91), but this number includes very unrealistic strawman. For example many papers (including one by the current first author (Keogh and Smyth, 1997)) compare times for an indexing method to sequential scan where both are preformed *in main memory*. However, it is well understood sequential scan enjoys a tenfold speed up when performed on disk because

any indexing technique must perform costly random access, whereas sequential scan can take advantage of an optimized linear traverse of the disk (Keogh et al., 2001). The limited number of rival methods is particularly troubling for papers that introduce a novel similarity measure. Although 29 of the papers surveyed introduce a novel similarity measure, only 12 of them compare the new measure to any strawman. The average number of rival similarity measures considered is only 0.97.

### 2.3. Number of different test datasets

Although the small sizes of the test databases and the relatively scarcity of comparisons with rival methods is by itself troublesome, the most interesting finding concerns the number of datasets used in the experimental evaluation. On average, each contribution is tested on 1.85 datasets (1.26 real and 0.59 synthetic). This numbers are astonishingly low when you consider that new machine learning algorithms are typically evaluated on at least a dozen datasets (Cohen, 1993, Kibler and Langley, 1988).

In fact, we feel that the numbers above are optimistic. Of the 30 papers that use two or more datasets, a very significant fraction (64%), use both stock market data and random walk data. However, we strongly believe these really should be counted as the same dataset. It is well known that random walk data can perfectly model stock market data in terms of all statistical properties, including variance, autocorrelation, stationarity etc (Faloutsos et al., 1994, Simon, 1994).

Work by the late Julian L. Simon suggested that humans find it impossible to differentiate between the two (1994). To confirm this finding we asked 12 professors at UCRs Anderson Graduate School of Management to look at Figure 1 and determine which three sequences are random walk, and which three are real S&P500 stocks. The confusion matrix is show in Table 1.

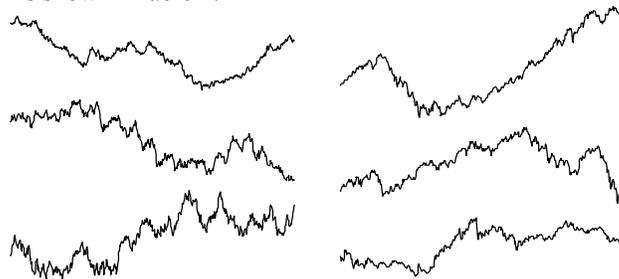


Figure 1. Six time series, three are random walk data, and three are real S&P500 stocks. Experiments suggest that humans cannot tell real and synthetic stock data apart (all the sequences on the right are real).

Table 1. The confusion matrix for human experts in attempting to differentiate between random walk data and stock market data.

		Predicted	
		S&P Stock	Random Walk
Actual	S&P Stock	20	16
	Random Walk	16	20

The accuracy of the humans was 55.6%, which does not differ significantly from random guessing.

Given the above, if we consider stock market and random walk data to be the same, each paper in the survey is tested on average on only 1.28 different datasets. This number might be reasonable if the contribution had been claimed for only a single type of data (Gavrilov et al., 2000, Lavrenko et al., 2000), or it had been shown that the choice of dataset has little influence on the outcome. However, the choice of dataset has a huge effect on the performance of time series algorithms. We will demonstrate this fact in the next 3 sections of this work.

### 3. Indexing (Query by Content)

Similarity search in time series databases has emerged as an area of active interest since the classic first paper by Agrawal et al. (1993). More than 68% of the indexing approaches surveyed here use the original GEMINI framework (Faloutsos et al., 1994), but suggest a different approach to the dimensionality reduction stage. The proposed representations include the Discrete Fourier Transform (DFT) (Agrawal et al., 1993, Chu and Wong, 1999, Faloutsos et al., 1997, Kahveci et al., 2002, Rafiei and Mendelzon, 1998, Rafiei, 1999), several kinds of Wavelets (DWT) (Chan and Fu, 1999, Kahveci and Singh, 2001, Popivanov and Miller, 2002, Shahabi et al., 2000, Wang and Wang, 2000b, Wu et al., 2000b), Singular Value Decomposition (Keogh et al., 2001, Korn et al., 1997), Adaptive Piecewise Constant Approximation (Keogh et al., 2001), Inner Products (Ferhatosmanoglu et al., 2001) and Piecewise Aggregate Approximation (PAA) (Yi and Faloutsos, 2000). The majority of work has focused solely on performance issues, however some authors have also considered other issues such as supporting non Euclidean distance measures (Keogh et al., 2001, Rafiei, 1999, Yi and Faloutsos, 2000) and allowing queries of arbitrary length (Keogh et al., 2001, Loh et al., 2000, Yi and Faloutsos, 2000).

#### 3.1. Implementation bias

Since most time series indexing techniques use the same indexing framework, and achieve the claimed speedup solely with the choice of representation, it is important to compare techniques in a manner that is free of implementation bias.

**Definition:** *Implementation bias* is the conscious or unconscious disparity in the quality of implementation of a proposed approach, vs. the quality of implementation of the competing approaches.

Implementing fairly complex indexing techniques allows many opportunities for implementation bias. For example, suppose you hope to demonstrate that DWT is superior to DFT. With shift-normalized data (Chu and Wong, 1999, Kahveci et al., 2002) the first DWT coefficient is zero so you could take advantage of that fact by indexing the 2<sup>nd</sup> to N+1<sup>th</sup> coefficients, rather than the 1<sup>st</sup> to N<sup>th</sup> coefficients. However, you might neglect doing a similar optimization for DFT, whose first real coefficient is also zero for normalized data. Another possibility is that you might use the simple  $O(n^2)$  DFT algorithm rather than spend the time to code the more complex  $O(n \log n)$  radix 2

algorithm (Keogh et al., 2001). In both these cases DFT's performance would be artificially deflated relative to DWT.

One possible solution to the problem of implementation bias is extremely conscientious implementations of all approaches, combined with diligent explanations of the experimental process. Another possibility, which we explain below, is to design experiments that are free from the possibility of implementation bias.

Since all the exact indexing techniques use the same basic framework, the efficiency of indexing depends only on how well the dimensionality-reduced approximation can model the distances between the original objects. We can measure this by calculating the tightness of the lower bounds for any given representation.

**Definition:** The *tightness of the lower bound* (denoted  $T$ ) for any given representation is the ratio of the estimated distance between two sequences under that representation, over the true distance between the same two sequences.

Note that  $T$  is in the range  $[0,1]$ . A value of 1 would allow a constant time search algorithm, and a value of 0 would force the indexing structure to degrade to sequence scan. In fact, because sequential scan can take advantage of a linear traverse of the disk, whereas any indexing scheme must make wasteful random disk accesses, it is well understood that  $T$  must be significantly greater than 0 if we are to use the representation to beat sequential scan (Keogh et al., 2001). Since one can always create artificial data for any representation that will give an arbitrary value of  $T$ , it should be estimated *for a particular dataset* by random sampling. Note that the value of  $T$  for any given dimensionality reduction technique depends only on the data and is independent of any implementation choices such as page size, buffer size, computer language, hardware platform, seek time etc. A handful of papers in the survey already make use of a similar measure to compare the quality of representations (Chan and Fu, 1999, Keogh et al., 2001).

This idea of an implementation free evaluation of performance is by no means new. In artificial intelligence, researchers often compare search algorithms by reporting the number of nodes expanded, rather than the CPU times (Kibler and Langley, 1988). The problem of implementation bias is also well understood in other computer science domains, including parallel processing (Bailey, 1991).

### 3.2. Data bias

As mentioned above, the tightness of the lower bound can be estimated by random sampling of a dataset. However we have not yet addressed the importance of *which* dataset(s) are sampled. The indexing papers included in this survey tested their approach on a median of 1 datasets. This would be reasonable if the utility of the approach was only being claimed for a single type of data, for example "*More Efficient Indexing of ECG Time Series*" or "*A New Approach to Indexing Stock Market Data*". However, none of the papers make such a limited claim. The papers are implicitly or explicitly claiming to be improvements over the state of the art on *any* time series data. In fact, the choice of test data has a great effect on the experimental results, and virtually all papers surveyed suffer from data bias.

**Definition:** *Data bias* is the conscious or unconscious use of a particular set of testing data to confirm a desired finding.

There does not appear to be a simple cure for data bias. One possibility is to limit the scope of the claim for a new approach to that which has actually been demonstrated, e.g. “*Faster indexing of Stock Market Data*”. Another possibility, which we favor, is to test the algorithms on a large, heterogeneous set of time series. Ideally this set should include data that covers the spectrum of time series properties; stationarity/ non-stationarity, noisy/ smooth, cyclical/ non-cyclical, symmetric/ asymmetric, etc.

### 3.3. Empirical demonstration of implementation and data Bias

To demonstrate the need for an implementation-free measure of the quality of indexing technique, and the absolute necessity of testing new algorithms on several datasets, consider the following contradictory claims made with regard the relative indexing abilities of DFT and DWT (wavelets):

- “*Several wavelets outperform the Haar wavelet (and DFT)*” (Popivanov and Miller, 2002).
- “*DFT-based and DWT-based techniques yield comparable results in similarity search*” (Wu et al., 2000b).
- “*Haar wavelets perform slightly better than DFT*” (Kahveci and Singh, 2001).
- “*DFT filtering performance is superior to DWT*” (Kawagoe and Ueda, 2002)

Which, if any, of these statements are we to believe? Because of the problems of implementation bias and the limited number of test datasets we feel little credence can be given to any of the claims. To demonstrate this we have performed a comprehensive series of experiments that show that the variance due to implementation bias and testing on different data can far outweigh the improvements claimed in the literature.

We calculated the value of  $T$  for both DFT and DWT. To ensure that we obtained good estimates we averaged over 100,000 randomly chosen subsequences from each dataset. For fairness we used the same 100,000 subsequences for each approach. To ensure randomness in our sampling technique we used true random numbers that were created by a quantum mechanical process (Walker, 2001).

#### 3.3.1. demonstration of data bias

The three papers listed above experimented on a maximum of 3 datasets. If we use that number of datasets we can demonstrate essentially any finding we wish. For example, by working with the Powerplant, Infrasound and Attas datasets we can find that DFT outperforms the Haar wavelet, as shown in Figure 2.

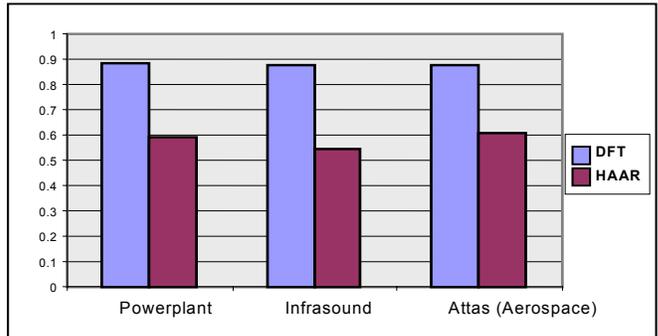


Figure 2. Experiments on the Powerplant, Infrasound and Attas datasets “demonstrate” that DFT outperforms DWT-Haar for indexing time series.

In contrast if we worked with the Network, ERPdata and Fetal EEG datasets we could conclude that there is no real difference between DFT and Haar, as suggested by Figure 3.

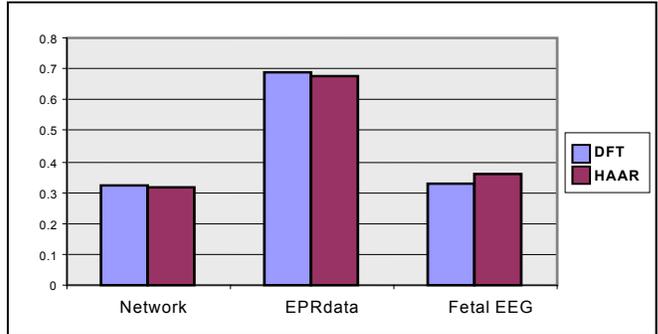


Figure 3. Experiments on the Network, ERPdata and Fetal EEG datasets “demonstrate” that DFT and DWT-Haar have the same performance for indexing time series.

Finally had we had chosen the Chaotic, Earthquake and Wind datasets we could use the graphs in Figure 4 to demonstrate “convincingly” that Haar is superior to DFT.

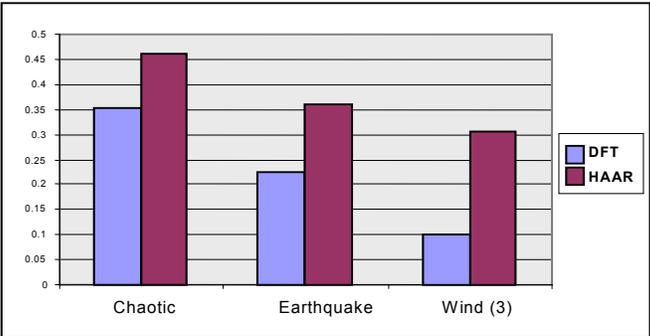


Figure 4. Experiments on the Chaotic, Earthquake and Wind datasets “demonstrate” that DWT-Haar outperforms DFT for indexing time series.

Although we used the value of  $T$  to demonstrate the problem, we also confirmed the findings on an implemented system, using an R-tree running on AMD Athlon 1.4 GHZ processor, with 512 MB of physical memory and 57.2 GB of secondary storage. The results were essentially identical, so we omit the graphs for brevity.

Note that we are not claiming any duplicity by the authors of the excellent papers listed above. We are merely demonstrating that the limited number of datasets used in the typical indexing paper severely limits the claims one can make.

### 3.3.2 demonstration of implementation bias

The vast majority of papers on indexing that do use a strawman comparison use the simplest possible one, sequential scanning. Here we will demonstrate the potential for implementation bias with sequential scanning performed in main memory.

The Euclidian distance function is shown in Eq. 1.

$$D(Q, C) \equiv \sqrt{\sum_{i=1}^n (q_i - c_i)^2} \quad (1)$$

The basic sequential search algorithm is shown in Table 2.

Table 2. The Sequential Search Algorithm

<b>Algorithm</b> sequential_scan(data, query)
<pre> best_so_far = inf; for every item in the database   if euclidian_dist(data<sub>i</sub>, query) &lt; best_so_far     pointer_to_best_match = i;     best_so_far = euclidian_dist(data<sub>i</sub>, query);   end; end;</pre>

One possibility implementation, which we call *Naïve*, is to calculate the Euclidian distance as shown in Eq. 1 with a loop to accumulate all the partial sums, followed by the taking of the square root. A possibility for optimization, which we call *Opt1*, is to neglect taking the square root. Since the square root function is monotonic, the ranking of the nearest neighbors will be identical under this scheme (Yi and Faloutsos, 2000). Finally we consider another optimization, which we call *Opt2*, which is simply to keep comparing the *best\_so\_far* variable to the partial sums at each iteration of the loop. If the partial sum ever exceeds the value of *best\_so\_far* we can admissibly abandon that calculation, since the partial distance can never decrease. To test the effect of these minor implementation details we performed 1-nearest neighbor searches in a random walk dataset, with a query length of 512 for increasingly larger datasets. The results are shown in Figure 5.

It is obvious that these very minor implementation details can produce large differences. If we are comparing a novel algorithm to sequential scan, and omit details of sequential scan implementation, it would be very hard to gauge the merit of our contribution. Note that for simplicity we only considered a main memory search. If we consider a disk-based search, there are a myriad of other implementation details that could effect the performance of sequential scan by at least an order of magnitude.

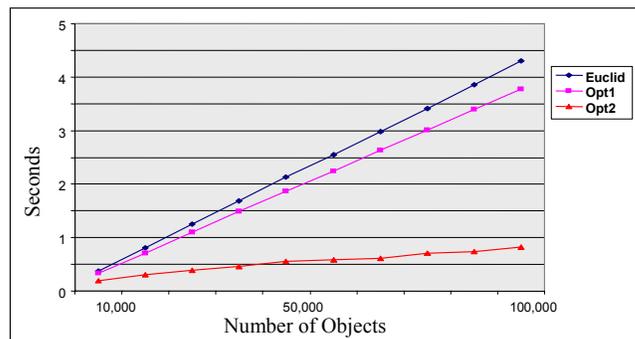


Figure 5. The affect of minor implementation details on the performance of sequential scan, for increasing large databases.

It is easy to find examples of data bias in the literature, it is much more difficult to know the scale of the problem for implementation bias. By its very nature, it is almost impossible to know what fraction of a claimed improvement should be credited to the proposed approach, and what fraction may be due to implementation bias. However, there are a handful of examples where this is clear. For example, one paper included in the survey finds a significant performance gap between the indexing abilities of Haar wavelets and Piecewise Aggregate Approximation (PAA) (Popivanov and Miller, 2002). However it was proved by two independent groups of researchers that these two approaches have exactly the same tightness of lower bounds when the number of dimensions is a power of two (and very little difference when the number of dimensions is not a power of two) (Keogh et al., 2001, Yi and Faloutsos, 2000). We empirically confirmed this fact 4,000,000 times during the experiments in Section 3.3.1. While there may be small differences in the CPU time to deal with the two representations, the order in which the original sequences are retrieved from disk by the index structure should be the same for both approaches, and disk time completely dominates CPU for time series indexing under Euclidean distance. We strongly suspect the spurious result reported above was the result of implementation bias, so we conducted an experiment to demonstrate how a simple implementation detail could produce an effect which is larger than the approximately 11% difference claimed.

We began our experiment by performing a fair comparison of the tightness of lower bounds for Haar and PAA on each of our 50 datasets, with a query length of 256 and 8 dimensions. Rather than estimate  $T$  with 100,000 random samples as in Section 3.1.1, we averaged over 100 samples as in the paper in question.

We repeated the experiment once more; this time neglecting to take advantage of the fact the first Haar coefficient is zero for normalized data. In other words, we wastefully index a value that is a constant zero. Once again we estimated  $T$  by averaging over 100 samples for each dataset.

For each dataset we calculated the ratio of the correct implementation's value of  $T$  to the poor implementation's value of  $T$ . The 50 results are plotted as a histogram in Figure 6.

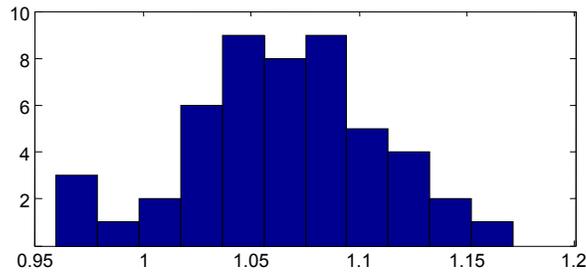


Figure 6. The distribution of the ratios of the results of correctly implemented experiments to experiments that have a slight implementation bias.

It is surprising to note that sometimes implementation bias that should favor an approach can actually hurt it, as happened 4 times out of the 50 experiments. However we must remember that the values of  $T$  for each dataset were only estimated from 100 samples, and the finding is not statistically significant. What is clear from the experiment however is that a simple minor implementation detail can produce effects that are as large as the claimed improvement of the proposed approach.

### 3.3.3 A real world case study illustrating the problems of implementation and data bias

In this section we will carefully deconstruct a recent paper to illustrate how data bias and implementation bias can render the contribution of a paper nil. We will consider a paper by Kim et al. (2001). Note that we chose this paper simply because the clear writing makes it easy to deconstruct and the detailed exposition of the method allows exact reimplementations. We do not mean to imply anything about the quality of the generally excellent work by these authors. Indeed the paper in question should be commended for including a rigorous proof of the proposed method, a rarity in the data mining literature. Once again, our point is simply to illustrate that even experienced researchers can be misled by data and implementation bias to make extravagant and unwarranted claims. In particular the claim made is that the proposed technique can allow similarity search under Dynamic Time Warping (DTW) that is up to 720 times faster than sequential search. As we shall show, the correct claim for the approach is that it can produce meaningless results quickly, or meaningful results very slowly (slower than sequential scan), but it *cannot* produce meaningful results quickly.

The paper is concerned with the laudable idea of indexing DTW. It has been forcibly demonstrated that in some domains the Euclidean distance is a brittle distance measure, because it is very sensitive to small distortions in the time axis (Keogh 2002, Berndt and Clifford, 1996, Yi et al. 1998). Dynamic Time Warping is a technique that measures the distance between two time series after first aligning them in the time axis. The drawback of DTW from a practical point of view is that it is  $O(n^2)$ , where  $n$  is the length of the time series of interest, in addition it cannot be indexed by standard spatial access methods which require the distance measure to be a metric. By contrast, Euclidean distance is merely  $O(n)$ , and is trivially indexable (Agrawal et al., 1993). The core contribution of the paper by Kim et al. (2001) is to introduce a novel linear time lower bounding technique for DTW. The lower bounding technique allows faster linear search, by

allowing one to prune off a fraction of the database by noting that the lower bound estimate of the distance to a candidate sequence is greater than the currently known *best-so-far* match. However, an additional claimed feature of the lower bounding estimate is that it is indexable, thus allowing one to retrieve candidate sequences in “most promising first” order. While a complete explanation of the technique is too detailed to present here, we have included a visual intuition of the lower bounding function in Figure 7 for completeness (See the original paper or Keogh, 2002. for a more detailed explanation). The lower bounding function works by extracting a 4-tuple-feature vector from each sequence. The features are the first and last elements of the sequence, together with the maximum and minimum values. The maximum squared difference of corresponding features is reported as the lower bound.

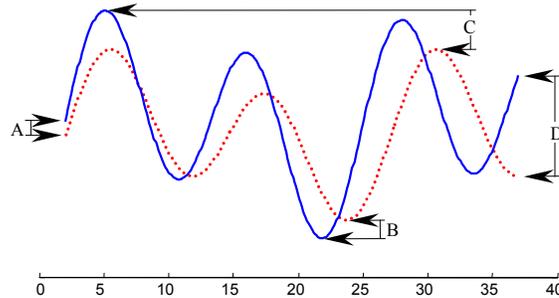


Figure 7. A visual intuition of the lower bounding measure introduced by Kim et al. The maximum of the squared difference between the two sequence’s first (A), last (D), minimum (B) and maximum points (C) is returned as the lower bound.

Before detailing the problems with the contribution, we must review an important preprocessing step that is necessary before attempting to match two time series under Euclidean distance, Dynamic Time Warping or any other distance measure. Consider the two time series shown in Figure 8. Although they appear to have similar shapes, they occur at different offsets in the Y-Axis. The Euclidean distance between the two sequences is proportional to the length of the gray hatch lines. We can see that shifting the two sequences to the same offset will greatly reduce the reported distance and reveal the true similarity between the sequences. The optimal shifting can be quickly achieved by simply subtracting the mean of a sequence from the entire sequence. This step is known as “normalizing” the data<sup>1</sup>. Note that although the Euclidean distance is shown here for simplicity, identical remarks apply to DTW or any other distance measure.

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<sup>1</sup> There are some other steps that may be included in the normalizing step, such as rescaling the time series and removing linear trend. We will ignore these steps for simplicity, since they don’t affect the argument that follows.

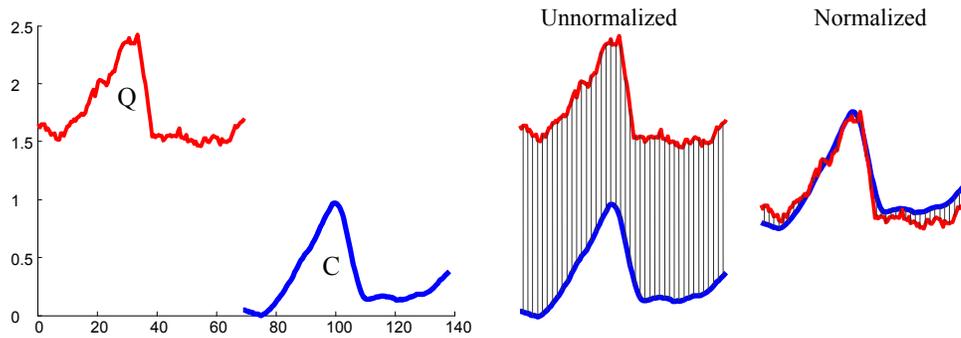


Figure 8. A visual intuition of the necessity to normalize time series before measuring the distance between them. The two sequences Q and C, on the left appear to have approximately the same shape, but have different offsets in the Y-Axis. The Euclidean distance between two sequences is proportional to the length of the gray hatch lines shown connecting the two sequences in the two graphics on the right. In the first case the data is unnormalized, and the reported distance greatly overstates the subjective dissimilarity. Normalizing the data reveals the true similarity of the two time series.

What would happen if we didn't normalize the time series before measuring the similarity? We can illustrate by clustering some Space Shuttle telemetry data with and without normalization as in Figure 9.

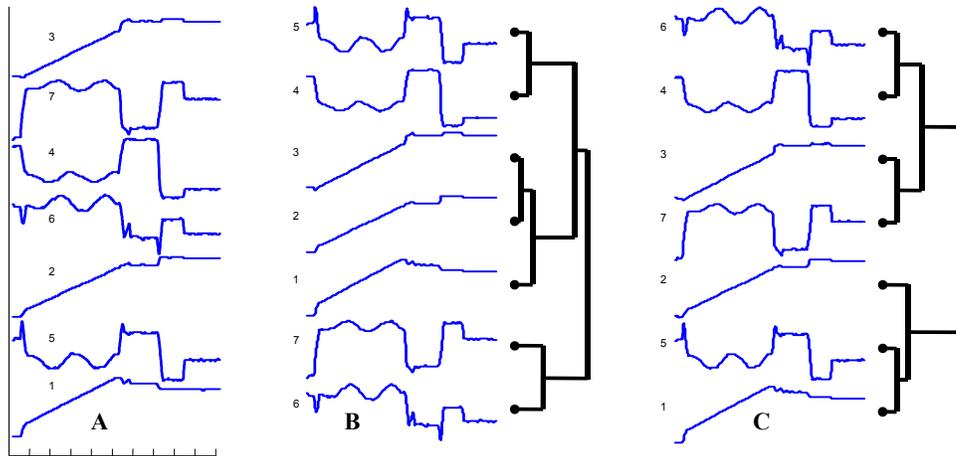


Figure 9. A visual explanation of the need to normalize time series before measuring their similarity. **A)** Seven time series obtained from inertial sensors on board Space Shuttle mission STS-57. **B)** The clustering obtained when we first normalize the signals, the clustering appears very natural and subjectively correct. **C)** The clustering obtained if we do *not* normalize the signals, the clustering appears essentially random.

The results are quite startling, without normalization time series similarity has essentially no meaning. More concretely, very small changes in offset rapidly dwarf any information about the shape of the two time series in question. For example, if we clustered just the mean values of the time series, the resulting dendrogram would be almost indistinguishable from the one shown in Figure 9.C. It is important to realize that although Figure 9 shows time series with greatly differing offsets for clarity, even

relatively small differences in offset, a tiny fraction of the variance of the signals, is enough to render the measurement of similarity meaningless.

Having reviewed the need for normalization, we are now in a position to demonstrate our earlier claim that the work of Kim et al. (2001) cannot produce meaningful results quickly. The authors report only experiments with no normalization (Park 2001). If the dataset used was already normalized, or had extremely low variance in offset this might not matter, however, this is not the case; the variance of offsets is greater than that shown in Figure 9, and the similarity measurements are equally meaningless (although it depends on the length of the query, we found that the variance of offsets is typically orders of magnitude larger than the variance of the signals on the dataset in question).

An obvious reaction is “so what”, can’t we just normalize the data and still get the 3 orders of magnitude speedup claimed? The answer is no! If we normalize the data the lower bound becomes extremely weak, a tiny fraction of the true distance, and thus the pruning power of the lower bound is greatly reduced. In essence the problem is this, the lower bounding function is mostly measuring information about how far apart the two time series are, and not measuring information about shape. If the time series are far apart, the lower bound is very tight (although the similarity is meaningless), but when the time series are normalized, the lower bound is so weak and only a small handful of candidate time series can be pruned, certainly not enough to justify the computational overhead.

To demonstrate our claim we tested the approach on 33 different datasets, using query lengths of 256, 512 and 1,042. In keeping with the author’s experiments, we compared the approach to sequential scan. We measured the normalized CPU cost (Keogh et al., 2001), which takes into account the fact that sequential scan enjoys a tenfold speed up when performed on disk.

**Definition:** *The Normalized CPU cost:* The ratio of average CPU time to execute a query using the index to the average CPU time required to perform a linear (sequential) scan. The normalized cost of linear scan is 1.0.

The justification being that any indexing technique must perform costly random access, whereas sequential scan can take advantage of an optimized linear traverse of the disk. Rather than summarize all the results with graphs, we will simply note that the technique never once beat sequential scan on any one of our 111,375 test queries.

Once again we must restate that we are not singling out this work for criticism. Only 57% of the papers in the survey explicitly state their normalization policy. For the papers that do not, we cannot be sure if their results are meaningful. The point of this section is to show that implementation bias, (the lack of normalization which helps the proposed technique while not affecting sequential scan), combined with data bias, (choosing a dataset with large offset variance), can result in work which appears to give dramatic speedup, but is actually worse than doing nothing at all.

#### **4. Classification and Clustering**

Classification and clustering problems have been the subject of active research for decades (Cohen, 1993, Kibler and Langley, 1988). However the unique structure of time series means that most classic machine learning algorithms do not work well for time series. In particular the high dimensionality, very high feature correlation, and the (typically) large amounts of noise that characterize time series data have been viewed as an interesting research challenge. Most of the contributions focus on providing a new similarity measure as a subroutine to an existing classification or clustering algorithm, so for simplicity we shall only consider the contribution of the suggested similarity measure. How well do these similarity measures capture the true similarity of time series? There are two ways to answer this question, subjectively and objectively, we consider both below.

#### *4.1 Subjective evaluation of similarity*

Since a goal of data mining is often to find patterns that map onto human intuition, one possible way to judge the utility of a similarity measure is to show examples of time series that the proposed measure found to be similar/dissimilar. Surprisingly, many of the papers included in the survey, whose main contribution was to introduce a new similarity measure, fail to show even one example of a matching pair of time series (André-Jönsson and Badal, 1997, Bozkaya et al., 1997, Gavrilov et al., 2000, Goldin and Kanellakis, 1995, Huang and Yu, 1999, Indyk et al., 2000, Kim et al., 2000, Lam and Wong, 1998, Lee et al., 2000, Park et al., 2001, Park et al., 1999, Qu et al., 1998, Wang and Wang, 2000b). Moreover, showing some examples of matching time series is of little utility unless some strawman comparison is used. Many papers ask us to consider the quality of their proposed similarity measure without a single comparison to another technique (Agrawal et al., 1995b, André-Jönsson and Badal, 1997, Bozkaya et al., 1997, Huang and Yu, 1999, Keogh and Smyth, 1997, Lee et al., 2000, Li et al., 1998, Park et al., 2000, Park et al., 2001, Pratt and Fink, 2002, Wang and Wang, 2000b). This is particularly surprising since the most obvious strawman, Euclidian distance, is trivial to implement (For example, in the Matlab programming language it requires only 19 characters: `sqrt(sum((q-c).^2))`).

We believe that one of the best (subjective) ways to evaluate a proposed similarity measure is to use it to create a dendrogram of several time series from the domain of interest (Keogh and Pazzani, 1998).

Additional dendrograms can be created using other measures then plotted side by side with the proposed approach. Figure 10 shows an example.

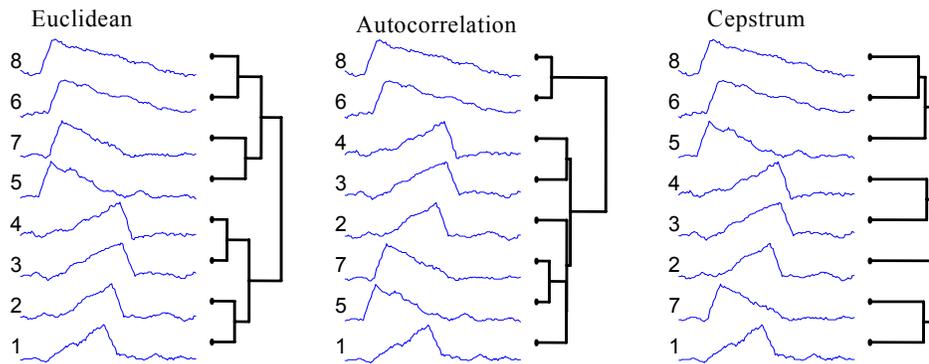


Figure 10. Dendrograms can be used to visually assess the usefulness of a similarity measure. Above a dataset of 8 objects is clustered using the single linkage method, with 3 different distance measures. Euclidean distance is a decade old strawman. The other two approaches have recently been proposed in data mining papers.

Dendrograms are particularly attractive since a clustering of  $M$  objects summarizes  $O(M)$  measurements, however other possibilities of visualizing the quality of a similarity measure included projecting the time series into 2 dimensional space (via MDS or SOMs for example (Debregeas and Hebrail, 1998)).

#### 4.2. Objective evaluation of similarity

Given a database of labeled time series, objective measurements of the quality of a proposed similarity measure can be readily obtained by running simple classification experiments. Although a few such databases do exist, very few advocates of a new similarity measure have chosen to demonstrate their contribution in this manner. The work by Geurts is a notable exception (2001). To repair this omission, we have undertaken an experimental comparison of many of the techniques included in the survey. We tested on two publicly available datasets:

- **Cylinder-Bell-Funnel:** This synthetic dataset has been in the literature for 8 years, and has been cited at least a dozen times (Geurts, 2001). It is a 3-class problem; we create 128 examples of each class for these experiments.
- **Control-Chart:** This synthetic dataset has been freely available for the UCI Data Archive since June 1998 (Bay, 1999). It is a 6-class problem, with 100 examples of each class.

Note that for both problems, informal experiments suggest humans can achieve an error rate of zero. For simplicity we use the 1-Nearest Neighbor algorithm, evaluated using “leaving-one-out”. We compare the proposed methods to the simplest strawman, Euclidean distance. This measure is well-known (Agrawal et al., 1993, Chan and Fu, 1999, Chu and Wong, 1999, Das et al., 1997, Das et al., 1998, Faloutsos et al., 1997, Faloutsos et al., 1994, Ferhatosmanoglu et al., 2001, Kahveci and Singh, 2001, Keogh et al., 2001, Korn et al., 1997, Lam and Wong, 1998, Loh et al., 2000, Popivanov and Miller, 2002, Rafiei and Mendelzon, 1998, Rafiei, 1999, Wu et al., 2000b, Yi and

Faloutsos, 2000, Yi et al., 1998), parameterless, trivial to implement and predates data mining by several decades.

We originally intended to implement every proposed similarity measure in our survey, but several of the papers do not include a detailed enough description to allow reimplementing (Li et al., 1998, Qu et al., 1998). We contented ourselves with reimplementing 11 measures. Some of the measures require the user to set some parameters. In these cases we wrapped the classification algorithm in a loop for each parameter, searched over all possible parameters and reported only the *best* result.

Table 3 summarized the results.

Table 3. The error rates for various similarity measures

Approach	Cylinder-Bell-Funnel	Control-Chart
<i>Euclidean Distance</i>	<i>0.003</i>	<i>0.013</i>
Aligned Subsequence [Park et al. 2001]	0.451	0.623
Piecewise Normalization [Indyk et al. 2002]	0.130	0.321
Autocorrelation Functions [Wang & Wang 2000b]	0.380	0.116
Cepstrum [Kalpakis et. al. 2001]	0.570	0.458
String (Suffix Tree) [Huang & Yu 1999]	0.206	0.578
Important Points [Pratt & Fink 2002]	0.387	0.478
Edit Distance [Bozkaya et al.1997]	0.603	0.622
String Signature [Jonsson & Badal 1997]	0.444	0.695
Cosine Wavelets [Huntala et al. 1999]	0.130	0.371
Hölder [Struzik & Siebes]	0.331	0.593
Piecewise Probabilistic [Keogh & Smyth 1997]	0.202	0.321

The results are quite surprising. None of the proposed techniques can beat the simple strawman. Their error rates are an order of magnitude worse than Euclidean distance. Several of the techniques have error rates close to the default rate (i.e. the same error you would get randomly guessing). Although the inability to perform well on these two objective tests does not necessarily mean the similarity measures in question are without any merit (there may exist datasets on which they have reasonable accuracy), one has to wonder about the contribution of a new similarity measure which fails to demonstrate its utility on *any* objective or subjective test<sup>2</sup>.

## 5. Segmentation

A large fraction of the papers in the survey either introduce a segmentation algorithm as their main contribution, or utilize a segmentation algorithm as a subroutine. Although the segments created could be polynomials of an arbitrary degree, the most common representation of the segments are linear functions. Intuitively a Piecewise Linear

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<sup>2</sup> Once again we wish to note that the current first author introduced one of the poorly performing measures.

Representation (PLR) refers to the approximation of a time series  $Q$ , of length  $n$ , with  $K$  straight lines. Figure 8 contains an example.

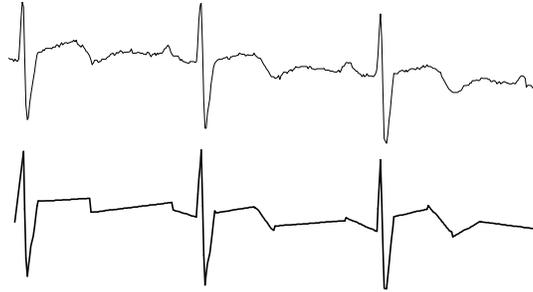


Figure 11. An example of a time series with its piecewise linear representation.

Because  $K$  is typically much smaller than  $n$ , this representation makes the storage, transmission and computation of the data more efficient. Specifically, in the context of data mining, piecewise linear representation has been used to:

- Support novel distance measures for time series, including “fuzzy queries” (Shatkay and Zdonik, 1996), weighted queries (Keogh and Pazzani, 1998), multiresolution queries (Li et al., 1998, Qu et al., 1998), dynamic time warping (Park et al., 2001, Pratt and Fink, 2002), autocorrelation queries (Wang and Wang, 2000b) and relevance feedback (Keogh and Pazzani, 1998).
- Support concurrent mining of text and time series (Lavrenko et al., 2000).
- Support novel clustering and classification algorithms (Keogh and Pazzani, 1998).
- Support change point detection (Ge and Smyth, 2000, Guralnik and Srivastava, 1999).

Surprisingly, in spite of the ubiquity of this representation, with the exception of the work by Shatkay and Zdonik, (1996), there has been little attempt to understand and compare the algorithms that produce it.

Although appearing under different names and with slightly different implementation details, most time series segmentation algorithms can be grouped into one of the following three categories.

- **Sliding-Windows (SW)**: A segment is grown until it exceeds some error bound. The process repeats with the next data point not included in the newly approximated segment.
- **Top-Down (TD)**: The time series is recursively partitioned until some stopping criteria is met.
- **Bottom-Up (BU)**: Starting from the finest possible approximation, segments are merged until some stopping criteria is met.

We can measure the quality of a segmentation algorithm in several ways, the most obvious of which is to measure the reconstruction error for a fixed number of segments. The reconstruction error is simply the Euclidean distance between the original data and the segmented representation.

### 5.1 Data bias in segmentation

Given that we have 3 algorithms to produce a segmented version of a time series, it is natural to ask which is best. The papers in the survey that use a segmentation algorithm test on a median of 1 dataset. However, if we use only one dataset we can demonstrate any finding we wish! There are 3 different algorithms, therefore  $3! = 6$  possible rankings. We tested the algorithms on our 50 fifty datasets, asking each algorithm to reduce a 1,024 datapoint time series to 64 segments. Amazingly, we found every possible ranking of the 3 algorithms as shown in Table 4.

Table 4. The 3 algorithms under consideration, ranked by reconstruction error (shown in brackets), on 6 datasets

Dataset	Best Algorithm	Second-Best Algorithm	Third-Best Algorithm
Soiltemp	TD (522.6)	SW (538.0)	BU (538.1)
Darwin	TD (575.2)	BU (821.0)	SW (833.9)
pHdata	SW (3.590)	TD (4.013)	BU (4.037)
Winding	SW (6.883)	BU (113.0)	TD (117.6)
Balloon	BU (168.1)	TD (224.5)	SW (234.1)
Network	BU (11.02)	SW (13.62)	TD (891.4)

Note that the fact that we could easily find datasets to demonstrate any ranking we wish does not preclude us from making a meaningful evaluation of the algorithms. In fact the Bottom-Up algorithm is significantly better than the other two approaches<sup>3</sup>. Our point, once again, is simply that little credence can be given to experimental results obtained from testing on a single dataset.

## 6. Conclusions and Recommendations

In this work we have conducted a comprehensive survey of recent work on time series data mining. We have shown that because of several kinds of experimental flaws, in particular data bias and implementation bias, many of the results claimed in the literature have very little generalizability to real world problems. We have demonstrated our claim with the most comprehensive set of time series experiments ever undertaken.

Once again we would like to note that we view this work as a “call to arms” to the data mining community, and not a criticism of the many wonderful and original papers cited here. The intended spirit of this paper is similar to the ironically titled work by Bailey, “*Twelve ways to fool the masses when giving performance results on parallel computers*” (1991). The author later noted that few, if any researchers set out to deliberately mislead the academic community, but unless greater effort is made to meaningfully compare rival

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<sup>3</sup> Bottom-Up outperformed Top-Down on 47 of 69 datasets, and it outperformed Sliding Windows on 58 of 69 datasets.

approaches, the entire field is in danger of being viewed with suspicion. This current work is an echo of that sentiment for the time series data mining community.

We conclude this paper with concrete suggestions for researchers working on time series data mining.

- Algorithms should be tested on a wide range of datasets, unless the utility of the approach is only been claimed for a particular type of data. If possible, one subset of the datasets should be used to fine tune the approach, then a different subset of the datasets should be used to do that the actual testing. This methodology is widely used in the machine learning community to help prevent implementation and data bias (Cohen, 1993).
- Where possible, experiments should be designed to be free of the possibility of implementation bias. Note that this does not preclude the addition of extensive implementation testing.
- Novel similarity measures should be compared to simple strawmen, such as Euclidian distance or Dynamic Time Warping. Some subjective visualization, or objective experiments should justify their introduction.
- Where possible, all data and code used in the experiments should be made freely available to allow independent duplication of findings (Bay, 1999).

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All papers except (Bailey, 1991, Bay, 1999, Cohen, 1993, Keogh 2002, Kibler and Langley, 1988, Prechelt, 1995, Simon, 1994), are included in the survey. The excluded papers contain background information.

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