

## Entropies for detection of Term/Preterm delivery:A Review

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### Abstract

*Impending preterm delivery is likely associated with a change in uterine cell excitability, favoring conduction of electrical activity across the uterine muscles, and myometrium. Analysis of the electrohysterogram (EHG), which is the noninvasive measurement of the uterine electrical activity, has therefore been investigated as a potential method for pregnancy monitoring and early diagnosis of preterm birth. Various Entropies are signal features derived from information theory. They reflect the regularity of a time series and have been extensively used for analysis and characterization of physiological signals. The sample entropy and approximate entropy can detect the difference between term and preterm delivery and performance of the system can further be improved after using various entropies.*

**Keywords:** *Electrohysterogram, Non Communicable Diseases, Approximate Entropy*

### 1.Introduction:

According to World Health Organization report “Born Too Soon”, there are about 15 million babies born prematurely in the world among a total of about 150 million births per year. Families all around the world are affected as more than 1 in 10 babies are born preterm. Over 1 million children die each year due to complications of preterm birth. Most of the children that survives face a lifetime of disability like chronic lung disease, learning disabilities, cerebral palsy, intellectual impairment, visual along with hearing problems and they have a strong chances of developing Non Communicable Diseases (NCD) like hypertension, diabetes and some other crucial health conditions later in life. They are physically not ready to face the world and often require special care [1]. In a systematic analysis and implications, India was given the highest rank with the highest numbers of preterm births in year 2010 with approximately 1 in 8 babies being preterm. In general most preterm births occur after 32 computed weeks of gestation. More than 90% of babies born before 28 weeks of gestation survive in high income countries but in low-income settings, only 10% of these babies or less survive [2]. Current approaches to prevention and treatment of preterm labor have been shown to be disappointingly unsuccessful [3]. Preterm Labor (PL) defined as labor before completing the 37th week of gestation is the main cause of newborn morbidity and mortality [4]. On the basis of previous studies it is known that even babies born at 34–37 weeks have an increased risk of immediate complications [5]. Lots of research is going on in this field but we are still unable to diagnose, prevent and treat preterm labor. Checking efficacy of involvements that would allow this is largely influenced by the inability to accurately identify true labor with the currently used crude technology. In the case of progestin treatment for prevention of preterm birth, uterine EMG and cervical LIF are essential tools to obtain the critically needed comparative data on effectiveness of various progestin formulations and their routes of administration in different patients at high risk for preterm delivery. To find an effective prevention and treatment for preterm labor, we must find a method that will allow focusing the treatment only to patients who would, if not treated, really deliver preterm [6]. ApEn

was developed by Pincus as a measure of regularity to quantify levels of complexity within a time series. The ability to discern levels of complexity within biological data sets has become increasingly important.

## 2. Previous Work:

Steven M. Pincus et al. (1990) proposed techniques to determine changing system complexity from data. Convergence of a frequently used correlation dimension algorithm to a finite value does not necessarily imply an underlying deterministic model or chaos. Analysis of a recently developed family of formulas and statistics, approximate entropy (ApEn), suggests that ApEn can classify complex systems, given at least 1000 data values in diverse settings that include both deterministic chaotic and stochastic processes. The capability to discern changing complexity from such a relatively small amount of data holds promise for applications of ApEn in a variety of contexts.[7]

Joshua S. Richman et al. (2000) have developed a new and related complexity measure, sample entropy (SampEn) and have compared ApEn and SampEn by using them to analyze sets of random numbers with known probabilistic character. They have also evaluated cross-ApEn and cross-SampEn, which use cardiovascular data sets to measure the similarity of two distinct time series. SampEn agreed with theory much more closely than ApEn over a broad range of conditions. The improved accuracy of SampEn statistics should make them useful in the study of experimental clinical cardiovascular and other biological time series. They have developed and characterized SampEn, a new family of statistics measuring complexity and regularity of clinical and experimental time series data and compared it with ApEn, a similar family. SampEn statistics agree much better than ApEn statistics with theory for random numbers with known probabilistic character over a broad range of operating conditions, maintain relative consistency where ApEn statistics do not, and have residual bias for very short record lengths, in a large part because of non-independence of templates. Furthermore, cross-SampEn is a more consistent measure of joint synchrony of pairs of clinical cardiovascular time series. The difficulties of ApEn analysis to the practice of counting self-matches and of cross-ApEn to the problem of unmatched templates resulting in undefined probabilities is also discussed. The differences are that SampEn does not count templates as matching themselves and does not employ a template-wise strategy for calculating probabilities. SampEn statistics provide an improved evaluation of time series regularity and should be a useful tool in studies of the dynamics of human cardiovascular physiology.[8]

Fele-Zorg et al. (2008) suggested that Sample entropy gives good results. The problem with sample entropy, lies in its extreme susceptibility to the parameter settings. If the matching pattern length,  $m$ , is too large, or if the sensitivity margin,  $r$ , is too low, within some time signals, no pattern matches can be found. The sample entropy in these cases depends only on the length of the signal and is much higher than usual. The outliers caused by this anomaly also affected the Student's  $t$ -test, producing a low  $p$ . To avoid such erroneous results, they limited their search for suitable  $m$  and  $r$  values to those that were found to be safe while using any of the three preprocessing filters. Noticeable differences were found between the groups of records recorded early and later. The technique indicated a difference between the term and pre-term delivery groups of records. As the time of gestation progresses, the average sample entropy values for term and pre-term delivery records drop indicating higher predictability of the signals as the delivery approaches. The average sample entropy values are lower for both early and later pre-term delivery records and indicate that the signals of pre-term delivery records exhibit higher predictability than the signals of term delivery records. The signal of the pre-term delivery record shows higher predictability than that of the term delivery record.[9]

Molina-Picó et al. (2011) proposed that there is an ongoing research effort devoted to characterize the signal regularity metrics approximate entropy (ApEn) and sample entropy (SampEn) in order to better

interpret these results in the context of biomedical signal analysis. They address the influence of abnormal spikes (impulses) on ApEn and SampEn measurements. A set of test signals consisting of generic synthetic signals, simulated biomedical signals, and real RR records was created. These test signals were corrupted by randomly generated spikes. ApEn and SampEn were computed for all the signals under different spike probabilities and for 100 realizations. The effect of the presence of spikes on ApEn and SampEn is different for test signals with narrowband line spectra and test signals that are better modeled as broadband random processes. In the first case, the presence of extrinsic spikes in the signal results in an ApEn and SampEn increase. In the second case, it results in an entropy decrease. For real RR records, the presence of spikes, often due to QRS detection errors, also results in an entropy decrease. They demonstrated that both ApEn and SampEn are very sensitive to the presence of spikes. Abnormal spikes should be removed, if possible, from signals before computing ApEn or SampEn. Otherwise, the results can lead to misunderstandings or misclassification of the signal regularity. The ApEn and SampEn performance in the context of biomedical signal analysis and the influence of spikes is evaluated. Neither ApEn nor SampEn should be sensitive to spikes, especially if they are infrequent. Theoretically,  $B_i$  and  $A_i$  should be big enough to cope with small variations due to outliers, that is to say, their relative changes should be very small. However, our experimental study demonstrated that spikes do influence the results of both regularity metrics. Mathematically, this is due to the fact that  $B_i$  and  $A_i$  are usually smaller than expected. For instance, in a sinusoidal random process, for  $B_i$ , although many matches are expected due to signal regularity, matches only account for 12% of the points approximately (for  $r = 0.2$ ), and in Gaussian white noise, for ten times less. The same applies to  $A_i$ , which is even lower than  $B_i$ . Thus, any change in the number of matches below the threshold  $r$ , may have a significant relative influence on the final result, and therefore, these metrics are sensitive to outliers. Additionally,  $A_i$  and  $B_i$  can be affected in a different scale. It can be seen that ApEn and SampEn are influenced by the presence of spikes, spikes increase the ApEn and SampEn in regular quasiperiodic signals but the opposite happens in the case of irregular broadband random processes and the capability to discern among types of signals based on features not apparent, seems not to be affected by the presence of spikes, provided all the classes are affected in the same manner and amount. Baseline wandering also influences the results of ApEn and SampEn since it reduces the number of matches, regardless of the signal regularity. It is very important to screen the signals before computing ApEn or SampEn in order to assure that abnormal spikes are not present. An improvement for ApEn or SampEn in the form of algorithm modification or new distance measure is necessary to obtain robustness against spikes and it may be advisable for researchers to apply spike-removal algorithms such as rank-order filters prior to the application of ApEn and SampEn in order to prevent erroneous signal regularity results. [10]

Mahmoud Hassan et al. (2011) have compared between three nonlinear methods, approximate entropy, correlation entropy and time reversibility was done on linear, nonlinear stationary and nonlinear nonstationary signals in order to choose the best method to apply on real EHG signals. Indeed these signals are thought to exhibit nonlinear as well as nonstationary characteristics. Detection of nonlinearity should be the first step before any analysis of nonlinearity or nonlinear behavior in biological signal. For the choice of best method characteristics of the signals under investigation are used to compare three methods widely used in nonlinearity detection: approximate entropy, correlation entropy and time reversibility. The false alarm rates with the numbers of surrogates for the three methods were computed on linear, nonlinear stationary and nonlinear nonstationary signals. The results indicate the superiority of time reversibility over the other methods for detecting linearity and nonlinearity in different signal types. The application of time reversibility on uterine electromyographic signal showed very good performance in classifying pregnancy and labor signals. The evolution of FAR of each method was computed with different surrogate numbers. The comparison demonstrated the superiority of time reversibility in the detection of linearity and nonlinearity of the different signals. Authors have tested EHG signals for their time reversibility property. The results indicate that uterine contractions during pregnancy are reversible, whereas labor contractions are temporally irreversible. They have shown that time reversibility could be a powerful tool to differentiate between pregnancy and labor contractions. It should be pointed out that this

nonlinearity measure, like other statistical nonlinearity measures, is based on the comparison with surrogate data. It is not a standalone measure needing the generation of surrogates. Therefore, the measure would fail if the generation of proper surrogates fails. Since non-Gaussianity is a characteristic common to both pregnancy and labor signals, the choice of the time reversibility parameter is made empirically, in order to obtain the best results on the synthetic signals and for the classification of pregnancy and labor signals. It will be important to investigate more closely the influence of this parameter on the detection of linearity and nonlinearity for the different synthetic signals. The results on the real data suggest that the property of time irreversibility is a strong characteristic for contraction measured on women in labor. This suggests that, for a more complete characterization of such recordings, additional nonlinear analysis techniques should be applied. The results provide a very powerful method for differentiating between pregnancy and labor contractions. [11]

Jennifer M. Yentes et al. (2012) proposed that Approximate entropy (ApEn) and sample entropy (SampEn) are mathematical algorithms created to measure the repeatability or predictability within a time series and both algorithms are extremely sensitive to their input parameters:  $m$  (length of the data segment being compared),  $r$  (similarity criterion), and  $N$  (length of data, especially for very short datasets,  $N \geq 200$ ). We suggest using  $N$  larger than 200, an  $m$  of 2 and examine several  $r$  values before selecting your parameters. There is no established consensus on parameter selection in short data sets, especially for biological data. They examined the robustness of these two entropy algorithms by exploring the effect of changing parameter values on short data sets. Data with known theoretical entropy qualities as well as experimental data from both healthy young and older adults was utilized. Based on their current findings, they proposed that SampEn is more reliable for short data sets. SampEn was less sensitive to changes in data length and demonstrated fewer problems with relative consistency. [12]

Peng Ren et al. (2015) discussed that preterm delivery increases the risk of infant mortality and morbidity, and therefore developing reliable methods for predicting its likelihood are of great importance. However, to date attempts at utilizing computational approaches to achieve sufficient predictive confidence, in terms of area under the curve (AUC) values, have not achieved the high discrimination accuracy that a clinical application requires. They proposed a new analytical approach for assessing the risk of preterm delivery using EMG recordings which firstly employs Empirical Mode Decomposition (EMD) to obtain their Intrinsic Mode Functions (IMF). Next, the entropy values of both instantaneous amplitude and instantaneous frequency of the first ten IMF components are computed in order to derive ratios of these two distinct components as features. Discrimination accuracy of this approach compared to those proposed previously was then calculated using six differently representative classifiers. Finally, three different electrode positions were analyzed for their prediction accuracy of preterm delivery in order to establish which uterine EMG recording location was optimal signal data. They have shown a clear improvement in prediction accuracy of preterm delivery risk compared with previous approaches, achieving an impressive maximum AUC value of 0.986 when using signals from an electrode positioned below the navel. This provides a promising new method for analyzing uterine EMG signals to permit accurate clinical assessment of preterm delivery risk. In this paper, we have demonstrated that it is possible to improve analysis of uterine EMG records to discriminate pregnant women at risk of preterm delivery with sufficient accuracy for potential clinical purposes. They have classified preterm and term delivery records based on the entropy ratios of the instantaneous amplitude and the instantaneous frequency of each two IMFs of uterine EMG signals using an EMD approach. Six different classifiers were implemented which revealed a mean AUC value of 0.89 and a maximum value of 0.986. A more detailed analysis of EMG recordings from specific abdominal locations revealed that the position with the strongest discrimination accuracy was the one below the navel. The study suggests that analysis of uterine EMG signals using the above approach is a very accurate method for discriminating women at risk of preterm delivery and may be important in clinical use. [13]

Massimo Mischi et al. (2017) suggested that preterm birth is a large-scale clinical problem involving over 10% infants. Diagnostic means for timely risk assessment are lacking and the underlying physiological mechanisms unclear. To improve the evaluation of pregnancy before term, they have introduced dedicated entropy measures derived from a single-channel electrohysterogram (EHG). The estimation of Approximate Entropy (ApEn) and Sample Entropy (SampEn) is adjusted to monitor variations in the regularity of single-channel EHG recordings, reflecting myoelectrical changes due to pregnancy progression. In particular, modifications in the tolerance metrics are introduced for improving robustness to EHG amplitude fluctuations. An extensive database of 58 EHG recordings with 4 monopolar channels in women presenting with preterm contractions was manually annotated and used for validation. The methods were tested for their ability to recognize the onset of labor and the risk of preterm birth. Comparison with the best single-channel methods according to the literature was performed. Results: The reference methods were outperformed. SampEn and ApEn produced the best prediction of delivery, although only one channel showed a significant difference ( $p < 0.04$ ) between labor and non labor. The modified ApEn produced the best prediction of preterm delivery, showing statistical significance ( $p < 0.02$ ) in 3 channels. The results were also confirmed by the area under the receiver operating characteristic curve and 5-fold cross-validation. Conclusion: The use of dedicated entropy estimators improves the diagnostic value of EHG analysis earlier in pregnancy. The changes in the EHG might manifest early in pregnancy, providing relevant prognostic opportunities for pregnancy monitoring by a practical single-channel solution. Entropy measures have been revisited and adjusted with the objective of achieving noninvasive prediction of delivery and diagnosis of preterm birth by analysis of single-channel EHG signals. On an extended database of in-house preterm EHG measurements, the estimated ApEn and SampEn have shown the best prediction of delivery, while the modified ApEn has shown a clear advantage for early diagnosis of delivery. These results prove the value of EHG entropy analysis as a tool for early, prognostic evaluation of pregnancy. [14]

### 3. Conclusion:

The use of dedicated entropy estimators improves the diagnostic value of EHG analysis earlier in pregnancy. The changes in the EHG might manifest early in pregnancy, providing relevant prognostic opportunities for pregnancy monitoring. The value of EHG entropy analysis can be used as a tool for early, prognostic evaluation of pregnancy

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