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# Approximate and Sample Entropy of Center of Pressure in Unperturbed Tandem Standing: Contribution of Embedding Dimension and Tolerance

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# Approximate and Sample Entropy of Center of Pressure in Unperturbed Tandem Standing:

# Contribution of Embedding Dimension and Tolerance

Jayla Wesley

A Thesis Submitted to the Graduate

# Faculty of GRAND VALLEY

# STATE UNIVERSITY

In

Partial Fulfillment of the

Requirements For the

Degree of

Master of Science in Engineering, Electrical and Computer Engineering

Padnos College of Engineering and Computing

April 2023

**Thesis Approval Form** 



The signatories of the committee below indicate that they have read and approved the thesis of Jayla Mashae Wesley in partial fulfillment of the requirements for the degree of Master of Science in Engineering, Electrical/Computer Engineering.

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Accepted and approved on behalf of the Accepted and approved on behalf of the Padnos College of Engineering and Computing Graduate Faculty

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#### **Abstract**

Approximate entropy (ApEn) and sample entropy (SampEn) are statistical methods designed to quantify the regularity or predictability of a time series. Although ApEn has been a prominent choice for use, it is currently unclear as to which method and parameter selection combination is optimal for its application in biomechanics. The goal of this thesis was to examine the difference between ApEn and SampEn related to center of pressure (COP) data during standing balance tasks, while also refining tolerance *r*, to determine entropy optimization. Six participants completed five 30-second, feet together and tandem standing, trials under eyes-open and eyesclosed conditions. Ground reaction force platform data (1200 Hz) was collected and downsampled to provide a 60 Hz COP time series. ApEn and SampEn were calculated using a constant pattern length, i.e., *m* = 2, and multiple values of *r* (tolerance). Four separate one-way analysis of variance analyses (ANOVA) were conducted for ApEn and SampEn in the anterior posterior (AP) and medial lateral (ML) directions. Dunnett's intervals were applied to the one-way ANOVA analyses to determine which conditions differed significantly. ApEn and SampEn provided comparable results in the predictability of patterns for different stability conditions, with increasing instability being associated with greater unpredictability. The selection of *r* had a relatively consistent effect on mean ApEn and SampEn values across *r =* 0.15 – 0.25\*SD, where both entropy methods tended to decrease as *r* increased. Mean SampEn values were generally lower than ApEn values. The results suggest that both ApEn and SampEn indices were equally effective in quantifying the level of center of pressure signal regularity during quiet tandem standing postural balance tests.

**Keywords:** approximate entropy, sample entropy, center of pressure, complexity, tolerance, downsampling

## **Abbreviations**

**ApEn** – approximate entropy

**COP** – center of pressure

**EC** – eyes closed

**EO** – eyes open

**FT** – feet together

**SampEn** – sample entropy

**SD** – standard deviation

**TanDB** – tandem dominant foot in back

**TanDF** – tandem dominant foot in front

#### **Introduction**

Two commonly used methods for quantifying physiological data are approximate entropy (ApEn) and sample entropy (SampEn). Entropy is defined as the loss of information in a time series or signal, i.e., it quantifies the amount of uncertainty regarding the order of an output signal<sup>10</sup>. Within the past twenty years, the use of entropy methods to define predictability or regularity in human physiological and biomechanical data has become quite prevalent<sup>47</sup>. Entropy quantifies the likelihood of the next state of a system, based on what is known about the present state of a time series, and has been used to quantify physiological changes with aging<sup>2,41</sup>, as well as cardiovascular<sup>14,40</sup> and respiratory pathology<sup>8,10,13</sup>. ApEn and SampEn are both also useful tools for understanding more about the function of changes in postural control system $6,16,21,36,310$  such as human gait mechanics<sup>50,51</sup> and standing balance<sup>5,6,7,36,47,48</sup>.

In 1991, Pincus developed approximate entropy as a mathematical instrument for measuring regularity to quantify levels of complexity within a time series<sup>31</sup>. It was meant to be a statistical measure of regularity whose foundations are similar and correctly quantifies finite data series<sup>47</sup>. ApEn was devised to quantify the rate of regularity in time data series, motivated by applications for relatively short, noisy data sets. Specifically, ApEn (and SampEn) have been shown to demonstrate changes in the complexity of various physiological signals such as chronic stroke<sup>28</sup>, electrocardiograms (ECG)<sup>4</sup>, electroencephalograms (EEG)<sup>49</sup>, heart rate variability<sup>30</sup>, and neural respiration signals $8,19$ . Complex systems such as these do not equate to being unmeasurable. Information entropy has been used to quantify complex systems where a time series with repeated patterns and less randomness will produce small entropy values whereas a time series with more randomness will equate with larger entropy values<sup>29,40</sup>. Equally, lower ApEn values reflect that a system is very persistent, repetitive, and predictive, with apparent patterns that repeat themselves throughout the series, while greater values of entropy mean independence between the data and greater randomness. So, it is more appropriate to use terms like probability, predictability, regularity, when describing the nature of a measurable complex system. In summary, the use of ApEn (and SampEn) was not meant to comprehensively analyze complex systems, but to statistically analyze the dynamics of time series related to complex systems<sup>12</sup>.

When this understanding of entropy is applied to postural sway in human stability, changes in entropy may provide insight into the control of static and dynamic balance. Cavanaugh et al.<sup>7</sup> evaluated the effect of cognitive task performance on postural control during quiet standing, revealing changes in ApEn as changes in the randomness of COP data occurred. ApEn showed the evolution of its complexity, providing meaningful comparisons to the detection of subtle influences on postural control after cerebral concussion in the alteration of the complex nature of motor control. Similarly, Ramdani et al.<sup>36</sup> analyzed the irregularity of postural sway during quiet standing comparing two sensory conditions using SampEn and concluded that the ability to successfully discriminate between levels of complexity, e.g., eyes open to eyes closed conditions, may provide insight toward characterizing the postural effects of aging and diseases. Hence, entropy provided researchers with the ability to quantify complexity within relatively short data sets based on meaningful experimental comparisons to control groups<sup>47</sup>.

Approximate entropy is calculated over a scale of time. Both ApEn and SampEn methods utilize three input parameters: *N* is the data length, *m* is the length of the window of the different vector comparisons, and *r* is the tolerance, i.e., function criterion of similarity or type of signal filter. Given the input parameters, ApEn  $(m, r, and N)^{30}$  is denoted by:

$$
ApEn(m, r, N) = -\frac{1}{N-m} \sum_{i=1}^{N-m} \log \frac{A_i}{B_i}
$$
 (1)

ApEn measures the logarithmic probability that nearby pattern runs remain close in the next incremental comparison<sup>7</sup>.  $B_i(r)$  is the probability that two sequences are similar for *m* points with self-counting and  $A_i(r)$  is probability that two sequences are similar for  $m + 1$  matches with selfcounting. Self-counting suggests that given one template, that segment in the sequence is compared to all the blocks in the sequence, including itself. For ApEn, self-counting is needed in the calculation of conditional probabilities to ensure the logarithms remain finite. Statistically, selecting *m* and *r* as input parameters would be the equivalent of dividing the space of states into cells of width *r*, to estimate the conditional probabilities of the *m*-th order7 . Higher *m* and smaller r describe details of sharper, more probabilistic parameters<sup>7</sup>. However, when dealing with stochastic processes, the analysis of conditional probabilities causes large values of *m* or minimal values of *r* to produce statistically low estimates. Ultimately, the value of the estimate depends on *m* and *r*. Pincus<sup>29</sup> suggested taking *m* as 2 and *r* as  $0.2*SD_x$  where  $SD_x$  is the standard deviation of the original data  $\langle x(n) \rangle$ , i.e.,

$$
SD_x = \sqrt{\frac{1}{N-1} \sum_{n=1}^{N} \left[ x(n) - \frac{1}{N} \sum_{n=1}^{N} x(n) \right]^2}
$$
(2)

Pincus<sup>29</sup> suggested that one of the advantages of ApEn is that the algorithm is finite for stochastic, noisy deterministic, and composite processes, i.e., models for complicated biological systems. It can differentiate between different mixed methods of deterministic and random components occurring with a different probability. ApEn is also robust to outliers because the pattern formed by wild points will rarely be repeated in the waveform<sup>32</sup>. An increasing ApEn corresponds to intuitively increasing process complexity in a biological modeling platform. However, the

limitations of ApEn include that the relative consistency is not guaranteed, and depending on the value of  $r$ , the ApEn values will change<sup>10</sup>. Additionally, the value of ApEn depends on the length of the data series. Lastly, the self-counting portion of the algorithm implies a statistical bias that is important in situations with small data sets, where when only a few or even no matches are present, the entopic result is biased toward zero $10$ .

Although the development of ApEn made a significant contribution to the understanding of complex physiological time series, its shortcomings may be significant. Richman and Moorman<sup>40</sup> introduced SampEn as an algorithm to counteract the limitations of ApEn, claiming that SampEn, as a statistical alternative, solved the self-counting problem to eliminate that bias. Eliminating self-counting is justified given that entropy is conceived as a measure of the rate of information production. ApEn uses the whole series to determine its value, needing only a that a template vector find a match of length  $m + 1$  to be defined<sup>7</sup>. SampEn contrasts with ApEn, where each template vector must find a match to be determined<sup>47</sup>. SampEn  $(m, r, and N)$  is defined as the negative value of the logarithm of the conditional probability that two similar sequences of *m* points remain identical at the next point  $m + 1$ , counting each vector over all the other vectors except on itself<sup>7</sup>.

$$
SampEn(m, r, N) = -log \frac{A^m(r)}{B^m(r)}
$$
\n(3)

 $B^m(r)$  is the probability that two sequences are similar for *m* points,  $A^m(r)$  is the probability that two sequences are similar for  $m + 1$  matches, the ratio is a conditional probability. The use of

SampEn appears to quantify regularity more effectively and eliminates many of the problems associated with ApEn40. **S**ampEn maintains the relative consistency and is also mostly independent

of the length of the series<sup>40</sup>. SampEn was created to address the bias and inconsistencies of ApEn, yet both methods retain similarities<sup>40</sup>. However, the literature is unclear about which method is preferable.

Important consideration must be given to parameter selection, as these choices may have the greatest impact on the final entropy value even in the presence of noise<sup>52</sup>. Given a time series with *N* data points, the calculation of entropy requires *a priori* determination of two unknown parameters, embedding dimension  $m$  and threshold  $r^{20}$ . Multiple pairings of parameter selections allow one to examine relative consistency where a better discrimination capacity can be accomplished. Incorrect parameter choice, and lack of due diligence in selecting *m, r*, *N,* can undermine the entropy results.

The parameter N is the length of the data series. According to Yentes et al.<sup>51</sup>, data sets larger than  $N = 200$  points are recommended over data lengths that consist of less than 200 points. It appears that both ApEn and SampEn stabilize around 2000 points. When the sampling rate is too high, i.e., frequency collection rates greater than 1000 Hz, too much redundancy likely exists within the data, which tends to artificially decrease entropy values<sup>48</sup>. Redundancy, i.e., repetitiveness of or repeating values, results in a reduced entropy and more signal regularity secondary to the counting of repeated matches. The over-redundant data problem can be solved by downsampling overly redundant time series data sets. Although downsampling removes real data, sensitivity analyses have demonstrated that the loss of data does not impact the subsequent application of the revised data set<sup>21,50</sup>. Rhea et al.<sup>38</sup>, for instance, examined the effect of downsampling on metrics that measure the magnitudes and structure of COP displacement and velocity variability. The results suggested that excessive downsampling, e.g., to 25 Hz, artificially

altered standing center of pressure displacement and velocity SampEn values. When analyzing changes in CoP variability, it is therefore essential to differentiate between those caused by the neuromotor system and those caused by data processing methods.

The parameter *m* determines the length of the sequences to be compared, and an estimation of its selection can be obtained by calculating the false closest neighbor<sup>20</sup>. Based on previous studies that include clinical applications,  $m = 1$  or  $m = 2$  appeared to produce good statistical validity for entropy calculations, where  $m = 2$  was the most popular<sup>[30](#page-79-0),52</sup>.

The third parameter, *r*, is the tolerance level for allowing similar patterns between two segments. It has been suggested that it be within 0.1-0.25 times the standard deviation (SD) for both deterministic and stochastic processes in order to be clinically useful<sup>25</sup>. These recommendations were largely based on its applications in heart rate analysis'4,19, neural processes as it relates to cognitive behaviors<sup>6</sup>, and long gait datasets<sup>50</sup>. Theoretically, with a greater *r* value, more randomized data are accepted, which produces a lower entropy value. With a smaller value of *r,* more similar data are rejected, only counting matches within a criterion thus producing a higher entropy value. Tipton's<sup>48</sup> work exposed the use of an unconventional  $r$  value. In the Tipton study, center of pressure data were analyzed at 1200 Hz, i.e., a very high sampling rate for purposes of estimating entropy. Because of the high sampling rate used by Tipton, a non-traditional method to determine *r* was used. Given the recommended range of *r*, authors<sup>4,19,20,50,51</sup> have shown that for ApEn values vary significantly even within the defined range of  $r = 0.1 - 0.25*SD$ , suggesting that additional studies are needed to accurately tighten this range.

The purpose of this master's thesis was to compare the analysis methods, ApEn and SampEn of the center of pressure data, to determine which entropy measure is less biased and most consistent. Center of pressure data were chosen because it represents a single measurement of the complex postural control mechanism used to maintain balance and incorporates all somatosensory and neuromotor inputs that influence stability<sup>48</sup>. ApEn and SampEn methods of analysis quantify these data regularity and unpredictability in its fluctuations over time and will determine an estimated baseline signal for different stability conditions. The goal was to examine distinctions in bias and consistency between ApEn and SampEn. We hypothesized that ApEn would be lower than SampEn for all calculations overall. Further, this study examined downsampling methods of the standing balance data to remove redundancy contained in high sampling rates, and to evaluate the effect of altering input tolerance *r* to determine the best *r* value for optimization.

#### **Materials and Methods**

#### *Participants*

Eight participants, between  $18 - 34$  years of age, participated in this research study after voluntarily providing their signed informed consent. Center of pressure force plate data were collected from participants but participants whose COP data contained any signal dropout were omitted. Therefore, six participants' data from the original cohort were included in the analysis. All participants were in good health and with no history of neurological or muscular disorders or injuries<sup>23</sup>. Before data collection commenced, foot dominance for each subject was determined based on the leg with which they preferred to kick a ball. This study was approved by the Grand

Valley State University Institutional Review Board (18-246-H), and data from a previous data collection were used in this study for the purpose of extending a prior analysis.

#### *Instrumentation*

Marker trajectories were captured at 120 Hz using Vicon Nexus v2.8 motion capture software (Vicon Motion System Ltd., Oxford Metrics, UK) and Vicon 16 MX and T40 cameras. Reflective markers were affixed to anatomical landmarks using two-sided hypoallergenic tape to track the movement trajectories of a modified Full-Body Plug-in-Gait model. Two floor embedded AMTI (Advanced Mechanical Technology Inc., Watertown, MA) force plates were used to synchronously collect ground reaction force data (1200 Hz). Data from six MA-411 surface preamplifiers, using the 16-channel MA300-XVI patient unit acquisition system (Motion Lab Systems Inc., Baton Rouge, LA) were synchronized with motion and ground force data to measure the electromyographic (EMG) signals of the medial gastrocnemius, soleus, and tibialis anterior bilaterally at a 1200 Hz sampling frequency. The MA300 has a fixed  $10 - 1000$  Hz (-3dB) bandwidth and uses a 500 Hz low-pass anti-aliasing filter.

Only force plate data were analyzed for this study. The force plates were oriented with one directly in front of the other (Figure 1). Center pressure data were extracted using Vicon NEXUS motion capture software v2.8 (Oxford Metrics, Oxford, UK) and exported to Excel for later analysis.

#### *Experimental Procedure*

Data for each trial were collected at 1200 Hz per second for 30 seconds until 5 successful trials were completed per stability condition (Table 1). Therefore, the time series of data collected totaled 36,000 data points. The standing postural condition of feet together with eyes open (EOFT) was defined as the most stable and hence was used as a baseline for all entropy comparisons. The subject was asked to hold this quiet standing position for thirty seconds without moving their body or stepping out of position. Balance tasks were performed barefoot with the arms positioned with the index finger pointed towards the shoulder on the same side of the body and the elbows pulled in and the knees extended. The subject then progressed through increasingly unstable balance conditions by changing visual status, with 2-minute breaks between each. Conditions included eyes open or closed, and changing foot position, i.e., feet together on force plate 5 or tandem stance using force plate 3 and 5 shown in Figure 1.



Figure 1. Force plate foot placement for feet together and feet tandem standing balance conditions, where the x-axis (anterior posterior (AP) direction) y-axis ( medial lateral (ML) direction define the center of pressure orientation. Note:  $D =$ dominant foot;  $ND =$ non dominant foot; DF = dominant foot forward; and DB = dominant foot back<sup>23,48</sup>





#### *Determining total body center of pressure from two force plates*

The extracted COP data files were analyzed using nonlinear analysis in the time and frequency domains. The tandem trial output data sets differed from the feet together trials in that two separate COP signals for tandem balance conditions were produced (one for data from each of the force plates), while feet together resulted in data from a single force plate. The two-column tandem trials were combined into one resultant COP to be directly compared to the feet together conditions using Equation 4.

$$
COP_{net} = COP_L \frac{F_{zL}}{F_{zL} + F_{zR}} + COP_R \frac{F_{zR}}{F_{zL} + F_{zR}}
$$
(4)

where  $COP<sub>L</sub>$  and  $COP<sub>R</sub>$  are the values of the COP signal from the left and right foot, respectively, and  $F_{zL}$  and  $F_{zR}$  are the vertical forces exerted on the force plates under the left and right foot, respectively<sup>41</sup>. The magnitude of the x and y axes of each subject and corresponding balance conditions were populated into curated databases. These data were then used for further estimation

of COP in the AP and ML directions using Approximate Entropy (ApEn) and Sample Entropy (SampEn).

#### *Downsampling and sensitivity analysis*

The total number of data points, *N*, required revaluation based on previous estimation by Tipton<sup>48</sup>. The original COP datasets, recorded at 1200 Hz for 30 seconds, produced 36,000 data points. Sampling data beyond 1000 Hz has been shown to lead to redundant information<sup>48</sup>. MATLAB's built-in downsample function was used to decrease the 1200 Hz sample rate by

keeping the first sample, and then every  $n<sup>th</sup>$  sample, i.e., 20, after the first. Consequently, *N* was downsampled from 36,000 data points by a factor of 20 to 1,800 data points. Simple downsampling typically exposes a system to aliasing; however, aliasing was not an issue because a fourth-order, zero-lag, low-pass Butterworth filter was applied with a cutoff frequency of 6 Hz via Nexus motion capture software v2.8 (Oxford Metrics, Oxford, UK) during preprocessing to eliminate any noise present in the signal that could not be attributed to each participant's postural control mechanism<sup>48</sup>. Simple sensitivity analyses as illustrated in Figures 2 and 3 and Figures 4 and 5 demonstrate the change in the time series before (Figs. 2 and 4) and after (Figs. 3 and 5) downsampling from a representative participant from an eyes open feet together and eyes closed tandem standing postures, respectively. Observation of the figures suggest that elimination of data points by downsampling did not impact the revised data set. These results are due to signal preprocessing and elements of eliminating redundancy, as disclosed by sensitivity analysis.



Figure 2. Representative time series for raw center of pressure (COP) data of Subject 1 eyes open, feet together (EOFT) Trial 4, where *N* = 36,000 datapoints, in the anterior posterior (AP) and medial lateral (ML) directions, respectively.



Figure 3. Representative time series for downsampled center of pressure (COP) data of Subject 1 eyes open, feet together (EOFT) Trial 4, where *N* = 1,800 datapoints, in the anterior posterior (AP) and medial lateral (ML) directions, respectively.



Figure 4. Representative time series for raw center of pressure (COP) data of Subject 1 eyes closed, feet tandem, dominant foot forward (ECTDF) Trial 29, where *N* = 36,000 datapoints, in the anterior posterior (AP) and medial lateral (ML) directions, respectively.



Figure 5. Representative time series for downsampled center of pressure (COP) data of Subject 1 eyes closed, feet tandem, dominant foot forward (ECTDF) Trial 29, where *N* = 1,800 datapoints, in the anterior posterior (AP) and medial lateral (ML) directions, respectively.

#### *Determination of Approximate and sample entropy*

Approximate (ApEn) and SampEn were determined for all five trials under each condition using custom MATLAB® (The MathWorks, Natick, MA) code. Given each COP time series, where  $N = 1800$  datapoints, a sequence of  $m = 2$  length vectors was formed. Comparisons were then made against each data segment that was 2 numbers long. Vectors were considered alike if vector components fell within a tolerance level,  $\pm r * SD^{29}$ . The similarity criteria were evaluated over a range of  $r = 0.05 - 0.3*SD$ . The total number of like vectors' logarithm sum was divided by  $N - m + 1$  to get the total number of like vectors, including a template comparison to itself<sup>51</sup>.

Looking one vector higher, *m* was raised by 1, i.e.,  $(m + 1)$ , the procedure was repeated. By deducting the conditional probabilities of  $m + 1$  from  $m$ , ApEn was calculated.

$$
ApEn(m, r, N) = -\frac{1}{N-m} \sum_{i=1}^{N-m} \log \frac{A_i}{B_i}
$$
 (5)

SampEn uses a different approach. It uses the whole series together, requiring only that a template vector find a match of length  $m + 1$  to be defined<sup>33</sup>. This contrasts with ApEn where each template vector (including itself) must find a match to be defined. So, the use of SampEn eliminates many of the problems associated with ApEn, in that it is useful to quantify regularity in a system more effectively<sup>33</sup>. The same input parameters as ApEn were used, and vectors were deemed similar if both their tail and head fell under the predetermined tolerance level. The sum of the total number of like vectors for *m* points was divided by  $N - m + 1$  and defined as  $B^m(r)$ . Further, SampEn defined  $A^m(r)$  as the subset of  $B^m(r)$  that two sequences are similar for  $m +$  $1^{51}$ . SampEn is then calculated as the conditional probability  $-\ln(A^m(r)/B^m(r))$ .

$$
SampEn(m, r, N) = -log \frac{A^m(r)}{B^m(r)}
$$
\n(6)

#### *Statistical Analysis*

Trials and stability conditions were independent of one another due to the breaks given between each of them<sup>23</sup>. The subject's ability to rest and then reset allows for this assumption. Four separate one-way analysis of variance (ANOVA) tests were performed for ApEn and SampEn in each of the AP and ML axial directions. The ANOVA testing revealed that each one-way ANOVA assessed for a difference of means between the baseline and increasingly difficult stability conditions within each subject. Additionally, the one-way ANOVA tests were run with Dunnett's

intervals to determine which conditions varied significantly. This test compared the baseline EOFT condition to each increasingly less stable condition.

#### **Results**

### *Comparison of Approximate and Sample Entropy Time Series*

Approximate and sample entropy were determined for all trials and conditions, yet it was important to first examine the ApEn and SampEn values over the 30-second time series for  $m = 2$ ,  $r = 0.02*SD$  for the purpose of establishing the fidelity of the data. Note that since  $N = 1800$  data points each second of COP data consisted of 60 points. Figures 6 and 7 illustrate these data for one trial of an eyes open, feet together condition and one trial of an eyes closed feet tandem condition for one representative participant. Visual inspection of each plot suggest that the ApEn and SampEn magnitude of the values were comparable to each other and the spikes over the 30-second time series appeared similar. Having established this for a representative participant and trial it seemed appropriate to determine the mean ApEn and SampEn values for the purpose of further statistical analysis.



Figure 6. Representative time series for ApEn and SampEn calculated over thirty 60-point segments of Subject 1 eyes open, feet together (EOFT) Trial 03 COP data; where  $N = 1,800$ datapoints,  $m = 2$ , and  $r = 0.02$ <sup>\*</sup>SD; in the anterior posterior (AP) and medial lateral (ML) directions, respectively.



Figure 7. Representative time series for ApEn and SampEn calculated over thirty 60-point segments of Subject 1 eyes closed, feet tandem, dominant foot back (ECTDB) Trial 19 COP data; where  $N = 1,800$  datapoints,  $m = 2$ , and  $r = 0.02$ <sup>\*</sup>SD; in the anterior posterior (AP) and medial lateral (ML) directions, respectively.

#### *Examination of ApEn and SampEn under different postural conditions*

Mean ApEn and SampEn values were obtained for all trials and conditions. Since one purpose of this study was to compare differences in the analysis methods in conjunction with the differing *r* parameter chosen, compilations of one-way ANOVAs with Dunnett's Test outputs for each condition were compared to baseline. These 144 analyses were condensed into summary graphics where Figures  $8 - 12$  depict subjects by column Average ApEn and SampEn values, charted in orange and blue respectively, are plotted in either row, where groups were separated by

data in the medial lateral (ML) and anterior posterior (AP) direction, based on if the entropy value was significantly different from baseline. The upper "No" row demonstrate that differences in entropy, with respect to the confidence interval, were not significantly different from baseline. The lower "Yes" row demonstrates that difference in entropy values were significantly different from baseline. The stature of 'significance' is based on an equivalent of 720 tests at  $\alpha = 0.05$ , where on average 36 "Yeses" are to be expected even if there are no statistically significant differences between the data. Notably there are a lot more than 36 significant instances found.



Figure 8. One-way ANOVA with Dunnett's Tests Outputs summary graphic, comparing the difference of eyes close, feet together (ECFT) trials of mean ApEn(orange) and SampEn(blue) values to baseline condition eyes open, feet together (EOFT), in the medial lateral (ML) and anterior posterior (AP) directions; where  $N = 1,800$  datapoints,  $m = 2$ , and  $r$  ranges from 0.05-0.3\*SD, for all subjects.



Figure 9. One-way ANOVA with Dunnett's Tests Outputs summary graphic, comparing the difference of eyes open, feet tandem, dominant foot forward (EOTDF) trials of mean ApEn(orange) and SampEn(blue) values to baseline condition eyes open, feet together (EOFT), in the medial lateral (ML) and anterior posterior (AP) directions; where  $N = 1,800$  datapoints, *m*  $= 2$ , and *r* ranges from 0.05-0.3\*SD, for all subjects.



Figure 10. One-way ANOVA with Dunnett's Tests Outputs summary graphic, comparing the difference of eyes closed, feet tandem, dominant foot forward (ECTDF) trials of mean ApEn(orange) and SampEn(blue) values to baseline condition eyes open, feet together (EOFT), in the medial lateral (ML) and anterior posterior (AP) directions; where  $N = 1,800$  datapoints, *m*  $= 2$ , and *r* ranges from 0.05-0.3\*SD, for all subjects.



Figure 11. One-way ANOVA with Dunnett's Tests Outputs summary graphic, comparing the difference of eyes open, feet tandem, dominant foot back (EOTDB) trials of mean ApEn(orange) and SampEn(blue) values to baseline condition eyes open, feet together (EOFT), in the medial lateral (ML) and anterior posterior (AP) directions; where  $N = 1,800$  datapoints,  $m = 2$ , and  $r$ ranges from 0.05-0.3\*SD, for all subjects.



Figure 12. One-way ANOVA with Dunnett's Tests Outputs summary graphic, comparing the difference of eyes closed, feet tandem, dominant foot back (ECTDB) trials of mean ApEn(orange) and SampEn(blue) values to baseline condition eyes open, feet together (EOFT), in the medial lateral (ML) and anterior posterior (AP) directions; where  $N = 1,800$  datapoints, *m*  $= 2$ , and *r* ranges from 0.05-0.3\*SD, for all subjects.

Assessment of each plot compares the "Yes" rows to the "No" rows in evaluating significance in the differences from baseline. Meaningful deviations from baseline conditions were found in the ML direction in cases of ECFT, EOTDB, ECTDB, EOTDF, and ECTDF. Considerable deviations were also found in the AP direction in cases of EOTDB, ECTDB, EOTDF, and ECTDF. The ECFT trials compared to baseline in the AP direction display majority insignificant results, implying there exists no tangible evidence that conditions were substantially different from baseline. The EOTanDB trails in both ML and AP directions, notably had the highest difference in entropy, and greatest amount of significances difference from baseline across all values of *r*. The consistent number of "Yeses" in Tandem trials suggest differences from baseline can be quantified for tandem standing. Overall, all trial mean entropy data assessed and

output into an ANOVA summary graphic, show that the significance of the difference from baseline is the equivalent, using either ApEn or SampEn approximation. Having determined that the significance of using either entropy calculation is equivalent when assessing the difference from baseline for all tandem trial data, it made sense to examine mean ApEn and SampEn entropy across varying *r* values.

#### *Examination of the changes in tolerance level, r, on ApEn and SampEn; bias and consistency*

Selection of parameters is the most important factor when evaluating entropy, and given *r* was determined using trial and error until reasonable values of ApEn were found in a previous study<sup>48</sup>, it was important to examine the effect of the *r* parameter for each entropy method. Figure 13 displays mean ApEn and SampEn values for all stability conditions, where each column separated by participant numbers, rows depicted by each ML and AP axial direction, and ranged over values of  $r = 0.05 - 0.3*SD$ . Figure 14 illustrates participants by method are separated into each column, grouping all subjects. See Appendix B. ANOVA Data Tables for participants by method and conditions by method mean entropy values.



Figure 13. Summary plot of all mean ApEn (orange) and SampEn (blue) values of all trials of stability conditions, separated by participant, in the medial lateral (ML) and anterior posterior (AP) directions; where  $N = 1,800$  datapoints,  $m = 2$ , across  $r$  range from 0.05-0.3\*SD.



Figure 14. Summary plot of all mean ApEn (orange) and SampEn (blue) values of all participants, separated by trials of stability conditions, in the medial lateral (ML) and anterior posterior (AP) directions; where  $N = 1,800$  datapoints,  $m = 2$ , across  $r$  range from 0.05-0.3\*SD.

Visual examinations of each plot indicate that ApEn and SampEn tended to decrease as *r* increased, with relative directionality evident between  $r = 0.15 - 0.25$  SD. Outliers that distorted this line of regression were most prominent in the range  $r = 0.05 - 0.1$  *\*SD*. Participants 4 and 5 revealed notable outliers in the AP directions within this range as calculated by the SampEn. Coincidingly, less rejected matches revealed themselves in the lowest entropy values, where *r = 0.3\*SD*. Despite comparing mean entropy values across all values of *r,* it does not appear that one method, i.e., ApEn and SampEn, is superior.

#### *Comparison of the magnitude of ApEn and SampEn*

Though the significance of the differences from baseline are consistent using either entropy calculation, differences in ApEn and SampEn values were apparent. This gave the purpose to directly compare outputs by taking the differences between the data derived from the ApEn and SampEn calculations in both medial lateral and anterior posterior directions. This comparison is shown in Figure 15 as a histogram of the SampEn and ApEn relationship of all the data, where the difference in value of the two calcuations are distributed across the x-axis, and the total number of calculations for all trials and conditions conducted are displayed on the y-axis.



Figure 15. SampEn versus ApEn histogram of each COP data timeseries of all subjects and stability conditions, including both the medial lateral (ML) and anterior posterior (AP) directions; where  $N = 1,800$  datapoints,  $m = 2$ , across  $r$  range from 0.05-0.3\*SD.

Evidence of a right skewed histogram indicates that the sample-approximate relationship is primarily negative, i.e., SampEn on average is smaller than ApEn. Given the bias in the ApEn calculation, ApEn estimates were expected to converge closer to 0; therefore, this result contradicted initial hypotheses. The sample-approximate relationship is anticipated to be positive based on previous research; however, these results are unique to this study's data set and should be reported.

#### **Discussion**

Understanding how a healthy brain responds to different stability conditions and utilizing the best method to measure those conditions is important when accurately comparing how a healthy brain responds versus how an injured brain responds to the same conditions. Previous research that has examined children and adults with mild traumatic brain injury, e.g., concussion, have

suggested that the use of the center of pressure (COP) data can be useful in delineating a normal from an abnormal response to the perturbations of static and dynamic balance<sup>5,6,35,36,37,46</sup>. Since it has been shown that the COP time series is non-linear, traditional methods of assessing various COP parameters, e.g., statistical use of means and standard deviation, have not been effective<sup>3</sup>. Previous work using non-linear methods, such as approximate and sample entropy, to study normal and pathological balance has been useful<sup>1,2,26,42</sup>. However, a consensus on the best methodological use of ApEn and SampEn has not yet been established. For example, previous work<sup>48</sup> in our laboratory used ApEn to examine healthy college-aged participants to describe changes in ApEn under more and less stable standing positions. Based on the need described in the literature for more work using ApEn and SampEn, and the methodological limitations of the previous research in our laboratory, the purpose of this study was to compare ApEn and SampEn under various stability conditions, and when altering tolerance (i.e., *r*) values. The results revealed that even though SampEn tend to yield lower mean values than ApEn, both indices were equally effective in quantifying the regularity of a COP signal given that the significance of using either entropy calculation is equivalent when assessing the difference from baseline. Further, the selection of *r* had a relatively consistent effect with both entropic statistical analyses, as the similarity criterion was increasingly tightened, more similarities were rejected, and therefore larger entropy values.

Tipton's<sup>48</sup> study was the most similar to ours in terms of the methodologies used to investigate conditions. Downsampling the COP data in this study was necessary because previously, entropy values were an artifact of quantifying an oversampled COP signal with an *r* value that was found through trial and error. Yentes et al.<sup>51</sup> recommend, as best practice, when using ApEn or SampEn for the analysis of human gait data that *N* not exceed sampling data beyond 1000 Hz because it's

over this frequency that may lead to redundant information. In the current study, previous entropy estimates at 1200 Hz yielded low entropy values between 0.005 and 0.030, where  $N = 36,000$ consisted of many closely spaced data points with identical values. In a different study, Lubetsky et al.<sup>21</sup> conducted sample entropy on COP data for prolonged standing tasks on normal and compliant surfaces. The study found that while down-sampling had increased SampEn values, it had an insignificant effect on the comparisons to the original datasets; however, if such procedures are performed, they should be well justified. In the present study, the waveforms were virtually identical when comparing the raw COP trial data to the downsampled trial data. Downsampling was necessary for the processing of our entropy results. Utilizing 1,800-point arrays, the entropy values in this study ranged from 0.08 to 0.90. However, it is suggested that it is essential for researchers to evaluate the influence that downsampling has on their particular dataset.

Incorrect parameter selection of the vector length, *m*, threshold, *r*, or data length, *N*, can undermine the ApEn and SampEn discrimination capacity<sup>9</sup>. The embedding dimension,  $m = 2$ , and dataset length, *N,* were fixed input parameters in this study. Many approaches of calculating *r* have been suggested, including utilizing the standard deviation (SD) of the whole time series<sup>26,30</sup>, the standard error of the entropy values<sup>18</sup>, predefined tolerance levels<sup>15,45</sup> and using a heuristic stochastic model<sup>9</sup>. Typically, it is suggested that for clinical data, *r* is to be set 0.02\*SD when utilizing an entropy algorithm. In Tipton's study (which use the same data set as the present study), only one value of *r* value was used. Therefore, one of the focusses of this study was placed on the filter parameter, r, to determine which or which are optimal for the objectives of this study. Center of pressure data sets demonstrated significant differences between the range of *r* values, *r =* 0.05,  $r = 0.1$ ,  $r = 0.15$ ,  $r = 0.2$ ,  $r = 0.25$ , and  $r = 0.3$  times standard deviation. Overall, the variations in

entropy values by ApEn and SampEn calculations as  $r$  changed imply that the usual choice of  $r =$ 0.2\*SD allows for contemplation. The stability of a metric is referred to as relative consistency. As r increased, ApEn and SampEn values decreased, where both calculations exhibited relative directional consistency; therefore, *r* had the same effect on both entropy calculations. In a different study Yentes et al.<sup>50</sup>, found that for walking trials of gait data SampEn decreased as *r* increased, which is consistent with our findings. However, for ApEn, different *r* values resulted in some variability<sup>50</sup>. For the current study, since ApEn showed relative consistency in a similar manner as SampEn, it is suggested that *r* be selected based on the individual study's criteria for parameters, and then examine the relative consistency across  $r = 0.15, 0.2$ , and 0.25 times the standard deviation.

Outliers and spikes were apparent in SampEn values for  $r = 0.05$  and  $r = 0.1*SD$  due to the overly stringent conditions. In a study that evaluated the impact of abnormal spikes on the interpretation of entropy results in the context of biosignal analysis, Molina-Picó et al.<sup>25</sup> suggested removing these results, as they can misrepresent the signal regularity. Therefore, it is illogical to assume that the anomalies exhibited by SampEn represent data that ApEn was incapable of detecting in any way. The obvious deviation of entropy values from the overall trend line of the data, within the range of  $r = 0.05$  and  $r = 0.1$ \*SD, suggests that such small entropy values should not be used, removing  $r = 0.05$  and  $r = 0.1$  SD from the list of recommendations for use.

The performance of ApEn and SampEn calculations of standing balance condition-based time series were able to differentiate between circumstances that were different from the baseline EOFT condition, in that the tandem stance trials indicated higher entropy values. ECFT was not consistent enough in indicating entropy values that were significantly higher than the baseline
entropy values in this study. In a similar study assessing balance by altering visual conditions, Ramdani<sup>36</sup> used SampEn to analyze human postural sway and was able to distinguish between eyes open and eyes closed conditions of participants standing on a single force plate. Importantly, in conditions with eyes closed, SampEn was lower than in conditions with eyes open. Some concur with these findings<sup>11</sup>, while others find them contradictory<sup>43</sup>. Given the equal significance in distinguishing between eyes closed, feet together (ECFT) and eyes open, feet together (EOFT) in our present study, it is suggested that ECFT condition not be a method primarily used to differentiate ECFT from baseline conditions.

All the participants showed a consistent significant difference from baseline under tandem conditions in both AP and ML directions. As expected, the tandem was the least stable condition, under both eyes open and eyes closed conditions. With tandem standing, placement of the dominant foot in the back or front did not seem to affect the entropy values. In contrast to other participants for tandem standing, participants 4 and 5 repeatedly exhibited insignificances when comparing ECTDF, EOTDB, and ECTDB trials to baseline in the one-way ANOVA with Dunnett's Tests Outputs summary graphics; however, previously discussed outliers may contribute to this. Though clear significances are found for differentiating from baseline in both ApEn and SampEn, which entropic measure is "better"? Few studies have compared ApEn and SampEn for standing balance data by altering only the tolerance parameter. However, Yentes<sup>50</sup> investigated postural control of walking trials and evaluated various combinations of *N, m*, and *r* and concluded that SampEn appeared to be more trustworthy for brief data sets and demonstrated fewer difficulties with relative consistency. As opposed to Yentes et al.'s<sup>50</sup>findings that generally mean ApEn values were lower than mean SampEn values in the analysis of gait data, in the present study it was found that mean SampEn values were lower. Given Our findings in that ApEn and SampEn are almost equally significant to distinguish from baseline for standing balance data, additional research needs to re-examine differences between ApEn and SampEn across similar values of *m* and *r*.

There are several limitations to this study beyond its use of a small sample of healthy participants. First, differentiating comparisons between ApEn and SampEn might be feasible utilizing alternative statistical practices. Future studies may propose the use of significantly more sophisticated statistical methods that permit testing across subjects concerning ANOVA analysis. Researchers may also consider comparing the time frequency analysis of the anterior posterior and medial lateral directions of ApEn and SampEn when considering directional differences. Secondly, additional methodologies exist for determining an optimal *r*. In addition to a few methodological propositions of  $r$ , such as using the standard error of the entropy values<sup>18</sup> and employing fixed tolerance values<sup>15, 45</sup>, I would propose comparing ApEn and SampEn using a method proposed by Chon et  $a^{\dagger}$ . In this study, the authors implement a heuristic stochastic model employing equations that autonomously determine the ApEn<sub>max</sub> value, such that the accuracy of the entropic output is unaffected by the different data lengths. Lastly, the outcomes of this study may be influenced by the resolution of the selected downsampling technique. Typically, downsampling by selecting every nth point, rather than averaging across each window, introduces aliasing into the system. However, a low-pass Butterworth filter was implemented during preprocessing to mitigate this effect. Though the resulting COP waveforms of the unprocessed and downsampled data appeared to be identical in this study, the results should be compared to other sampling techniques, such as

the use of the MATLAB decimate function rather than downsample. Or evaluate the effect of obtaining a windowed average by downsampling an average of 20, 30, 40, etc. points at a time.

This study's objective was to investigate the impact of varying the input parameter *r*, the similarity criterion, required for the calculations of both ApEn and SampEn, in order to determine the optimal choice. Choosing a suitable input *r* value is necessary to ensure relative consistency. The selection of *r* had a relatively consistent effect on both statistical analyses, as the similarity criterion was increasingly tightened, more similarities were rejected, resulting in larger entropy values. For optimal results, values between  $r = 0.15$  and  $0.25$ <sup>\*</sup>SD are deemed optimal. It is these ranges of *r* values that the entropic data is less susceptible to outliers while maintaining a critical threshold in lieu of retaining low experimental error. Further, the purpose of this master's thesis was to compare the analysis methods, ApEn and SampEn of center of pressure balance data, to determine which entropy measure is most consistent. ApEn and SampEn analysis methods accurately quantify the varying stability conditions, i.e., both methods could accurately decipher between the stability conditions appropriately. It can be concluded that both systems are highly predictable over time and share equivalent significance in terms of their ability to differentiate from baseline. Consequently, one estimation technique was not particularly "better" than the other. Since both methods of statistical data analysis produce the same overall pattern of results, it can be concluded that both indices were equally effective in quantifying the level of center of pressure signal regularity during quiet tandem standing postural balance tests and it is thereby suitable to use either methodology. However, it was discovered that mean SampEn values were typically lesser than ApEn values. In order to substantially advance the determination of the optimal technique for clinically diagnosing concussions, additional research will be required to demonstrate what this really implies for our data.

### **Appendix A. ANOVA with Dunnett's Test Figures**

Supplementary data associated with this master's thesis can be found below. Participant 1 serves as the representative for all subjects' one-way ANOVA statistics with Dunnett's tests in Appendix A. The data from graphics such as those in Appendix A were compiled into summary graphics for methods section, Figures  $8 - 12$ .



Figure A.1. One-way ANOVA, mapping the 5 mean ApEn trial values in the anterior posterior (AP) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where  $N = 1,800$ datapoints,  $m = 2$ , and  $r = 0.15*SD$ .



Figure A.2. One-way ANOVA, mapping the 5 mean ApEn trial values in the medial lateral (ML) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where *N* = 1,800 datapoints,  $m = 2$ , and  $r = 0.15*SD$ .



Figure A.3. One-way ANOVA, mapping the 5 mean ApEn trial values in the anterior posterior (AP) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where  $N = 1,800$  datapoints,  $m = 2$ , and  $r = 0.2$ <sup>\*</sup>SD.



Figure A.4. One-way ANOVA, mapping the 5 mean ApEn trial values in the medial lateral (ML) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where *N* = 1,800 datapoints,  $m = 2$ , and  $r = 0.2$ <sup>\*</sup>SD.



Figure A.5. One-way ANOVA, mapping the 5 mean ApEn trial values in the anterior posterior (AP) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where  $N = 1,800$  datapoints,  $m = 2$ , and  $r = 0.25$ <sup>\*</sup>SD.



Figure A.6. One-way ANOVA, mapping the 5 mean ApEn trial values in the medial lateral (ML) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where  $N = 1,800$ datapoints,  $m = 2$ , and  $r = 0.25$ <sup>\*</sup>SD.

*T*



Figure A.7. One-way ANOVA, mapping the 5 mean SampEn trial values in the anterior posterior (AP) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where  $N = 1,800$ datapoints,  $m = 2$ , and  $r = 0.15*SD$ .



Figure A.8. One-way ANOVA, mapping the 5 mean SampEn trial values in the medial lateral (ML) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where  $N = 1,800$  datapoints,  $m = 2$ , and  $r = 0.15$ <sup>\*</sup>SD.



Figure A.9. One-way ANOVA, mapping the 5 mean SampEn trial values in the anterior posterior (AP) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where  $N = 1,800$  datapoints,  $m = 2$ , and  $r = 0.2$ <sup>\*</sup>SD.



Figure A.10. One-way ANOVA, mapping the 5 mean SampEn trial values in the medial lateral (ML) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where  $N = 1,800$  datapoints,  $m = 2$ , and  $r = 0.2$ <sup>\*</sup>SD.



Figure A.11. One-way ANOVA, mapping the 5 mean SampEn trial values in the anterior posterior (AP) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where  $N = 1,800$  datapoints,  $m = 2$ , and  $r = 0.25$ <sup>\*</sup>SD.



Figure A.12. One-way ANOVA, mapping the 5 mean SampEn trial values in the medial lateral (ML) direction for each stability condition; with Dunnett's tests, comparing eyes closed, feet together (ECFT); eyes open, feet tandem, dominant foot back (EOTDB); eyes closed, feet tandem, dominant foot back (ECTDB); eyes open, feet tandem, dominant foot forward (EOTDF); and eyes closed, feet tandem, dominant foot forward (ECTDF); to baseline (EOFT); where  $N = 1,800$  datapoints,  $m = 2$ , and  $r = 0.25$ <sup>\*</sup>SD.

# **Appendix B. ANOVA Data Tables**

Supplementary data associated with this master's thesis can be found below. Table B.1 and Table B.2 depict the associated ApEn and SampEn values plotted in Figures 13 and 14. Each mean entropy value is a result of a compilation of each of the columns, i.e., value of *r*, axis direction, participant, stability condition, and entropy estimation method.

	r	<b>Axis</b>	Subject	<b>Method</b>	mean	std
$\boldsymbol{l}$	0.05	AP		Approximate	0.552923336033609	0.0488963051243597
$\overline{\mathbf{2}}$	0.05	AP	$\mathbf{1}$	Sample	0.513363004523101	0.0754283729720997
$\overline{\mathbf{3}}$	0.05	AP	$\overline{2}$	Approximate	0.573654693622424	0.0271048578781064
$\overline{\boldsymbol{4}}$	0.05	AP	$\overline{2}$	Sample	0.555312510534061	0.0462103755086795
$\overline{5}$	0.05	AP	$\overline{4}$	Approximate	0.773897907699981	0.183724893881889
6	0.05	AP	$\overline{4}$	Sample	0.789084817867085	0.202248702651556
$\overline{7}$	0.05	AP	5	Approximate	<b>NA</b>	<b>NA</b>
8	0.05	AP	5	Sample	<b>NA</b>	<b>NA</b>
9	0.05	AP	6	Approximate	0.543356457836966	0.0490237753617992
10	0.05	AP	6	Sample	0.496973395612704	0.0769347412887044
11	0.05	AP	8	Approximate	0.574055805788667	0.0383543544455246
12	0.05	AP	8	Sample	0.55617824887009	0.0681921869237874
13	0.05	ML	$\mathbf{1}$	Approximate	0.532937654547443	0.153309748577886
14	0.05	ML	$\mathbf{1}$	Sample	0.513974937202097	0.194325970349666
15	0.05	ML	$\overline{c}$	Approximate	0.594209164205528	0.0864580077985201
16	0.05	ML	$\overline{2}$	Sample	0.598869208020955	0.14416650843452
17	0.05	ML	$\overline{4}$	Approximate	0.914593728118234	0.208024389784852
18	0.05	ML	$\overline{4}$	Sample	1.05959078000325	0.334259652183477
19	0.05	ML	5	Approximate	<b>NA</b>	<b>NA</b>
20	0.05	ML	5	Sample	<b>NA</b>	<b>NA</b>
21	0.05	ML	6	Approximate	0.573649856971166	0.0648231736059189
22	0.05	ML	6	Sample	0.552868221749671	0.103898283656662
23	0.05	ML	8	Approximate	0.619926536346686	0.0824860324099032
24	0.05	ML	8	Sample	0.625694031364102	0.135668768455753
25	0.1	AP	$\mathbf{1}$	Approximate	0.448215894627326	0.0969965564832068
26	0.1	AP	$\mathbf{1}$	Sample	0.375800257292317	0.0928149762734519
27	0.1	AP	$\overline{c}$	Approximate	0.488664106635178	0.0598657503255546
28	0.1	AP	$\overline{2}$	Sample	0.415247057735097	0.0603946849773664

*Table B.1. All Subjects by Axis and Method Mean and Standard Deviation Data* 





115	0.25	ML	5	Approximate	<b>NA</b>	<b>NA</b>
116	0.25	ML	5	Sample	<b>NA</b>	<b>NA</b>
117	0.25	ML	6	Approximate	0.207186730634017	0.0980794582018656
118	0.25	ML	6	Sample	0.176988769866324	0.0782941305485083
119	0.25	ML	8	Approximate	0.244696727592634	0.124487145447648
120	0.25	ML	8	Sample	0.207408219078263	0.101614949747247
121	0.3	AP	$\mathbf{1}$	Approximate	0.147413586034671	0.0518672659890784
122	0.3	AP	$\mathbf{1}$	Sample	0.131962574805219	0.0459571907503602
123	0.3	AP	$\overline{2}$	Approximate	0.16388583641981	0.0390441019200517
124	0.3	AP	$\overline{2}$	Sample	0.146801678879389	0.0337274095388648
125	0.3	AP	$\overline{4}$	Approximate	0.194206631964223	0.0476272709196575
126	0.3	AP	$\overline{4}$	Sample	0.180739110430945	0.0481953037370043
127	0.3	AP	5	Approximate	<b>NA</b>	<b>NA</b>
128	0.3	AP	$\overline{5}$	Sample	<b>NA</b>	<b>NA</b>
129	0.3	AP	6	Approximate	0.134879717826831	0.0416876733790984
130	0.3	AP	6	Sample	0.12168185097051	0.0374015752515928
131	0.3	AP	8	Approximate	0.164974361215884	0.0477887043158255
132	0.3	AP	8	Sample	0.148274547353833	0.0438802271151863
133	0.3	ML	1	Approximate	0.156971276857867	0.110095250592128
134	0.3	ML	$\mathbf{1}$	Sample	0.133732045614251	0.092158732785613
135	0.3	ML	$\overline{2}$	Approximate	0.216016175947946	0.113356088509414
136	0.3	ML	$\overline{2}$	Sample	0.17941797306955	0.091253195428536
137	0.3	ML	$\overline{4}$	Approximate	0.328574546195869	0.129810217731111
138	0.3	ML	$\overline{4}$	Sample	0.278120656533204	0.10725972063412
139	0.3	ML	5	Approximate	<b>NA</b>	<b>NA</b>
140	0.3	ML	$\overline{5}$	Sample	<b>NA</b>	<b>NA</b>
141	0.3	ML	6	Approximate	0.16591360343697	0.0799861047818259
142	0.3	ML	6	Sample	0.143981036209407	0.0646059827742168
143	0.3	ML	8	Approximate	0.199078641904989	0.107871255326046
144	0.3	ML	8	Sample	0.169858737868292	0.0878226919358317

*Table B.2. Conditions by Axis and Method Mean and Standard Deviation Data*











## **Appendix C. Code**

#### C.1 *Downsampling COP Data*

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
% Title: Downsample_COP.m
% Author: Jayla Wesley
% Notes: this script file loads subject data into a usable structure for 
% further analysis
   1. Cx is the Anterior-Posterior direction
% 2. Cy is the Medial Lateral direction<br>% 3. Downsampling by a factor of 20, 12
   3. Downsampling by a factor of 20, 1200Hz to 60Hz
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
clc; close all; clear;
% load trial condition
filtdata = readmatrix('C:\Users\jmwesley\OneDrive - Newkirk Electric Associates, 
Inc\Documents\GVSU Masters 22\EGR 695 Masters Thesis\Subject 01\Subject 01 Trial Data 
Raw\SB01 Trial04.csv');
% separate COP data into axial vectors
Cx filt = filtdata(6:36005,18);Cy filt = filtdata(6:36005,19);
Cz filt = filtdata(6:36005,20);
t1 = 1inspace(1,30,36000);
% downsample by factor of 20
Cx filtds = downsample(Cx filt, 20);
Cy filtds = downsample(Cy filt, 20);
Cz filtds = downsample(Cz filt, 20);
t2 = 1inspace(1,30,1800);
% plot generation
figure(1)
\texttt{subplot}(1,2,1), \texttt{plot}(t1,Cx_filt)xlabel('Time(s)','FontSize', 12), ylabel('AP Distance from Origin (mm)','FontSize', 
12);
subplot(1,2,2), plot(t1, Cy filt)
xlabel('Time(s)','FontSize', 12), ylabel('ML Distance from Origin 
(mm)','FontSize',12);
figure(2)
subplot(1,2,1), plot(t2,Cx filtds)
xlabel('Time(s)','FontSize', 12), ylabel('AP Distance from Origin (mm)','FontSize', 
12);
subplot(1,2,2), plot(t2, Cy filtds)
xlabel('Time(s)','FontSize', 12), ylabel('ML Distance from Origin 
(mm)','FontSize',12);
```
## C.2 *Approximate Entropy Function*

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
function [AE] = ApEntr( data, dim, r )
% Title: ApEntr(data,m,R): returns the approximate entropy value, function
% Adapted from Jenna Yentes [44].
% inputs - data, single column (transpose COP) time series
```

```
% - m, length of vectors to be compared<br>% - R. filter for accepting matches (as
% - R, filter for accepting matches (as a proportion of the
           standard deviation)
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
% variable declaration
r = r*std(data); % tolerance, r, times the standard deviation
N = length(data); % data length
phim = zeros(1,2); \frac{1}{2} phi m and phi m + 1
% ApEn calculation
for j = 1:2<br>m = dim+j-1;
                                         % define vector length
    phi = zeros(1, N-m+1);dataMat = zeros(m, N-m+1);for i = 1:mdataMat(i,:) = data(i:N-m+i); % divide data series into blocks
    end<br>for i = 1:N-m+1% calculate the conditional probability of each
vector
        tempMat = abs(dataMat - repmat(dataMat(:,i),1, N-m+1));
        AorB = any( (tempMat > r), 1); % count m and m + 1 matches within tolerance
r*SD
        phi(i) = sum(\alphaAorB)/(N-m+1); % sum of natural log for ea cond. proab m and m
+ 1
     end
    phim(j) = sum(log(phi))/(N-m+1); % sum of natural log for ea cond. prob divide by
N - m + 1 and N-mend
AE = \text{phim}(1) - \text{phim}(2); % \text{phi } m - \text{phi } m + 1End
```
## C.3 *Sample Entropy Function*

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
function [SE] = SampEntr(data, m, R)
% Title: SampEntr(data,m,R): returns the sample entropy value, function
% Adapted from Jenna Yentes [44].
% inputs: - data, single column (transpose COP) time series
% - m, length of vectors to be compared<br>% - R. filter for accepting matches (as
          - R, filter for accepting matches (as a proportion of the
% standard deviation)
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
% variable declaration
r = R * std(data); % tolerance, r, times the standard deviation
N = length(data); % data length
dij=zeros(N-m,m+1);
dj=zeros(N-m,1);
dj1=zeros(N-m,1);Bm=zeros(N-m,1);
Am=zeros(N-m,1);
% SampEn calculation
for i = 1:N-mfor k = 1:m+1dij(:,k) = abs(data(1+k-1:N-m+k-1)-data(i+k-1)); end
    dj = max(dij(:,1:m),[], 2); % divide series into m blocks
```

```
dj1 = max(di;[],2); % divide series into m + 1 blocks<br>d = find(dj<=r); % count m matches within toleranc
      d = find(dj <= r);<br>
d = \frac{1}{\sqrt{3}} = find(dj 1 <= r);<br>
\frac{1}{\sqrt{3}} = find(dj 1 <= r);<br>
\frac{1}{\sqrt{3}} = count m + 1 matches within tolerance
                                              \frac{1}{2} count m + 1 matches within tolerance r*SD
     nm = length(d)-1; \frac{1}{1} % subtract the self-match<br>Bm(i) = nm/(N-m); \frac{1}{1} % number of similar vector
      Bm(i) = nm/(N-m); % number of similar vector for m points<br>
nm1 = length(d1)-1; % subtract the self-match
     nm1 = length(d1)-1; \frac{1}{1} \frac{1}{1} \frac{1}{2} subtract the self-match<br>Am(i) = nm1/(N-m); \frac{1}{2} s number of similar vector
                                               % number of similar vector for m + 1 matches
end
Bmr = sum(Bm)/(N-m); % sum of natural log for cond. prob two sequences
                                 % are similar for m points divided by N - m
\text{Amr} = \text{sum(Am)/(N-m)}; \text{\$ sum of natural log for cond. prob two sequences} % are similar for m + 1 matched divided by N - m
SE = -log(Amr/Bmr); % negative natural log of A/Bend
```
### C.4 *ApEn and SampEn over Time Series*

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
% Title: ApEn SampEn TimeSeries.m
% Author Jayla Wesley
% Notes: this script file loads subject data into a usable structure for 
% further analysis
  1. Cx is the Anterior-Posterior direction
 2. Cy is the Medial Lateral Direction
% 3. Downsampling by a factor of 20, 1200Hz to 60Hz
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
close all; clear all; clc;
filtdata = readmatrix('C:\Users\jmwesley\OneDrive - Newkirk Electric Associates, 
Inc\Documents\GVSU Masters 22\EGR 695 Masters Thesis\Subject 01\Subject 01 Trial Data 
Raw\SB01 Trial19.csv');
Cx filt = filtdata(6:36005,27);Cy filt = filtdata(6:36005,28);
Cz filt = filtdata(6:36005,29);
% downsample the data 1200Hz to 60Hz
Cx filtds = downsample(Cx filt, 20);
Cy filtds = downsample(Cy filt, 20);
Cz filtds = downsample(Cz filt, 20);
%% Center of Pressure (COP) Calculation %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
% Get COP data by window - choose filtered because overtime difference is 
insignificant.
% 1 second %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x1 = Cx filtds(1:60); \frac{1}{2} second of data 1-60
y1 = Cy filtds(1:60);
z1 = Cz filtds(1:60);
% 2 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x2 = Cx filtds(61:120); \frac{1}{2} second of data 61-120
y2 = Cy filtds(61:120);
z2 = Cz filtds(61:120);
% 3 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
```

```
x3 = Cx filtds(121:180); % 1 second of data 121-180
y3 = Cy_{tiltds}(121:180);z3 = Cz filtds(121:180);
% 4 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x4 = Cx filtds(181:240); % 1 second of data 180-240
y4 = Cy_{\text{filds}}(181:240);z4 = Cz filtds (181:240);
% 5 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x5 = Cx filtds(241:300); \frac{1}{6} 1 second of data 240-300
y5 = Cy^-filtds(241:300);
z5 = Cz filtds(241:300);
% 6 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x6 = Cx filtds(301:360); \frac{6}{5} 1 second of data 300-360
y6 = Cy filtds(301:360);
z6 = Cz filtds(301:360);
% 7 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x7 = Cx filtds(361:420); % 1 second of data 361-420
y7 = Cy_filtds(361:420);
z7 = Cz_filtds(361:420);
% 8 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x8 = Cx filtds(421:480); % 1 second of data 421-480
y8 = Cy filtds(421:480);
z8 = Cz filtds (421:480);
% 9 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x9 = Cx filtds(481:540); % 1 second of data 481-540
y9 = Cy filtds(481:540);
z9 = Cz filtds (481:540);
% 10 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x10 = Cx filtds(541:600); % 1 second of data 541-600
y10 = Cy_filtds(541:600);
z10 = Cz_filtds(541:600);
% 11 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x11 = Cx filtds(601:660); % 1 second of data 601-660
y11 = Cy_{\text{filds}}(601:660);z11 = Cz filtds (601:660);
% 12 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x12 = Cx filtds(661:720); % 1 second of data 661-720
y12 = Cy_{tiltds}(661:720);z12 = Cz filtds (661:720);
% 13 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x13 = Cx filtds(721:780); % 1 second of data 721-780
y13 = Cy_filtds(721:780);
z13 = Cz_filtds(721:780);
% 14 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x14 = Cx filtds(781:840); % 1 second of data 781-840
y14 = Cy_{tiltds}(781:840);z14 = Cz filtds(781:840);
% 15 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x15 = Cx filtds(841:900); % 1 second of data 841-900
```

```
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```

```
y15 = Cy filtds(841:900);
z15 = Cz filtds (841:900);
% 16 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x16 = Cx filtds(901:960); % 1 second of data 901-960
y16 = Cy\_filtds(901:960);z16 = Cz_filtds(901:960);
% 17 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x17 = Cx filtds(961:1020); % 1 second of data 961-1020
y17 = Cy\_filtds(961:1020);z17 = Cz filtds (961:1020);
% 18 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x18 = Cx filtds(1021:1080); % 1 second of data 1021-1080
y18 = Cy_{\text{filds}}(1021:1080);z18 = Cz filtds(1021:1080);
% S<sub>3</sub> 8 seconds % S<sub>3</sub> % S<sub>4</sub> % S<sub>5</sub> % S<sub>6</sub> % S<sub>7</sub> % S<sub>8</sub> % S<sub>7</sub> % S<sub>8</sub> % S<sub>8x19 = Cx filtds(1081:1140); % 1 second of data 1081-1140
y19 = Cy_filtds(1081:1140);
z19 = Cz_filtds(1081:1140);
% 20 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x20 = Cx filtds(1141:1200); % 1 second of data 1141-1200
y20 = Cy filtds(1141:1200);
z20 = Cz filtds(1141:1200);
% 21 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x21 = Cx filtds(1201:1260); % 1 second of data 1201-1260
y21 = Cy_{tiltds}(1201:1260);z21 = cz filtds(1201:1260);
% 22 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x22 = Cx filtds(1261:1320); % 1 second of data 1261-1320
y22 = Cy_{\text{filds}}(1261:1320);
z22 = Cz filtds(1261:1320);
% 23 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x23 = Cx filtds(1321:1380); % 1 second of data 1321-1380
y23 = Cy filtds(1321:1380);
z23 = Cz filtds (1321:1380);
% 24 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x24 = Cx filtds(1381:1440); % 1 second of data 1381-1440
y24 = Cy filtds(1381:1440);
z24 = Cz filtds(1381:1440);
% 25 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x25 = Cx filtds(1441:1500); % 1 second of data 1441-1500
y25 = Cy_filtds(1441:1500);
z25 = Cz_filtds(1441:1500);
% S<sup>8</sup> 26 seconds % S<sup>8</sup> % S<sup>x26 = Cx filtds(1501:1560); % 1 second of data 1501-1560
y26 = Cy filtds(1501:1560);
z26 = Cz filtds(1501:1560);
% 27 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x27 = Cx filtds(1561:1620); % 1 second of data 15611-1620
y27 = Cy filtds(1561:1620);
```

```
z27 = Cz filtds(1561:1620);
% 28 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x28 = Cx filtds(1621:1680); % 1 second of data 1621-1680
y28 = Cy filtds(1621:1680);
z28 = Cz filtds (1621:1680);
% 29 seconds %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x29 = Cx filtds(1681:1740); % 1 second of data 1681-1740
y29 = Cy filtds(1681:1740);
z29 = Cz filtds (1681:1740);
% S<sup>8</sup> 30 seconds % S<sup>8</sup> % S<sup>x30 = Cx filtds(1741:1800); \frac{1}{8} 1 second of data 1741-1800
y30 = Cy filtds(1741:1800);
z30 = Cz filtds (1741:1800);
%% Entropy Variable Declaration
dim = 2; % embedded dimension = 2
r = 0.2; % tolerance
% Sample Entropy(SampEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%%%%
% AP direction
SE1 AP = SampEntr(x1',dim,r);
SE2 AP = SampEntr(x2',dim,r);
SE3<sup>_</sup>AP = SampEntr(x3',dim,r);
SE4 AP = SampEntr(x4',dim,r);
SE5 AP = SampEntr(x5',dim,r);SE6 AP = SampEntr(x6',dim,r);
SE7 AP = SampEntr(x7',dim,r);
SE8<sup>-</sup>AP = SampEntr(x8',dim,r);
SE9<sup>-</sup>AP = SampEntr(x9',dim,r);
SE10 AP = SampEntr(x10',dim,r);
SE11<sup>AP</sup> = SampEntr(x11',dim,r);
SE12<sup>_</sup>AP = SampEntr(x12',dim,r);
SE13<sup>-</sup>AP = SampEntr(x13',dim,r);SE14<sup>-</sup>AP = SampEntr(x14',dim,r);SE15 AP = SampEntr(x15', dim, r);
SE16 AP = SampEntr(x16',dim,r);
SE17<sup>-</sup>AP = SampEntr(x17',dim,r);SE18 AP = SampEntr(x18',dim,r);
SE19 AP = SampEntr(x19',dim,r);
SE20 AP = SampEntr(x20',dim,r);
SE21 AP = SampEntr(x21',dim,r);
SE22 AP = SampEntr(x22',dim,r);
SE23<sup>AP</sup> = SampEntr(x23',dim,r);
SE24<sup>-</sup>AP = SampEntr(x24',dim,r);SE25<sup>-</sup>AP = SampEntr(x25',dim,r);SE26<sup>AP</sup> = SampEntr(x26',dim,r);
SE27_AP = SampEntr(x27',dim,r);
SE28<sub>_</sub>AP = SampEntr(x28',dim,r);
SE29<sub>_</sub>AP = SampEntr(x29',dim,r);
SE30AP = Samplentr(x30',dim,r);
%ML direction
SE1 ML = SampEntr(y1',dim,r);
SE2 ML = SampEntr(y2',dim,r);
SE3 ML = SampEntr(y3',dim,r);
```

```
SE4 ML = SampEntr(y4',dim,r);
```

```
SE5 ML = SampEntr(y5',dim,r);
SE6 ML = SampEntr(y6', dim, r);
SE7<sup>-ML</sup> = SampEntr(y7',dim,r);
SE8 ML = SampEntr(y8',dim,r);
SE9 ML = SampEntr(y9', dim, r);
SE10 ML = SampEntr(y10',dim,r);
SE11ML = SampEntr(y11', dim, r);
SE12_ML = SampEntr(y12',dim,r);SE13_ML = SampEntr(y13',dim,r);
SE14 ML = SampEntr(y14',dim,r);
SE15<sup>-ML</sup> = SampEntr(y15',dim,r);
SE16<sup>ML</sup> = SampEntr(y16',dim,r);
SE17<sup>ML</sup> = SampEntr(y17', dim, r);
SE18ML = SampEntr(y18', dim, r);
SE19 ML = SampEntr(y19',dim,r);
SE20 ML = SampEntr(y20',dim,r);
SE21 ML = SampEntr(y21',dim,r);
SE22<sup>-ML</sup> = SampEntr(y22',dim,r);
SE23_ML = SampEntr(y23',dim,r);
SE24 ML = SampEntr(y24',dim,r);
SE25_ML = SampEntr(y25',dim,r);
SE26_ML = SampEntr(y26',dim,r);
SE27<sup>-ML</sup> = SampEntr(y27',dim,r);
SE28_ML = SampEntr(y28',dim,r);SE29 ML = SampEntr(y29',dim,r);
SE30 ML = SampEntr(y30',dim,r);
```
SampEn2 AP = [SE1 AP SE2 AP SE3 AP SE4 AP SE5 AP SE6 AP SE7 AP SE8 AP SE9 AP SE10 AP SE11\_AP SE12\_AP SE13\_AP SE14\_AP SE15\_AP SE16\_AP SE17\_AP SE18\_AP SE19\_AP SE20\_AP SE21 AP SE22 AP SE23 AP SE24 AP SE25 AP SE26 AP SE27 AP SE28 AP SE29 AP SE30 AP]; tS  $AP = 1$ : length (SampEn2 AP);

SampEn2\_ML = [SE1\_ML SE2\_ML SE3\_ML SE4\_ML SE5\_ML SE6\_ML SE7\_ML SE8\_ML SE9\_ML SE10\_ML SE11\_ML SE12\_ML SE13\_ML SE14\_ML SE15\_ML SE16\_ML SE17<sup>T</sup>ML SE18\_ML SE19\_ML SE20\_ML SE21\_ML SE22\_ML SE23\_ML SE24\_ML SE25\_ML SE26\_ML SE27\_ML SE28\_ML SE29\_ML SE30\_ML]; tS  $ML = 1$ : length (SampEn2 ML);

% Approximate Entropy(ApEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%%%% % AP direction AP1  $AP =$  ApEntr(x1',dim,r);  $AP2^{\top}AP = APEntr(x2',dim,r);$  $AP3$ <sup>-</sup>AP = ApEntr(x3',dim,r);  $AP4$ <sup> $A$ </sup> $P$  = ApEntr(x4',dim,r);  $AP5^{\top}AP = ApEntr(x5',dim,r);$ AP6 AP = ApEntr( $x6'$ ,dim,r); AP7 AP = ApEntr(x7',dim,r); AP8  $AP = ApEntr(x8',dim,r);$ AP9  $AP =$  ApEntr(x9',dim,r); AP10 AP = ApEntr(x10',dim,r);  $API1$ <sup>-</sup> $AP$  = ApEntr(x11',dim,r);  $AP12$ <sup> $AP$  = ApEntr(x12',dim,r);</sup> AP13\_AP = ApEntr( $x13'$ ,dim,r); AP14\_AP = ApEntr(x14',dim,r); AP15\_AP = ApEntr(x15',dim,r); AP16\_AP = ApEntr(x16',dim,r); AP17\_AP = ApEntr(x17',dim,r);  $AP18$ <sup>-</sup> $AP$  = ApEntr(x18',dim,r);  $AP19$ <sup>-</sup> $AP$  = ApEntr(x19',dim,r); AP20 AP = ApEntr(x20',dim,r); AP21 AP = ApEntr(x21',dim,r); AP22 AP = ApEntr(x22',dim,r);

```
AP23 AP = ApEntr(x23',dim,r);
AP24 AP = ApEntr(x24',dim,r);
AP25 AP = ApEntr(x25',dim,r);
AP26 AP = ApEntr(x26',dim,r);
AP27 AP = ApEntr(x27',dim,r);
AP28 AP = ApEntr(x28',dim,r);
AP29 AP = ApEntr(x29',dim,r);
AP30 AP = ApEntr(x30', dim, r);
%ML direction
AP1 ML = ApEntr(y1',dim,r);
AP2<sup>-</sup>ML = ApEntr(y2',dim,r);
AP3 ML = ApEntr(y3',dim,r);
AP4<sup>_ML</sup> = ApEntr(y4', dim, r);
AP5 ML = ApEntr(y5',dim,r);
AP6 ML = ApEntr(y6',dim,r);
AP7 ML = ApEntr(y7',dim,r);
AP8 ML = ApEntr(y8',dim,r);
AP9<sup>_ML</sup> = ApEntr(y9', dim, r);
AP10ML = ApEntr(y10',dim,r);
AP11<sup>ML</sup> = ApEntr(y11',dim,r);
AP12_ML = ApEntr(y12',dim,r);
AP13_ML = ApEntr(y13',dim,r);
AP14_ML = APEntr(y14',dim,r);AP15_ML = ApEntr(y15',dim,r);
AP16_ML = ApEntr(y16', dim, r);
AP17<sup>-</sup>ML = ApEntr(y17',dim,r);AP18<sup>ML</sup> = ApEntr(y18',dim,r);
AP19 ML = ApEntr(y19',dim,r);
AP20 ML = ApEntr(y20',dim,r);
AP21 ML = ApEntr(y21',dim,r);
AP22 ML = ApEntr(y22',dim,r);
AP23 ML = ApEntr(y23',dim,r);
AP24<sup>-</sup>ML = ApEntr(y24',dim,r);
AP25_ML = ApEntr(y25',dim,r);
AP26_ML = ApEntr(y26',dim,r);
AP27_ML = APEntr(y27',dim,r);AP28_ML = ApEntr(y28',dim,r);
AP29_ML = ApEntr(y29',dim,r);
AP30 ML = ApEntr(y30',dim,r);
```
ApEn2\_AP = [AP1\_AP AP2\_AP AP3\_AP AP4\_AP AP5\_AP AP6\_AP AP7\_AP AP8\_AP AP9\_AP AP10\_AP AP11\_AP AP12\_AP AP13\_AP AP14\_AP AP15\_AP AP16\_AP AP17\_AP AP18\_AP AP19\_AP AP20\_AP AP21\_AP AP22\_AP AP23\_AP AP24\_AP AP25\_AP AP26\_AP AP27\_AP AP28\_AP AP29\_AP AP30\_AP]; tA  $AP = 1$ : length (ApEn2 AP);

ApEn2\_ML = [AP1\_ML AP2\_ML AP3\_ML AP4\_ML AP5\_ML AP6\_ML AP7\_ML AP8\_ML AP9\_ML AP10\_ML AP11\_ML AP12\_ML AP13\_ML AP14\_ML AP15\_ML AP16\_ML AP17\_ML AP18\_ML AP19\_ML AP20\_ML  $AP21$ ML AP22 $-ML$  AP23 $-ML$  AP24 $-ML$  AP25 $-ML$  AP26 $-ML$  AP27 $-ML$  AP28 $-ML$  AP29 $-ML$ ]; tA  $ML = 1$ : length (ApEn2 ML);

%% PLOT GENERATION %% figure(1) subplot $(2,2,1)$ , plot $(tA$  AP, ApEn2 AP) xlabel('Time (s)','FontSize', 12), ylabel('ApEn (AP)','FontSize', 12),ylim([0 .6]); subplot(2,2,2), plot(tA\_ML, ApEn2\_ML) xlabel('Time (s)','FontSize', 12), ylabel('ApEn (ML)','FontSize', 12),ylim([0 .6]); subplot(2,2,3), plot(tS\_AP, SampEn2\_AP) xlabel('Time (s)','FontSize', 12), ylabel('SampEn (AP)','FontSize', 12),ylim([0 .6]); subplot $(2, 2, 4)$ , plot $(tSML, SampEn2ML)$ 

```
xlabel('Time (s)','FontSize', 12), ylabel('SampEn (ML)','FontSize', 12),ylim([0 
.6]);
```
## C.5 *ApEn and SampEn Averages*

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
% Title: ApEn SampEn Averages.m
% Author Jayla Wesley
% Notes: this script file loads subject data into a usable structure for 
% further analysis. estimates the average of entire data set EOFT: Trials 2 - 6
% 1. Cx is the Anterior-Posterior direction<br>% 2. Cy is the Medial Lateral Direction
% 2. Cy is the Medial Lateral Direction
  3. Downsampling by a factor of 20, 1200Hz to 60Hz
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
close all; clear; clc;
%% Trial 02 %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
filtdata2 = readmatrix('C:\Users\jmwesley\OneDrive - Newkirk Electric Associates, 
Inc\Documents\GVSU Masters 22\EGR 695 Masters Thesis\Subject 04\Subject 04 Trial Data 
Raw\SB04 Trial02.csv');
Cx filt = filtdata2(4:36003,9);Cy filt = filtdata2(4:36003,10);
Cz filt = filtdata2(4:36003,11);
% downsample the data 1200Hz to 60Hz
Cx filtds = downsample(Cx filt, 20);
Cy filtds = downsample(Cy filt, 20);
Cz filtds = downsample(Cz_filt,20);
x = Cx _{ 11tds(1:1800);y = Cy_filtds(1:1800);z = Cz filtds(1:1800);
% Calculate Center of Pressure
data COPAP = x'; % get transpose for time series AP calculation
data COPML = y'; % get transpose for time series ML calculation
% Entropy Variable Declaration Trial 2
dim = 2; % embedded dimension = 2
r = 0.2; % tolerance
% Sample Entropy(ApEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%%%%%%
% ranging r values
SampEntropy22ML = SampEntr(data COPML,dim,r);
SampEntropy22AP = SampEntr(data COPAP, dim, r);
\text{m s} r = .05
r = 0.05; % tolerance
SampEntropy052ML = SampEntr(data COPML,dim,r);
SampEntropy052AP = SampEntr(data COPAP,dim,r);
\text{m s} r = .1
r = 0.1; % tolerance
SampEntropy12ML = SampEntr(data COPML,dim,r);
SampEntropy12AP = SampEntr(data_COPAP,dim,r);
\text{m s} r = .15
r = 0.15; % tolerance
SampEntropy152ML = SampEntr(data COPML,dim,r);
SampEntropy152AP = SampEntr(data COPAP,dim,r);
```

```
\text{m s} r = .25
r = 0.25; % tolerance
SampEntropy252ML = SampEntr(data COPML,dim,r);
SampEntropy252AP = SampEntr(data COPAP,dim,r);
\text{m s} r = .3
r = 0.3; % tolerance
SampEntropy32ML = SampEntr(data COPML,dim,r);
SampEntropy32AP = SampEntr(data COPAP, dim, r);
% Approximate Entropy(ApEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%
% ranging r values
r = 0.2; % tolerance
AproxEntropy22ML = ApEntr(data COPML,dim,r);
AproxEntropy22AP = ApEntr(data-COPAP, dim, r);\text{m s} r = .05
r = 0.05; % tolerance
AproxEntropy052ML = Aperbr(data COPML,dim,r);AproxEntropy052AP = ApEntr(data_COPAP,dim,r);
\text{m s r = .1}r = 0.1; %tolerance
AproxEntropy12ML = ApEntr(data COPML, dim,r);AproxEntropy12AP = ApEntr(data COPAP, dim,r);\text{m s r} = .15r = 0.15; % tolerance
AproxEntropy152ML = ApEntr(data_COPML,dim,r);
AproxEntropy152AP = ApEntr(data COPAP, dim,r);\text{m s} r = .25
r = 0.25; % tolerance
AproxEntropy252ML = Aperint(data COPML, dim,r);AproxEntropy252AP = ApEntr(data_COPAP,dim,r);
\text{m s} \cdot \text{r} = .3r = 0.3; %tolerance
AproxEntropy32ML = ApEntr(data COPML,dim,r);
AproxEntropy32AP = ApEntr(data-COPAP, dim,r);%% Trial 03 %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
filtdata3 = readmatrix('C:\Users\jmwesley\OneDrive - Newkirk Electric Associates, 
Inc\Documents\GVSU Masters 22\EGR 695 Masters Thesis\Subject 04\Subject 04 Trial Data 
Raw\SB04 Trial03.csv');
Cx filt = filtdata3(4:36003,9);CY filt = filtdata3(4:36003,10);
Cz filt = filtdata3(4:36003,11);Cx filtds = downsample(Cx filt, 20);
Cy_filtds = downsample(Cy_filt,20);cz<sup>-filtds = downsample(Cz_filt,20);</sup>
% Center of Pressure (COP) Calculation %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x = Cx filtds(1:1800); \frac{1}{6} 1 second of data 1-60
y = Cy_filtds(1:1800);
z = Cz_filtds(1:1800);
% Calculate Center of Pressure
data COPAP = x'; % get transpose for time series AP calculation
data COPML = y'; % get transpose for time series ML calculation
% Entropy Variable Declaration Trial 3
dim = 2; % embedded dimension = 2
```

```
% Sample Entropy(ApEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%%%%%%
% range of r values
SampEntropy23ML = SampEntr(data COPML,dim,r);
SampEntropy23AP = SampEntr(data COPAP, dim, r);
\text{m s} r = .05
r = 0.05; % tolerance
SampEntropy053ML = SampEntr(data COPML,dim,r);
SampEntropy053AP = SampEntr(data COPAP,dim,r);
\text{m s} \cdot r = .1r = 0.1; % tolerance
SampEntropy13ML = SampEntr(data_COPML,dim,r);
SampEntropy13AP = SampEntr(data COPAP, dim, r);
\text{m s} r = .15
r = 0.15; % tolerance
SampEntropy153ML = SampEntr(data COPML,dim,r);
SampEntropy153AP = SampEntr(data_COPAP,dim,r);
\text{m s} r = .25
r = 0.25; % tolerance
SampEntropy253ML = SampEntr(data COPML,dim,r);
SampEntropy253AP = SampEntr(data COPAP,dim,r);
\text{m s} \cdot r = .3r = 0.3; % tolerance
SampEntropy33ML = SampEntr(data COPML,dim,r);
SampEntropy33AP = SampEntry (data COPAP, dim,r);% Approximate Entropy(ApEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%
% range of r values
r = 0.2; % tolerance
AproxEntropy23ML = ApEntr(data COPML,dim,r);AproxEntropy23AP = ApEntr(data COPAP, dim,r);\text{m s} r = .05
r = 0.05; % tolerance
AproxEntropy053ML = ApEntr(data COPML,dim,r);
AproxEntropyO53AP = ApEntr(data COPAP,dim,r);\text{m s} \cdot r = .1r = 0.1; %tolerance
AproxEntropy13ML = ApEntr(data COPML,dim,r);
AproxEntropy13AP = ApEntr(data-COPAP, dim, r);% r = .15r = 0.15; % tolerance
AproxEntropy153ML = Aperint(data COPML, dim,r);AproxEntropy153AP = ApEntr(data COPAP, dim,r);\text{m s} r = .25
r = 0.25; % tolerance
AproxEntropy253ML = ApEntr(data COPML, dim, r);
AproxEntropy253AP = Aperb}(dataCORAP,dim,r);\text{m s} \cdot r = .3r = 0.3; %tolerance
AproxEntropy33ML = ApEntr(data COPML,dim,r);AproxEntropy33AP = ApEntr(data COPAP,dim,r);
```
 $r = 0.2$ ; % tolerance

%% Trial 04 %% filtdata4 = readmatrix('C:\Users\jmwesley\OneDrive - Newkirk Electric Associates, Inc\Documents\GVSU Masters 22\EGR 695 Masters Thesis\Subject 04\Subject 04 Trial Data Raw\SB04 Trial04.csv');

```
Cx filt = filtdata4(4:36003,9);Cy filt = filtdata4(4:36003,10);
Cz filt = filtdata4(4:36003,11);Cx filtds = downsample(Cx filt, 20);
Cy_filtds = downsample(Cy_filt,20);Cz filtds = downsample(Cz filt,20);
% Center of Pressure (COP) Calculation %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x = Cx filtds(1:1800); % 1 second of data 1-60
y = Cy filtds(1:1800);
z = Cz filtds(1:1800);
data COPAP = x'; % get transpose for time series AP calculation
data<sup>-</sup>COPML = y'; \frac{1}{2} get transpose for time series ML calculation
% Entropy Variable Declaration Trial 4
dim = 2; % embedded dimension = 2
r = 0.2; % tolerance
% Sample Entropy(ApEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%%%%%%
% ranges of r
SampEntropy24ML = SampEntr(data COPML,dim,r);
SampEntropy24AP = SampEntr(data COPAP,dim,r);
\text{m s} r = .05
r = 0.05; % tolerance
SampEntropy054ML = SampEntr(data_COPML,dim,r);
SampEntropy054AP = SampEntr(data_COPAP,dim,r);
\text{m s} \cdot r = .1r = 0.1; % tolerance
SampEntropy14ML = SampEntr(data COPML,dim,r);
SampEntropy14AP = SampEntr(data COPAP,dim,r);
\text{m s r} = .15r = 0.15; % tolerance
SampEntropy154ML = SampEntr(data COPML,dim,r);
SampEntropy154AP = SampEntr(data COPAP,dim,r);
\text{er } = .25r = 0.25; % tolerance
SampEntropy254ML = SampEntr(data COPML,dim,r);
SampEntropy254AP = SampEntr(data COPAP,dim,r);
\text{m s} r = .3
r = 0.3; % tolerance
SampEntropy34ML = SampEntr(data COPML,dim,r);
SampEntropy34AP = SampEntr(data COPAP,dim,r);
% Approximate Entropy(ApEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%
% ranges of r
r = 0.2; % tolerance
AproxEntropy24ML = ApEntr(data_COPML,dim,r);
AproxEntropy24AP = ApEntr(data COPAP,dim,r);
\text{m s r} = .05r = 0.05; % tolerance
AproxEntropy054ML = ApEntr(data COPML,dim,r);
AproxEntropyO54AP = ApEntr(data COPAP, dim,r);\text{m s} r = .1
r = 0.1; %tolerance
AproxEntropy14ML = Apentr(data COPML,dim,r);
AproxEntropy14AP = Apentr(data COPAP,dim,r);
\frac{15}{6} r = .15
```
```
r = 0.15; % tolerance
AproxEntropy154ML = ApEntr(data COPML,dim,r);AproxEntropy154AP = ApEntr(data COPAP, dim,r);\text{m s} \cdot r = .25r = 0.25; % tolerance
AproxEntropy254ML = ApEntr(data COPML,dim,r);
AproxEntropy254AP = Aperort(data COPAP, dim,r);\text{m s} \cdot r = .3r = 0.3; %tolerance
AproxEntropy34ML = ApEntr(data COPML,dim,r);AproxEntropy34AP = ApEntr(data_COPAP,dim,r);
%% Trial 05 %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
filtdata5 = readmatrix('C:\Users\jmwesley\OneDrive - Newkirk Electric Associates, 
Inc\Documents\GVSU Masters 22\EGR 695 Masters Thesis\Subject 04\Subject 04 Trial Data 
Raw\SB04 Trial05.csv');
Cx filt = filtdata5(4:36003,9);Cy<sup>-filt</sup> = filtdata5(4:36003,10);
Cz filt = filtdata5(4:36003,11);
Cx_filtds = downsample(Cx_filt,20);Cy_filtds = downsample(Cy_filt,20);
Cz_filtds = downsample(Cz_filt,20);
% Center of Pressure (COP) Calculation %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x = Cx filtds(1:1800); % 1 second of data 1-60
y = Cy filtds(1:1800);
z = Cz filtds(1:1800);
data COPAP = x'; % get transpose for time series AP calculation
data COPML = y'; % get transpose for time series ML calculation
% Entropy Variable Declaration Trial 5
dim = 2; % embedded dimension = 2r = 0.2; % tolerance
% Sample Entropy(SampEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%%%%%%
% ranges of r
SampEntropy25ML = SampEntr(data COPML,dim,r);
SampEntropy25AP = SampEntr(data_COPAP,dim,r);
\text{m s} r = .05
r = 0.05; % tolerance
SampEntropy055ML = SampEntr(data COPML,dim,r);
SampEntropy055AP = SampEntr(data COPAP,dim,r);
\text{m s} r = .1
r = 0.1; % tolerance
SampEntropy15ML = SampEntr(data_COPML,dim,r);
SampEntropy15AP = SampEntr(data COPAP,dim,r);
\text{m s r} = .15r = 0.15; % tolerance
SampEntropy155ML = SampEntr(data COPML,dim,r);
SampEntropy155AP = SampEntr(dataCOPAP,dim,r);
\text{m s} r = .25
r = 0.25; % tolerance
SampEntropy255ML = SampEntr(data COPML,dim,r);
SampEntropy255AP = SampEntr(data COPAP,dim,r);
\text{m s} \cdot r = .3r = 0.3; % tolerance
```

```
SampEntropy35ML = SampEntr(data COPML,dim,r);
SampEntropy35AP = SampEntr(data COPAP, dim, r);
% Approximate Entropy(ApEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%
% ranges of r
r = 0.2; % tolerance
AproxEntropy25ML = ApEntr(data_COPML, dim, r) ;
AproxEntropy25AP = Apentr(data COPAP, dim,r);\text{m s} r = .05
r = 0.05; % tolerance
AproxEntropyO55ML = ApEntr(data COPML,dim,r);AproxEntropy055AP = Aperort(data-COPAP, dim,r);\text{m s r = .1}r = 0.1; %tolerance
AproxEntropy15ML = Aptntr(data COPML,dim,r);AproxEntropy15AP = ApEntr(data COPAP, dim,r);\text{m s r} = .15r = 0.15; % tolerance
AproxEntropy155ML = ApEntr(data COPML,dim,r);
AproxEntropy155AP = ApEntr(data COPAP, dim, r);\text{m s} r = .25
r = 0.25; % tolerance
AproxEntropy255ML = Aperint(data COPML,dim,r);AproxEntropy255AP = ApEntr(data COPAP, dim,r);\text{m s} \cdot r = .3r = 0.3; %tolerance
AproxEntropy35ML = ApEntr(data_COPML,dim,r);
AproxEntropy35AP = ApEntr(data-COPAP, dim, r);%% Trial 06 %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
filtdata6 = readmatrix('C:\Users\jmwesley\OneDrive - Newkirk Electric Associates, 
Inc\Documents\GVSU Masters 22\EGR 695 Masters Thesis\Subject 04\Subject 04 Trial Data 
Raw\SB04 Trial6.csv');
Cx filt = filtdata6(4:36003,9);Cy_filt = filtdata6(4:36003,10);Cz filt = filtdata6(4:36003,11);
Cx filtds = downsample(Cx filt, 20);
Cy_filtds = downsample(Cy_filt,20);Cz_filtds = downsample(Cz_filt,20);
% Center of Pressure (COP) Calculation %%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
x = Cx filtds(1:1800); % 1 second of data 1-60
y = Cy filtds(1:1800);
z = Cz_{\text{flits}}(1:1800);data COPAP = x'; % get transpose for time series AP calculation
data COPML = y'; % get transpose for time series ML calculation
% Entropy Variable Declaration Trial 6
dim = 2; % embedded dimension = 2
r = 0.2; % tolerance
% Sample Entropy(SampEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%%%%%%
% ranges of r
SampEntropy26ML = SampEntr(data COPML,dim,r);
SampEntropy26AP = SampEntr(data COPAP,dim,r);
```

```
\text{m s} r = .05
r = 0.05; % tolerance
SampEntropy056ML = SampEntr(data COPML,dim,r);
SampEntropy056AP = SampEntr(data COPAP,dim,r);
\text{m s} r = .1
r = 0.1; % tolerance
SampEntropy61ML = SampEntr(data COPML,dim,r);
SampEntropy61AP = SampEntr(data COPAP, dim, r);
\text{ m s r = .15}r = 0.15; % tolerance
SampEntropy156ML = SampEntr(data COPML,dim,r);
SampEntropy156AP = SampEntry1 (data COPAP, dim, r);
\text{m s} r = .25
r = 0.25; % tolerance
SampEntropy256ML = SampEntr(data COPML,dim,r);
SampEntropy256AP = SampEntr(data COPAP, dim, r);
\text{m s} \cdot r = .3r = 0.3; % tolerance
SampEntropy36ML = SampEntr(data COPML,dim,r);
SampEntropy36AP = SampEntr(data COPAP,dim,r);
% Approximate Entropy(ApEn) Calculation (Function) %%%%%%%%%%%%%%%%%%%%%%%%
% ranges of r
r = 0.2; % tolerance
AproxEntropy26ML = ApEntr(data COPML, dim,r);AproxEntropy26AP = ApEntr(data COPAP, dim,r);\text{m s} r = .05
r = 0.05; % tolerance
AproxEntropyO56ML = ApEntr(data COPML, dim,r);AproxEntropyO56AP = ApEntr(data COPAP, dim,r);\text{m s} \cdot r = .1r = 0.1; %tolerance
AproxEntropy61ML = ApEntr(data COPML,dim,r);
AproxEntropy61AP = ApEntr(data COPAP, dim,r);\text{m s} r = .15
r = 0.15; % tolerance
AproxEntropy156ML = Aperint(data COPML, dim,r);AproxEntropy156AP = Aperb(data^COPAP,dim,r);\text{m s} r = .25
r = 0.25; % tolerance
AproxEntropy256ML = ApEntr(data_COPML,dim,r);
AproxEntropy256AP = Aperb(data COPAP, dim,r);\text{m s} \cdot r = .3r = 0.3; %tolerance
AproxEntropy36ML = Aplentr(data COPML,dim,r);AproxEntropy36AP = ApEntr(data COPAP, dim,r);
```
## %% Results %%

## format long

[AproxEntropy052ML AproxEntropy053ML AproxEntropy054ML AproxEntropy055ML AproxEntropy056ML AproxEntropy12ML AproxEntropy13ML AproxEntropy14ML AproxEntropy15ML AproxEntropy61ML AproxEntropy152ML AproxEntropy153ML AproxEntropy154ML AproxEntropy155ML AproxEntropy156ML AproxEntropy22ML AproxEntropy23ML AproxEntropy24ML AproxEntropy25ML AproxEntropy26ML AproxEntropy252ML AproxEntropy253ML AproxEntropy254ML AproxEntropy255ML AproxEntropy256ML AproxEntropy32ML AproxEntropy33ML AproxEntropy34ML

AproxEntropy35ML AproxEntropy36ML]

[AproxEntropy052AP AproxEntropy053AP AproxEntropy054AP AproxEntropy055AP AproxEntropy056AP AproxEntropy12AP AproxEntropy13AP AproxEntropy14AP AproxEntropy15AP AproxEntropy61AP AproxEntropy152AP AproxEntropy153AP AproxEntropy154AP AproxEntropy155AP AproxEntropy156AP AproxEntropy22AP AproxEntropy23AP AproxEntropy24AP AproxEntropy25AP AproxEntropy26AP AproxEntropy252AP AproxEntropy253AP AproxEntropy254AP AproxEntropy255AP AproxEntropy256AP AproxEntropy32AP AproxEntropy33AP AproxEntropy34AP AproxEntropy35AP AproxEntropy36AP]

[SampEntropy052ML SampEntropy053ML SampEntropy054ML SampEntropy055ML SampEntropy056ML SampEntropy12ML SampEntropy13ML SampEntropy14ML SampEntropy15ML SampEntropy61ML SampEntropy152ML SampEntropy153ML SampEntropy154ML SampEntropy155ML SampEntropy156ML SampEntropy22ML SampEntropy23ML SampEntropy24ML SampEntropy25ML SampEntropy26ML SampEntropy252ML SampEntropy253ML SampEntropy254ML SampEntropy255ML SampEntropy256ML SampEntropy32ML SampEntropy33ML SampEntropy34ML SampEntropy35ML SampEntropy36ML]

[SampEntropy052AP SampEntropy053AP SampEntropy054AP SampEntropy055AP SampEntropy056AP SampEntropy12AP SampEntropy13AP SampEntropy14AP SampEntropy15AP SampEntropy61AP SampEntropy152AP SampEntropy153AP SampEntropy154AP SampEntropy155AP SampEntropy156AP SampEntropy22AP SampEntropy23AP SampEntropy24AP SampEntropy25AP SampEntropy26AP SampEntropy252AP SampEntropy253AP SampEntropy254AP SampEntropy255AP SampEntropy256AP SampEntropy32AP SampEntropy33AP SampEntropy34AP SampEntropy35AP SampEntropy36AP]

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