

1 **Temporal Changes of Neocortical High Frequency Oscillations in Epilepsy**

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Abstract

High frequency (100-500 Hz) oscillations (HFOs) recorded from intracranial electrodes are a potential biomarker for epileptogenic brain. HFOs are commonly categorized as ripples (100-250 Hz) or fast ripples (250-500 Hz), and a third class of mixed frequency events has also been identified. We hypothesize that temporal changes in HFOs may identify periods of increased likelihood of seizure onset. 86,151 HFOs from five patients with neocortical epilepsy implanted with hybrid (micro + macro) intracranial electrodes were detected using a previously validated automated algorithm run over all channels of each patient’s entire recording. HFOs were characterized by extracting quantitative morphologic features and divided into four time epochs (interictal, preictal, ictal, and postictal) and three HFO clusters (ripples, fast ripples, and mixed events). We used supervised classification and nonparametric statistical tests to explore quantitative changes in HFO features before, during, and after seizures. We also analyzed temporal changes in the rates and proportions of events from each HFO cluster during these periods. We observed patient-specific changes in HFO morphology linked to fluctuation in the relative rates of ripples, fast ripples, and mixed frequency events. These changes in relative rate occurred in pre- and postictal periods up to thirty minutes before and after seizures. We also found evidence that the distribution of HFOs during these different time periods varied greatly between individual patients. These results suggest that temporal analysis of HFO features has potential for designing custom seizure prediction algorithms and for exploring the relationship between HFOs and seizure generation.

Key Words: Epilepsy, HFO, oscillation, machine learning, classifier

61 **Introduction**

62 High frequency oscillations (HFOs) have received increasing interest as a promising
63 biomarker for epileptogenic tissue. HFOs are discrete electrophysiological events that stand out
64 from background activity and are seen in normal as well as epileptic tissue (Engel et al. 2009).
65 They occur in the 100-500 Hz frequency range and generally last on the order of tens of
66 milliseconds. HFOs have conventionally been separated into two classes by frequency: ripples
67 (100-250 Hz) and fast ripples (250-500 Hz), though a third class of mixed frequency events has
68 been automatically identified in human neocortical epilepsy (Blanco et al. 2010). Ripples were
69 first identified during normal brain functions such as memory consolidation (Buzsaki 1998;
70 Grenier et al. 2001; Siapas and Wilson 1998) but were later identified in epileptic tissue as well
71 (Bragin et al. 2002a).

72 The study of HFOs may help elucidate mechanisms of seizure generation (Demont-
73 Guignard et al. 2012; Jefferys et al. 2012). Many investigations have focused on the mechanisms
74 generating ripples (Bragin et al. 2004; Bragin et al. 2007; Ylinen et al. 1995) and fast ripples
75 (Dzhala and Staley 2004; Foffani et al. 2007; Ibarz et al. 2010; Menendez de la Prida and
76 Trevelyan 2011). Fast ripples have been more closely linked to pathological activity and localize
77 to the seizure onset zone (Bragin et al. 2002b; Urrestarazu et al. 2007). However, investigations
78 of human intracranial recordings indicate that HFOs in both frequency ranges increase in
79 epileptogenic brain regions (Jacobs et al. 2010; Worrell et al. 2008). Like seizures, HFOs
80 increase after medication reduction, indicating a close link between the two phenomena
81 (Zijlmans et al. 2009). These studies investigated HFOs primarily as a spatial biomarker, as they
82 appear to have great potential for delineating epileptogenic brain. Removal of HFO-generating
83 tissue has been shown to correlate with better outcomes of resective surgery (Akiyama et al.

84 2011; Jacobs et al. 2010). However, one of the primary challenges in HFO research remains how
85 to distinguish between normal and abnormal HFOs (Engel et al. 2009; Traub 2003); merely
86 identifying fast ripples and ripples is not specific enough, as all types of HFOs are present even
87 in normal human brain tissue (Blanco et al. 2011).

88

89 One method of characterizing HFOs that has received relatively little attention is a
90 detailed analysis of their temporal properties in the periods during and surrounding seizures. The
91 seminal research on this subject has been encouraging. An in-vitro study demonstrated increased
92 ripple and fast ripple activity before seizures (Khosravani et al. 2005). Later human studies
93 showed increased HFO activity 10 seconds before seizure onset (Zijlmans et al. 2011) and
94 increased high-frequency power 8 seconds before seizure onset (Khosravani et al. 2009).
95 Another investigation analyzing fifteen minutes preceding seizures found preictal changes in
96 HFO rates and HFO-band power in all patients. However, the magnitude and direction of these
97 changes were variable, and there were no clear systematic trends across patients and seizures
98 (Jacobs et al. 2009). The majority of these analyses focused on counting the number of HFOs
99 and determining the peak spectral content to classify them as ripples or fast ripples. We
100 hypothesize that quantifying other features of HFO signals and evaluating them with more robust
101 statistics provides information critical to characterizing HFOs. This information could lead to
102 new algorithms capable of predicting seizures or identifying abnormal areas of brain tissue.

103

104 The current study evaluates HFOs as a temporal biomarker for seizures, analyzing 86,151
105 events from five patients with neocortical epilepsy. HFOs were detected using an automated
106 algorithm and grouped with unsupervised clustering (Blanco et al. 2010). This method allows

107 for processing massive datasets in a manner that minimizes human bias (Gardner et al. 2007).
108 Using this dataset, we quantified waveform morphology using several signal processing features.
109 These features were the basis for comparing the dynamics of HFOs in the epochs surrounding
110 seizures, using several supervised classifiers and nonparametric statistical tests. The temporal
111 changes in the rates of each type of HFO were also evaluated. We find that each patient has
112 unique, statistically significant temporal changes in HFO rates and features in the 30 minute
113 period before and after seizures.

114

115 **Methods**

116 *Patient Selection and Data Acquisition*

117 We used a previously published HFO dataset (Blanco et al. 2010). Nine patients
118 diagnosed with medically refractory epilepsy were implanted with subdural electrodes (Ad-Tech
119 Medical Instruments, Racine, WI). As the goal of the analysis was to characterize HFOs at
120 different times relative to seizure onset, for inclusion in the study we required the presence of at
121 least one HFO during an electrographic seizure. Five of the nine patients met this study criterion.
122 Three of those excluded did not have any seizures, and the fourth did experience any HFOs
123 during the single seizure. Table 1 shows the clinical characteristics and electrode type for each of
124 the five included patients. The experimental protocol involved medication taper but not sleep
125 staging.

126

127 Electrodes were modified versions of standard grid and strip electrodes with added arrays
128 of non-penetrating, 40 μm platinum-iridium “micro” wires (Van Gompel et al. 2008). Besides
129 intracranial electrodes, all patients received a limited montage of standard gold scalp electrodes,

130 as well as other electrodes placed on the chin and anterior surface of the tibialis muscle to record
131 electromyographic activity. Stainless steel surgical sutures (Ethicon, Somerville, NJ) located at
132 the vertex region of the head served as the reference and ground for the intra- and extracranial
133 electrodes. Data were acquired on a Digital Lynx Data Acquisition System (Neuralynx,
134 Bozeman, MT) continuously at 32,556 samples per second with 18-bit resolution in up to 144
135 channels per patient, with a 9-kHz low-pass anti-aliasing filter.

136
137 HFOs were previously extracted from raw EEG data, as described in (Blanco et al.
138 2010). Briefly, we used a well-known HFO detector (Staba et al. 2002) to identify candidate
139 events, then clustered them using an unsupervised algorithm that did not presuppose the number
140 of subpopulations of HFOs. This algorithm identified four clusters of HFOs, corresponding
141 roughly to ripples, fast ripples, mixed events, and artifact. Events classified as artifacts were
142 removed, and all remaining detected HFOs in each patient were used in the analysis. It is
143 important to point out that the clustering algorithm removed filtering artifacts (Benar et al. 2010)
144 and other nonphysiological waveforms with a success rate that was similar to trained human
145 reviewers, who reviewed over 4700 individual HFOs in raw and filtered form. The algorithm
146 was used on the entire dataset without modification for ictal versus interictal period. The present
147 work is comprised of all HFOs that were detected and validated in the previous paper (Blanco et
148 al. 2010), excluding the artifacts. The EEG data, as well as the HFO markings, are freely
149 available at www.ieeg.org (Appendix A).

150

151 *Feature extraction*

152 Eight quantitative measures were calculated from each of the 86,151 remaining HFOs, a

153 process known as feature extraction. The features were: 1) fast ripple/ripple band power ratio; 2)
154 spectral centroid; 3) spectral peak; 4) line length after spectral equalization; 5) bandpassed line
155 length; 6) zero-crossings per sample length; 7) maximum amplitude; and 8) number of peaks per
156 sample. The first four features characterize the frequency content of each event. The remaining
157 four features capture elements of waveform morphology that distinguish HFO classes in
158 published literature (see Appendix B). After calculating all features, we found that the data from
159 features 7 and 8 were highly correlated with others (correlation > 0.84), but that the remaining 6
160 provided unique information. To reduce computational complexity, we used the reduced set of 6
161 features for all analyses except the initial classification (next paragraph).

162

163 *HFO Ictal/Non-ictal Classification*

164 We used three standard supervised machine-learning techniques to attempt to distinguish
165 between HFO events occurring during ictal versus non-ictal periods: logistic regression, k-
166 nearest neighbors (k-NN), and a support vector machine (SVM). Each of these techniques uses a
167 different algorithm to classify data. The goal is to determine how to label HFOs automatically in
168 subsequent data. We used two labels (ictal and non-ictal) in the first test to assess whether
169 classification was feasible, and if more complex classification experiments aimed at seizure
170 prediction would be possible. Information from all eight features was used to inform the
171 classifiers, and we performed Principal Components Analysis (PCA) to lessen the computational
172 burden on the learning algorithms. This reduced the eight features to five components while
173 retaining 96.3% of the data variance. We ran the experiments on the aggregate patient data and
174 on an individual patient basis. However, it is important to note that conclusive analyses were
175 performed on the full feature set, rather than just PCA data (see below). We split each set of

176 samples into equally-sized sets of training and testing data.

177

178 To further simplify the initial test of the classifiers, we first trained them on data from
179 each patient individually, with a segregated partition of testing data. We also trained and tested
180 on data from all patients in aggregate. In each case, we created ten random partitions from the
181 training and testing data sets, reserving 25% of the samples in each data set for cross-validation.
182 Appendix C describes the details of the cross-validation for the three methods. We assessed
183 classifier performance on the testing data using the F_1 measure (Eq. 1), the harmonic mean of
184 sensitivity and precision:

$$S = \frac{TP}{TP+FN}$$

185

(1)

$$P = \frac{TP}{TP+FP}$$

186

$$F_1 = \frac{2*S*P}{S+P}$$

187 (S : sensitivity, P : precision, TP : true positive, TN : true negative, FP : false positive)

188

189 We used a permutation test to compare the testing set results to those generated by randomly
190 assigned ictal/non-ictal labels. This tested the null hypothesis that the classification was no better
191 than randomly assigning the labels. After permuting the labels of the training data, we retrained
192 the classifiers, then reclassified the testing data. We repeated this procedure for 10,000 trials for
193 each patient and for 10,000 trials of the aggregate of all five patients. We compared the original
194 F_1 performance of the classifiers to the expected performance under the null hypothesis (the
195 distribution of F_1 from the random permutations) at a significance level $\alpha = 0.05$ (Bonferroni
196 corrected $\alpha = 0.001$).

197

198 *Temporal distribution of HFO features*

199 After calculating all features, each HFO can be represented as a unique point in “feature
200 space” that has as many dimensions as the number of features. We divided the HFOs into four
201 time epochs: preictal, ictal, postictal, and interictal. Pre- and postictal windows were defined as
202 10 minutes before and after a seizure, respectively. Interictal was defined as greater than 10
203 minutes from a seizure for this test. We tested whether the distribution of HFOs within the
204 feature space in each epoch was non-random. The null hypothesis was that the scattered
205 distribution of features of all HFOs in each epoch was no different than if the HFOs were
206 randomly assigned to an epoch. The centroid of features 1-6 was calculated (7-8 were not
207 included due to high correlation with the other features). We measured the Euclidean distance of
208 each HFO’s features to the centroids, grouped HFOs according to epoch, and used the median
209 distance as the measure of dispersion for that epoch. We first calculated the dispersion using the
210 original labels. We then randomly permuted the epoch labels, determined a new centroid for each
211 of the four epochs, and recalculated the dispersion of all relabeled HFOs within that epoch,
212 repeating this procedure over 10,000 trials. The probability of the null hypothesis (i.e. that the
213 observed dispersion in each epoch was that of a random sample of all HFOs) was the proportion
214 of permutations with dispersion values more extreme than the real data, tested with a
215 significance level of 0.05. Similar analyses were done with pre/postictal periods of several other
216 durations from 2-120 minutes; results of this analysis were similar.

217

218 *Temporal evolution of HFO features*

219 We assessed changes in the morphology features before and after seizures using two non-
220 parametric statistical tests that are resistant to outliers and/or skewed data. We first performed

221 Principal Components Analysis (PCA) in each patient to reduce the set of HFO features to two
222 dimensions for the purpose of visualization. We divided the four epochs above into smaller 5- to
223 10-minute “stages” (10 total) between 0-30 minutes before and after seizures. Interictal for these
224 analyses was defined as greater than 30 minutes from a seizure. We made scatter plots of the
225 HFOs in the first two PCA dimensions for each stage. The Kruskal-Wallis Test of the first two
226 PCA components determined whether the 2-dimensional PCA distributions from the different
227 stages were unique versus samples of the same distribution. We then evaluated temporal changes
228 of the six individual morphology features (see previous paragraph) using Spearman’s rank
229 correlation, the nonparametric equivalent of the Pearson correlation. This test, which used 6
230 features explicitly, served as a more robust analysis of the conclusions in the PCA data. This test
231 determines whether there is a monotonic relationship between time and each feature. HFOs prior
232 to the first seizure or after the last were excluded from this analysis, as their relationship to
233 previous or subsequent seizures could not be determined.

234

235 *Stereotyped responses in different seizures*

236 We evaluated whether the temporal changes above were consistent from seizure to
237 seizure within each patient with an analysis of variance (ANOVA) on the first two PCA
238 components in each of the ten stages. We calculated the variance of each component in each
239 temporal stage for each seizure, then computed the ANOVA across all seizures in each patient.
240 The ANOVA p-values were averaged across each patient and across each stage: high p-values
241 indicate that the distributions are similar from seizure to seizure, i.e. that the response is
242 stereotyped.

243

244 *Temporal evolution of HFO rate and class*

245 In order to analyze temporal changes in the rates and proportions of each of the three
246 HFO classes (ripples, fast ripples, mixed events (Blanco et al. 2010)), we calculated the average
247 rate (events/minute) and percentage of total events from each HFO class within each of the 5- to
248 10-minute stages described above. To quantify the relationship between HFO class and latency
249 to seizure, we performed a Kruskal-Wallis test comparing the rank order of each class with time
250 to nearest seizure, aggregating all seizures. This was done twice, evaluating time to next and time
251 to previous seizure. A significant p-value (<0.05) indicates that the three clusters have different
252 distributions, in other words that each class tends to occur at different times in relation to the
253 nearest seizure. We tested whether the preictal rates of each class change significantly from
254 interictal baseline with a chi-square test, comparing the proportion of each type of HFO during
255 interictal versus the combination of all preictal stages (0-30 minutes).

256

257 **Results**

258 *HFO Ictal/Non-ictal Classification*

259 The first analysis tested three classifiers to determine whether they could distinguish
260 HFOs occurring during seizures (ictal) from those occurring at other times (non-ictal, including
261 interictal, postictal, and preictal). Figure 1 shows how well each classifier was able to separate
262 the morphology of ictal and non-ictal HFOs, which we quantified using the F_1 measure. The
263 logistic regression classifier is not included in the figure because it did not make a single true
264 positive identification. A higher F_1 score indicates a larger number of true positives relative to
265 false positives and false negatives. The highest possible score is 1, resulting from perfect
266 sensitivity and precision. The classifiers had limited success in separating the ictal and non-ictal

267 HFOs. This is in part due to the fact that the proportion of non-ictal HFOs was much greater than
268 ictal in the training data. The k-NN and SVM classifiers were statistically superior to random
269 classification (0.05 significance, Bonferroni correction $p \ll 0.001$) in patients A and B, and the
270 k-NN was superior to random classification in patient D and in the aggregate of all 5 patients.
271 Thus, in certain patients it was possible to distinguish ictal from non-ictal HFOs better than
272 random; however, the actual F1 scores were all < 0.2 , suggesting the classification was not very
273 sensitive nor precise.

274

275 *Temporal distribution of HFO features*

276 To evaluate whether any feature characteristics varied over time, we further subdivided
277 the non-ictal period into postictal, preictal, and interictal epochs. Table 2 displays the number of
278 HFOs in each epoch. Seizures were a mean of 120 minutes apart (range 9.5 to 839 min). In three
279 patients, some seizures occurred less than one hour apart, causing pre- and postictal periods to
280 overlap. HFOs occurring during these overlaps were counted as both pre- and post-ictal events,
281 which may introduce bias into some of the following analyses. However, this involved a small
282 number of HFOs (15% of HFOs in patient A, none in B and C, and 1% in D and E), and
283 removing them from analysis did not change any of the conclusions (next section). In each
284 patient, we tested whether the distribution of HFOs in feature space was different during each
285 epoch than what would be expected from a random sample of HFOs from all epochs in that
286 patient. Figure 2 displays a plot of all HFOs from one patient, projected into two-dimensional
287 feature space via PCA, with raw data from several HFOs displayed to demonstrate the varied
288 appearance across the PCA space. Figure 3 displays the same data separated into time epochs,
289 showing that the HFO distribution in each epoch (i.e. the “shape” of the scattered HFOs) differs

290 from the shape of the population as a whole. Similar findings were found in the other patients,
291 though the distributions and rates of each HFO type were very different from this patient (see
292 Fig. 6). Though it cannot be displayed graphically, the HFO distributions using all six features
293 have similar temporal differences. When analyzing the data in six dimensions (one for each
294 feature), the distributions were significantly different than random: there were specific
295 characteristics during the different epochs, demonstrated by a difference in cluster dispersion as
296 measured by Euclidean distance ($p < 0.05$). We made similar temporal comparisons for each
297 feature individually and found great inter-patient variability. Table 3 shows the p-values for two
298 example features, the line length after spectral equalization and power band ratio. The other
299 features demonstrated similar inter-patient variability and were significant in some, but never all,
300 of the five patients. These results imply that HFO distributions vary during different peri-ictal
301 epochs and involve complex relationships between several features, but that these changes are
302 unique to individual patients. Thus, samples from different times, especially across different
303 patients, will likely have different feature distributions.

304

305 *Temporal evolution of HFO features*

306 The previous section tested whether each epoch had a different distribution from random.
307 We next sought to test the time-dependency of HFO features. The first step was to compare the
308 distributions in each stage in the 2-dimensional PCA plots (e.g. Fig. 3 for Patient C). The
309 Kruskal-Wallis Test demonstrated that for all patients there were highly significant differences
310 between stages ($p \ll 0.0001$). We then evaluated how the individual features changed
311 temporally, using time before or after seizure as a continuous variable rather than constraining
312 the analysis to categorical epochs. We used the Spearman correlation to assess whether there was

313 any time dependence for each of the six individual features. The Spearman correlation
314 determines strength and direction of any monotonic relationship between a feature and time; in
315 this case, it assesses how much of a measurable, progressive change the HFO features have as a
316 function of time prior or subsequent to seizure. Figure 4 shows the Spearman correlations for
317 each patient. Correlations closer to ± 1 indicate stronger relationships. In most cases, several
318 features demonstrated weak but statistically significant correlations with both time to next
319 seizure and time from previous seizure. Interestingly, certain features showed a significant
320 positive correlation with time in some patients and a significant negative correlation with time in
321 others. These analyses were repeated in patients A, D, and E with the overlapping HFOs
322 removed, and there were no substantive changes to the conclusions: Kruskal Wallis tests were
323 still all significant, and there were minor changes to the Spearman correlations (Fig. 4B).

324 In order for these temporal changes to be helpful in the development of customized
325 seizure prediction algorithms, they must be consistent from seizure to seizure. Looking at each
326 patient individually, we evaluated the seizure to seizure variability by running ANOVA on the
327 first two PCA components in each peri-ictal stage across all seizures. In this analysis, when the
328 p-value is “insignificant” (> 0.05), it corresponds to a similar distribution of the PCA values in
329 subsequent seizures, i.e., the response is stereotyped and thus the ANOVA fails to find a
330 difference between them. Conversely, a “significant” p-value indicates there are differences
331 across seizures, so the distributions of PCA are not stereotyped. An example of each case is
332 shown in Fig. 5. These plots demonstrate that some patients are stereotyped while others are not,
333 but do not capture all of the temporal changes described in previous paragraphs. The HFO
334 distributions are quite similar from different seizure periods for example, there are a large
335 number of HFOs 20-30 minutes before seizure 3 with different morphology, so it would be

336 difficult to predict seizure onset in this patient based upon these data. We calculated the ANOVA
337 of the PCA distributions across different seizures in every stage for every patient to determine if
338 the changes were stereotyped. We found that three of the patients (A, B, E) had stereotyped
339 responses across all stages ($p > 0.05$) (Fig. 5C). In the other two patients (C, D), the ANOVA
340 found that at least one of the PCA components was significantly different between different
341 seizures, meaning it would be hard to predict how the distribution would look in later seizures.
342 Taken together, these data demonstrate that several HFO features are correlated with time, that
343 these changes are often stereotyped, but that there is significant inter-patient variability.

344

345 *Temporal evolution of HFO rate and class*

346 Although it is clear from these analyses that a wide set of features is helpful to
347 describe HFOs and their temporal dynamics, the data also suggest that the relative rate of each
348 HFO cluster (ripple, fast ripple, mixed event) may differ between different stages. Constraining
349 each HFO to one of 3 cluster labels removes most information about its dynamic features, but is
350 also the most common way they are currently studied. In order to evaluate any changes
351 descriptively, we first plotted the relative rates of each HFO cluster, as well as the total HFO
352 rate, for each patient in each stage (Fig. 6). We observed distinct patient-specific trends in the
353 evolution of rates and proportions of each cluster before and after seizures. In three of the
354 patients (A, B, D), the total number of HFOs is relatively constant until an increase during the
355 ictal period, an effect less pronounced in patient B. But the individual HFO clusters behave quite
356 differently in these three patients. In A, there is a subtledrop in the proportional of ripples just
357 before the seizures. B has a similar effect postictally. In D, mixed events dominate preictally,
358 but disappear during the seizures. Observing the datapoints from individual seizures shows that

359 similar findings are seen at each seizure. Patients C and E have much smaller changes in total
360 HFO rate, sometimes having fewer HFOs during seizures. The only consistent change in patient
361 C is increased mixed events and loss of ripples during seizures. Patient E is difficult to evaluate
362 with only 2 seizures, but tends to have a predominance of ripples preictally and mixed events
363 immediately postictally. We tested whether the HFO rates changed significantly from baseline
364 by comparing the proportion of each HFO cluster in the interictal stage with the aggregate of the
365 4 preictal stages. Patients C, D and E had very significant changes from baseline, while A and B
366 did not ($p < 0.0001$, chi-square). In order to quantify the relationships between each type of HFO
367 and time to seizure in Fig. 6, we compared the firing times of every HFO with a Kruskal-Wallis
368 test using time to the previous and subsequent seizure as continuous variables. This assesses by
369 rank order whether there are trends in the temporal distribution of each HFO cluster in each
370 patient. A significant result ($p < 0.05$) indicates that the firing times of the three clusters are
371 unlikely to be randomly distributed—that at least one type of HFO (ripple, fast ripple, mixed
372 event) is more likely to occur at specific times between seizures. We found that HFO times both
373 before and after seizures were significantly different in all patients and times (Table 4, Kruskal-
374 Wallis Test, $p < 0.05$). Thus, in each patient specific types of HFOs became more or less likely to
375 occur in the pre- and post-ictal periods. These findings demonstrate the strong temporal
376 relationships HFOs have with seizures as well as the disparate patterns found in different
377 patients.

378

379 **Discussion**

380 *Peri-ictal HFO distributions vary greatly among patients*

381 Our current analysis evaluates properties of three classes of HFOs in four epochs (inter-,

382 pre-, postictal, and ictal), collecting a large number of events in 5 patients. This allowed a
383 thorough comparison of the statistical characteristics of HFOs during these epochs. Our use of
384 three classes of HFO is a slight departure from previous work on these types of oscillations in
385 patients with epilepsy. The first two classes are very similar to ripple and fast ripples, which are
386 commonly recognized in the literature and were originally identified based upon their peak
387 oscillation frequency. In this work, we have included mixed events that were identified by an
388 unsupervised algorithm (Blanco et al. 2010) as a third class, as their power spectra contain
389 features of both ripples and fast ripples

390

391 Our results show that the relative and absolute rates of each class of HFO changed in the
392 periods before and after seizures, and that these effects varied greatly among patients (Fig. 6).
393 The ictal period was most variable in terms of the number and type of HFOs. The total HFO rate
394 increased in three of the patients during seizures, while in the other two patients it decreased in
395 the preictal period. Such disparities in preictal HFO rates have been identified previously
396 (Jacobs et al. 2009); to our knowledge this is the first work evaluating the postictal and interictal
397 periods as well. Interestingly, not all patients exhibited their highest total HFO rate during
398 seizures. Fast ripple and mixed event rate increased beginning from 20-30 minutes prior to
399 seizures in all patients but B. Fast ripple rates increased postictally in patients B-E. Ripples were
400 predominant in 3 patients, though the ratio of ripples to fast ripples changed significantly in the 5
401 minutes before and after seizures. The time between seizures was highest in patients B and C,
402 but there are no clear patterns that distinguish these two patients from the others. It is notable
403 that there were no trends consistent among all 5 patients. With such a small patient sample size
404 identifying so many differences between individuals, it is likely a common occurrence for

405 patients to have unique HFO distributions. The cause of these findings is not clear. There are two
406 main possibilities. The first is that behavior of HFOs during, between and after periods of
407 seizure generation may be unique to a patient's epileptic network or mechanism of epilepsy. Our
408 analysis of PCA features (Fig. 5) revealed that at least three of the patients had unique responses
409 that were stereotyped in different seizures, suggesting this may be true in some patients. Another
410 possibility is that these individualized patterns may be due in part to sampling error, as there is
411 no uniform implant strategy for all patients, and the microwires which detected many of the
412 HFOs are dispersed unevenly in the implanted grids. These questions could be explored further
413 by a larger sample with different implantation strategies that maximize high resolution electrode
414 coverage in the brain, or perhaps by guiding electrode placement through intraoperative mapping
415 of HFO generating regions to augment standard electrode placement strategies. Whatever the
416 cause, our findings suggest that each patient must be evaluated individually when using HFO
417 rates as an electrical seizure biomarker.

418 One intriguing possibility for future work lies in comparing the results from patients B
419 and D, which had the most similar ictal responses in Fig. 6 (increased rate of HFO and ripples
420 during seizure). These two patients both had frontal cortical dysplasia (Table 1). Although this
421 sample size is small, this result suggests that future work should aim to stratify patients by
422 epileptic pathology, and perhaps common trends could be identified. Adding patients with other
423 types of epilepsy (e.g. mesial temporal sclerosis) would also provide crucial information. Such
424 work would likely also benefit from additional features that were not present in the current
425 analysis.

426

427 *HFO features during the ictal period, and seizure prediction*

428 We tested three machine learning algorithms with a large dataset of automatically
429 detected HFOs to determine whether supervised classification could be used to identify
430 differences between ictal and nonictal HFOs. Using the measurements described in Appendix B,
431 none of the algorithms obtained an F1 score higher than 0.2, which is fairly poor performance.
432 This finding suggests that, at least with these tools and features, it is difficult to distinguish any
433 HFO signal features that are specific to the ictal period.

434 Despite the low performance, two methods did beat random chance, so we further
435 investigated the temporal changes in HFOs in the peri-ictal period to determine the source of the
436 differences. The first step was to analyze the HFO features in “feature space.” When the HFOs
437 were grouped together based upon stage, each patient’s HFO distribution had a unique shape in
438 feature space that changed over time. We found that these changes continued to evolve as much
439 as 30 minutes before or after seizures. We tested these changes in several different ways:
440 comparing the distributions of their PCA-reduced features, comparing their 6-dimensional
441 distributions to the distributions of random samples, and determining the Spearman correlation
442 of each feature with time before and after seizures. In each case, there was strong statistical
443 evidence of measureable temporal changes in HFO features.

444 These findings have significant implications for developing automated seizure prediction
445 or warning algorithms. Although this analysis did not attempt to develop such an algorithm, it
446 was performed on an unbiased sample of continuous EEG, thus simulating the “real-world”
447 conditions necessary for such an algorithm to be successful. Additionally, it is well known that
448 classification algorithms used for seizure detection and prediction usually perform better with
449 input from multiple orthogonal features; most of our analyses were based upon single or PCA-
450 reduced features. Future attempts to develop seizure prediction algorithms based upon

451 combinations of these and other features may be very useful, but will need to be rigorously tested
452 using independent testing and training sets with prolonged continuous intracranial EEG,
453 comparing against chance predictors using established methods (Mormann et al. 2007; Snyder et
454 al. 2008). Furthermore, these data also suggest that these algorithms will be most effective if
455 they are personalized for individual patients based upon their unique features (Stacey and Litt
456 2008). For example, certain features might be relevant for one patient but not another, or the
457 correlation of certain features with seizures might differ in strength or direction between patients.
458 This will require a thorough analysis of HFO data on a per-patient basis before adjusting and
459 applying the prediction algorithm. It will be critical to identify appropriate patients for such
460 algorithms, and likewise not to abandon these techniques if they fail in some patients.

461

462 *Identifying abnormal, epileptic HFOs*

463 As HFOs have been identified in both normal and abnormal brain, it is imperative that
464 any clinical decisions based upon them are able to distinguish whether they are markers of
465 abnormal activity or not (Engel et al. 2009). To date, the primary clinical strategies proposed are
466 to identify the electrodes with the highest number of HFOs (Jacobs et al. 2010) or to stratify
467 HFOs into different categories based upon peak frequency (i.e. ripples 100-250 Hz, fast ripples >
468 250 Hz) (Akiyama et al. 2011). While such strategies are promising, they were based upon
469 selected retrospective data, and there is considerable risk in basing prospective clinical decisions
470 upon such data. The primary concern is that HFOs also occur outside the seizure onset zone,
471 even in patients who do not have epilepsy at all (Blanco et al. 2011). It is quite possible that any
472 patient can have a region with a relatively higher HFO density, but that the increase could be due
473 to local edema, physiological variation, or various other non-epileptic causes. It is therefore

474 crucial to develop a method of characterizing HFOs that are true markers of epileptic tissue.

475 One characteristic of HFOs that has received relatively little attention is their temporal
476 variation. A recent study evaluated HFOs and sharp waves with respect to seizures and found
477 that, unlike sharp waves, HFO incidence changes similar to seizures (Zijlmans et al. 2009). This
478 finding is intriguing in light of recent work investigating the mechanisms of abnormal HFOs and
479 seizures. Ripples were originally described in normal tissue (Ylinen et al. 1995), and are
480 considered to be the product of fast synchronous firing of inhibitory potentials. Conversely, later
481 work in epileptic tissue has shown that “abnormal” HFOs are comprised of population spikes
482 (Bragin et al. 2011). This difference has led to a great deal of research to determine how
483 epileptic HFOs are made (Jefferys et al. 2012). But detailed physiological studies into these
484 phenomena are extremely difficult with current technology. Therefore, computational modeling
485 has been used to confirm the role of population spikes in ripples (Stacey et al. 2011; Stacey et al.
486 2009; Wendling et al. 2012) and fast ripples (Demont-Guignard et al. 2012; Ibarz et al. 2010;
487 Roopun et al. 2010). In each case, epileptic pathologies (i.e. increased excitation, decreased
488 inhibition, abnormal coupling) cause the abnormal HFO to occur. Wendling et al. demonstrated
489 that HFO characteristics change on a continuum from normal activity to seizure, depending upon
490 the parameters (Wendling et al. 2012). These realistic modeling studies suggest that epileptic
491 pathologies will alter the appearance of HFOs, and predict that a wide range of parameters
492 (Demont-Guignard et al. 2012; Stacey et al. 2009) can be responsible. Our data demonstrate
493 clinical evidence supporting this: that there are temporal changes in HFO morphology leading
494 into seizures. These data likely represent a change in underlying physiology that is not yet
495 quantified, which suggests that a robust feature analysis of HFOs may be able to illuminate
496 future research into the underpinnings of this phenomenon.

497 If one assumes that there is an underlying process that alters the network parameters
498 before and after seizures, and that it affects both seizures and HFOs, it follows that HFOs may
499 have different features in the time surrounding seizures. Until technology is able to characterize
500 these network phenomena on smaller scales, our best method is to analyze EEG data. In this
501 study, we analyze an unbiased sample of intracranial EEG and find that there are indeed
502 temporal changes in HFO characteristics. However, the differences were relatively small and
503 were highly variable between patients. Further work evaluating different signal features,
504 recording EEG on different spatial and temporal scales, and comparing HFOs from “normal” and
505 “abnormal” regions may be better able to distinguish these changes and lead to understanding of
506 the mechanisms involved in generating both HFOs and seizures.

507

508 **Conclusion**

509 With increasing interest in and evidence for HFOs as an electrical seizure biomarker, it is
510 crucial to develop automated methods to identify and classify them that are not operator
511 dependent, and that can be applied to a wide range of patients as well as animal models. Using
512 data from an automated HFO detection algorithm, this study provides statistical evidence that
513 some patients demonstrate temporal changes in the distribution of HFOs in the periods before
514 and after seizures. Initially, these findings can be the basis of further experimental algorithms to
515 explore temporal phenomena such as seizure generation, prediction, and epileptogenesis. With
516 further validation, they may also provide a basis for algorithms capable of identifying periods of
517 increased risk of seizure onset. Perhaps the most important finding from this study is that
518 individual patients have vastly different HFO patterns related to seizures; these findings suggest
519 that any clinical decision based upon HFO analysis must be customized to each individual

520 patient.

521

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523

524 **Figure Legends**

525 Figure 1: Classifier performance for the aggregate patient data and for individual patients as
526 measured by the F_1 score, the harmonic mean of sensitivity and precision (max=1). Statistical
527 significance at the level of $\alpha = 0.05$ (Bonferroni corrected $\alpha = 0.001$) is indicated by an asterisk.
528 Results from the logistic regression classifier are not shown because the F_1 score was zero in
529 every trial. SVM: support vector machine; k-NN: k-Nearest Neighbor.

530

531 Figure 2: Scatter plot of all HFOs in Patient C in the first two dimensions of the PCA space.
532 Each dot represents one HFO and is colored according to HFO cluster as determined by (Blanco
533 et al. 2010). Localization of each dot corresponds to the first and second PCA components
534 plotted in x-y coordinates. Representative waveforms from different regions of the PCA feature
535 space provide examples of the morphology at each location.

536

537 Figure 3: Outside: Scatter plots of the HFOs in each time stage for Patient C in the first two
538 dimensions of the PCA space. Color indicates HFO cluster as in Fig. 2. Bar graphs show the
539 average rate (events per minute) for each cluster. Center: Scatter plot of all HFOs from Patient C.

540

541 Figure 4: A: Bar graphs showing the Spearman correlations for morphology features and time to
542 next seizure (left) and time from previous seizure (right). Statistically significant correlations (α
543 = 0.05) are indicated by an asterisk. B: Repeat analysis when overlapping HFOs are removed.
544 Patient B and C had none so the analysis was unchanged (*italics*). Black circles indicate subtle
545 differences from A, most notably that two values in patients A and D become significant
546 ($p < 0.05$). Kruskal Wallis tests all remained $p < 0.001$ when overlappers were removed (not

547 shown).

548

549

550 Figure 5: Temporal HFO distributions are stereotyped across different seizures in some patients.

551 HFOs are displayed by their first two principle components and stratified by seizure. Plot is

552 similar in organization to Fig. 3, but coloring corresponds to nearest seizure. A: In patient C,

553 there are distinct patterns seen before or after certain seizures, such as 30-20 minute preictally in

554 seizure 3. This patient does not have stereotyped HFO distributions ($p < 0.05$). B: Patient A has

555 very similar distribution across all seizures in every stage, a stereotyped response ($p > 0.1$). All

556 plots have same axes and scale, shown in lower left (bars: 1 unit). C: P-values for the ANOVA

557 of the first two PCA components. Bar graphs of average p-value across all stages except

558 interictal in each patient. Bars indicate one standard deviation ($n=9$ stages). Asterisks indicate

559 statistical significance ($p < 0.05$) Insignificant p-values correspond to similarity in the

560 distributions.

561

562

563 Figure 6: Graphs of the relative rates of HFOs in each frequency cluster (orange, blue, and pink

564 lines; left axis) and the total HFO rate (dotted dark line; right axis). Rates of each HFO cluster

565 during individual seizures are indicated with indicated symbols; lines indicate mean across all

566 seizures. Error bars: standard deviation of total rate. Lack of any consistent trends among

567 different patients implies that there is marked inter-patient variability in HFO rates before and

568 after seizures. Clustering of datapoints from different seizures indicates stereotyped responses,

569 most prominent in patients A, B and D. #- Chi-square testing indicated that proportions of each

570 HFO cluster are significantly different from interictal in the patients C, D, and E. Parentheses
571 next to patient name indicate number of seizures in each patient.

572

573

574

575 **Tables**

576 Table 1: Patient summary

Subject	Age	Sex	Pathology	Electrodes	Electrode Placement
A (SZ01)	35	F	Frontal neocortical oligodendroglioma	Standard: 6x6 Micro: 1x8	Right: frontal, motor cortex
B (SZ02)	24	M	Frontal cortical dysplasia	Standard: 6x6 Micro: 4x6	Left: frontal, frontoparietal
C (SZ04)	39	F	Temporal neocortical gliosis	Standard: 6x6, 1x8, 1x8, 1x4, 1x4, 1x4 Micro: 1x4	Left: temporal, inferior frontal, superior frontal, anterior temporal, posterior temporal, anterior, posterior
D (SZ07)	42	F	Frontal cortical dysplasia	Standard: 6x8, 3x8, 1x4 Micro: 1x4, 1x8	Right: frontal, temporal, anterior temporal, inferior temporal, posterior temporal
E (SZ08)	38	F	Temporal neocortical gliosis	Standard: 6x6 Standard depth: 1x4 Micro depth: 1x4, 1x4	Left: temporal, anterior temporal, middle temporal, posterior temporal

577

578 Summary of patient clinical characteristics and electrode placement. Parentheses: labels used in
579 prior study, see Table 1 in (Blanco et al. 2011) for further details.

580

581

582

583 Table 2: HFO and seizure counts
584

Patient	Total seizures	Total HFOs	Inter-ictal	Preictal	Ictal	Post-ictal
A	9	2552	1487	314	371	380
B	3	9905	9237	280	139	249
C	4	12417	11585	329	317	186
D	7	53015	49519	1787	433	1276
E	2	8262	7210	493	46	513
All	25	86151	79038	3206	1306	2604

585
586 Counts of HFOs in each time epoch for dispersion analysis. Preictal and postictal epochs were
587 defined as 10 minutes before and after seizures, respectively.

588

589

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591
592

Table 3: Dispersion p-values from 2 features

Feature 3 (Line length)						
	Pt A	Pt B	Pt C	Pt D	Pt E	All
Interictal	0.0018	*	*	*	0.0011	*
Preictal	0.3713	0.0615	*	*	0.0002	0.2242
Ictal	0.0426	0.0007	*	0.3148	0.0155	*
Postictal	0.0983	*	*	0.0038	*	*

593

Feature 4 (Power Band Ratio)						
	Pt A	Pt B	Pt C	Pt D	Pt E	All
Interictal	*	0.2285	*	*	*	0.1398
Preictal	0.0442	0.4916	*	*	0.4208	*
Ictal	*	0.0001	*	*	*	*
Postictal	*	0.1059	*	*	*	0.3285

594
595

596 Table 3: P-values from the epoch dispersion analysis. Values were calculated by dividing the
597 number of permutation test trials with results more extreme than the true values by the total
598 number of permutation test trials (10,000). Bold text indicates values that were not statistically
599 significant. The combination of all (six) features was significant in all patients. Each patient had
600 different trends with the individual features. Asterisks: highly significant ($p < 1e-6$)

601
602

603 Table 4: P-values of temporal rank
604

	Time to Next	Time From Previous	Closest cluster (next)
Pt A	<.0001*	<.0001*	FR, M
Pt B	0.013*	<.0001*	M, R
Pt C	<.0001*	<.0001*	FR, R
Pt D	<.0001*	<.0001*	FR, R
Pt E	<.0001*	<.0001*	R

605
606

607 Table 4: P-values from the second Kruskal-Wallis Test, used to assess whether an HFO near a
608 seizure was more likely to be of a specific cluster. Third column indicates which clusters were
609 closest to the next seizure (R: ripple, M: mixed, FR: fast ripple). Asterisks: statistical
610 significance.

611
612

613 **Appendix A: Online access to data at www.ieeg.org**

614 The EEG data and HFO markings used in this study have been posted by the authors of (Blanco
615 et al. 2011) for viewing and analysis on the IEEG-Portal (<https://www.ieeg.org>). This free online
616 database allows the user to view, download, and annotate EEG data, and has been established to
617 allow datasharing and collaboration. The five studies used in this manuscript can be accessed
618 with the following names:

619 Patient A- I001_P034_D01; Patient B- I001_P011_D01; Patient C- I001_P015_D01; Patient
620 D- I001_P014_D01; Patient E- I001_P017_D02.

621
622

623 **Appendix B: Feature Equations**

624

625 1) Power Band Ratio (250-500 Hz)/(100-200 Hz): The ratio of estimated power in the
626 hypothesized “fast-ripple” band to that in the hypothesized “ripple” band. Computed on the
627 band-passed data; frequency-input domain.

628

$$\text{Power band ratio} = \hat{P}_{[250,500]} / \hat{P}_{[100,200]}$$

629

where

$$\hat{P}_{[a,b]} = \sum_{\{k \in Z \mid \zeta(a) \leq k \leq \zeta(b)\}} |M_k|^2, (a < b) \in (0, \frac{F_s}{2})$$

630 and M_k is a multitaper power spectral density estimate. For more details, see Blanco et al. 2010.

631

632 2) Spectral Centroid: The frequency corresponding to the “center of mass” of the spectrum.
633 Computed on the band-passed data; frequency-input domain.

634

$$\text{Spectral centroid} = \frac{\sum_{k=0}^{N/2} \frac{k}{NT} |M_k|^2}{\sum_{k=0}^{N/2} |M_k|^2}$$

635

636 3) Spectral Peak: The frequency corresponding to the peak of the estimated power spectral
637 density. Computed on raw data; frequency-input domain.

638

$$\text{Spectral peak} = \frac{1}{NT} \underset{k}{\operatorname{argmax}} |M_k|^2, k \in \{0, 1, \dots, \frac{N}{2}\}$$

639

640 4) Line length after spectral equalization: Detections are first detrended and energy normalized
641 by dividing by their Euclidean lengths and by sample length. Computed on raw data after first-
642 order backward differencing; time-domain input.

$$\text{Line length} = \frac{1}{L} \sum_{i=0}^{L-2} |x_{i-1} - x_i|$$

643

644

645 5) Line length (bandpassed) per sample length: line length of the 100-500 Hz bandpassed HFO
646 waveform divided by the total length of the sample

$$\text{Line length (bandpassed) per sample length} = \frac{1}{N} \sum_{i=2}^N |x_i - x_{i-1}|$$

647 6) Zero-crossings per sample length: the number of times the sample crosses its mean divided by
648 the total length of the sample, which is zero-meaned

649

$$\text{Zero crossings per sample} = \frac{1}{N} \sum_{i=2}^N \text{sign}(x_{i-1}) \neq \text{sign}(x_i)$$

650

651 7) Maximum amplitude range: the difference between a sample's maximum and minimum
652 values

$$\text{Max. amplitude range} = \max(x) - \min(x)$$

653

654 8) Number of peaks per sample length: the total number of local maxima in a sample divided by
655 the total length of the sample

$$\text{No. peaks per sample} = \frac{1}{N} \sum_{i=2}^{N-1} (x_i > x_{i-1} \wedge x_i > x_{i+1})$$

656

657 **Appendix C: Classifier Cross Validation**

658 The k-NN classifier requires specification of the value for k , the number of nearest neighbors to
659 use. We determined the optimal k by calculating the average performance of the classifier for a
660 range of k values (from 1 to 20) over the ten partitions of training and validation data and
661 selecting the input parameter that corresponded to the best average performance. We similarly
662 obtained optimal parameters for the support vector machine (SVM) classifier, which requires
663 specification of a) a cost parameter, C , that controls the tradeoff between reducing training errors
664 and controlling the complexity of the model, and b) a parameter, γ , that controls the width of the
665 Gaussian kernel. The latter two classifiers were re-trained on the augmented training and
666 validation data using these optimal parameters. We did not optimize any parameters for the
667 logistic regression classifier.

668

669

670

671

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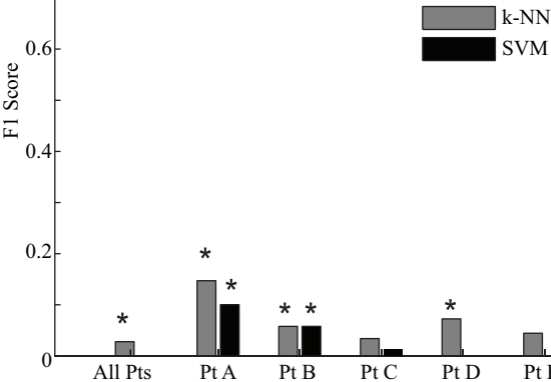
784 2011) for sharing their EEG data and HFO markings on the IEEG-Portal (www.ieeg.org).

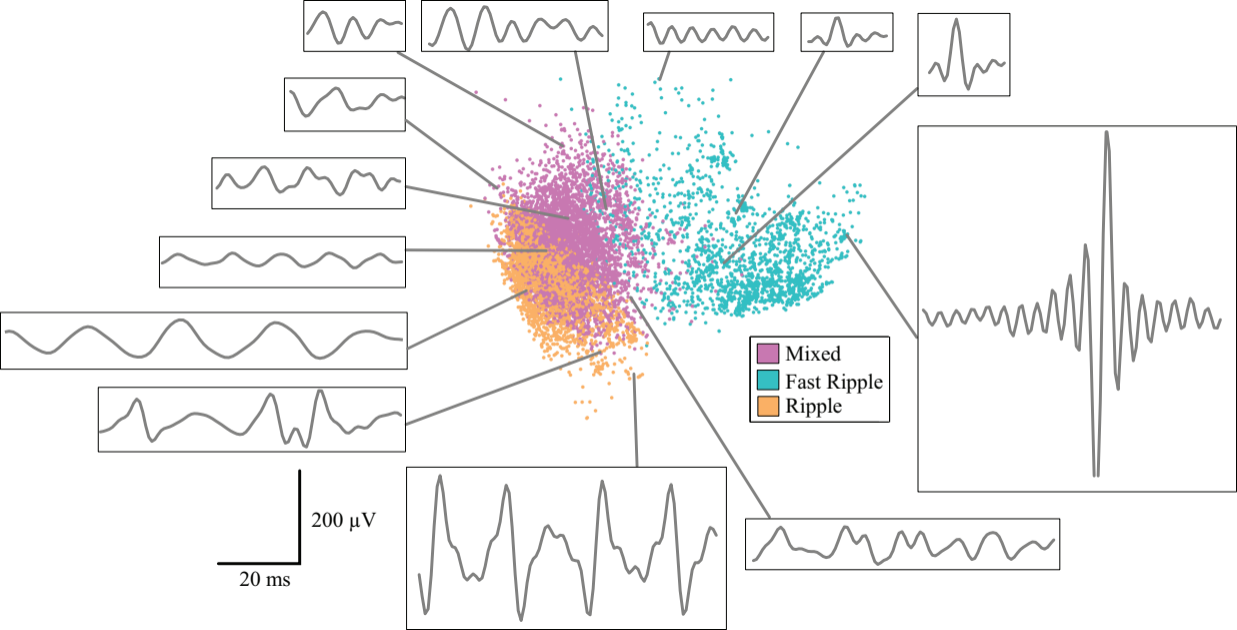
785

786

787

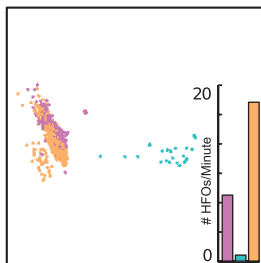
Classifier Performance per Patient



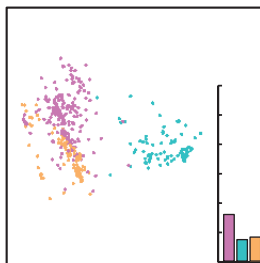


Preictal

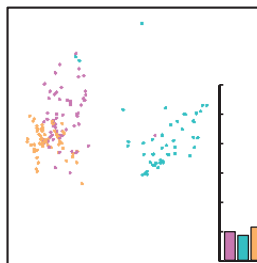
30-20 min
(4 stages combined=40 min total)



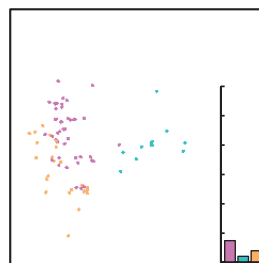
20-10 min
(4 stages)



10-5 min
(4 stages=20 min)

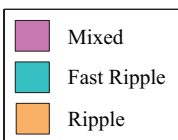
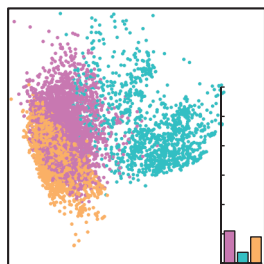


5-0 min
(4 stages)



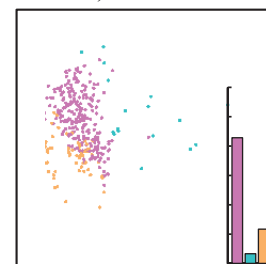
Interictal

7.28 hours total

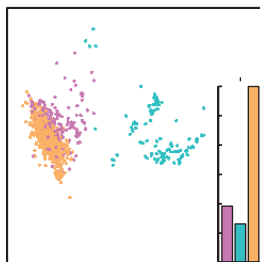


Ictal

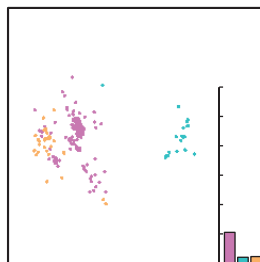
4 seizures, 17.25 minutes total



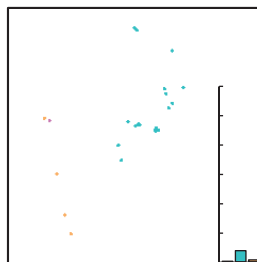
30-20 min
(4 stages)



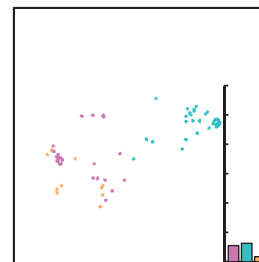
20-10 min
(4 stages)



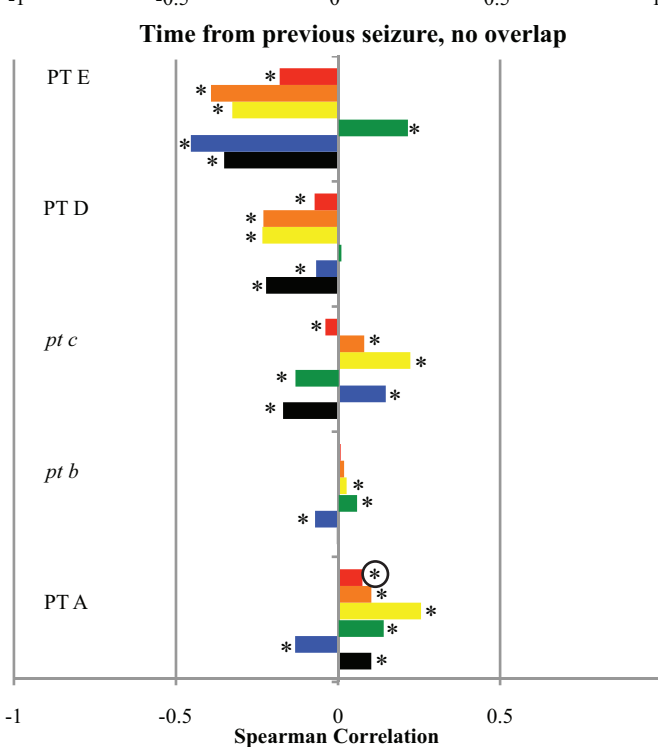
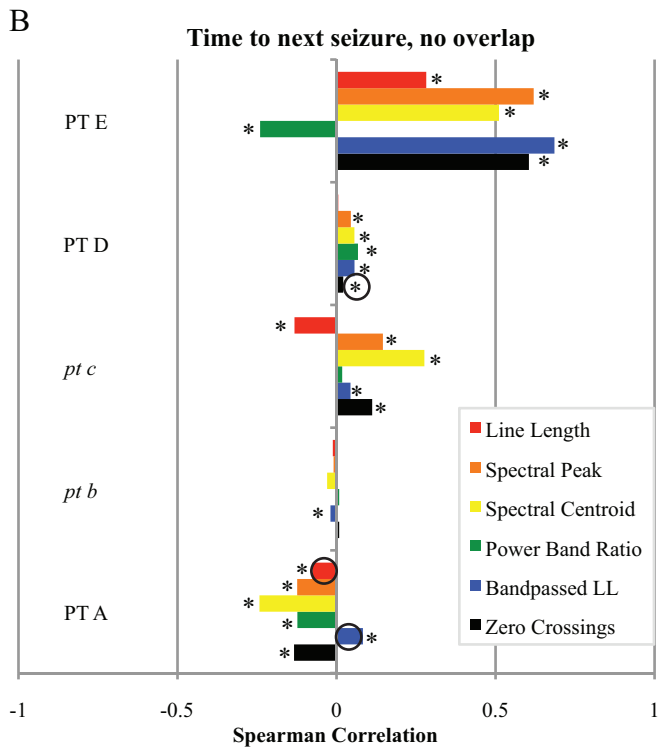
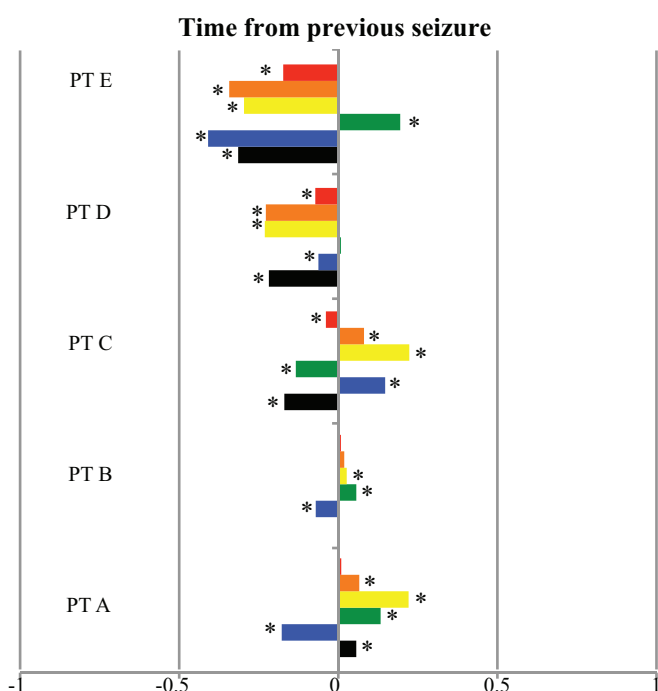
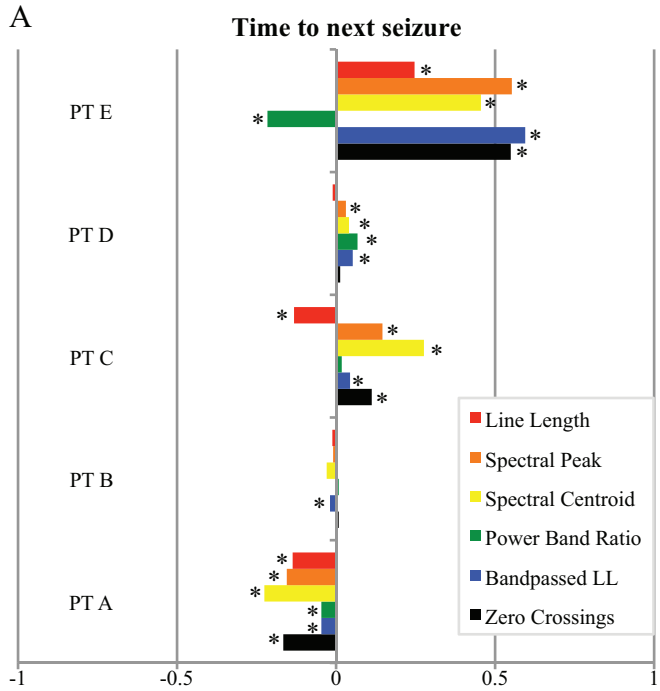
10-5 min
(4 stages)

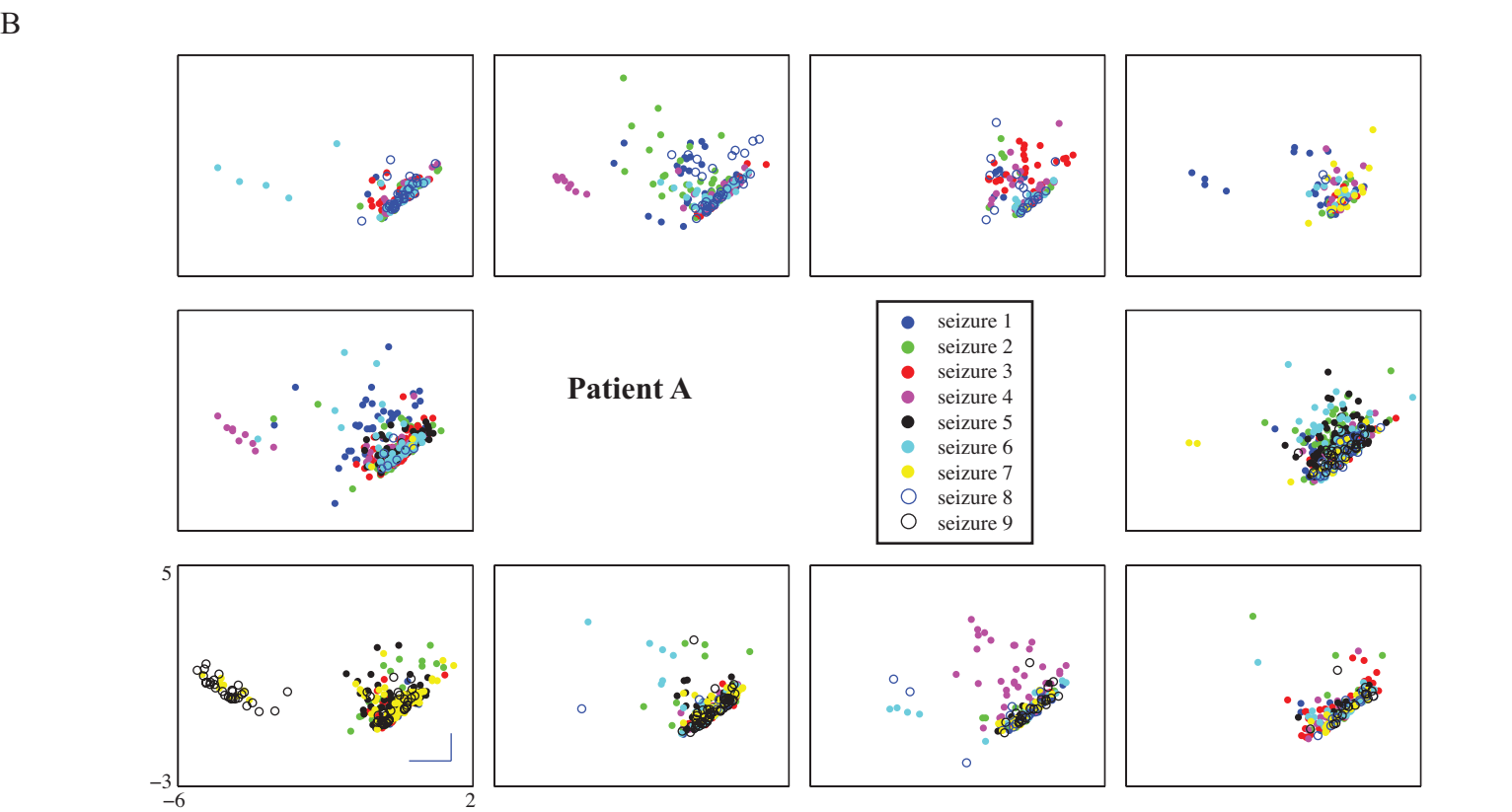
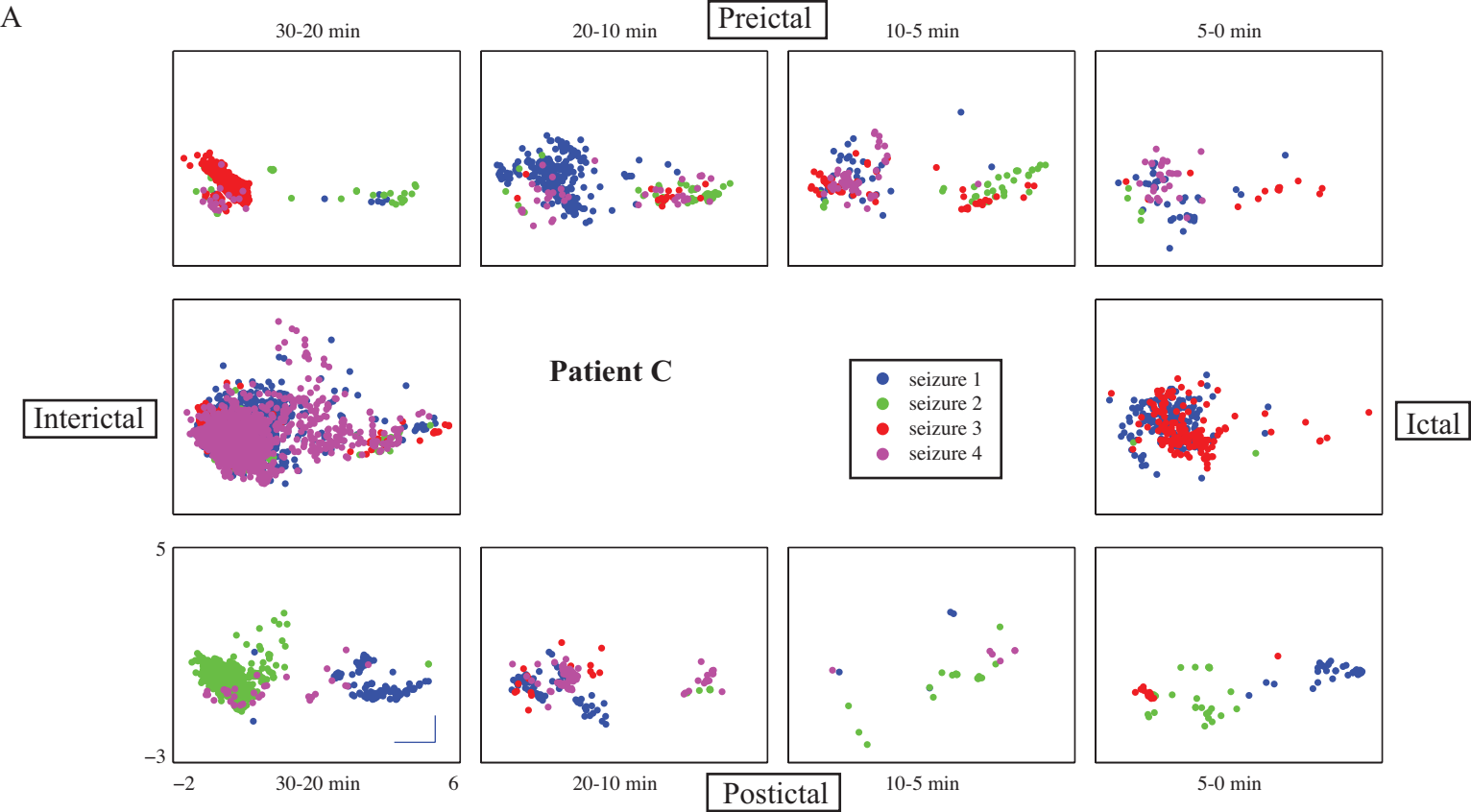


5-0 min
(4 stages)

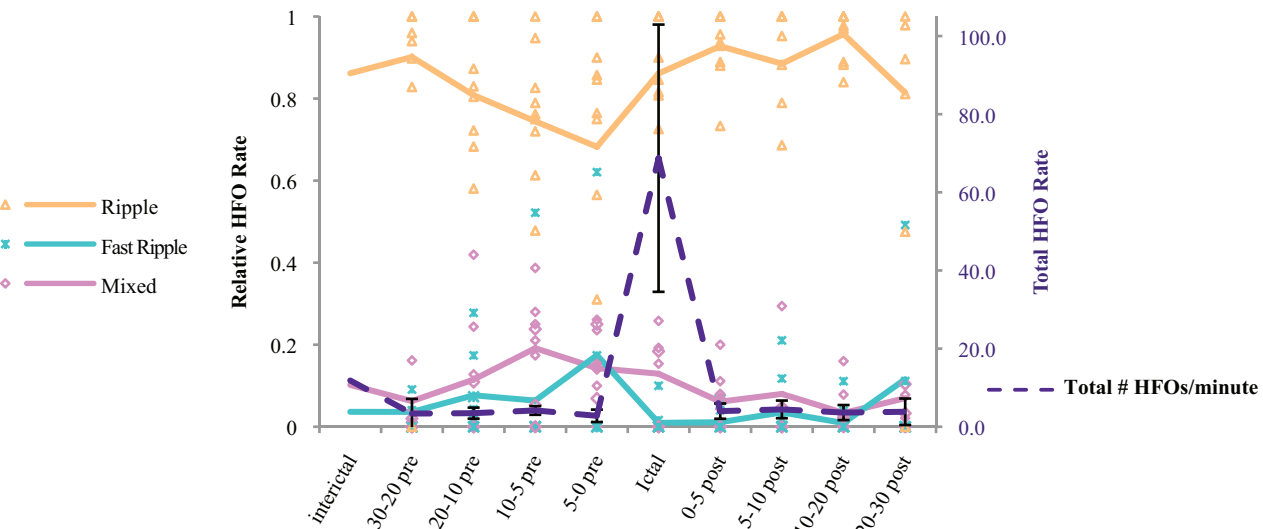


Postictal

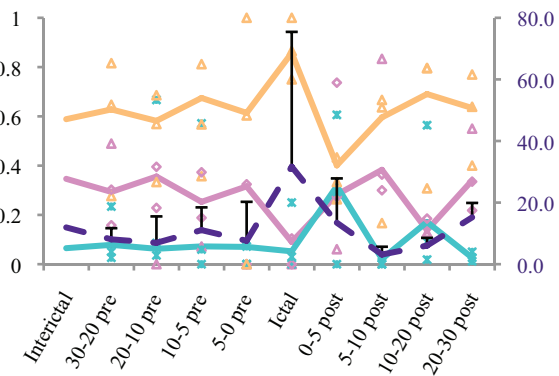




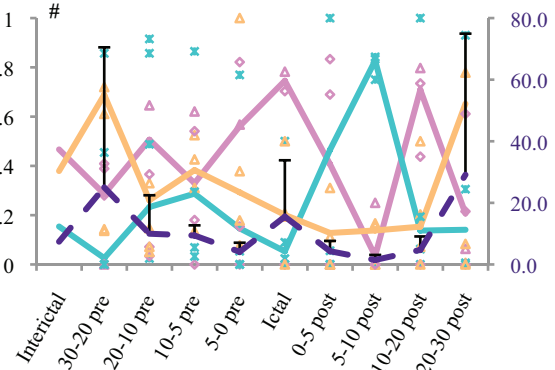
Patient A (9 seizures)



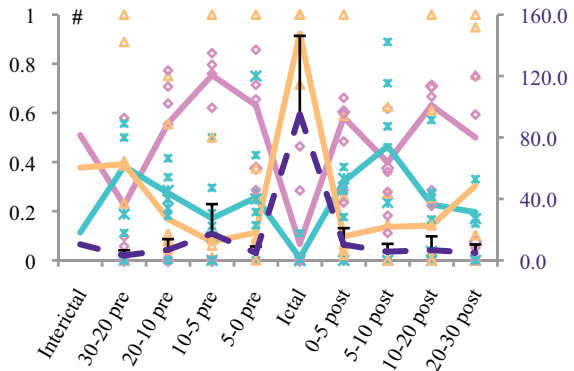
Patient B (3)



Patient C (4)



Patient D (7)



Patient E (2)

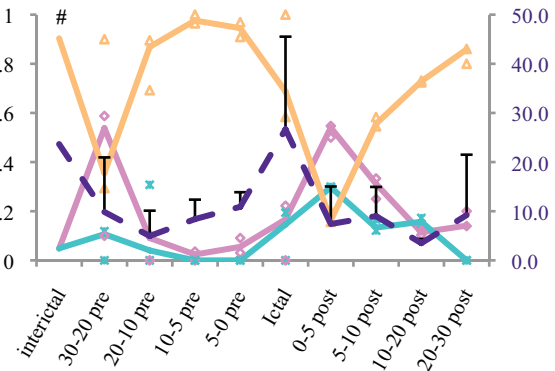


Table 1: Patient summary

Subject	Age	Sex	Pathology	Electrodes	Electrode Placement
A (SZ01)	35	F	Frontal neocortical oligodendroglioma	Standard: 6x6 Micro: 1x8	Right: frontal, motor cortex
B (SZ02)	24	M	Frontal cortical dysplasia	Standard: 6x6 Micro: 4x6	Left: frontal, frontoparietal
C (SZ04)	39	F	Temporal neocortical gliosis	Standard: 6x6, 1x8, 1x8, 1x4, 1x4, 1x4 Micro: 1x4	Left: temporal, inferior frontal, superior frontal, anterior temporal, posterior temporal, anterior, posterior
D (SZ07)	42	F	Frontal cortical dysplasia	Standard: 6x8, 3x8, 1x4 Micro: 1x4, 1x8	Right: frontal, temporal, anterior temporal, inferior temporal, posterior temporal
E (SZ08)	38	F	Temporal neocortical gliosis	Standard: 6x6 Standard depth: 1x4 Micro depth: 1x4, 1x4	Left: temporal, anterior temporal, middle temporal, posterior temporal

Summary of patient clinical characteristics and electrode placement. Parentheses: labels used in prior study, see Table 1 in (Blanco et al. 2011) for further details.

Table 2: HFO and seizure counts

Patient	Total seizures	Total HFOs	Inter-ictal	Preictal	Ictal	Post-ictal
A	9	2552	1487	314	371	380
B	3	9905	9237	280	139	249
C	4	12417	11585	329	317	186
D	7	53015	49519	1787	433	1276
E	2	8262	7210	493	46	513
All	25	86151	79038	3206	1306	2604

Counts of HFOs in each time epoch for dispersion analysis. Preictal and postictal epochs were defined as 10 minutes before and after seizures, respectively.

Table 3: Dispersion p-values from 2 features

	Feature 3 (Line length)					
	Pt A	Pt B	Pt C	Pt D	Pt E	All
Interictal	0.0018	*	*	*	0.0011	*
Preictal	0.3713	0.0615	*	*	0.0002	0.2242
Ictal	0.0426	0.0007	*	0.3148	0.0155	*
Postictal	0.0983	*	*	0.0038	*	*

	Feature 4 (Power Band Ratio)					
	Pt A	Pt B	Pt C	Pt D	Pt E	All
Interictal	*	0.2285	*	*	*	0.1398
Preictal	0.0442	0.4916	*	*	0.4208	*
Ictal	*	0.0001	*	*	*	*
Postictal	*	0.1059	*	*	*	0.3285

Table 3: P-values from the epoch dispersion analysis. Values were calculated by dividing the number of permutation test trials with results more extreme than the true values by the total number of permutation test trials (10,000). Bold text indicates values that were not statistically significant. The combination of all (six) features was significant in all patients. Each patient had different trends with the individual features. Asterisks: highly significant ($p < 1e-6$)

Table 4: P-values of temporal rank

	Time to Next	Time From Previous	Closest cluster (next)
Pt A	<.0001*	<.0001*	FR, M
Pt B	0.013*	<.0001*	M, R
Pt C	<.0001*	<.0001*	FR, R
Pt D	<.0001*	<.0001*	FR, R
Pt E	<.0001*	<.0001*	R

Table 4: P-values from the second Kruskal-Wallis Test, used to assess whether an HFO near a seizure was more likely to be of a specific cluster. Third column indicates which clusters were closest to the next seizure (R: ripple, M: mixed, FR: fast ripple). Asterisks: statistical significance.

