

pdCluster: Partial Discharges Clustering

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```
> library(pdCluster)
```

The set of examples will use a dataset which is loaded with:

```
> load("~/Investigacion/PD/Datos/20100922/DescargasRAW.RData")
```

1 Feature generation

1.1 Prony

A clean partial discharge signal can be regarded as a finite combination of damped complex exponentials. Under this assumption, the so-called Prony's method allows for the estimation of frequency, amplitude, phase and damping components of the signal [KT82, HDS90, KTS84].

Let's use some signals from the dataset (figure 1). The signals contain zeros at the beginning and at the end. The no0 function can remove these parts (figure 2).

```
> signals <- lista[1:25]
```

With these cleaned signals the Prony's method can provide their components (figure 3). Since the number of components must be fixed *a priori*, the function compProny allows the comparison of different numbers (figure 4)

1.2 Feature generation

The pdCluster includes several functions for feature generation. The analysis functions comprises all of them. The results for our example signal are:

```
> analysis(signal)

  RefMax      W1      W2      W3      W4      range   N
1  154 1.585665e-07 2.80655e-05 0.004660942 0.2967786 0.0009498277 323
     energy    nZC   freq1   freq2  damp1  damp2
1 8.301006e-06 0.04643963 3071675 1704751 1140735 3385223
```

This function can be used with a list of signals in order to obtain a matrix of features:

```
> analysisList <- lapply(lista[1:10], analysis)
> pdData <- do.call(rbind, analysisList)
```

Now we need the angle and reflection information, available from another different file. In order to safely share the information, both data frames must be reordered by their energy values:

```
> pdSummary <- read.csv("~/Investigacion/PD/Datos/20100922/descargas.csv")[1:10,
+ ]
> idxOrderSummary = order(pdSummary$sumaCuadrados)
> idxOrderData = order(pdData$energy)
> pdDataOrdered = cbind(pdData[idxOrderData, ], pdSummary[idxOrderSummary,
+   c("angulo", "separacionOriginal")])
```

Later, the data frame to be used with the clustering algorithm has to ordered by time. Thus the samples of the clara method will be random.

```
> idx <- do.call(order, pdSummary[idxOrderSummary, c("segundo",
+   "inicio")])
> pdDataOrdered <- pdDataOrdered[idx, ]
```

We can now construct a PD object¹.

¹The pdCluster package is designed with S4 classes and methods. Two classes have been defined: PD and PDCluster.

```

> p <- xyplot(signals, y.same = NA, FUN = function(x) {
+   xyplot(ts(x))
+ })
> print(p)

```

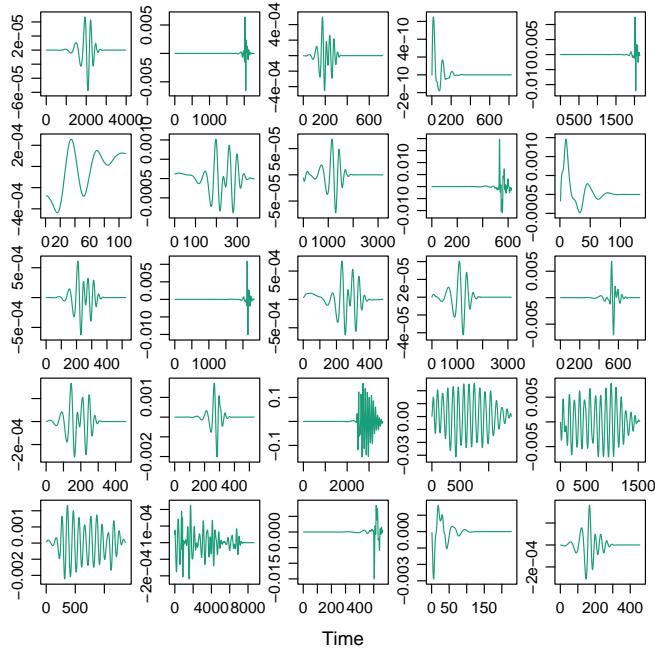


Figure 1: Partial discharge signals

```

> p <- xyplot(signals, y.same = NA, FUN = function(x) {
+   xyplot(ts(no0(x)))
+ })
> print(p)

```

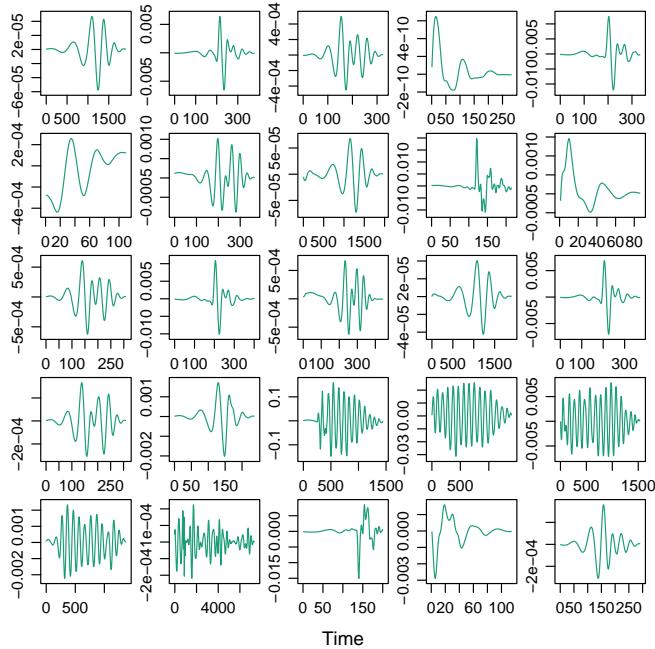


Figure 2: Partial discharge signals after a threshold cleaning

```

> signal <- signals[[3]]
> pr <- prony(signal, M = 10)
> print(xyplot(pr))

```

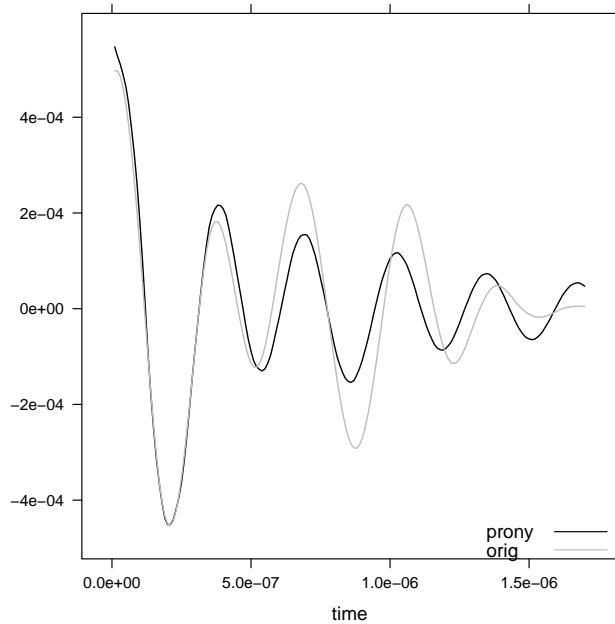


Figure 3: Prony's method results

```

> p <- compProny(signal, M = c(10, 20, 30, 40))
> print(p)

```

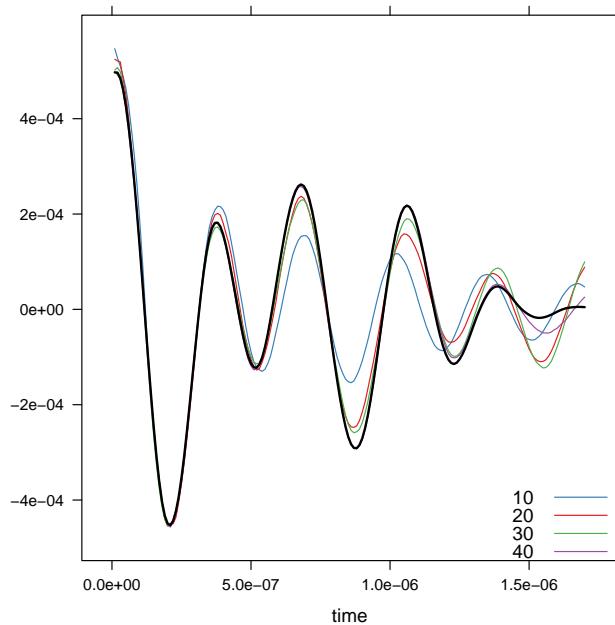


Figure 4: Comparison of different Prony decompositions

```

> pd <- df2PD(pdDataOrdered)
> pd

Object of class PD

Source of measurements:
Number of observations: 10
Filtered?: FALSE
Filter: '<undef>'()
Transformed?: FALSE

Data:
  RefMax      W1      W2      W3
Min. : 10.0 Min. :8.417e-14 Min. :1.487e-11 Min. :2.875e-09
1st Qu.: 56.5 1st Qu.:3.521e-08 1st Qu.:6.808e-06 1st Qu.:1.158e-03
Median : 176.0 Median :2.128e-07 Median :2.981e-05 Median :4.565e-03
Mean   : 320.0 Mean   :2.423e-04 Mean   :6.606e-03 Mean   :3.205e-02
3rd Qu.: 212.0 3rd Qu.:5.609e-07 3rd Qu.:9.860e-05 3rd Qu.:9.841e-03
Max.   :1152.0 Max.   :2.234e-03 Max.   :6.465e-02 Max.   :2.749e-01
      W4      range      N      energy
Min. :0.0000000 Min. :8.272e-10 Min. : 86.0 Min. :9.544e-18
1st Qu.:0.0007306 1st Qu.:2.792e-04 1st Qu.: 230.8 1st Qu.:2.038e-06
Median :0.2051125 Median :1.428e-03 Median : 328.0 Median :1.208e-05
Mean   :0.1893751 Mean   :7.392e-03 Mean   : 594.7 Mean   :6.944e-04
3rd Qu.:0.3037326 3rd Qu.:1.021e-02 3rd Qu.: 374.5 3rd Qu.:6.760e-04
Max.   :0.5386891 Max.   :3.070e-02 Max.   :1954.0 Max.   :3.282e-03
      nZC      freq1      damp1
Min. :0.005249 Min. : 353312 Min. : 175314
1st Qu.:0.020226 1st Qu.:1370276 1st Qu.: 679159
Median :0.030888 Median :2731837 Median :2755076
Mean   :0.033111 Mean   :2210122 Mean   :2320301
3rd Qu.:0.044445 3rd Qu.:3066133 3rd Qu.:3635957
Max.   :0.070093 Max.   :3113590 Max.   :4715804
Number of reflections: 6

```

The results of analysis to the whole dataset are available with:

```

> load("~/Investigacion/PD/Datos/20100922/dfHibr17112010.RData")
> dfHibr <- df2PD(dfHibr)
> dfHibr

Object of class PD

Source of measurements:
Number of observations: 9955
Filtered?: FALSE
Filter: '<undef>'()
Transformed?: FALSE

Data:
  RefMax      W1      W2      W3
Min. : 0.0 Min. :0.000e+00 Min. :0.000e+00 Min. :0.000e+00
1st Qu.: 57.0 1st Qu.:6.838e-14 1st Qu.:1.228e-11 1st Qu.:2.378e-09
Median : 205.0 Median :8.011e-13 Median :1.250e-10 Median :1.730e-08
Mean   : 559.4 Mean   :1.197e-03 Mean   :2.366e-03 Mean   :1.352e-02
3rd Qu.: 896.0 3rd Qu.:1.104e-07 3rd Qu.:1.740e-05 3rd Qu.:2.009e-03
Max.   :34079.0 Max.   :2.973e-01 Max.   :1.165e-01 Max.   :5.869e-01
      W4      range      N      energy
Min. :0.000e+00 Min. :0.000e+00 Min. : 0 Min. :0.000e+00
1st Qu.:3.341e-07 1st Qu.:2.265e-05 1st Qu.: 172 1st Qu.:2.833e-08
Median :1.103e-06 Median :1.064e-04 Median : 417 Median :5.665e-07
Mean   :7.648e-02 Mean   :5.624e-03 Mean   : 992 Mean   :4.023e-02
3rd Qu.:3.686e-02 3rd Qu.:1.026e-03 3rd Qu.: 1568 3rd Qu.:1.307e-05
Max.   :1.430e+00 Max.   :3.108e-01 Max.   :34234 Max.   :5.459e+00
      nZC      freq1      damp1
Min. :0.0000000 Min. : 0 Min. : 0
1st Qu.:0.003629 1st Qu.: 299379 1st Qu.: 178107
Median :0.005682 Median : 384840 Median : 427577
Mean   :0.013704 Mean   : 1250969 Mean   : 1807198
3rd Qu.:0.021283 3rd Qu.: 1849583 3rd Qu.: 1625622
Max.   :0.147059 Max.   :50000000 Max.   :205053894
Number of reflections: 5626

```

2 Transformations

Prior to the clustering algorithm, the feature matrix has to be filtered:

```
> dfFilter <- filterPD(dfHibr)
```

and transformed [BC64]:

```

> dfTrans <- transformPD(dfFilter)
> dfTrans

Object of class PD

Source of measurements:
Number of observations: 3695
Filtered?: TRUE
Filter: N > 10 & nZC > 0 & W4 > 0 & freq1 > 1 & freq1 < 2e+07 & !refl
Transformed?: TRUE

Data:
    RefMax          W1          W2          W3
Min.   : 0.000  Min.  :-53.341  Min.  :-38.720  Min.  :-27.9936
1st Qu.: 6.809  1st Qu.: 45.027  1st Qu.: 29.386  1st Qu.: 20.6000
Median : 7.896  Median : -26.477  Median : -18.205  Median : -13.0398
Mean   : 8.387  Mean   : -30.009  Mean   : -19.236  Mean   : -12.7681
3rd Qu.:10.407 3rd Qu.: -18.402  3rd Qu.: -10.554  3rd Qu.: -5.1754
Max.   :18.979  Max.   : -1.496  Max.   : -2.249  Max.   : -0.5339
      W4        range          N          energy
Min.  :-20.5419  Min.  :-31.952  Min.  : 3.381  Min.  :-59.712
1st Qu.: -14.3313 1st Qu.: -12.589  1st Qu.: 3.934  1st Qu.: 19.287
Median : -7.3826  Median : -10.500  Median : 4.171  Median : 16.453
Mean   : -8.1065  Mean   : -9.551  Mean   : 4.217  Mean   : 14.776
3rd Qu.: -1.2579  3rd Qu.: -6.206  3rd Qu.: 4.561  3rd Qu.: 8.933
Max.   : 0.1203  Max.   : -1.210  Max.   : 5.339  Max.   : 1.650
      nZC       freq1       damp1
Min.  :-12.686  Min.  : 4.996  Min.  : 6.581
1st Qu.: -9.468  1st Qu.: 5.131  1st Qu.: 10.800
Median : -8.107  Median : 5.226  Median : 11.327
Mean   : -7.424  Mean   : 5.216  Mean   : 11.527
3rd Qu.: -5.080  3rd Qu.: 5.313  3rd Qu.: 12.434
Max.   : -3.268  Max.   : 5.433  Max.   : 14.677
Number of reflections: 0

```

The figure 5 compares the datasets after and before of the transformations:

```

> nZCbefore <- as.data.frame(dfFilter)$nZC
> nZCafter <- as.data.frame(dfTrans)$nZC
> comp <- data.frame(After = nZCafter, Before = nZCbefore)

```

The filterPD method is a wrapper for the general subset method. With `subset` it is possible to extract a group of samples based on a condition and select only certain columns. The figure 6 shows a scatterplot matrix with `splom` of this subset. The `splom` graphical tool is explained in the next section.

```

> dfTransSubset <- subset(dfTrans, subset = (angle >= 90 & angle <=
+     180), select = c(energy, W1, nZC))
> dfTransSubset

Object of class PD

Source of measurements:
Number of observations: 624
Filtered?: TRUE
Filter: N > 10 & nZC > 0 & W4 > 0 & freq1 > 1 & freq1 < 2e+07 & !refl
Transformed?: TRUE

Data:
    energy          W1          nZC
Min.  :-42.785  Min.  :-52.106  Min.  :-12.154
1st Qu.: -19.589  1st Qu.: -45.646  1st Qu.: -9.568
Median : -17.837  Median : -41.668  Median : -9.184
Mean   : -15.985  Mean   : -34.488  Mean   : -8.000
3rd Qu.: -12.644  3rd Qu.: -20.409  3rd Qu.: -6.064
Max.   :  1.650  Max.   : -1.743  Max.   : -3.896
Number of reflections: 0

```

2.1 Graphical tools

The pdCluster packages includes a set of graphical exploratory tools, such as a scatterplot matrices with hexagonal binning [CLNL87] (figure 7), density plots (figure 8), histograms (figure 9) or phase resolved partial discharge patterns, both with partial transparency (figure 10) or hexagonal binning (figure 11).

3 Clustering

The filtered and transformed object can now be used with the clustering algorithm [SHR97]. The results are displayed with a phase resolved pattern with clusters in separate panels in the figure 12. The colors encode the distance of each point to the *medoid* of its cluster. The figure 13 displays the same pattern with superposed clusters. Here the colors encode the membership to a certain cluster, and transparency is used to denote density of points in a region.

```

> h <- histogram(~After + Before, data = comp, scales = list(x = list(relation = "free"),
+   y = list(relation = "free", draw = FALSE)), breaks = 100,
+   col = "gray", xlab = "", strip.names = c(TRUE, TRUE), bg = "gray",
+   fg = "darkblue")
> print(h)

```

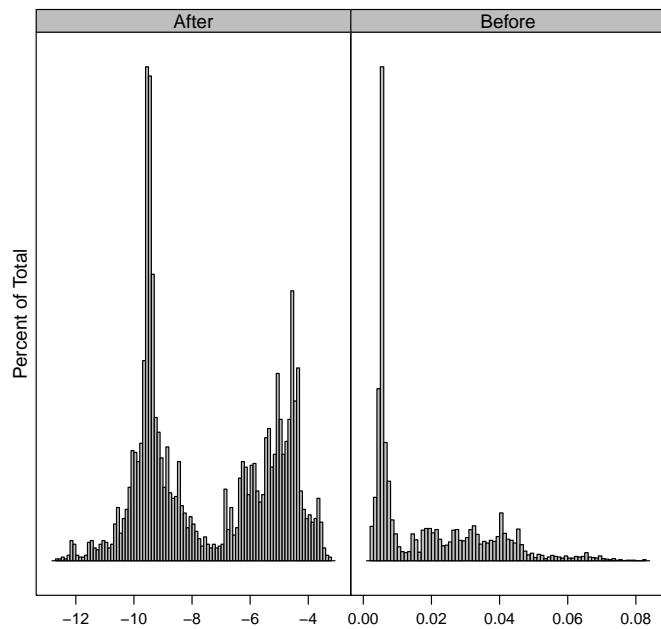


Figure 5: Histogram of a collection of partial discharges

```

> p <- splom(dfTransSubset)
> print(p)

```

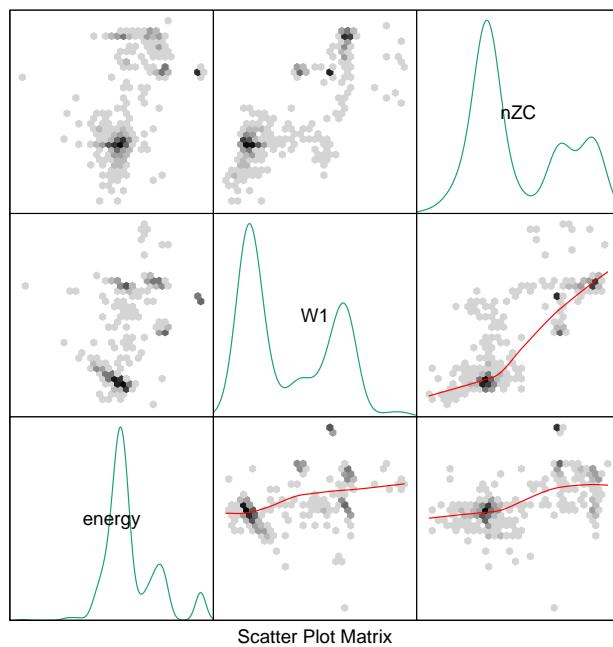


Figure 6: Scatterplot matrix of a subset of dfTrans.

```
> p <- splom(dfTrans)
> print(p)
```

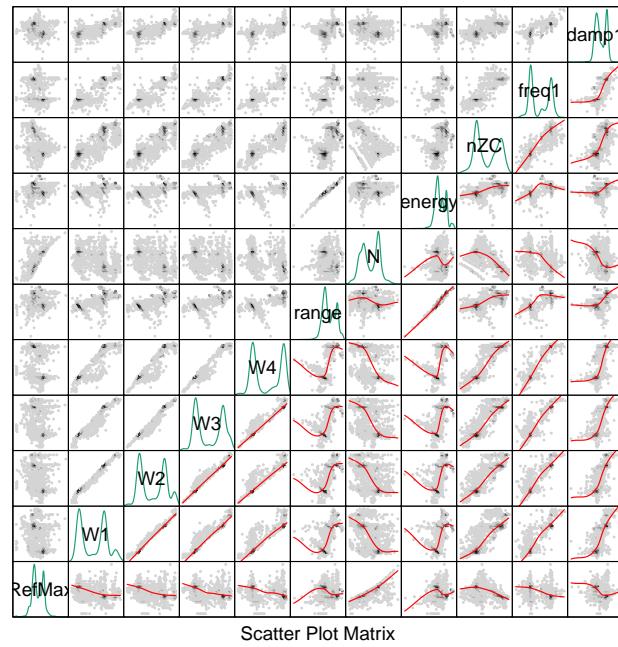


Figure 7: Scatterplot matrix of a collection of partial discharges

```
> p <- densityplot(dfTrans)
> print(p)
```

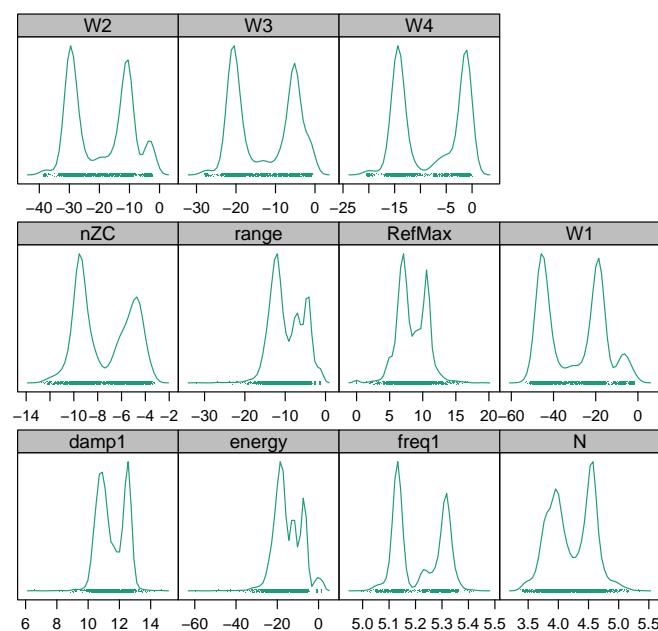


Figure 8: Density plot of a collection of partial discharges

```
> p <- histogram(dfTrans)
> print(p)
```

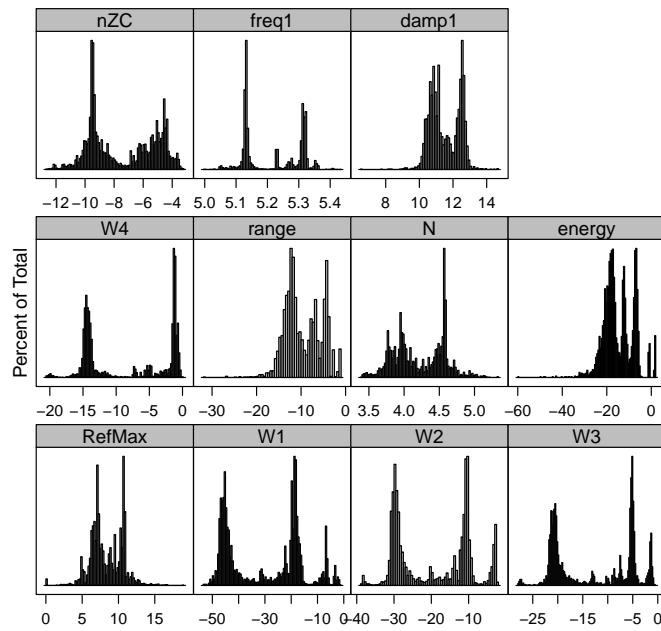


Figure 9: Histograms of a collection of partial discharges

```
> p <- xyplot(dfTrans)
> print(p)
```

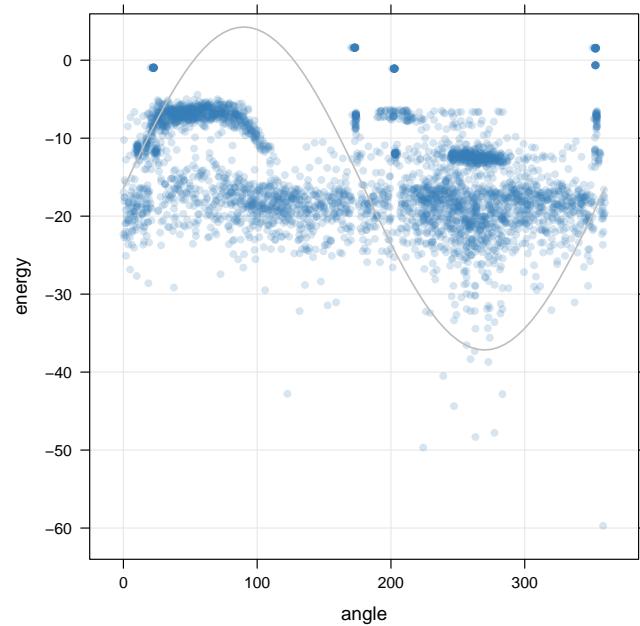


Figure 10: Partial discharge phase resolved pattern.

The results can be easily understood with the density plots of each cluster and feature (figure 14) or with the histograms (figure 15).

```
> dfTransCluster <- claraPD(dfTrans, noise.level = 0.7, noise.rm = TRUE)
> dfTransCluster

Object of class PD

Source of measurements:
Number of observations: 2643
Filtered?: TRUE
Filter: N > 10 & NZC > 0 & W4 > 0 & freq1 > 1 & freq1 < 2e+07 & !refl
Transformed?: TRUE

Data:
  RefMax          W1          W2          W3
Min.   : 1.972  Min.  :-53.341  Min.  :-38.720  Min.  :-27.994
1st Qu.: 7.081  1st Qu.:-45.517  1st Qu.:-29.706  1st Qu.:-20.887
Median : 9.025  Median :-35.255  Median :-24.944  Median :-19.066
Mean   : 8.841  Mean   :-31.585  Mean   :-20.194  Mean   :-13.379
3rd Qu.:10.612 3rd Qu.:-18.438  3rd Qu.:-10.554  3rd Qu.:-5.171
Max.   :16.753  Max.   :-6.286   Max.   :-2.715   Max.   :-1.162
      W4          range          N          energy
Min.  :-20.54193  Min.  :-18.600  Min.  :3.381  Min.  :-30.428
1st Qu.:-14.48318 1st Qu.:-12.492  1st Qu.:3.943  1st Qu.:-19.013
Median :-13.66034  Median :-10.973  Median :4.421  Median :-16.647
Mean   :-8.43715  Mean   :-9.288  Mean   :4.272  Mean   :-14.145
3rd Qu.:-1.23210  3rd Qu.:-4.736  3rd Qu.:4.569  3rd Qu.:-7.572
Max.   :-0.09173  Max.   :-1.210  Max.   :5.202  Max.   : 1.650
      NZC          freq1          damp1
Min.  :-12.192  Min.  :5.005  Min.  : 6.581
1st Qu.: -9.523  1st Qu.:5.131  1st Qu.:10.706
Median : -8.822  Median :5.139  Median :11.135
Mean   : -7.414  Mean   :5.211  Mean   :11.460
3rd Qu.:-4.954  3rd Qu.:5.312  3rd Qu.:12.475
Max.   : -3.268  Max.   :5.329  Max.   :13.115
Number of reflections: 0
Number of clusters: 7
Number of elements per cluster:
  1   2   3   4
1383 696 420 144

Metric: manhattan
Number of simulations: 25
Noise level: 0.7
Distances
  dist          distRel          distFactor
Min.   : 0.1020  Min.  :0.0000  Min.  : 1.000
1st Qu.: 2.1595  1st Qu.:0.7387  1st Qu.: 3.000
Median : 4.9140  Median :0.8760  Median : 5.000
Mean   : 6.8265  Mean   :0.8294  Mean   : 4.977
3rd Qu.:11.0155  3rd Qu.:0.9463  3rd Qu.: 7.000
Max.   :50.0681  Max.   :0.9975  Max.   : 9.000
NA's    :263.000
```

References

- [BC64] G. E. P. Box and D. R. Cox. An analysis of transformations. *Journal of the Royal Statistical Society. Series B (Methodological)*, 26(2):pp. 211–252, 1964.
- [CLNL87] D. B. Carr, R. J. Littlefield, W. L. Nicholson, and J. S. Littlefield. Scatterplot matrix techniques for large n. *Journal of the American Statistical Association*, 82(398):pp. 424–436, 1987.
- [HDS90] J.F. Hauer, C.J. Demeure, and L.L. Scharf. Initial results in prony analysis of power system response signals. *Power Systems, IEEE Transactions on*, 5(1):80 –89, feb 1990.
- [KT82] R. Kumaresan and D. Tufts. Estimating the parameters of exponentially damped sinusoids and pole-zero modeling in noise. *Acoustics, Speech and Signal Processing, IEEE Transactions on*, 30(6):833 – 840, dec. 1982.
- [KTS84] R. Kumaresan, D.W. Tufts, and L.L. Scharf. A prony method for noisy data: Choosing the signal components and selecting the order in exponential signal models. *Proceedings of the IEEE*, 72(2):230 – 233, feb. 1984.
- [SHR97] Anja Struyf, Mia Hubert, and Peter Rousseeuw. Clustering in an object-oriented environment. *Journal of Statistical Software*, 1(4):1–30, 2 1997.

```
> p <- hexbinplot(dfTrans)
> print(p)
```

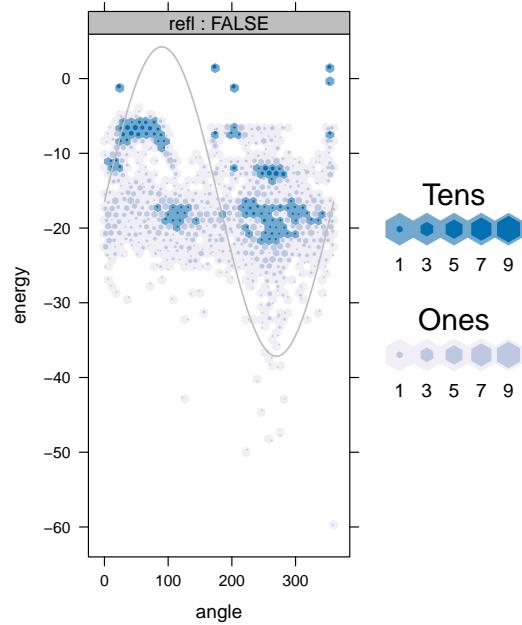


Figure 11: Partial discharge phase resolved pattern with hexbinplot

```
> p <- xyplot(dfTransCluster)
> print(p)
```

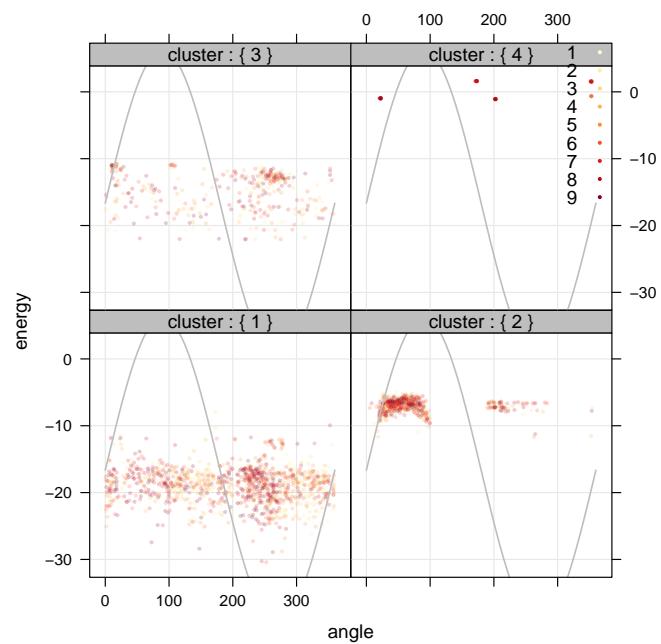


Figure 12: Partial discharge phase resolved pattern with clusters in separate panels.

```
> p <- xyplot(dfTransCluster, panelClust = FALSE)
> print(p)
```

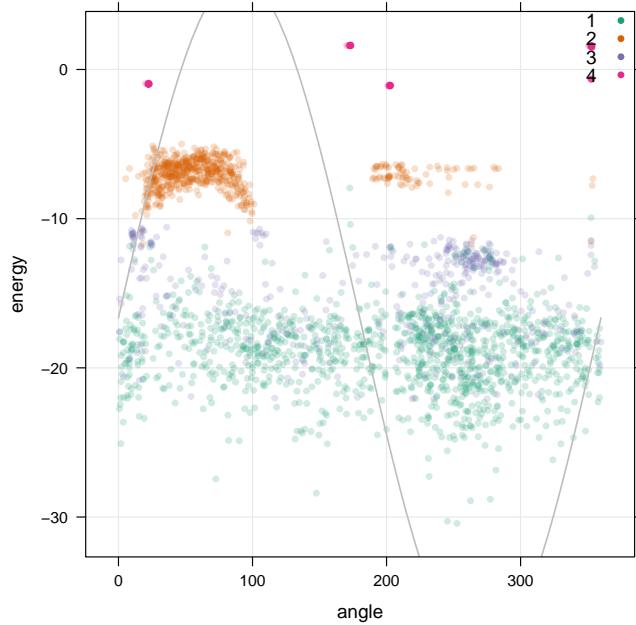


Figure 13: Partial discharge phase resolved pattern with clusters marked with colors.

```
> p <- densityplot(dfTransCluster)
> print(p)
```

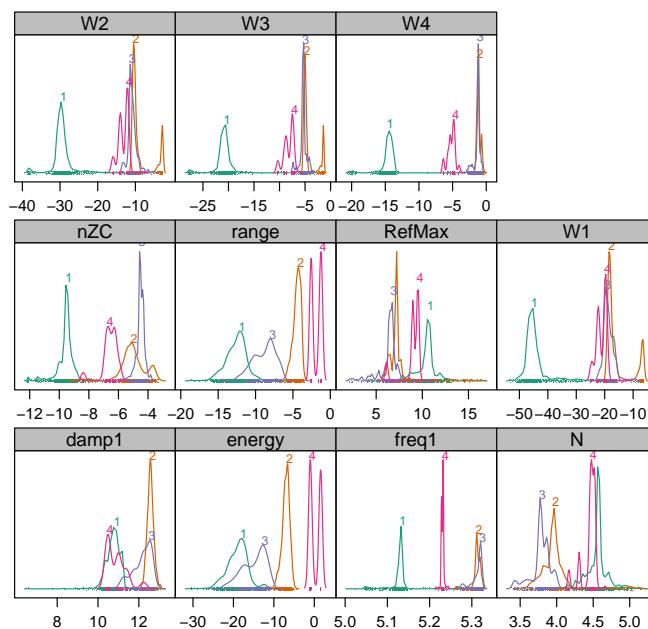


Figure 14: Density plot of the clusters of partial discharges

```
> p <- histogram(dfTransCluster)
> print(p)
```

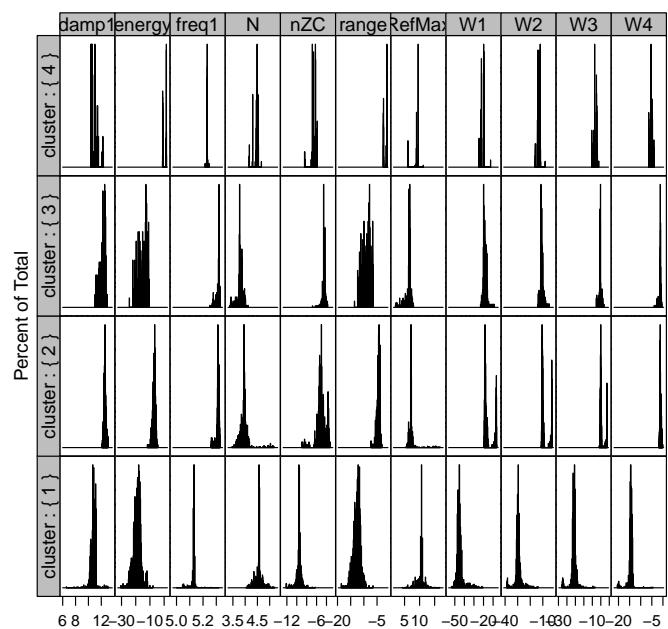


Figure 15: Histograms of the clusters of partial discharges