

advanced biosignal processing and analysis

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## To the Reader

Welcome to g.tec's world of medical and electrical engineering! Discover the only professional biomedical signal processing platform under MATLAB and Simulink. Your ingenuity finds the appropriate tools in the g.tec elements and systems. Choose and combine flexibly the elements for biosignal amplification, signal processing and stimulation to perform even real-time feedback.

Our team is prepared to find the better solution for your needs.

Take advantage of our experience!

Dr. Christoph Guger

Dr. Guenter Edlinger

### **Researcher and Developer**

Reduce development time for sophisticated real-time applications from month to hours. Integrate g.tec's open platform seamlessly into your processing system. g.tec's rapid prototyping environment encourages your creativity.

### Scientist

Open new research fields with amazing feedback experiments. Process your EEG/ECG/EMG/EOG data with g.tec's biosignal analyzing tools. Concentrate on your core problems when relying on g.tec's new software features like ICA, AAR or online Hjorth's source derivation.

### Study design and data analysis

You are planning an experimental study in the field of brain or life sciences? We can offer consultation in experimental planning, hardware and software selection and can even do the measurements for you. If you have already collected EEG/ECG/EMG/EOG, g.tec can analyze the data starting from artifact control, do feature extraction and prepare the results ready for publication.

# Preface

This section includes the following topics:

**Required Products** 

<u>Using This Guide</u> - Suggestions for reading the handbook

<u>Conventions</u> - Text formats in the handbook

# **Required Products**

g®.CLASSIFYtoolbox uses:

**g®.BSanalyze** – the advanced biosignal analysis software package from g.tec

 $\ensuremath{\textbf{MATLAB}}\xspace -$  as basic matrix operation platform

Signal Processing Toolbox - to give access to standard signal analysis tools

# Using This Guide

"<u>Generating a Feature Matrix</u>" shows how to extract specific features that are used for the classification task.

Chapter "<u>Generating a Classifier</u>" demonstrates the set-up of linear and non-linear classifiers and shows how to classify the feature matrix. Linear Discriminant Analysis (LDA), Minimum Distance Classifier (MDC), Multi-layer Perceptron (MLP), Radial Basis Function (RBF) and Distinction Sensitive Learning Vector Quantization (DSLVQ) are explained.

"<u>Receiver Operator Curve</u>" demonstrates the computation of the sensitivity and specificity and presentation as ROC curve.

Chapter "KMEANS Clustering" demonstrates the clustering on a three-class problem.

"DSLVQ Feature Weighting" explains the necessary steps for feature selection based on the DSLVQ algorithm.

"<u>Using the Classifier</u>" explains how to apply already calculated classifiers on new data and how to test a classifier on new data.

"<u>Data Access</u>" shows how to access the feature matrix and classifier objects from the MATLAB command line.

Chapter "<u>Help</u>" explains the usage of the on-line help, the printable documentation and the function help.

Chapter "<u>Batch-Mode</u>" shows how to use the g.BSanalyze commands from the MATLAB command line.

# Conventions

| Item             | Format          | Example                              |
|------------------|-----------------|--------------------------------------|
| MATLAB code      | Courier         | to start simulink, type<br>simulink  |
| String variables | Courier italics | <pre>set(P_C, 'PropertyName',)</pre> |
| Menu items       | Boldface        | Select Save from the File menu.      |

# Hardware and Software Requirements

For Hardware and Software Requirements see the g.BSanalyze manual.

# **Generating a Feature Matrix**

The first step in generating a feature matrix is to calculate specific features (e.g. bandpower values) with the Parameter Extraction methods of g.BSanalyze. There are three options available:

- Trial attributes extract the features of trials with specific attributes
- Time points extract the features of the signals at specific time points
- Time segments extract the features of the signals of specific segments

Perform the following steps:

1. After starting MATLAB and setting the correct path, type:

gbsanalyze

into the MATLAB command line

| 📣 Command Window | - | × |
|------------------|---|---|
| fx >> gbsanalyze |   | ۲ |
|                  |   |   |
|                  |   |   |
|                  |   |   |
|                  |   |   |
|                  |   |   |
|                  |   |   |
|                  |   |   |
|                  |   |   |
|                  |   |   |

g.BSanalyze starts with a blank data window

2. Select Load Data under the File menu and open the file session1234triggered.mat from the following directory:

Documents\gtec\gBSanalyze\testdata\BCI

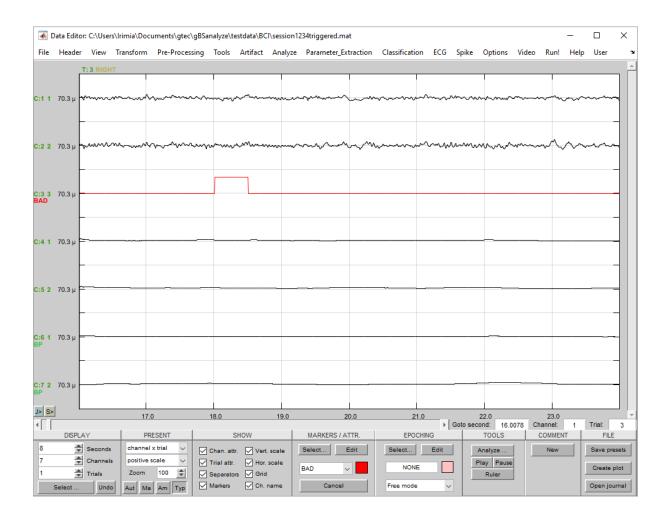
The Data Editor shows a brain-computer interface (BCI) experiment data-set with 2 EEG channels and 1 trigger channel. The first channel was recorded from channel C3, the second channel from channel C4. The paradigm is described in detail in the g.BSanalyze documentation in chapter **Data-sets** – **Movement Imagination**.

3. To extract relevant information out of the raw EEG data select **Bandpower** under the **Parameter Extraction** menu. Select the ALPHA filter to calculate the bandpower between 7 and 13 Hz and set the **Length** of the averaging window to 128 samples with an **Overlap** of 127 samples.

| 承 Bandpower  |   |                   |                   | _              |                   | ×   |
|--|---|-------------------|-------------------|----------------|-------------------|-----|
| Calculate the bandpower b<br>data.   |   |                   | raging over a spe | ecific wind    | dow of the        | •   |
| _ Select CHANNELS:   |   |                   |                   | Select         | channels          |     |
| □ Specify METHOD and PARAM   | Meters:                                   |                   |                   |                |                   |     |
| Choose a filter for the estimation:  | ALPHA / BP / 7 / 13<br>name / type / f (k |                   | ∼<br>order        | Design         | new filter        | ·   |
| Specify averaging window:  | Length: 1000<br>128                       | [ms]<br>[samples] | Overlap           | 992.187<br>127 | 5 [ms]<br>[sample | es] |
|  |   |                   |                   |                |                   |     |
| Result procedure: <ul> <li>Add new</li> <li>Replace</li> <li>Save res</li> </ul> | all channels                              | Auto<br>Filename: | matic treemaker i | is: enat       |                   | ~   |
|  |   | Help              | Cancel            |                | Start !           | !   |

- 4. Press the **Select channels** button and select only the two EEG channels 1 and 2 for the operation
- 5. Chose Add new channels to append the bandpower values to the raw data
- 6. Press **Start** to perform the operation
- 7. Repeat steps 4 to 7 with the BETA-3 filter (14 to 20 Hz)

8. After finishing the calculation the Data Editor visualizes the newly created features. Channels 4 and 5 represent the bandpower in the alpha range and channels 6 and 7 the bandpower in the beta range.



# **Trial Attributes**

Trial Attributes allows to extract features of trials which correspond to a specific class. The class is assigned with trial attributes in g.BSanalyze (such as right or left hand movement imagination).

1. Open the Feature Matrix window from the Classification menu

| 承 Feature Matrix      |  |              |           |          |             |              | _          |             | ×  |
|-----------------------|--|--------------|-----------|----------|-------------|--------------|------------|-------------|----|
|                       | feature matrix as input f<br>ch attribute corresponds<br>s to a class. |              |           |          |             |              |            |             | nt |
| C Specify CLASSIF     | ICATION INTERVAL   | L:           |           |          |             |              |            |             |    |
| Start at: 1000        | [ms] Step:   | 1000 [       | [ms]      | Stop at: | 8000        | [ms]         | Merge      | e time poin | ts |
| 128                   | [samples]  | 128 [        | [samples] |          | 1024        | [samples]    |            |             |    |
| - Specify CLASS L     | ABELS / TIME POI   | NT:          |           |          |             |              |            |             | _  |
| Select class:         | ARTIFACT   | ^            |           | Select   | time point: | 1000         |            |             |    |
|                       | REMOVE<br>LEFT   | - 11         |           |          |             | 2000<br>3000 |            |             |    |
|                       | RIGHT  |              |           |          |             | 4000         |            |             |    |
|                       |  |              |           |          |             | 5000         |            |             |    |
|                       |  |              |           |          |             | 6000<br>7000 |            |             |    |
|                       |  | ¥            |           |          |             | 8000         |            | `           |    |
| - Select FEATURE      | CHANNELS:  |              |           |          |             |              |            |             |    |
|                       |  |              |           |          |             | Select fe    | eature cha | innels      |    |
| Choose METHOD         | and OPTIONS:   |              |           |          |             |              |            |             | _  |
| Classification method | d: Linear Discriminant A   | nalvsis (LDA | 0         | ~        |             |              |            |             |    |
|                       |  |              | •/        |          |             |              |            |             |    |
| Randomly permuta      | ate the matrix   |              |           |          |             |              |            |             |    |
|                       |  |              |           |          |             |              |            |             |    |
| Result procedure:     | ✓ Classify data  |              |           |          | Automat     | ic treemaker | is: enable | ed N        |    |
|                       | Save results   |              | Filenam   | e:tdat   | a\BCI\featu | rematrix\ban | dpower.ma  | at          | ]  |
|                       |  |              |           | Hel      | p           | Cancel       |            | Start       |    |

- 2. The CLASSIFICATION INTERVAL should Start at 1000 ms with a Step size of 1000 ms and Stop at 8000 ms. These settings extract for each time point (1000, 2000, ... 8000 ms) the corresponding features.
- 3. Chose classes LEFT and RIGHT to extract only trials with these trial attributes

4. Press the **Select features channels** ... button and select the 4 bandpower channels 4, 5, 6 and 7

| Select channels                                 |   | _                                   |      |
|---|---|-------------------------------------|------|
| and channel name on the left<br>right list box. | processing. Currently available c<br>t. Calculations are applied only o |                                     |      |
| Available channels:                             |   | Selected channels:<br>number / name |      |
| 1 1<br>2 2<br>3 3                               |   | 4 1<br>5 2<br>6 1<br>7 2            |      |
|   | Add to list -><br><- Remove from list                                   |                                     |      |
|   | Select all ->> < Remove all   |                                     |      |
| ·   |   |                                     | ~    |
|   | Help  | Cancel                              | OK ! |

- 5. Under **Classification method** it is possible to select a specific method for the classification of the currently generated feature matrix. If the **Classify data** checkbox is enabled the selected classification window will be started.
- 6. Uncheck the **Randomly permutate the matrix** checkbox because the trials with the left and right class labels are already randomly permutated. If this is not the case enable the box to generate a randomly permutated feature matrix.
- 7. Uncheck the **Classify data** box (if the box is checked the Linear Discriminant Analysis (LDA) window is immediately opened for further processing)
- 8. Check **Save results** to store the generated feature matrix. The automatic treemaker generates a subdirectory under the current data path with the name featurematrix. Enter as filename bandpower.mat.
- 9. Press the **Start** button to generate the feature matrix.

To perform the example demonstrated above from the MATLAB command line use the following code:

#### %Load Data

```
P_C=data;
File=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\session1234triggered.mat'];
P_C=load(P_C,File);
```

#### % Bandpower

```
ChannelExclude = [3];
Filter.Name = 'ALPHA';
Filter.Type = 'BP';
Filter.f_low = [7];
Filter.f_high = [13];
Filter.Realization = 'fft';
Filter.Order = [0];
IntervalLength = 128;
Overlap = 127;
Replace = 'add channels';
FileName = '';
ProgressBarFlag = 0;
P_C = gBSbandpower(P_C, ChannelExclude, Filter, IntervalLength,...
Overlap, Replace, FileName, ProgressBarFlag);
```

#### % Bandpower

```
ChannelExclude = [3 4 5];
Filter.Name = 'BETA-3';
Filter.Type = 'BP';
Filter.f_low = [14];
Filter.f_high = [20];
Filter.Realization = 'fft';
Filter.Order = [0];
IntervalLength = 128;
Overlap = 127;
Replace = 'add channels';
FileName = '';
ProgressBarFlag = 0;
P_C = gBSbandpower(P_C, ChannelExclude, Filter, IntervalLength,...
Overlap, Replace, FileName, ProgressBarFlag);
```

#### %Feature Matrix

```
Interval=[128 128 1024];
AttributeName={
    'LEFT'
    'RIGHT'
};
ChannelExclude=[1 2 3];
Permutate=0;
MergeTimePoints=0;
FileName=[ 'C:\Users\' getenv('USERNAME')
    '\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\bandpower.mat'];
ProgressBarFlag=[0];
F_O=gBSfeaturematrix(P_C,Interval,AttributeName,Permutate,...
MergeTimePoints,ChannelExclude,FileName,ProgressBarFlag);
```

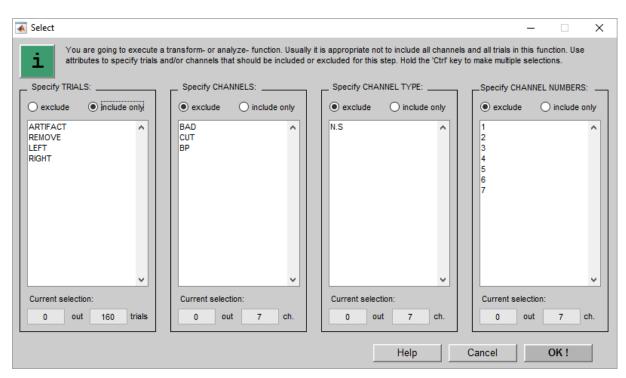
# Time Points

To extract features of the signal at different time points to investigate e.g. an initial state and an active state perform the following steps.

1. Load the data-set session1234bp.mat that was created in the previous section from

Documents\gtec\gBSanalyze\testdata\BCI\session1234bp.mat

2. Open **Cut Trials Channels** from the **Transform** menu and click the **Select trials/chan.** button. In **Specify TRIALS** this dialog click the **Include only** radio button and select only RIGHT trials for further operation. This allows inspecting changes between the inactive and active state of the BCI experiment for all right hand movement imagination trials.



- 3. After finishing the settings click **OK** to close the **Select** dialog.
- 4. Click **Start** to perform the action with the settings provided above.

|  | 5. | Open the | <b>Feature Matrix</b> | window from | n the Classification | menu |
|--|----|----------|-----------------------|-------------|----------------------|------|
|--|----|----------|-----------------------|-------------|----------------------|------|

| 承 Feature Matrix       |  |                |          |                            |             |  | -         |             | $\times$ |
|------------------------|--|----------------|----------|----------------------------|-------------|--|-----------|-------------|----------|
|                        | ature matrix as input f<br>n attribute correspond<br>to a class. |                |          |                            |             |  |           |             | ıt       |
| C Specify CLASSIFI     | CATION INTERVA   | L:             |          |                            |             |  |           |             |          |
| Start at: 1000         | [ms] Step:   | 1000 [m        | is]      | Stop at:                   | 8000        | [ms]   | Merg      | e time poin | ts       |
| 128                    | [samples]  | 128 [s         | amples]  |                            | 1024        | [samples]  |           |             |          |
| Specify CLASS LA       |  | NT-            |          |                            |             |  |           |             |          |
| R                      | RTIFACT<br>EMOVE<br>EFT<br>IGHT                                  | NI:            |          | <ul> <li>Select</li> </ul> | time point: | 1000<br>2000<br>3000<br>4000<br>5000<br>6000<br>7000<br>8000 |           |             |          |
| - Select FEATURE       | CHANNELS:  |                |          |                            |             | Select fe  | eature ch | annels      |          |
| - Choose METHOD        | and OPTIONS: -   |                |          |                            |             |  |           |             |          |
| Classification method: | Linear Discriminant A  | Analysis (LDA) |          | ~                          |             |  |           |             |          |
| Randomly permuta       |  |                |          |                            |             |  |           |             |          |
|                        |  |                |          |                            |             |  |           |             |          |
| Result procedure:      | Classify data  |                |          |                            | Automat     | ic treemaker   | is: enab  | ed ~        |          |
|                        | Save results   |                | Filename | e:ata\                     | BCNfeature  | ematrix\tpban  | dpower.m  | at          |          |
|                        |  |                |          | Help                       | p           | Cancel   |           | Start       |          |

- 6. Set the CLASSIFICATION INTERVAL Start at box to 1000 ms, the Step box to 1000 ms and the Stop at box to 8000 ms. These settings populate the Select time point listbox with specific time points.
- 7. Select 2000 ms and 7000 ms to extract only the features at these time points
- 8. Check the **Save results** box to store the features under tpbandpower.mat. The automatic treemaker generates the path.
- 9. Press **Start** to perform the operation.
- 10. The **Linear Classifier** dialog will open, see corresponding chapter for further description.

To perform the example demonstrated above from the MATLAB command line use the following code:

#### %Load Data

```
P_C=data;
File= ['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\session1234bp.mat'];
P_C=load(P_C,File);
```

#### %Select Trials and Channels

```
trial_id=[4];
channel_id=[];
type_id=[];
channelnr_id=[];
flag_tr='tr_inc';
flag_ch='ch_exc';
flag_type='type_exc';
flag_nr='nr_exc';
[TrialExclude, ChannelExclude]=gBSselect(P_C,trial_id,...
flag_tr,channel_id,flag_ch,type_id,flag_type,channelnr_id,flag_nr);
```

P\_C=gBScuttrialschannels(P\_C,TrialExclude,ChannelExclude);

#### %Feature Matrix

```
Interval=[256 896];
AttributeName={};
ChannelExclude=[];
Permutate=0;
MergeTimePoints=0;
FileName= ['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\tpbandpower.mat'];
ProgressBarFlag=[0];
F_O=gBSfeaturematrix(P_C,Interval,AttributeName,Permutate,...
MergeTimePoints,ChannelExclude,FileName,ProgressBarFlag);
```

# Time Segments

Time Segments allows to extract multiple segments of a trial to a feature matrix. Each segment of a trial receives its own class label. Therefore, if three segments are extracted the feature matrix contains three classes.

Follow these steps to generate a time segment feature matrix:

1. Load the data file Data.mat from

 ${\tt Documents \gtec \gBS analyze \test data \Self Paced}$ 

into the Data Editor. The Data Editor shows one ECoG channel and one trigger impulse channel.

2. To detect each trigger impulse start the **Eventfinder** from the **Artifact** menu. Check **Mark overflows** to search for trigger impulses which exceed 90 % of the maximum of the channel.

| Eventfinder   |                                       | _      |              |
|---|---------------------------------------|--------|--------------|
| The eventfinder searches for overflows, zero epochs and/or markers. Trial attributes can be | · · · · · · · · · · · · · · · · · · · |        | be marked by |
| _ Specify EVENTS:   |                                       |        |              |
| Mark overflows (amplitude criterion)  |                                       |        |              |
|   | Value exeeds:                         | 200    | μV ~         |
| Show epoching areas   | Value exeeds level:                   | 90     | [% of max.]  |
| Set start-marker: OR1 Set end-marker: OR2   |                                       |        |              |
|   |                                       | Select | trials/chan. |
| Assign trial attribute: OVERRUN V   |                                       | Select | unais/chain. |
| Mark zerolines  |                                       |        |              |
|   | Value keeps below:                    | 5      | μV 🗸         |
| ✓ Show epoching areas   | Level keeps below:                    | 2      | [% of max.]  |
| Set start-marker: ZL1   | Slew rate below:                      | 3      | [µV/ms]      |
| Set end-marker: ZL2   | within a time window of:              | 500    | [ms]         |
| Assign trial attribute: ZERO $\sim$   |                                       | 100    | [samples]    |
|   |                                       |        |              |
|   |                                       | Select | trials/chan. |
|   |                                       |        |              |
| Mark event (slew rate criterion)  | O Slew rate below:                    | 50     | [µV/ms]      |
| ✓ Show epoching areas   | Slew rate above:                      | 50     | [µV/ms]      |
| Set start-marker: E1  | within a time window of:              | 500    | [ms]         |
| Set end-marker: E2  |                                       | 100    | [samples]    |
| Assign trial attribute: EVENT ~   |                                       |        |              |
|   |                                       | Select | trials/chan. |
|   |                                       |        |              |
| Expected events   | Number of events:                     | 0      |              |
| Expected events -> Number of overflows: 0   | Number of events.                     | 0      |              |
| Number of zerolines: 0  |                                       |        |              |
|   |                                       |        |              |
|   | Help Cance                            |        | Start !      |
|   |                                       |        |              |

- 3. Check Set-start marker to assign an OR1 marker to each rising edge on the trigger channel. Uncheck the Show epoching areas box.
- 4. To search only on the trigger impulse channel press the **Select trials /chan.** button and select channel 2
- 5. Press **Start** to search for the trigger impulses.

The Data Editor shows now 50 markers.

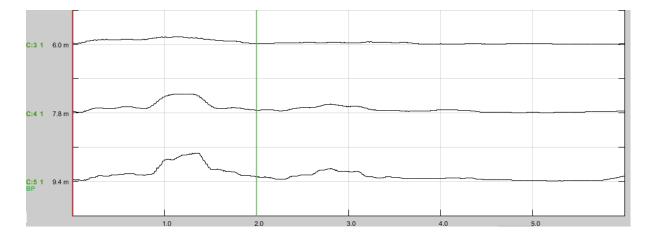
6. Open the **Trigger** window from the **Transform** menu and check the **Marker** radio button and select OR1 to extract 6 second trials around each OR1 marker

| 承 Trigger   |  | – 🗆 X                        |
|---|--|------------------------------|
| The trigger function splits your recorded data into trials channels or markers. The use of different markers or cl automatically to resulting trials. Channel attributes and m  | nannels allows you to ass  | ign attributes               |
| Time before trigger: 2000 [ms] Time after trigger:  | 4000 [ms]  | Accept incomplete            |
| 400 [samples]   | 800 [samples]  | last trial                   |
|   |  |                              |
| Specify TRIGGERS and ATTRIBUTES:     Physical channel: 1(CH1)     A Edge: rising  | 0  | 201.4                        |
| O Physical channel: 1(CH1)<br>2(CH2)  | <ul> <li>Threshold voltage:</li> <li>Threshold level:</li> </ul> | 201.1 [µV]<br>90 [% of max.] |
| ×   | <ul> <li>Slew rate:</li> </ul>                                   | 24.67 [µV/ms]                |
| Marker: OR1      Chan. (Market)   | )/ Name/ Edge/ Value/ Att  |                              |
| Assign attribute to resulting trials: mark/ OR1/ -/   | . 2  |                              |
| Accept overlap add to list -> <ul> <li>&lt;- remove from list</li> <li>apply changes -&gt;</li> </ul> <li>Change color: <ul> <li>red</li> <li>blue</li> <li>green</li> <li>yellow</li> <li>pink</li> <li>orange</li> <li>purple</li> <li>olive</li> <li>brown</li> <li>grey</li> </ul></li> |  | *                            |
| Generate LINED UP TRIALS:   |  |                              |
| O Line up trials (no trigger) Length of trials: 1000<br>200   | [ms] Overlap:<br>[samples]                                       | 0 [ms]<br>0 [samples]        |
| Select CHANNELS for the triggered file:   |  | Select channels              |
| H   | lelp Cance   | Start !                      |

- 7. Check the Accept Overlap box to allow overlapping trials
- 8. Press the **add to list** button to accept this trigger criterion
- 9. Press **Start !** to perform the operation. Now the Data Editor shows 49 trials with a trial length of 6 seconds. 2 seconds prior to the trigger impulse and 4 seconds after the impulse.
- 10. To calculate parameters from the ECoG channel open the **Bandpower** window from the **Parameter Extraction** menu

| 承 Bandpower                         |   |                      |                           | -          |             | ×   |
|-------------------------------------|---|----------------------|---------------------------|------------|-------------|-----|
| Calculate the bandpower data.       | by bandpass filtering                   | , squaring and aver  | aging over a spec         | ific windo | ow of the   |     |
| - Select CHANNELS:                  |   |                      |                           |            |             |     |
|                                     |   |                      |                           | Select c   | hannels     |     |
| Specify METHOD and PARA             | METERS:                                 |                      |                           |            |             |     |
| Choose a filter for the estimation: | ALPHA / BP / 7 / 13                     | / fft / 0            | ~                         | Design r   | new filter. |     |
|                                     | name / type / f (k                      | ) / f (hi) / real. / | order                     |            |             | -1  |
|                                     |   | .,                   |                           |            |             |     |
| Specify averaging window:           | Length: 500                             | [ms]                 | Overlap                   | 495        | [ms]        |     |
|                                     | 100                                     | [samples]            |                           | 99         | [sample     | es] |
| <u> </u>                            | v channels<br>all channels<br>sult data | Autor<br>Filename:   | natic treemaker is:<br>ei | enabl      |             | ~   |
|                                     |   | Help                 | Cancel                    |            | Start !     |     |

- 11. Select the Alpha filter to extract the bandpower from 7 to 13 Hz  $\,$
- 12. Press the Select channels button and chose only channel 1 for the calculation
- 13. Select **Add new channels** to append the new feature channel to the data in the Data Editor
- 14. Press Start to perform the calculation
- 15. Repeat steps 10 to 14 with the BETA-3 and with the BETA filter



### The Data Editor contain now 3 additional bandpower channels

- 16. Open the **Time Segment Feature Matrix** window from the **Classification** menu and define the classification segments. Enter under **Start at** 2000 ms and under **Stop at** 2200 ms and press the **Add** button. Then enter 5000 ms and 5200 ms and press again the **Add** button. These settings will extract two segments from each trial. The first segment will be the first class and the second segment the second class.
- 17. Press the **Select features channels ...** button and chose the bandpower channels 3, 4 and 5
- 18. Check the **Randomly permutate the matrix** box to generate a random permutation of the trials

19. Click on **Save results** and enter the name selffmseg.mat into the upcoming window to store the feature matrix.

| 承 Time Segment Featu                    | re Matrix                                   |                                       |                      | -                   | -          | ×       |
|---|---|---------------------------------------|----------------------|---------------------|------------|---------|
|   | eature matrix of spe<br>k can be used for c | cific data segments<br>lassification. | . Each segment       | forms its own clas  | ss. The re | sulting |
| C Specify CLASSIF                       | ICATION INTER                               | VAL:                                  |                      |                     |            |         |
| Start at: 5000<br>1000<br>Stop at: 5200 | [ms]<br>[samples]<br>[ms]                   | Add>                                  | 400 440<br>1000 1040 |                     | ^          |         |
| 1040                                    | [samples]                                   |                                       |                      |                     |            |         |
|   | _   |                                       |                      |                     | <u>×</u>   |         |
| - Select FEATURE                        | CHANNELS:                                   |                                       |                      |                     |            |         |
|   |   |                                       |                      | Select feat         | ure char   | inels   |
|   | and OPTIONS:                                |                                       |                      |                     |            |         |
| Classification method:                  | Linear Discriminar                          | nt Analysis (LDA)                     | $\sim$               |                     |            |         |
| Randomly permuta                        | te the matrix                               |                                       |                      |                     |            |         |
| Result procedure:                       | 🗹 Classify data                             |                                       | Auton                | natic treemaker is: | enabled    | ~       |
|   | Save results                                | Filename:                             | SelfPaced\fe         | eaturematrix\selffn | nseg.mat   |         |
|   |   |                                       | Help                 | Cancel              |            | Start   |

- 20. If the **Classify data** box is checked the **Linear Discriminant Analysis (LDA)** algorithm window would be opened for further processing, see chapter **Linear Classifier** for further description.
- 21. Press **Start** to perform the operation.

To perform the example demonstrated above from the MATLAB command line use the following code:

#### %Load Data

```
P_C=data;
File=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\SelfPaced\Data.mat'];
P_C=load(P_C,File);
```

#### %Select Trials and Channels

```
trial_id=[];
channel_id=[];
type_id=[];
channelnr_id=[2];
flag_tr='tr_exc';
flag_ch='ch_exc';
flag_type='type_exc';
flag_nr='nr_inc';
[TrialExclude, ChannelExclude]=gBSselect(P_C,trial_id,flag_tr,...
channel_id,flag_ch,type_id,flag_type,channelnr_id,flag_nr);
```

#### % Eventfinder (overflow)

```
MarkOverflow = 1;
showEpochingAreas over = 0;
setStartMarker over = 1;
setStopMarker over = 0;
AssignAttribute over = 0;
StartMarker over = 'OR1';
StopMarker over = 'OR2';
TrialAttribute over = 'OVERRUN';
Threshold over = 90;
getUnit over = '% of max';
TrialExclude over = [];
ChannelExclude over = [1];
ProgressBarFlag = 0;
[P C, PreviewOverflow, VecThreshold] = gBSoverflow...
        (P C, MarkOverflow, showEpochingAreas over,...
        setStartMarker over, setStopMarker over,...
        AssignAttribute over, StartMarker over, StopMarker over, ...
        TrialAttribute over, Threshold over, getUnit over,...
        TrialExclude_over, ChannelExclude_over, ProgressBarFlag);
```

#### %Trigger

```
New_tm{1}={3 1};
SamplesBefore=400;
SamplesAfter=800;
Uncomplete=0;
ChannelExclude=[];
P_C=gBStrigger(P_C,New_tm,SamplesBefore,SamplesAfter,Uncomplete,ChannelExcl
ude);
```

```
% Bandpower
ChannelExclude = [2];
Filter.Name = 'ALPHA';
Filter.Type = 'BP';
Filter.f low = [7];
Filter.f high = [13];
Filter.Realization = 'fft';
Filter.Order = [0];
IntervalLength = 100;
Overlap = 99;
Replace = 'add channels';
FileName = '';
ProgressBarFlag = 0;
P C = gBSbandpower(P C, ChannelExclude, Filter, IntervalLength,...
                   Overlap, Replace, FileName, ProgressBarFlag);
% Bandpower
ChannelExclude = [2]
                    31;
Filter.Name = 'BETA-3';
Filter.Type = 'BP';
Filter.f_low = [14];
Filter.f high = [20];
Filter.Realization = 'fft';
Filter.Order = [0];
IntervalLength = 100;
Overlap = 99;
Replace = 'add channels';
FileName = '';
ProgressBarFlag = 0;
P C = gBSbandpower(P C, ChannelExclude, Filter, IntervalLength,...
                   Overlap, Replace, FileName, ProgressBarFlag);
% Bandpower
ChannelExclude = [2 3 4];
Filter.Name = 'BETA';
Filter.Type = 'BP';
Filter.f low = [14];
Filter.f high = [32];
Filter.Realization = 'fft';
Filter.Order = [0];
IntervalLength = 100;
Overlap = 99;
Replace = 'add channels';
FileName = '';
ProgressBarFlag = 0;
P C = gBSbandpower(P C, ChannelExclude, Filter, IntervalLength,...
                   Overlap, Replace, FileName, ProgressBarFlag);
%Time Segment Feature Matrix
Interval=[
400 440
1000 1040
1;
ChannelExclude=[1 2];
Permutate=1;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\SelfPaced\featurematrix\selffmseg.mat'
1;
ProgressBarFlag=[0];
F O=gBStimesegmentfeaturematrix(P C,Interval,Permutate,...
ChannelExclude,FileName,ProgressBarFlag);
```

# **Generating a Classifier**

This section explains the classification and classifier generation of the feature matrix with the following methods:

Linear Classifier Multi-Class Linear Discriminant Analysis (LDA) Minimum Distance Classifier (MDC)

Neural Network Multi-Layer Perceptron (MLP) Radial Basis Function (RBF)

DSLVQ

Distinction Sensitive Learning Vector Quantization (DSLVQ)

All classification windows have certain control fields in common:

The **Load FEATURE MATRIX** field allows to **Browse** for a feature matrix file and shows the filename as well as the **Dimension** of the matrix. The number of features corresponds to the number of channels used for the feature matrix generation.

The number of trials depends on the feature matrix generation mode:

**Mode 1 – Trial Attribute** ... the number corresponds to the number of trials which have a certain trial attribute (e.g. left or right)

**Mode 2 – Time Point** ... the number corresponds to the number of selected time points (e.g. 2000 ms and 7000 ms) times the number of trials (e.g. 80)

**Mode 3 – Time Segment** ... the number corresponds to the number of samples in the time segments times the number of trials

| Load FEATURE MATRIX:         |                                   |
|------------------------------|-----------------------------------|
| Name of feature matrix file: | Browse                            |
| Dimension:                   | [features x trials x time points] |

**Time points** expresses the generation time points of the feature matrix. For each single time point the whole feature matrix is generated (e.g. 1000, 2000, ... 8000 ms).

The **Select FEATURE CHANNELS** field allows to specify the numbers (corresponds to the channel number when the feature matrix was generated) of the features that should be used for plotting a cloud of the feature matrix in gResult2d.

| Select FEATURE CHANNELS | ):       |
|-------------------------|----------|
| Map feature no.:        | against: |

**Choose METHOD and OPTIONS** allows to select the classification method and the training and test-sets:

**10 x 10 cross validation** ... The 10 times 10 fold cross validation mixes the data set randomly and divides it into 10 equally sized distinct partitions. Each partition is then used

| Choose METHOD a        | and OPTIONS:                       |   |                               |
|------------------------|------------------------------------|---|-------------------------------|
| Classification method: | Linear Discriminant Analysis (LDA) | ~ | Metric 🔘 Mahalanobis distance |
| Training / test-sets:  | 10 x 10 cross-validation           | ~ | O Euclidian distance          |

once for testing, the other partitions are used for training. This results in 10 different error rates, which are averaged. This is the error rate of a 10 fold cross validation. To further improve the estimate the procedure is repeated 10 times and again all error rates are averaged.

**Train 50 % - Test 50 %** ... uses the first 50 % of the feature matrix for training and the rest for testing

Train 100 % - Test 100 % ... uses all the data for training and testing

**Train 100 % - Test 0 %** ... uses all the data for training. This is useful to generate a classifier.

The **Result procedure** field allows to open gResult2d with the classification result and a cloud of the features. The **classifier window** option opens the MATLAB Editor with classification results and classifiers (weight vectors). **Save results** allows to store the classification result under a specific filename. If the **Automatic treemaker** is enabled a subdirectory under the current data directory is created. The result is stored into this subdirectory.

| Result procedure: | Show with Result2D     |           | Automatic treemaker is: enabled | ~ |
|-------------------|------------------------|-----------|---------------------------------|---|
|                   | Open classifier window |           |                                 |   |
|                   | Save results           | Filename: | enter filename                  |   |

## Linear Classifier

The Linear Classifier window allows to perform a linear discriminant analysis and minimum distance classifier analysis of multiple classes.

### **Multi-Class LDA**

#### Linear Discriminant of Fisher

An optimal decision rule for minimizing the probability of misclassification is based on the idea of discriminant functions. The simplest form consists of a linear combination of the inputs. The parameters are obtained with a learning algorithm from a set of training data. Fisher introduced a method that reduces the dimensionality before classification [Bishop 1995].

The dimension reduction is done by projecting the input data  $\mathbf{x}$  onto a value y with adjustable weights  $\mathbf{w}$ 

$$y = \mathbf{w}^T \mathbf{x} \tag{1}$$

Of course this leads to a loss of information but we chose  $\mathbf{w}$  in such a way that maximizes the class separation between class1 and class 2 (e.g. left and right finger movement). For class "left finger" the mean vector is

$$\mathbf{m1} = \frac{1}{N1} \sum_{n \in Classleftfinger} \mathbf{x}_{nleftfinger}$$
(2)

where N1 is the length of the input vector. For the class ,,right hand" **m2** is

$$\mathbf{m2} = \frac{1}{N2} \sum_{n \in Classrightfinger} \mathbf{x}_{nrightfinger}$$
(3)

The separation of the two classes is made by separating m1 and m2

$$m2 - m1 = \mathbf{w}^T (\mathbf{m2} - \mathbf{m1}) \tag{4}$$

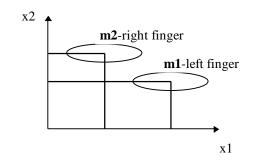
By choosing w arbitrary large the difference increases, therefore define

$$\sum_{i} \mathbf{w}_{i}^{2} = 1 \tag{5}$$

and after some calculation we obtain

$$w \propto (\mathbf{m} 2 - \mathbf{m} \mathbf{1}) \tag{6}$$

But a problem arises with this separation that is shown below.



Separation problem.

If we project m1 and m2 onto the x1 axes the difference is bigger than in the case of projecting onto the x2 axes. But there are within-class spreads that cause a better separation when m1 and m2 are projected onto x2.

Fisher proposed as solution

$$\mathbf{w} \propto \mathbf{S}_{w}^{-1} (\mathbf{m2} - \mathbf{m1}) \tag{7}$$

where  $S_w$  is the total within class covariance matrix

$$\mathbf{S}_{w} = \sum_{n \in Classieftfinger} (\mathbf{x}_{n} - \mathbf{m1}) (\mathbf{x}_{n} - \mathbf{m1})^{T} + \sum_{n \in Classrightfinger} (\mathbf{x}_{n} - \mathbf{m2}) (\mathbf{x}_{n} - \mathbf{m2})^{T}$$
(8)

Which is a projection rule for the data down to one dimension.

By choosing a threshold y0 we can classify a point to class 1 if it is greater zero or to class 2 otherwise.

Obviously the dimension reduction reduces the amount of information, but it can lead to improvements of the classifier performance [Bishop 1995].

#### **References:**

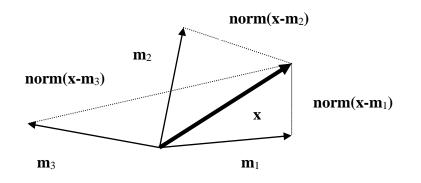
Bishop, C. M., Neural Networks for Pattern Recognition. Clarendon Press, Oxford, 1995.

## **Minimum Distance Classifier**

Let **x** be the feature vector for the unknown input. Vectors  $\mathbf{m}_1$ ,  $\mathbf{m}_2$ , ...,  $\mathbf{m}_c$  are the templates (i.e., perfect, noise-free feature vectors) for the c classes. Then the error in matching **x** against  $\mathbf{m}_k$  is given by

$$\|\mathbf{x} - \mathbf{m}_k\|$$
.

where  $|| \mathbf{u} ||$  is called the **norm** of the vector  $\mathbf{u}$ . The minimum-error classifier calculates  $|| \mathbf{x} - \mathbf{m}_k ||$  for k = 1 to c to find the class for which this error is minimum.  $|| \mathbf{x} - \mathbf{m}_k ||$  is also the distance from  $\mathbf{x}$  to  $\mathbf{m}_k$  and therefore the method is called **minimum-distance** classifier.



### Linear Classifier window:

**Classification method** can be Linear Discriminant Analysis (LDA) or Minimum Distance Classifier (MDC). In the case of MDC under Metric the Mahalanobis or Euclidian distance can be selected.

| 承 Linear Classifier  |   |                    | – 🗆 X               |  |  |
|--|---|--------------------|---------------------|--|--|
| Compute a classifier (weight vector) between groups of trials marked by class labels (trial attributes). The performance of the classifier can be estimated by cross-validation. The weight vector can be viewed, edited and stored for real-time processing with q.RTsys. |   |                    |                     |  |  |
| Load FEATURE M   | IATRIX:                                   |                    |                     |  |  |
| Name of feature matr   | ix file:                                  |                    | Browse              |  |  |
| Dimension: [features x trials x time points]   |   |                    |                     |  |  |
| - Select FEATURE   | CHANNELS:against:                         |                    |                     |  |  |
| Choose METHOD  | and OPTIONS:                              |                    |                     |  |  |
| Classification method:   | Linear Discriminant Analysis (LDA)        | ✓ Metric ○ Ma      | ahalanobis distance |  |  |
| Training / test-sets:  | 10 x 10 cross-validation                  | ✓ ○ Eu             | iclidian distance   |  |  |
| Result procedure:  | Show with Result2D Open classifier window | Automatic treemake |                     |  |  |
|  | Save results Filename:                    | enter mename       |                     |  |  |
|  |   | Help Cancel        | Start               |  |  |

### Example 1:

Perform the following steps to make a classification of an EEG-based brain-computer interface data-set:

- 1. Open the window Linear Classifier from the Classification menu
- 2. Press the **Browse** button and select the feature matrix file bandpower.mat that was created in the previous example and is stored under

Documents\gtec\gBSanalyze\testdata\BCI\featurematrix

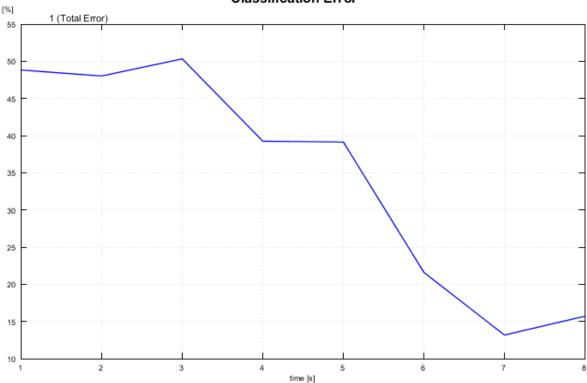
The featurematrix contains 4 features (2 bandpower values in the alpha range and 2 bandpower values in the beta range), 160 trials (80 right and 80 left) and 8 time points (1000, 2000, ... 8000 ms).

- 3. Under Select FEATURE CHANNELS select Map feature no 3 against 4 to plot a cloud of the feature matrix in gResult2d
- 4. Select Linear Discriminant Analysis (LDA) from the pull-down menu

| 承 Linear Classifier  |  | —             |          | $\times$      |  |
|--|--|---------------|----------|---------------|--|
| Compute a classifier (weight vector) between groups of trials marked by class labels (trial attributes). The performance of the classifier can be estimated by cross-validation. The weight vector can be viewed, edited and stored for real-time processing with q.RTsys. |  |               |          |               |  |
| Load FEATURE M   | ATRIX:   |               |          |               |  |
| Name of feature mate   | rix file:alyze\testdata\BCNfeaturematrix\bandpower.mat |               | se       |               |  |
| Dime   | nsion: 4 160 8 [features x trials x time points        | 3]            |          |               |  |
|  |  |               |          |               |  |
| C Select FEATURE   | CHANNELS:  |               |          |               |  |
| Map feature no.:   | 3 against: 4   |               |          |               |  |
| Choose METHOD  | and OPTIONS:   |               |          |               |  |
| Classification method:   | Linear Discriminant Analysis (LDA) V Metric O Mal      | halanobis o   | distance |               |  |
| Training / test-sets:  | 10 x 10 cross-validation V O Euc                       | clidian dista | ance     |               |  |
| Result procedure:  | Show with Result2D Automatic treemaker                 | is: enable    | ed       | $\overline{}$ |  |
|  | Open classifier window                                 |               |          |               |  |
|  | Save results Filename:stdata\BC\/featurematrix\/c      | LDAbp.ma      | at .     |               |  |
|  | Help Cancel  |               | Start    |               |  |

- 5. Select 10 x 10 cross-validation to randomly mix the training and testing data-set
- 6. Check the **Show with Result2D** and **Open classifier window** to open gResult2d and the MATLAB Editor to view the classification result and the weight vector (not available for 10 x 10 cross validation)
- 7. To store the classification result check **Save results** and enter the filename LDAbp.mat. The automatic treemaker generates the directory.

gResult2d opens with the classification result. The classification error is at the beginning around 50 % and drops down to 13 % at second 7.



Classification Error

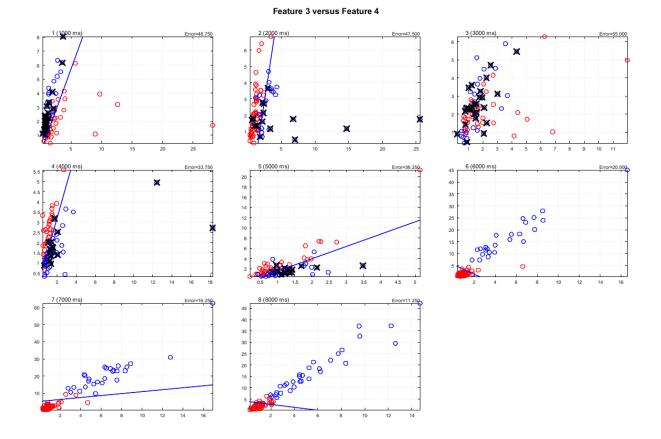
The MATLAB Editor shows the ASCII description of the classification. The first and second columns of the matrix shows the classification time point in seconds and samples. The third column represents the mean classification error followed by the standard deviation. The following columns give the classification errors of the 10 cross-validation runs.

Classification Method: Linear Discriminant Analysis Training- and testdata option: CV class 1: LEFT class 2: RIGHT Total Nr. of Trials: 160 Trials per selected Class 1: 80 Trials per selected Class 2: 80 Second/Sample/Mean Erkor/Std/[Total Error CV Runs] 1.000 128.000 48.3 2.6 48.8 46.3 46.9 50.0 50.6 50.0 52.5 44.4 45.0 48.8 2.000 256.000 46.8 1.4 44.4 46.9 46.3 48.1 46.3 47.5 48.8 46.3 45.0 48.1 3.000 384.000 49.6 1.8 50.6 50.0 49.4 52.5 48.1 50.6 48.8 47.5 51.2 46.9 4.000 512.000 39.1 1.2 39.4 41.3 39.4 39.4 38.8 37.5 38.1 39.4 40.0 37.5 5.000 640.000 39.4 1.1 40.6 39.4 40.0 40.6 39.4 39.4 36.9 38.8 39.4 39.4 6.000 768.000 21.3 0.6 21.9 21.3 21.9 20.6 21.3 20.6 21.3 20.6 21.3 22.5 7.000 896.000 13.0 0.4 12.5 12.5 13.1 12.5 13.1 13.1 13.1 13.1 13.8 13.1 8.000 1024.000 15.3 0.4 15.0 15.0 15.0 15.6 15.6 15.6 14.4 15.0 15.6 15.6

8. Repeat steps 1 to 7 but select under Training / test-sets Train 50 % - Test 50 %

gResult2d opens with the classification result on page 1. Change to page 2 of gResult2d to view the cloud of feature 3 versus feature 4. The red circles show the feature of a LEFT trial, the blue circles of a RIGHT trial. The blue line represents the weight vector of the linear discriminant analysis. The black crosses show the wrong classified trials.

At second 1 the classification error is 48,75 % and therefore the left and right classes can not be differentiated. But at second 8 the discrimination is possible with an error of 11.25 %. Note that the blue and red circles are clearly separated.



User Manual g.CLASSIFYtoolbox 5.16.02

Now the MATLAB Editor shows also the weight vectors for each calculated time point. Under Classifier 1 the bias value of the LDA classifier can be found. Classifier 2 shows the weight values for each feature channel in the same sequence as the features were extracted in the Data Editor.

```
Second/Sample: 8.000 1024.000
Classifier 1
0.565 |
Classifier 2
0.336 |
-0.098 |
-0.090 |
-0.139 |
```

Furthermore, the trial number of the wrong classified trials is given for each time point. In this case only trials of class 2 were wrong classified.

Second/Sample: 8.000 1024.000 Class Number/Nr Trials: Class 1: Class 2: 83 86 91 92 97 101 123 124 155 To perform the example demonstrated above from the MATLAB command line use the following code:

#### %Load FeatureMatrix

```
F_M=featurematrix;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\bandpower.mat'];
F_M=load(F_M,FileName);
```

#### %Linear Classifier

```
PlotFeatures=[1 2];
Method=['LDA'];
P.metric=[''];
TrainTestData=['CV'];
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\lc\LDAbp.mat'];
ProgressBarFlag=[0];
C_O=gBSlinearclassifier(F_M,Method,P,TrainTestData,PlotFeatures,...
FileName,ProgressBarFlag);
```

#### %Linear Classifier

```
PlotFeatures=[3 4];
Method=['LDA'];
P.metric=[''];
TrainTestData=['50:50'];
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\lc\LDAbp.mat'];
ProgressBarFlag=[0];
C_O=gBSlinearclassifier(F_M,Method,P,TrainTestData,PlotFeatures,...
FileName,ProgressBarFlag);
```

### Example 2:

This example shows the classification of a three class problem with the Minimum Distance Classifier (MDC).

1. Load the data file 3classes.mat into the Data Editor from

Documents\gtec\gBSanalyze\testdata\Classify

The data-set contains 150 trials with 2 feature channels. The channels contain artificial generated random numbers and the classes are separated by specific mean values. Trial 128 represents an outlier.

2. Open Feature Matrix from the classification menu and set the CLASSIFCATION INTERVAL to Start at 1000 ms, Step 1000 ms and Stop at 5000 ms.

| 承 Feature Matrix       |   |          |              |                           |              | —           |               | × |
|------------------------|---|----------|--------------|---------------------------|--------------|-------------|---------------|---|
|                        | eature matrix as input f<br>h attribute correspond<br>to a class. |          |              |                           |              |             |               |   |
| C Specify CLASSIFI     | CATION INTERVA  | L:       |              |                           |              |             |               | _ |
| Start at: 1000         | [ms] Step:  | 1000 [n  | ns] Stop at: | 5000                      | [ms]         | Merge       | e time points |   |
| 128                    | [samples]   | 128 [s   | amples]      | 640                       | [samples]    |             |               |   |
| - Specify CLASS LA     | BELS / TIME POI   | NT:      |              |                           |              |             |               | _ |
| Select class: A        | RTIFACT   | ^        | O Sele       | ct time point:            | 1000         |             | ^             |   |
| R                      | EMOVE   | _        |              |                           | 2000<br>3000 |             |               |   |
| 4                      |   |          |              |                           | 4000         |             |               |   |
| 5                      |   |          |              |                           | 5000         |             |               |   |
|                        |   | <b>_</b> |              |                           |              |             | <b>v</b>      |   |
|                        |   | •        |              |                           | L            |             | · ·           |   |
| - Select FEATURE       | CHANNELS:   |          |              |                           |              |             |               | _ |
|                        |   |          |              |                           | Select fe    | eature cha  | innels        |   |
|                        |   |          |              |                           |              |             |               |   |
| Choose METHOD          | and OPTIONS: _  |          |              |                           |              |             |               |   |
|                        |   |          |              |                           |              |             |               |   |
| Classification method: | Minimum Distance Cl   |          |              |                           |              |             |               |   |
|                        | Minimum Distance Cl   |          |              |                           |              |             |               |   |
| Classification method: | Minimum Distance Cl   |          |              |                           |              |             |               |   |
| Classification method: | Minimum Distance Cl   |          |              | Automati                  | c treemaker  | is: enable  | id 🗸          |   |
| Classification method: | Minimum Distance Cl<br>te the matrix<br>Classify data             |          | ~            |                           |              | enable      |               |   |
| Classification method: | Minimum Distance Cl   |          | ~            | Automati<br>ata\Classify\ |              | enable      |               |   |
| Classification method: | Minimum Distance Cl<br>te the matrix<br>Classify data             |          | ~            | ata\Classify\             |              | x\3clfm.mat |               |   |

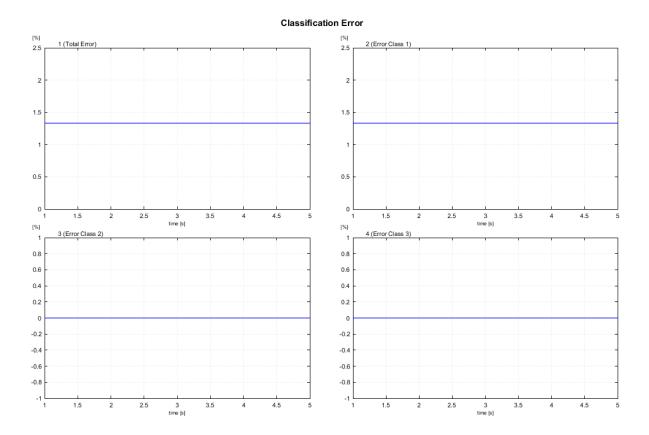
- 3. Select classes 3, 4 and 5 to separate all 3 classes
- 4. Chose Minimum Distance Classifier (MDC) under Classification method
- 5. Check Save results and enter 3clfm.mat as filename
- 6. Press **Start** to extract the feature matrix and to open the **Linear Classifier** window

The dimension of the feature matrix is 2 features (2 channels), 150 trials and 5 time points (1000, 2000,...5000ms).

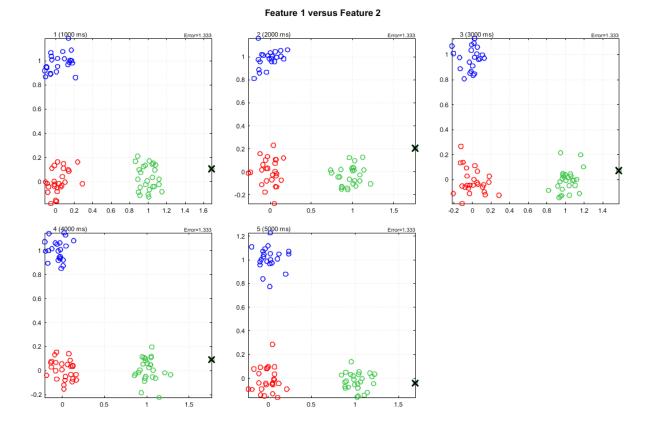
| 承 Linear Classifier                 |  | _              |            | × |
|-------------------------------------|--|----------------|------------|---|
| performance                         | assifier (weight vector) between groups of trials marked by class la<br>of the classifier can be estimated by cross-validation. The weight ve<br>ored for real-time processing with g.RTsys. |                |            |   |
| Load FEATURE M                      | IATRIX:  |                |            |   |
| Name of feature mate                | rix file:lyze\testdata\Classify\featurematrix\3clfm.mat  | Bro            | wse        |   |
| Dime                                | nsion: 2 150 5 [features x trials x time poin  | nts]           |            |   |
| Select FEATURE     Map feature no.: | CHANNELS:  |                |            |   |
|                                     | and OPTIONS:   |                |            |   |
| Classification method:              | Minimum Distance Classifier (MDC) V Metric 💿 M   | lahalanobi     | s distance |   |
| Training / test-sets:               | Train 50 % - Test 50 %   | uclidian dis   | stance     |   |
| Result procedure:                   | Show with Result2D Automatic treemake Open classifier window Save results Filename:ata\Classify\featuremate  | Cita           | bled       | ~ |
|                                     | Save results Filename:ata\Classify\featuremat  | Texticiael.III |            |   |
|                                     | Help Cance   |                | Start      | : |

- 7. Select Minimum Distance Classifier (MDC) under Classification method and chose Train 50 % Test 50 %
- 8. Press **Start** to perform the classification

Page 1 of gResult2d shows 4 classification error time courses. The first channel shows the **Total Error** of all classes. Channels 2 to 4 represent the error rates for each individual class. In this case only class 1 has an error rate which is not zero. This is the case because trial 128 is marked as class 1 trial but represents an outlier. Note that only 50 % of the trials are used for testing and therefore the total trial number is 75 trials.



The second page maps feature 1 versus feature 2. A color is assigned to each class and the wrong classified trial 128 is indicated by the black cross. Trial 128 was classified as green class but belongs to the red class. Therefore, the linear method is not able to correctly identify trial 128.



# The MATLAB Editor shows that 27 trials were selected of class 1, 22 of class 2 and 26 of class 3.

```
Classification Method: Linear Discriminant Analysis
Training- and testdata option: 50:50
class 1: 3
class 2: 4
class 3: 5
Total Nr. of Trials: 75
Trials per selected Class 1: 27
Trials per selected Class 2: 22
Trials per selected Class 3: 26
```

The weight vector is given for all three classes. The symbol "|" is used to separate the classifiers.

```
Second/Sample: 5.000 640.000

Classifier 1

1.000 0.000 | 1.000 0.000 | 1.000 0.000 |

0.000 1.000 | 0.000 1.000 | 0.000 1.000 |

Classifier 2

0.032 | 0.032 | 1.016 |

-0.010 | 0.997 | 0.006 |
```

To perform the example demonstrated above from the MATLAB command line use the following code:

```
%Load Data
P_C=data;
File=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\Classify\3classes.mat'];
P_C=load(P_C,File);
%Feature Matrix
Interval=[128 128 640];
AttributeName={
    '3'
```

```
'4'
'5'
};
ChannelExclude=[];
Permutate=0;
MergeTimePoints=0;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\Classify\featurematrix\3clfm.mat'];
ProgressBarFlag=[0];
F_O=gBSfeaturematrix(P_C,Interval,AttributeName,Permutate,...
MergeTimePoints,ChannelExclude,FileName,ProgressBarFlag);
```

#### %Load FeatureMatrix

```
F_M=featurematrix;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\Classify\featurematrix\3clfm.mat'];
F_M=load(F_M,FileName);
```

#### %Linear Classifier

```
PlotFeatures=[1 2];
Method=['MDC'];
P.metric=['Mahalanobis'];
TrainTestData=['50:50'];
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\Classify\featurematrix\lc\3cl.mat'];
ProgressBarFlag=[0];
C_O=gBSlinearclassifier(F_M,Method,P,TrainTestData,PlotFeatures,...
FileName,ProgressBarFlag);
```

### Neural Network

Neural Network allows to calculate a Multi-Layer Perceptron (MLP) or a Radial Basis Function (RBF).

Artificial Neural Networks (ANNs) can approximate the discriminant function by varying the connection strength (weight value) between the units. Such networks can have only one layer or multiple sequential layers (MLP). The MLP available in g.BSanalyze has 1 input layer, 1 hidden unit and 1 output layer.

Radial Basis Function networks have 2 layers. The first one performs a non-linear parameter transformation and the second layer makes a linear discrimination of the first layer parameters. The idea of RBF is based on the assumption that a complex pattern classification problem becomes linearly better separable when the parameters are non-linearly mapped to higher dimensionality.

The window has the following **OPTIONS** settings:

**No. inputs** ... number of input units. This number corresponds to the number of feature channels of the feature matrix

No. hidden units ... enter the number of hidden units

**No. outputs** ... number of output units. This number corresponds to the number of classes of the feature matrix

**Learning rate** ... specify the learning rate of the neural network. The learning rate controls the learning process. It gives the influence of the present example compared to all the past examples. A learning rate of 0 means that the neural network is not changed at all, while a learning rate of 1 would change the neural network according to the present example, independent of all previous examples.

Stop error ... enter the stop error rate

**Epochs** ... specify the number of training iterations

Momentum (only for MLP) ... enter the momentum value

Perform the following steps to classify a BCI experiment data-set:

1. Start Neural Network from the Classification menu and open the feature matrix file bandpower.mat from

Documents\gtec\gBSanalyze\testdata\BCI\featurematrix

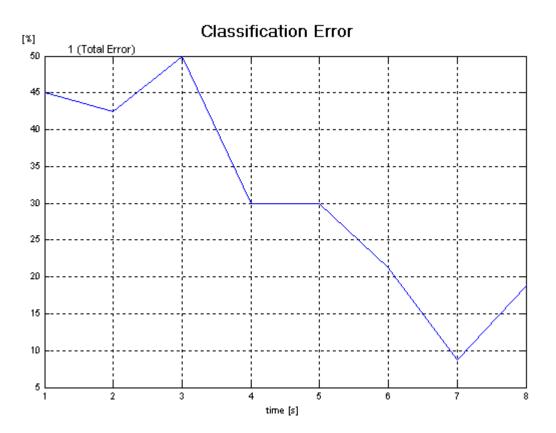
The feature matrix has 4 channels, 160 trials and was created for 8 time points (1000 ms, 2000 ms, ..., 8000 ms).

| 承 Neural Network  |   | – 🗆 X               |  |  |  |  |  |  |
|---|---|---------------------|--|--|--|--|--|--|
| Perform a classification of a feature matrix with a neural network. The neural net can be used for real-time analysis with g.RTsys. |   |                     |  |  |  |  |  |  |
|   | IATRIX:   |                     |  |  |  |  |  |  |
| Name of feature mate  | ix file:alyze\testdata\BC\featurematrix\bandpower.mat | Browse              |  |  |  |  |  |  |
| Dime  | nsion: 4 160 8 [features x trials x time p            | points]             |  |  |  |  |  |  |
|   |   |                     |  |  |  |  |  |  |
| No.   | 4 Learning 0.0002                                     | Epochs: 4000        |  |  |  |  |  |  |
| No. hidden units:   | 8 Stop 0.01   | Momentum: 0.98      |  |  |  |  |  |  |
| No. outputs:  | 2   |                     |  |  |  |  |  |  |
| - Select FEATURE  | CHANNELS:   |                     |  |  |  |  |  |  |
| Map feature no.:  | 3 against: 4  |                     |  |  |  |  |  |  |
| Choose METHOD   | and OPTIONS:  |                     |  |  |  |  |  |  |
| Classification method:  | Multi Layer Perceptron (MLP)                          |                     |  |  |  |  |  |  |
| Training / test-sets:   | Train 50 % - Test 50 %                                |                     |  |  |  |  |  |  |
| Result procedure:   |   | naker is: enabled 🗸 |  |  |  |  |  |  |
|   | Open classifier window                                |                     |  |  |  |  |  |  |
|   | Save results Filename:stdata\BCI\featurema            | trix\nn\NNbp.mat    |  |  |  |  |  |  |
|   | Help Can  | cel Start           |  |  |  |  |  |  |

- 2. Set the No. hidden units to 8 and the Epochs to 4000
- 3. Enter 3 under Map feature no. and 4 under against
- 4. Select Train 50 % Test 50 % to split the training and testing data in 50 % partitions

- 5. Check Save results and enter the filename NNbp.mat
- 6. Press **Start** to calculate the neural network

gResult2d opens automatically with the classification error time course. The minimum error of 8 % is reached at second 7.



The MATLAB Editor shows also the weight vectors of the generated MLP network. Classifier 1 shows the weights for the hidden layer (8 hidden layer nodes = 8 columns, 4 inputs layers + 1 bias = 5 rows). Classifier 2 represents the output layer (2 columns = 2 output nodes, 8 hidden layer nodes + 1 bias value = 9 rows).

```
Second/Sample: 7.000 896.000
Classifier 1
-1.211 -4.580 -1.104 0.576 0.845 0.307 0.084 0.110
-1.421 -2.675 -1.516 -1.390 1.171 0.171 -0.117 -0.947
-0.290 -1.677 -0.728 0.310 0.382 1.271 -0.422 1.423
1.071 3.122 -0.961 0.254 0.895 4.709 0.672 -1.460
0.781 2.537 -0.795 2.089 0.172 -0.107 -0.618 0.681
Classifier 2
0.871 -1.360
-2.710 2.593
1.158 -2.000
-0.977 0.010
0.169 0.903
-0.093 -0.157
2.087 0.033
-1.029 1.952
0.781 0.432
```

To perform the example demonstrated above from the MATLAB command line use the following code:

#### %Load FeatureMatrix

```
F_M=featurematrix;
FileName= ['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\bandpower.mat'];
F_M=load(F_M,FileName);
```

#### %Neural Network

```
P.ninput=[4];
P.nhidden=[8];
P.noutput=[2];
P.learningrate=[0.0002];
P.stoperror=[0.01];
P.epochs=[4000];
P.momentum=[0.98];
PlotFeatures=[3
                41;
Method=['MLP'];
TrainTestData=['50:50'];
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\nn\NNbp.mat'];
ProgressBarFlag=[0];
C_O=gBSneuralnetwork(F_M,Method,P,TrainTestData,PlotFeatures,...
FileName, ProgressBarFlag);
```

### DSLVQ

The method Distinction Sensitive Learning Vector Quantization is based on the individual testing of candidate feature subsets. The classifier is initially trained on all features and through implicit feature relevance analysis the system finds which features are not necessary and adapts the weighting of the features accordingly. Finally only the relevant features are used for the classification problem.

The window has the following **OPTIONS** settings:

| JSLVQ                  |                                       |                                  | – 🗆 X         |
|------------------------|---------------------------------------|----------------------------------|---------------|
| Perform a Dis          | tinction Sensitive Learning Vector Qu | antization.                      |               |
| Load FEATURE M         | IATRIX:                               |                                  |               |
| Name of feature mate   | rix file:testdata\SelfPaced\fea       | turematrix\selffmseg.mat         | Browse        |
| Dime                   | nsion: 3 4018 1                       | [features x trials x time points | \$]           |
|                        |                                       |                                  |               |
| Codebooks per          | 8 Alpha:                              | 0.05 Epochs:                     | 2000          |
| Map feature no.:       | CHANNELS:<br>2 against: 3             |                                  |               |
| Choose METHOD          | and OPTIONS:                          |                                  |               |
| Classification method: | DSLVQ                                 | ~                                |               |
| Training / test-sets:  | Train 50 % - Test 50 %                | ~                                |               |
| Result procedure:      | Show with Result2D                    | Automatic treemaker              | is: enabled V |
|                        | 🗹 Open classifier window              |                                  |               |
|                        | Save results Filename:                | enter filename                   |               |
|                        |                                       | Help Cancel                      | Start         |

Codebooks per class ... define the number of codebooks for each class

Alpha ... define the learning speed of the algorithm

**Epochs** ... define the number of iterations

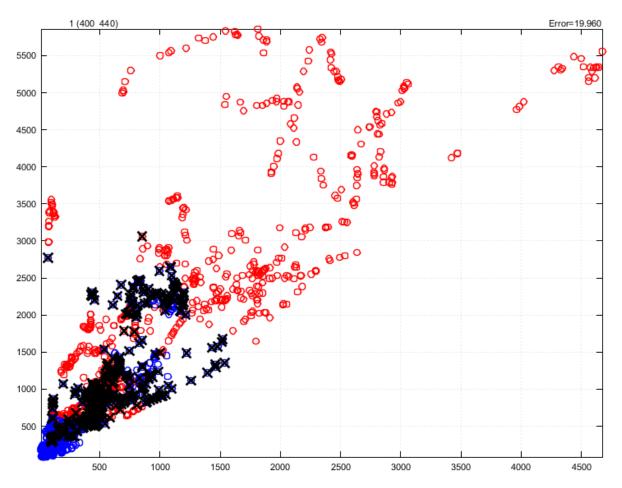
Perform the following steps to classify a finger movement ECoG experiment:

1. Open the **DSLVQ** window from the **Classification** menu and load the feature matrix file selffmseg.mat from

Documents\gtec\gBSanalyze\testdata\SelfPaced\featurematrix

- 2. Enter under Codebooks per class 8 and set the number of Epochs to 2000
- 3. Enter under Map feature no. 2 and under against 3
- 4. Chose the Train 50 % Test 50 % option
- 5. Press the **Start** button to train the DSLVQ

gResult2d opens with the classification result. The classification error is 15.978 % and the plot shows the distribution of features 2 versus feature 3. Red colors correspond to the first class (segment 2000-2200 ms) and blue colors correspond to the second segment (5000 - 5200 ms). Black crosses indicate wrong classified examples.



### Feature 2 versus Feature 3

To perform the example demonstrated above from the MATLAB command line use the following code:

```
%Load FeatureMatrix
F_M=featurematrix;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\SelfPaced\featurematrix\selffmseg.mat'
];
F_M=load(F_M,FileName);
```

#### %DSLVQ

```
P.CBperclass=[8];
P.alpha=[0.05];
P.epochs=[2000];
PlotFeatures=[2 3];
Method=['DSLVQ'];
TrainTestData=['50:50'];
FileName=[''];
ProgressBarFlag=[0];
C_O=gBSdslvq(F_M,Method,P,TrainTestData,PlotFeatures,FileName,...
ProgressBarFlag);
```

### Support Vector Machine Classifier

**Support Vector Machine Classifier** creates a classifier based on support vector machines [Cortes 1995].

To achieve good performance, it is required to scale the data appropriately, three scaling methods are implemented: **unit variance/zero mean**, **scaling to range [0 1]**, and **scaling to range [-1 1]**. Further, the hyperparameters have to be tuned for the problem. Use a linear SVM for linear problems, and tune the trade-off parameter **C-value** with **OPTIMIZATION**. For non-linear problems use a radial basis function (RBF) kernel, this adds an additional hyperparameter **Gamma-value** that can be also optimized.

Two procedures for the tuning of hyperparameters are implemented: **Grid-search** and **Random-search**. The latter is recommended if a larger search space is defined. **Grid-Search** also allows a fine tuning of the parameters over several levels. Both optimization procedures use cross-validation to determine the best parameters. The **Range**, of the tuned hyperparameters can be selected as well as the **Resolution** between the range and the **Scaling**.

### **References:**

Cortes, C. and Vapnik, V. "Support-Vector Networks", Machine Learning, 20, pp. 273-297, 1995.

| 承 Support Vector Machine Clas   | sifier  |   | – 🗆 X                            |  |  |  |  |
|---|---|---|----------------------------------|--|--|--|--|
| Compute a SVM-classifier between groups of trials marked by class labels (trial attributes). The performance of the classifier can be estimated by cross-validation. The classifier can be viewed, edited and stored for real-time processing with g.RTsys. |   |   |                                  |  |  |  |  |
| Name of feature matrix file:  | alyze\testdata\BC\\f  | eaturematrix\bandpower.mat  | Browse                           |  |  |  |  |
| Dimension:  | 4 160 8   | [features x trials x time points  |                                  |  |  |  |  |
| - OPTIONS:  |   |   |                                  |  |  |  |  |
| Classifier: Non-linear  | (RBF) V   |   | C-value: 2                       |  |  |  |  |
| Scaling: unit varian  | ce/zero mean 🗸 🗸  | Gam   | ma-value: 1                      |  |  |  |  |
| Training / test-sets: Train 50 %  | 5 - Test 50 % ✓   |   |                                  |  |  |  |  |
| - Optimization:   |   |   |                                  |  |  |  |  |
| Procedure:<br>Grid-search V<br>Levels: 2<br>Loops: 100  | C-value:<br>Range: 1<br>Scaling: logarithm<br>Resolution:       | to 10<br>ic v<br>20<br>Gamma-value<br>Range:<br>Scaling:<br>Resolution: | :<br>0.1 to 20<br>linear ~<br>10 |  |  |  |  |
|   | ow with Result2D<br>en classifier window<br>ve results Filename | Automatic treemaker   | is: enabled V                    |  |  |  |  |
|   | [   | Help Cancel   | Start                            |  |  |  |  |

### Example:

Perform the following steps to make a classification of an EEG-based brain-computer interface data-set:

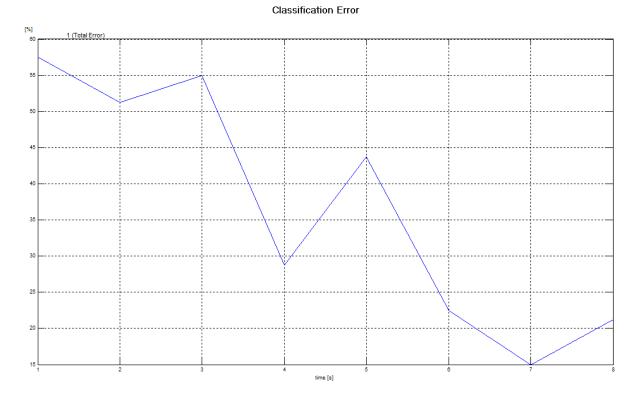
- 1. Open the window Support Vector Machine Classifier from the Classification menu
- 2. Press the **Browse** button and select the feature matrix file bandpower.mat that was created in the previous example and is stored under

Documents\gtec\gBSanalyze\testdata\BCI\featurematrix

The featurematrix contains 4 features (2 bandpower values in the alpha range and 2 bandpower values in the beta range), 160 trials (80 right and 80 left) and 8 time points (1000, 2000, ... 8000 ms)

- 3. Select Non-linear (RBF) for the Classifier to use a radial basis function as kernel
- 4. Select unit variance/zero mean as scaling method
- 5. Select **Train 50 % Test 50 %** to train the classifier on 50 % of the data and test it on the other 50 %
- 6. Select Grid-search for Optimization, using 2 Levels
- 7. Select for the **C-value** a **Range** between 1 and 10 with logarithmic **Scaling** and a **Resolution** of 20
- 8. Select for the **Gamma-value** a **Range** between 0.1 and 20 with linear **Scaling** and a **Resolution** of 10
- 9. Press the **Start** button

gResult2d opens with the classification result. The classification error is at the beginning around 50 - 55 % and drops down to 15 % at second 7.



The MATLAB Editor shows the ASCII description of the classifier. The kernel, the optimization method and the optimized C-values and Gamma-values for each sample are shown at the beginning:

Support Vector Machine Kernel: radial basis function Optimization method: Grid-search Number of levels: 2 Optimization of the C-value: Range: [1.0 10.0] Scaling: logarithmic Resolution: 20 Optimized C-values: Second/Sample/C-value:

1.000 128.000 2.0000 2.000 256.000 2.0000 3.000 384.000 2.0000 4.000 512.000 2.0000 5.000 640.000 2.0000 6.000 768.000 2.0000 7.000 896.000 2.0000 8.000 1024.000 2.0000 Optimization of the Gamma-value: Range: [0.1 20.0] linear Scaling: Resolution: 10 Optimized Gamma-values: Second/Sample/Gamma-value: 1.000 128.000 1.0000 2.000 256.000 1.0000 3.000 384.000 1.2222 4.000 512.000 8.0000 5.000 640.000 1.0000 6.000 768.000 1.0000 7.000 896.000 1.0000 8.000 1024.000 1.0000

The classification error is shown in the following table: The first and second columns of the matrix show the classification time point in seconds and samples, the third column shows the error.

```
Second/Sample/Total Error/[Error Class 1/Error Class 2...]

1.000 128.000 57.5

2.000 256.000 51.2

3.000 384.000 55.0

4.000 512.000 28.7

5.000 640.000 43.8

6.000 768.000 22.5

7.000 896.000 15.0

8.000 1024.000 21.3
```

The following code shows how to perform the example demonstrated above from the MATLAB command line.

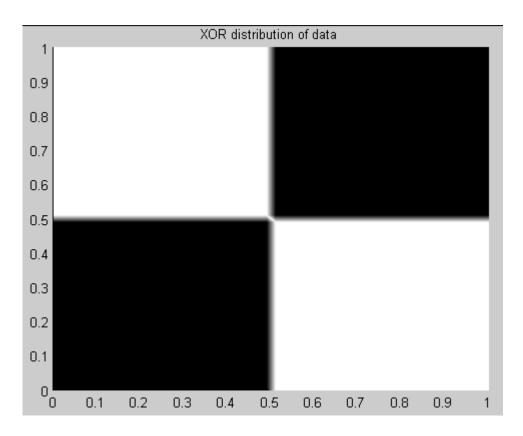
```
%Load FeatureMatrix
F M=featurematrix;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\bandpower.mat'];
F M=load(F M,FileName);
%Support Vector Machine Classifier
CLOption=[1];
Scaling=[1];
TrainTestData=['50:50'];
CParam=[10];
GParam=[0.5];
Optimization=[1];
Levels=[2];
Loops=[100];
CRange=[1.0 10];
CScaling=[1];
CResolution=[20];
GRange=[0.1 20];
```

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```
GScaling=[0];
GResolution=[10];
FileName=[''];
ProgressBarFlag=[1];
C_O = gBSsvmclassifier(F_M,CLOption,Scaling,TrainTestData,CParam,GParam,...
Optimization,Levels,Loops,CRange,CScaling,CResolution,GRange,GScaling,...
GResolution,FileName,ProgressBarFlag);
```

### Comparison of LDA and DSLVQ

This section compares the LDA and DSLVQ methods on the example of an XOR problem. The XOR distribution is demonstrated in the figure below. Basically the data-set consists of 2 channels with values between 0 and 1. If both channels have values below 0.5 or above 0.5 than the example belongs to class 0. If one channel is below 0.5 and the second channel is above 0.5 than the example belongs to class 1.



Perform the following steps:

1. Open the data-set XOR.mat from

Documents\gtec\gBSanalyze\testdata\Classify

into the Data Editor

2. Select the **Feature Matrix** window from the **Classification** menu to generate a feature matrix for a single time point.

| 🕢 Feature Matrix —  | × |
|---|---|
| Generate a feature matrix as input for the classification methods. Select class allows to select trials with a certain attribute. Each attribute corresponds to a class. Select time point allows to select specific time points. Each time point corresponds to a class. |   |
| Specify CLASSIFICATION INTERVAL:  | 7 |
| Start at:         600         [ms]         Step:         600         [ms]         Stop at:         600         [ms]         Merge time points   |   |
| 3 [samples] 3 [samples] 3   |   |
| Specify CLASS LABELS / TIME POINT:  | _ |
| Select class: ARTIFACT     Select time point: 600   |   |
| REMOVE<br>1   |   |
| 0   |   |
|   |   |
| ✓ ✓   |   |
| Select FEATURE CHANNELS:  |   |
| Select feature channels   |   |
| Choose METHOD and OPTIONS:  | _ |
| Classification method: Linear Discriminant Analysis (LDA)   |   |
|   |   |
| Randomly permutate the matrix   |   |
| Result procedure: Classify data Automatic treemaker is: enabled ~   |   |
| Filename:data\Classify\featurematrix\xorfm.mat  |   |
| Help Cancel Start   |   |

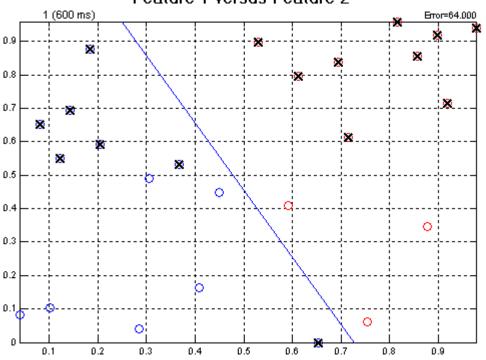
- 3. Set Start at, Step and Stop at to 600 ms and select classes 1 and 0 under Select class
- 4. Check the Save results box and enter the filename xorfm.mat
- 5. Press **Start** to save the feature matrix file

6. Open the **Linear Classifier** window under **Classification** and **Browse** for the feature matrix file xorfm.mat

| 承 Linear Classifier  |                                    |                                  | – 🗆 X              |  |  |  |  |  |
|--|------------------------------------|----------------------------------|--------------------|--|--|--|--|--|
| Compute a classifier (weight vector) between groups of trials marked by class labels (trial attributes). The performance of the classifier can be estimated by cross-validation. The weight vector can be viewed, edited and stored for real-time processing with g.RTsys. |                                    |                                  |                    |  |  |  |  |  |
| Name of feature mat  | rix file:lyze\testdata\Classify\f  | eaturematrix\xorfm.mat           | Browse             |  |  |  |  |  |
| Dime   | nsion: 2 50 1                      | [features x trials x time points | ;]                 |  |  |  |  |  |
| Select FEATURE   | CHANNELS:                          |                                  |                    |  |  |  |  |  |
| Choose METHOD  | and OPTIONS:                       |                                  |                    |  |  |  |  |  |
| Classification method:   | Linear Discriminant Analysis (LDA) | ✓ Metric ○ Mat                   | halanobis distance |  |  |  |  |  |
| Training / test-sets:  | Train 50 % - Test 50 %             | ✓ ○ Euc                          | clidian distance   |  |  |  |  |  |
| Result procedure:  | Show with Result2D                 | Automatic treemaker              | is: enabled V      |  |  |  |  |  |
|  | Save results Filename:             | enter filename                   |                    |  |  |  |  |  |
|  |                                    | Help Cancel                      | Start              |  |  |  |  |  |

7. Select Train 50 % - Test 50 % and press the Start button

gResult2d maps feature 1 versus feature 2 and shows the classification error of 64 %. The LDA is not able to discriminate the XOR problem.



Feature 1 versus Feature 2

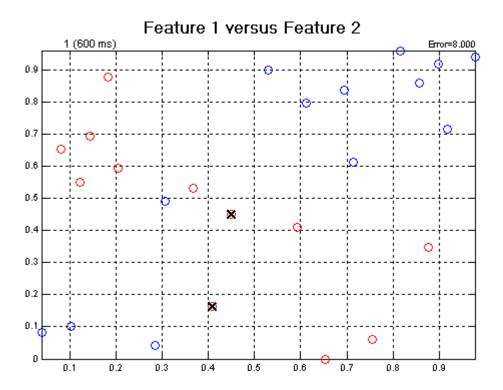
8. Close the Linear Classifier window

### 9. Open the **DSLVQ** window

- 10. Load again the xorfm.mat file and set the **Codebooks per class** to 8 and the **Epochs** to 4000
- 11. Select Train 50 % Test 50 % and press the Start button

| A DSLVQ - 🗆 🗙  |
|--|
| Perform a Distinction Sensitive Learning Vector Quantization.  |
| Load FEATURE MATRIX:   |
| Name of feature matrix file:lyze\testdata\Classify\featurematrix\xorfm.mat Browse  |
| Dimension: 2 50 1 [features x trials x time points]  |
| _ OPTIONS:   |
| Codebooks per 8 Alpha: 0.05 Epochs: 4000   |
| Select FEATURE CHANNELS:     Map feature no.: 1 against: 2   |
| Choose METHOD and OPTIONS:   |
| Classification method: DSLVQ ~   |
| Training / test-sets: Train 50 % - Test 50 %   |
| Result procedure:       Show with Result2D       Automatic treemaker is:       enabled          Open classifier window       Image: Comparison of the second sec |
| Save results Filename: enter filename  |
| Help Cancel Start  |

gResult2d shows now the classification result of the DSLVQ classifier with an error rate of 8 %. Note that only 2 trials (which are close to the border of 0.5) were wrongly classified. If the number of training examples is enhanced the error rate is reduced.



To perform the example demonstrated above from the MATLAB command line use the following code:

#### %Load Data

```
P_C=data;
File= ['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\Classify\XOR.mat'];
P_C=load(P_C,File);
```

#### %Feature Matrix

```
Interval=[3 3 3];
AttributeName={
    '1'
    '0'
};
ChannelExclude=[];
Permutate=0;
MergeTimePoints=0;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\Classify\featurematrix\xorfm.mat'];
ProgressBarFlag=[0];
F_O=gBSfeaturematrix(P_C,Interval,AttributeName,Permutate,...
MergeTimePoints,ChannelExclude,FileName,ProgressBarFlag);
```

#### %Load FeatureMatrix

```
F_M=featurematrix;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\Classify\featurematrix\xorfm.mat'];
F_M=load(F_M,FileName);
```

#### **%Linear Classifier**

```
PlotFeatures=[1 2];
Method=['LDA'];
P.metric=[''];
TrainTestData=['50:50'];
FileName=[''];
ProgressBarFlag=[0];
C_O=gBSlinearclassifier(F_M,Method,P,TrainTestData,PlotFeatures,...
FileName,ProgressBarFlag);
```

#### %Load FeatureMatrix

F\_M=featurematrix; FileName=['C:\Users\' getenv('USERNAME') '\Documents\gtec\gBSanalyze\testdata\Classify\featurematrix\xorfm.mat']; F\_M=load(F\_M,FileName);

#### %DSLVQ

```
P.CBperclass=[8];
P.alpha=[0.05];
P.epochs=[4000];
PlotFeatures=[1 2];
Method=['DSLVQ'];
TrainTestData=['50:50'];
FileName=[''];
ProgressBarFlag=[0];
C_O=gBSdslvq(F_M,Method,P,TrainTestData,PlotFeatures,FileName,...
ProgressBarFlag);
```

# **Receiver Operator Curve**

The sensitivity and specificity of a diagnostic test depends on more than just on the quality of the test. It depends also on the definition of what constitutes an abnormal test. In practice a threshold is selected to distinguish e.g. normal from disease. The threshold level determines the number of true positives, true negatives, false positives and false negatives. By moving the threshold to a higher level the sensitivity can be improved which makes the criterion for a positive test less strict. The specificity can be improved by moving the threshold to a lower value which makes the criterion for a positive test more strict. Thus, there is a tradeoff between sensitivity and specificity. Therefore, a Receiver Operator Characteristic curve (ROC) can be used to find the optimal threshold. The ROC curve plots the true positive rate against the false positive rate for different thresholds.

The ROC curve demonstrates the following:

- It shows the tradeoff between sensitivity and specificity an increase of the sensitivity is accompanied by a decrease of the specificity
- The test is less accurate if the ROC curve comes closer to the 45 degree diagonal line
- The test is more accurate if the curve comes close to the left and upper border
- The area under the curve is a measure of accuracy:
  - 0.9 1 excellent 1.8 - 0.9 good 0.7 - 0.8 fair 0.6 - 0.7 poor 0.5 - 0.6 fail

Perform the following steps:

- 1. Open the Receiver Operator Curve window from the Classification menu
- 2. Press the **Browse** button to load feature matrix file selffmseg.mat from

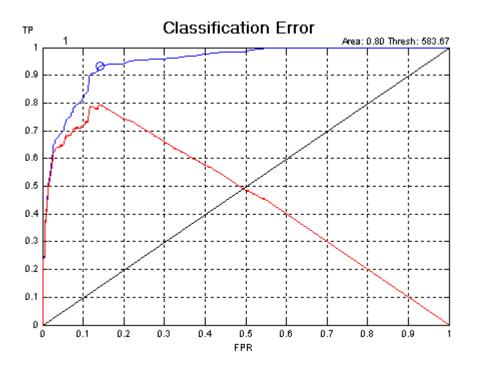
Documents\gtec\gBSanalyze\testdata\SelfPaced\featurematrix

The features matrix has 3 channels and 4018 trials

| 承 Receiver Operator Curve   | _         |           | ×  |
|---|-----------|-----------|----|
| Calculate a Reiceiver Operator Curve to find a threshold between two classes. The corresponds to the quality of classification. | e area be | low the R | ос |
| Load FEATURE MATRIX:  |           |           |    |
| Name of feature matrix file:testdata\SelfPaced\featurematrix\selffmseg.mat  | Bro       | wse       |    |
| Dimension: 3 4018 1 [features x trials x time points  | \$]       |           |    |
| OPTIONS:  | ~         |           |    |
| Result procedure: Show with Result2D Automatic treemaker  | is: ena   | bled      | ~  |
| Save results Filename: enter filename   |           |           |    |
| Help Cancel   |           | Start     |    |

- 3. Select feature 3
- 4. Press the **Start** button

gResult2d opens with the ROC curve. The true positive rate (TP) is plotted on the y-axis, the false positive rate (FPR) is plotted on the x axis. The area under the blue ROC curve is 0.8 which corresponds to a good accuracy and the optimal threshold level is 583.67 (indicated by the circle). The circle marks also the maximum of the red HF (hit-false) difference curve.



To perform the example demonstrated above from the MATLAB command line use the following code:

```
%Load Feature Matrix
```

```
F_O=featurematrix;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\SelfPaced\featurematrix\selffmseg.mat'
];
F_O=load(F_O,FileName);
```

#### %Receiver Operator Curve

```
FeatureNumber=[3];
NrBins=[100];
ClassNumber=[1];
FileName=[''];
ProgressBarFlag=[0];
D_O=gBSreceiveroperatorcurve(F_O,FeatureNumber,ClassNumber,...
NrBins,FileName,ProgressBarFlag);
```

# **DSLVQ Feature Weighting**

DSLVQ can be used to analyze the importance of specific features to a discrimination task.

The window has the following OPTIONS settings:

Codebooks per class ... define the number of codebooks for each class

Alpha ... define the learning speed of the algorithm

**Epochs** ... define the number of iterations

Bootstraps ... number of bootstrap repetition

Evaluation ... percentage of data used for testing'

Perform the following steps:

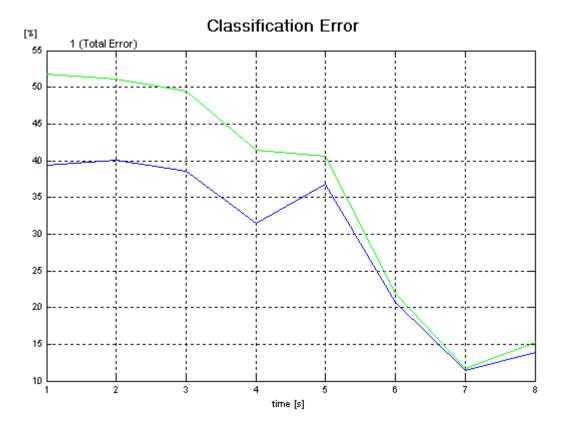
1. Open DSLVQ Feature Weighting from the Classification menu and Browse for the feature matrix bandpower.mat under

Documents\gtec\gBSanalyze\testdata\BCI\featurematrix

| 承 DSLVQ Feature Weigl  | nting      |                       |             |                       | _           |       | ×  |
|--|------------|-----------------------|-------------|-----------------------|-------------|-------|----|
| Weights each feature according to its importance to the classification task. |            |                       |             |                       |             |       |    |
|  | /IATRIX: - |                       |             |                       |             |       |    |
| Name of feature mat  | rix file:  | alyze\testdata\BCI\fe | aturematrix | bandpower.mat         | Br          | owse  |    |
| Dime   | nsion:     | 4 160 8               | [featu      | res x trials x time p | points]     |       |    |
| C OPTIONS:   |            |                       |             |                       |             |       |    |
| Codebooks per  | 2          | Alpha:                | 0.05        | Epochs                | s: 4000     |       |    |
| Bootstraps:  | 20         | Evaluation            | 50          |                       |             |       |    |
| Result procedure:  | Show v     | vith Result2D         |             | Automatic treem       | nakeris: en | abled | ~  |
|  | Save re    | sults Filenam         | e:Nfea      | turematrix\dslvqfv    | w\dslvqfwbj | p.mat |    |
|  |            | [                     | Help        | Can                   | icel        | Sta   | rt |

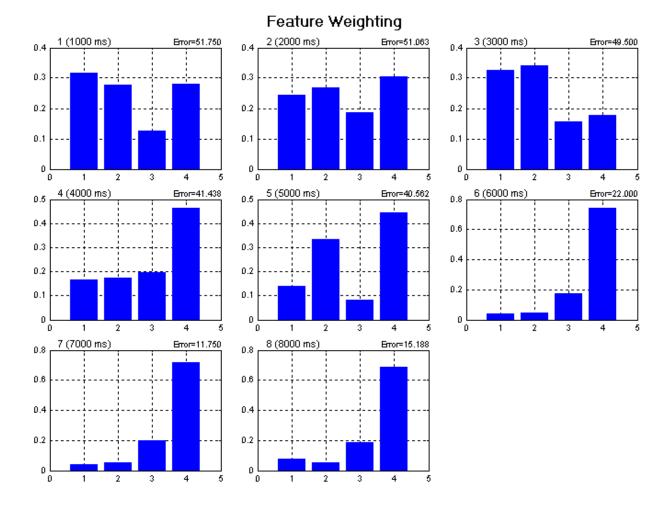
- 2. Set the number of **Epochs** of 4000
- 3. Check Save results and enter the filename dslvqfwbp.mat
- 4. Press **Start** to perform the feature weighting

Page 1 of gResult2d show the classification error of the training data (blue line) and of the testing data (green line). The minimum is reached at second 7. During the first 5 seconds the error rate of the testing data is higher as the training error.



The second page of gResult2d shows the feature weights for each time point. From second 1 to 3 the error is around 50 % and drops down to a minimum of 11.75 % at second 7.

The blue bars represent the importance of each feature to the discrimination task. At second 1 features 1 is the most important one, followed by 4, 2 and 3. But the classification error is 51,75 % and therefore the result is random. At second 7 the error is 11,75 % and therefore the feature weighting can be considered as reliable. Therefore, feature 4 (bandpower in the beta range of channel 2) is the most important one. The bar of feature 3 is much smaller but the feature can still be considered as important for the classification task. Features 1 and 2 are not important and should not be considered for the discrimination.



To perform the example demonstrated above from the MATLAB command line use the following code:

#### %Load FeatureMatrix

```
F_M=featurematrix;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\bandpower.mat'];
F_M=load(F_M,FileName);
```

#### %DSLVQ Feature Weighting

```
P.CBperclass=[2];
P.alpha=[0.05];
P.epochs=[4000];
P.bootstraps=[20];
P.evaluation=[50];
Method=['DSLVQ'];
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\dslvqfwbp.mat'];
ProgressBarFlag=[0];
C O=gBSdslvqfeatureweighting(F M,Method,P,FileName,ProgressBarFlag);
```

# **KMEANS Clustering**

A very common method to find the optimal position of codebook vectors is k-means. The codebook vectors are approximated iteratively.

Perform the following steps to perform an unsupervised clustering of the data:

- 1. Open KMEANS Clustering from the Classification menu
- 2. Click on the **Browse** button and search for the feature matrix file 3clfm.mat which is stored under

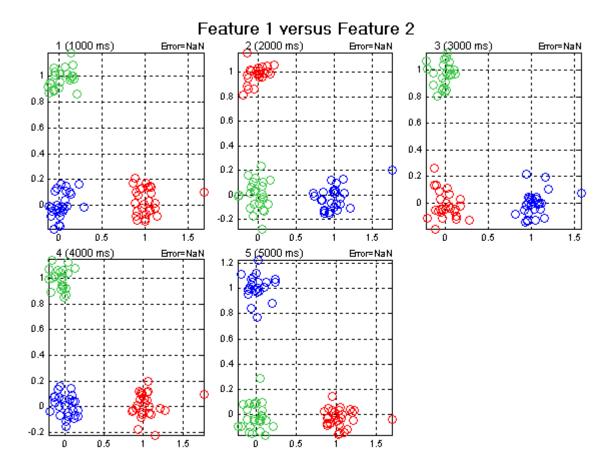
Documents\gtec\gBSanalyze\testdata\Classify\featurematrix

The feature matrix contains 2 feature channels with 150 examples and 3 classes. The feature matrix was calculated for 5 time points. The **No. of clusters** is set to the number of loaded classes.

3. Select Train 50 % - Test 50 % and press the Start button

| 🛦 KMEANS Clustering — 🗆 🗙   |   |
|---|---|
| Perfom a KMEANS clustering of the feature matrix. The method takes specific trials/samples for training an tests it on the remaining samples. | d |
| Load FEATURE MATRIX:  | ٦ |
| Name of feature matrix file:lyze\testdata\Classify\featurematrix\3clfm.mat Browse   |   |
| Dimension: 2 150 5 [features x trials x time points]  |   |
| _ OPTIONS:  | _ |
| No. of clusters: 3 Stop 0.98 Epochs: 1000   |   |
| Select FEATURE CHANNELS:       Map feature       1       against:       2   |   |
| Choose METHOD and OPTIONS:  | 7 |
| Classification method: Clustering ~   |   |
| Training / test-sets: Train 50 %-Test 50%   |   |
| Result procedure: Show with Result2D Automatic treemaker is: enabled ~  | 1 |
| Save results Filename: enter filename   |   |
| Help Cancel Start   |   |

gResult2d shows the clustering results for all 5 time points. The method used the first 50 % of the data for finding the codebook vectors and used the result to cluster the test data. Basically the k-means algorithm was able to find un-subervised the 3 classes. The outlier (trial 128) yielded to a wrong clustering result.



To perform the example demonstrated above from the MATLAB command line use the following code:

#### %Load FeatureMatrix

```
F_M=featurematrix;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\Classify\featurematrix\3clfm.mat'];
F_M=load(F_M,FileName);
```

#### %KMEANS Clustering

```
P.ncluster=[3];
P.stoperror=[0.98];
P.epochs=[1000];
PlotFeatures=[1 2];
Method=['KMEANS'];
TrainTestData=['50:50'];
FileName=[''];
ProgressBarFlag=[0];
C_O=gBSkmeansclustering(F_M,Method,P,TrainTestData,PlotFeatures,...
FileName,ProgressBarFlag);
```

# **Using the Classifier**

After generating a classifier with the linear or non-linear methods it is possible to use this classifier for the classification of new data.

There are two ways:

<u>Apply Classifier</u> – classifies data of the Data Editor with the classifier and generates a new channel in the Data Editor. This channel represents the output of the classification method.

<u>Test Classifier</u> – test the generated classifier on new data and produce an error rate curve and feature cloud in gResult2d

## Apply Classifier

Perform the following steps:

1. Load the data-set session1234bp.mat from

Documents\gtec\gBSanalyze\testdata\BCI

into the Data Editor. The Data Editor visualizes 2 EEG channels, 1 trigger channel and 4 bandpower channels.

2. Open the **Cut Trials Channels** window from the **Transform** menu and exclude channels 1, 2 and 3

| 承 Select                              |                           |   | – 🗆 X                                |
|---------------------------------------|---------------------------|---|--------------------------------------|
|                                       |                           | y it is appropriate not to include all channels<br>or excluded for this step. Hold the 'Ctrl' key |                                      |
| Specify TRIALS:                       | Specify CHANNELS:         | Specify CHANNEL TYPE:   | Specify CHANNEL NUMBERS:             |
| exclude      include only             | exclude      include only | exclude      include only   | exclude      include only            |
| ARTIFACT A<br>REMOVE<br>LEFT<br>RIGHT | BAD<br>CUT<br>BP          | N.S   | 1<br>2<br>3<br>4<br>5<br>6<br>7<br>7 |
| Current selection:                    | Current selection:        | Current selection:  | Current selection:                   |
| 0 out 160 trials                      | 0 out 7 ch.               | 0 out 7 ch.   | 3 out 7 ch.                          |
|                                       |                           | Help  | Cancel OK !                          |

3. Open Apply Classifier from the Classification menu and Browse for the classifier file NNbp.mat under

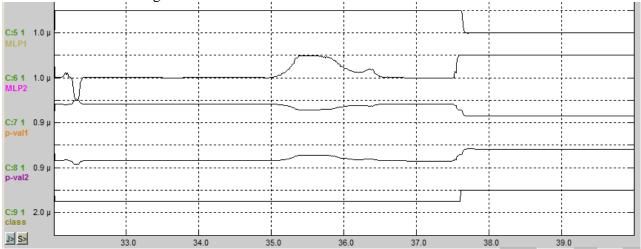
Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\nn

The listbox shows all classification time points with the corresponding classification error.

| Apply Classifier   | – 🗆 X   |
|--|---|
| This function applies a classifier (i.e. a weight vector) calculated befo<br>that the correct feature channels (same as for generation) are used.<br>Load and select CLASSIFIER:           | re to the new test data. Make sure,   |
| Load classifier:       Select classifier from list:        stdata\BC\/featurematrix\nn\\N\bp.mat          Method:       MLP         Add zero class       Confidence interval [%]       2.5 | 48.125   1000<br>57.5   2000<br>53.125   3000<br>38.125   4000<br>34.375   5000<br>21.25   6000<br>12.5   7000<br>13.125   8000 |
| Result procedure:       Add new channels       Automa         O Replace all channels       Filename:         Save result data  | tic treemaker is: enabled   |
| Help   | Cancel Start  |

- 4. Select the appropriate classifier (normally the best one) and check Add new channels
- 5. Press Start to classify the data

After classification the Data Editor shows five additional channels with the classification result of the neural network. The first two channels display the two outputs of the neural network. Channel 5 reaches at the end of the trial zero, channel 6 reaches at the end 1, therefore the trial belongs at the end of the trial to the second class.



Channel 7 and 8 represent the probabilities that the data would be assigned to one of the two classes. A value of 0 means the data does not belong to the corresponding class and a value of 1 means that it belongs to the corresponding class. From channel 7 it can be deduced that it is up to 80% probable that the trial belongs at the beginning to class 1 and with 72% to the

second class at the end of the trial. This fact is also indicated by channel 9 added by the **ApplyClassifier** function. A value of 1 means that this actual time point belongs to the first class and a 2 means that the time point belongs to the second class.

6. Repeat steps 1 to 5 but check the **Add zero class** and select a value of 40 for the **Confidence interval [%]** parameter

Now the channel 9 indicating the class assignment shows an additional zero class. The **ApplyClassifier** function assigns the data samples to this virtual class whenever the probability that the selected class assignment is wrong is higher than the value of the **Confidence interval [%]** parameter which was set to 40 %. This is the case in the middle of the trial between 35 s and 36 s and when the class assignment switches from class 1 to class to between 37.5 s and 38 s.

|             |   |      |     |    |     | i      |     |   |            |     |
|-------------|---|------|-----|----|-----|--------|-----|---|------------|-----|
|             |   |      |     |    |     |        | 1   |   |            |     |
| C-0 4 2 0   |   |      |     |    |     |        |     | ( |            |     |
| C:9 1 2.0 µ |   |      |     |    |     |        |     |   | <br>!<br>! |     |
| Class       |   |      |     | Ì  |     | 1      | 1   |   |            |     |
| J> S>       | 3 | 30 3 | 4.0 | 35 | 0 3 | 36.0 3 | 7.0 | 3 | 80 3       | 9.0 |

To perform the example demonstrated above from the MATLAB command line use the following code:

### %Load Data

```
P_C=data;
File=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\session1234bp.mat'];
P_C=load(P_C,File);
```

### %Select Trials and Channels

```
trial_id=[];
channel_id=[];
type_id=[];
channelnr_id=[1 2 3];
flag_tr='tr_exc';
flag_ch='ch_exc';
flag_type='type_exc';
flag_nr='nr_exc';
[TrialExclude, ChannelExclude]=gBSselect(P_C,trial_id,flag_tr,...
channel_id,flag_ch,type_id,flag_type,channelnr_id,flag_nr);
P_C=gBScuttrialschannels(P_C,TrialExclude,ChannelExclude);
```

### %Load Classifier

C\_O\_S=classifierobj; FileName=['C:\Users\' getenv('USERNAME') '\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\nn\NNbp.mat']; C\_O\_S=load(C\_O\_S,FileName);

#### %Apply Classifier

```
ClassifierNumber=[7];

Replace=['add channels'];

FileName=[''];

ProgressBarFlag=[0];

ConfidenceInterval=[];

P_C=gBSapplyclassifier(P_C,C_O_S,ClassifierNumber,Replace, ...

ConfidenceInterval,FileName,ProgressBarFlag );
```

### **Test Classifier**

After calculating a classifier it is possible to test the classifier on new data.

Perform the following steps:

1. Open Test Classifier from the Classification menu and in the Load FEATURE MATRIX section Browse to the feature matrix bandpower.mat from

 ${\tt Documents \gtec \gBS analyze \test data \BCI \feature matrix}$ 

2. In the Load and select CLASSIFIER section Browse for the classifier file NNbp.mat under

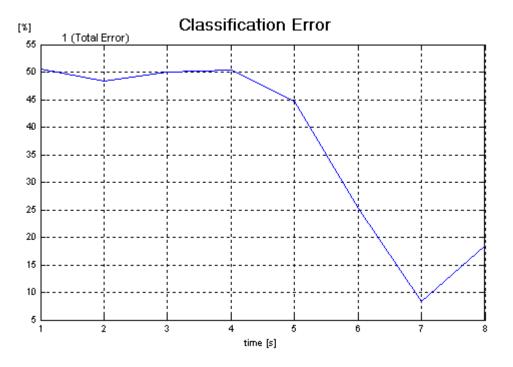
Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\nn

and select the classifier from second 7

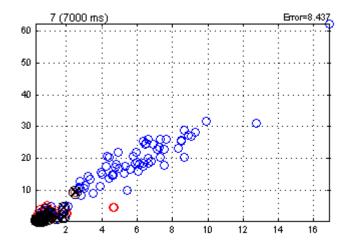
| Test Classifier   | – 🗆 X   |  |  |  |  |  |  |  |
|---|---|--|--|--|--|--|--|--|
| Test a classifier on a specific feature matrix.   |   |  |  |  |  |  |  |  |
| Load FEATURE MATRIX:  |   |  |  |  |  |  |  |  |
| Name of feature matrix file:alyze\testdata\BC\\featurematrix\bandpow  | er.mat Browse   |  |  |  |  |  |  |  |
| Dimension: 4 160 8 [features x trials   | s x time points]  |  |  |  |  |  |  |  |
| L Load and select CLASSIFIER:   |   |  |  |  |  |  |  |  |
| Select best classifier from list:<br>Load classifier:<br>stdata\BCI\featurematrix\nn\NNbp.mat<br>Method: MLP              | 48.13 %         1000.00 ms         57.50 %         2000.00 ms         53.13 %         3000.00 ms         38.13 %         4000.00 ms         34.38 %         5000.00 ms         21.25 %         6000.00 ms         12.50 %       7000.00 ms         13.13 %         8000.00 ms |  |  |  |  |  |  |  |
| Select FEATURE CHANNELS:         Map feature no.:       3         against:       4  |   |  |  |  |  |  |  |  |
| Result procedure:       Show with Result2D       Automa         Open validation window       Save results       Filename: | tic treemaker is: enabled ~   |  |  |  |  |  |  |  |
| Help  | Cancel Start  |  |  |  |  |  |  |  |

### 3. Press the **Start** button

gResult2d display the time course of the classification error. The minimum is reached at second 7 and is 8 %.



Page 2 of gResult2d displays the feature map of feature 3 versus feature 4.



To perform the example demonstrated above from the MATLAB command line use the following code:

#### %Load Feature Matrix

```
F_O=featurematrix;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\bandpower.mat'];
F_O=load(F_O,FileName);
```

#### %Load Classifier

```
C_O=classifierobj;
FileName=['C:\Users\' getenv('USERNAME')
'\Documents\gtec\gBSanalyze\testdata\BCI\featurematrix\nn\NNbp.mat'];
C O=load(C O,FileName);
```

#### **%Test Classifier**

```
ClassifierNumber=[7];
PlotFeatures=[3 4];
FileName=[''];
ProgressBarFlag=[0];
C_O=gBStestclassifier(F_O,C_O,ClassifierNumber,PlotFeatures,FileName,...
ProgressBarFlag);
```

### **Classification Output Mapping**

After applying a classifier to a dataset, it is possible to plot the classification outputs and calculate the classification error rates averaged over trials for the selected classes and channels.

Perform the following steps:

1. Load the data-set mi\_2class\_80trials.mat from

Documents\gtec\gBSanalyze\testdata\Classify\classifieddata

into the Data Editor. The Data Editor displays 5 channels with classification results from a two-class Motor Imagery BCI experiment, which was created with linear discriminant analysis (LDA). The first two channels display the two outputs of the LDA classification, channels 3 and 4 represent the probabilities that the data would be assigned to one of the two classes and channel 5 indicates the number of this class.

Open the Classification Output Mapping window from the Classification menu, select the RIGHT and LEFT classes in the Select Class listbox and select a value of 5 % for the Significance level (alpha) parameter.

| Classification Output Mapping  | – 🗆 X                               |  |  |  |  |  |  |
|--|-------------------------------------|--|--|--|--|--|--|
| Plots the classification outputs and calculates classification error rates averaged over trials for the selected classes and channels. |                                     |  |  |  |  |  |  |
| Select class: ARTIFACT   |                                     |  |  |  |  |  |  |
| Select CHANNELS:   | Select channels                     |  |  |  |  |  |  |
| Result procedure:       Show with Result2D       Automatic treen         Show with Result3D       Save results       Filename:         | naker is: enabled    enter filename |  |  |  |  |  |  |
| Help Car   | ncel Start                          |  |  |  |  |  |  |

3. Open the **Select channels** menu by clicking on the **Select channels** button. Then, add (into the **Selected channels** list) the first 2 channels displayed into the **Available channels** list, then press the **OK** button. The first channel is assigned to the first

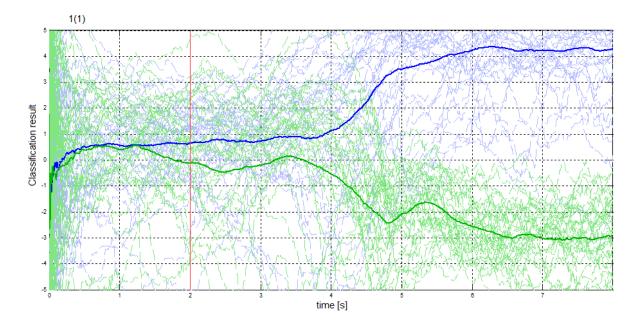
selected class (RIGHT), and the second channel is assigned to the second selected class (LEFT).

| Select channels   | – 🗆 X                               |  |  |  |  |  |  |  |
|---|-------------------------------------|--|--|--|--|--|--|--|
| Select channels for further processing. Currently available channels are listed by channel number and channel name on the left. Calculations are applied only on the selected channels shown in the right list box. |                                     |  |  |  |  |  |  |  |
| Select CHANNELS:  |                                     |  |  |  |  |  |  |  |
| Available channels:<br>number / name  | Selected channels:<br>number / name |  |  |  |  |  |  |  |
| 3       2         4       2         5       2         Add to list ->         <- Remove from list  | 1 2 2 1 2                           |  |  |  |  |  |  |  |
| · · · · · · · · · · · · · · · · · · ·   | · · · · · ·                         |  |  |  |  |  |  |  |
| Help  | Cancel OK !                         |  |  |  |  |  |  |  |

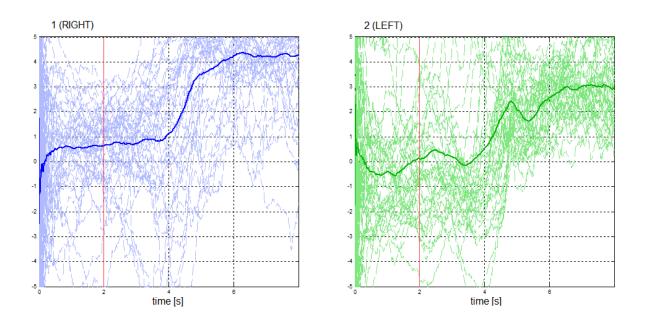
# 4. Check the **Show with Result2D** and **Save results** boxes, enter a filename to store the results and press the **Start** button.

ON the first page, gResult2D displays the trials (dashed lines) and trial averages (solid lines) for both classes. The assigned colors are blue for the first selected class (RIGHT) and green for the second selected class (LEFT). The red vertical line represents the time-point of the trigger. The y-axis shows the classification result (in this case the LDA distance), and the x-axis presents the timing of a single trial in seconds.

**Please note:** For an easier graphical comparison between the two classes, the second output channel (assigned to class LEFT) was inverted.

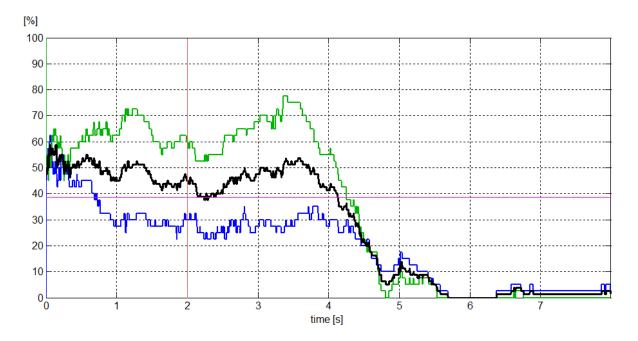


The second gResult2D page displays the trials and trials averages for each class in separate plots, as in the example below. This time the second output channel was not inverted.



The third gResult2D page displays the classification error time courses for individual classes (blue for class RIGHT and green for class LEFT) and the total error (black), averaged over all trials.

The violet horizontal line represents the upper border of the confidence interval. Based on it, the user can decide if the achieved error rates are statistically significant.

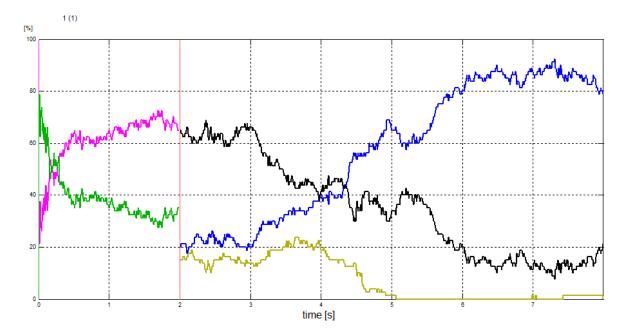


- 5. Repeat steps 1 to 4 but select channels 3 and 4 to map the classification probability outputs. The gResult2D will contain the same pages described before.
- 6. Repeat steps 1 to 4, but select channel 5 only.

If the **Apply Classifier** was done with the **Add zero class** parameter checked, the first two pages of gResult2D will contain the same features as presented before, page number 3 will display the statistical measures of the classification process, and page number 4 will contain the classification error rates.

The color assignment for the statistical features shown in the third page of gResult2D is presented below:

| — | True True Positive (TTP) – correct detections after the cue.  |  |  |  |  |
|---|---|--|--|--|--|
|   | False True Positive (FTP) – false detections after the cue.   |  |  |  |  |
|   | False Negative detections (FN) – all zero-class samples after the cue.                              |  |  |  |  |
|   | False Positive detections (FP) – all detections before the cue that are not assigned to zero-class. |  |  |  |  |
|   | True Negative detections $(TN)$ – all samples correctly assigned to zero-class before the cue.      |  |  |  |  |



To perform the example shown above from the MATLAB command line, please use the following code:

### %Load Data

```
P_C=data;
File=['C:\Users\' getenv('USERNAME') '\Documents\gtec\gBSanalyze' ...
'\testdata\Classify\classifieddata\mi_2class_80trials.mat'];
P_C=load(P_C,File);
```

#### %Classification Output Mapping

```
ClassIndex=[3 4];
ChannelExclude=[3 4 5];
TrialExclude=[];
FileName=['C:\Users\' getenv('USERNAME') '\Documents\gtec\gBSanalyze\'...
'testdata\Classify\Classifieddata\classificationoutputmapping.mat'];
SignificanceLevel=[5];
ProgressBarFlag=1;
V_O = gBSclassificationoutputmapping(P_C,ClassIndex,ChannelExclude,...
TrialExclude,FileName,SignificanceLevel,ProgressBarFlag);
result2D = CreateResult2D(V_O);
gResult2d(result2D);
```

### **Data Access**

g.BSanalyze stores the feature matrix and the classifier data in specific objects. To access both objects the get and set commands can be used:

### Using the get Command

The get method provides a way to access the object entries

### Syntax

```
get(C 0 S, 'PropertyName')
```

returns the value of the property 'PropertyName' of the object C O S

### Example

get(C O S, 'out err')

### Using the set Command

The set method provides a way to set object properties.

### **Syntax**

set(F 0 S, 'PropertyName', 'PropertyValue')

assigns the 'PropertyValue' to the specified 'PropertyName' of the object.

### Example

set(F O S, 'SamplingFrequency', 128)

### Accessing the Feature Matrix

g.BSanalyze stores feature matrix data in an object called F\_O of class featurematrix. If the data are stored to harddisk the name of the data object is changed to F\_O\_S.

| Entry              | Description                               |  |  |  |
|--------------------|---|--|--|--|
| Features           | Feature matrix of each time point         |  |  |  |
| <b>ClassLabels</b> | Class label for each sample. Each row     |  |  |  |
|                    | corresponds to a class                    |  |  |  |
| Interval           | Mode 1:                                   |  |  |  |
|                    | [Start Step End]                          |  |  |  |
|                    | Mode 2:                                   |  |  |  |
|                    | [Time Point 1 - Time Point 2]             |  |  |  |
|                    | Mode 3 :                                  |  |  |  |
|                    | [Segment 1 Start - Segment 1 End          |  |  |  |
|                    | Segment 2 Start - Segment 2 End           |  |  |  |
|                    | ]   |  |  |  |
| Classes            | Class names                               |  |  |  |
| ChannelExclude     | Number of channels which were excluded    |  |  |  |
| FileName           | Name of featurematrix file                |  |  |  |
| SamplingFrequency  | Sampling rate of data file                |  |  |  |
| Mode               | Mode=1 generated from trial attributes    |  |  |  |
|                    | Mode=2 generated from time points         |  |  |  |
|                    | Mode=3 generated from segments            |  |  |  |
| TrialNumber        | TrialNumber for each feature matrix trial |  |  |  |

### Accessing the Classifier Object

g.BSanalyze stores classifier data in an object called C\_O of class classifierobj. If the data are stored to harddisk the name of the object is changed to C\_O\_S.

| Entry               | Description   |  |  |  |  |
|---------------------|---|--|--|--|--|
| Out_err             | Stores the classification error and classification  |  |  |  |  |
|                     | time point.   |  |  |  |  |
|                     | First row: time point                               |  |  |  |  |
|                     | Second row: total error                             |  |  |  |  |
|                     | Third row: error of class 1                         |  |  |  |  |
|                     |   |  |  |  |  |
|                     | N-row: error of class N                             |  |  |  |  |
| Out_clssfyr         | Classifier or weight vector for each time point.    |  |  |  |  |
|                     | Each row corresponds to a time point.               |  |  |  |  |
| Out_erridx          | Trial id for wrong classified trials for each class |  |  |  |  |
|                     | and time point                                      |  |  |  |  |
| Out_clstest         | Classification result for each time point           |  |  |  |  |
| Out_rmse            | Root mean squared error (only for neural            |  |  |  |  |
|                     | networks)   |  |  |  |  |
| TPC                 | Trials per class                                    |  |  |  |  |
| Interval            | Mode 1: [Start Step End]                            |  |  |  |  |
|                     | Mode 2: [Time point 1 start - time point 1 end      |  |  |  |  |
|                     | Time point 2 start – time point 2 end               |  |  |  |  |
|                     |   |  |  |  |  |
|                     | ]   |  |  |  |  |
| AttrNr              | Class name  |  |  |  |  |
| Methods             | Classification method                               |  |  |  |  |
| TrainTestData       | CV, 100:100, 100:0, 50:50, 0:100                    |  |  |  |  |
| Metric              | Euclidian or Mahalanobis                            |  |  |  |  |
| InputObj            | Classifier method object                            |  |  |  |  |
| Features            | Features for each time point                        |  |  |  |  |
| <b>PlotFeatures</b> | Number of features to plot as cloud                 |  |  |  |  |

# Help

g.BSanalyze and the g.CLASSIFYtoolbox provide a printable documentation and a function help.

The printable documentation is stored under

Your MATLAB path\gtec\gBSanalyze\Help

as gCLASSIFYtoolbox.pdf. Use Acrobat Reader to view the documentation.

To view the function help type

help gBSfunctionname

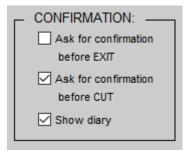
under the MATLAB command window.

To view all functions that are available in batch mode type

gBSfunctions

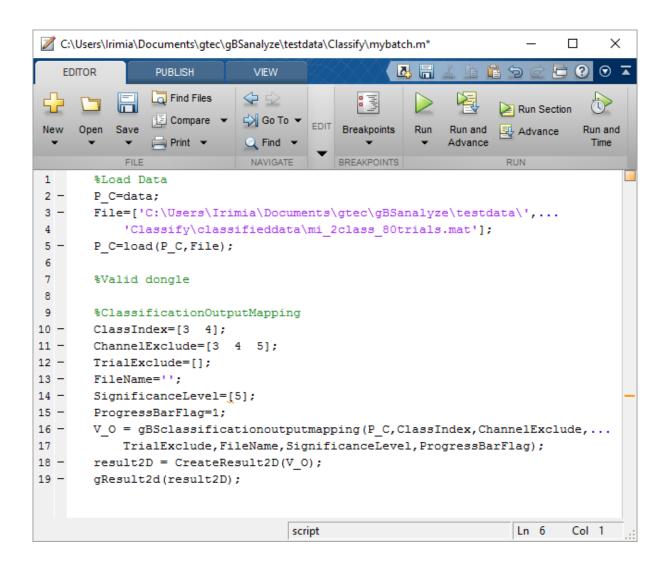
## **Batch Mode**

The easiest way to create a batch for data processing is to perform the analysis under the Data Editor with the graphical user interfaces. Make sure that the **Show diary** checkbox is enabled in **Appearance Settings** under the **Options** menu.



This forces g.BSanalyze to report all calculations in the MATLAB command window. After finishing the analysis open a **New Script** and copy and paste all commands into the file.

| 📣 MA               | TLAB     | R2015a                   |                |   |              |  |  |
|--------------------|----------|--------------------------|----------------|---|--------------|--|--|
| но                 | ME       | PLOTS                    | APPS           |   |              |  |  |
| New<br>Script      | New      |                          | mport Sa       | New Variable  New Variable  Open Variable  Ve  Space  VARIABLE                | Analyze Code |  |  |
| <                  |          | Script 🔓 Ctrl+N          |                | ocuments + gtec + gBSanalyz   |              |  |  |
| Curren             | fx       | Function                 | Command Window |   |              |  |  |
|                    | P        | Example                  |                | %Load Data<br>P C=data;   |              |  |  |
| ⊕ <mark> </mark> f | <u>}</u> | Class                    |                | File='C:\Users\Irin<br>P C=load(P C,File)                                     |              |  |  |
|                    | °-       | System Object >          |                | %Valid dongle   | ,            |  |  |
|                    |          | Figure                   |                | %ClassificationOut;   | nutManning   |  |  |
|                    |          | Graphical User Interface |                | ClassIndex=[3 4];   |              |  |  |
|                    | 7        | Command Shortcut         |                | ChannelExclude=[3<br>TrialExclude=[];   | 4 5];        |  |  |
|                    | SIMU     | ILINK                    |                | FileName='';<br>SignificanceLevel=  | [5];         |  |  |
|                    | ▶        | Simulink Model           |                | ProgressBarFlag=1;  |              |  |  |
|                    |          | Simulink Project >       |                | <pre>V_0 = gBSclassific<br/>result2D = CreateR<br/>gBooult2d (result2D)</pre> | esult2D(V_O) |  |  |
|                    |          |                          |                | gResult2d(result2D  | 1.           |  |  |



Save the batch in your own directory as  ${\tt mybatch.m}$  and start the batch under the MATLAB command window with

mybatch

For further data-sets just replace the input data file to perform the same analysis.

# **Product Page**

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- Additional demonstrations



# contact information

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