Matching pursuit decomposition in Svarog

Mp5 software for <u>matching pursuit</u> (MP) decomposition of (biomedical multivariate) time series was developed at the <u>University of Warsaw</u>, <u>Faculty of Physics</u>. The algorithm is based upon the seminal paper by Mallat and Zhang (<u>Mallat1993</u>) with modifications described partly by Durka et al. (<u>Durka2001</u>) and Kuś et al. (<u>Kuś2013</u>). Mp5 is designed for batch processing input signals and writing results to disk files called – after (<u>Mallat1993</u>) – *decomposition books*. These books (*.b) contain parameters of the functions selected for representation of the analyzed signal.

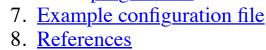
Svarog offers an interactive interface for MP decomposition, allowing for:

- 1. simple selection of the part of the signal for MP decomposition,
- 2. running mp5 decomposition according to selected parameters
- 3. visualization of the results in terms of interactive time-frequency maps of signal's energy density.

Some of the topics presented below are covered in a screencast available at <u>http://braintech.pl/svarog/screencast.html</u>. *Caveat*: default settings are optimized for speed rather than quality of the decomposition. This may be sufficient for the first try, but serious use requires some understanding of the procedure.

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Configuration

By default, Svarog looks for the file containing the mp5 binary in PATH. If you installed the system from .deb packages (see <u>http://deb.braintech.pl</u>) it should be already preconfigured. If that does not work, you should provide the correct path in the "Preferences" dialog (Edit/Preferences), tab Tools:

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If you downloaded mp5 together with Svarog in one archive, executables for different operating systems can be located under mp5 in subfolders linux, windows, and mac.

Selecting the signal epoch(s) for decomposition

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Using icons in the toolbar, mark in Svarog the part of the signal that you want to decompose and select Tools/MP decomposition from the menu

MP Decomposition configuration						
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Time fragment selection Whole signal Selected Marked Selection type Page selection Block selection Block selection Image: Free (channel) selection Selection parameters Start time 4.044 $\frac{1}{2}$ Length (seconds) 5.212 $\frac{1}{2}$	Channel selection Fp1 Fp2 F7 F3 Fz F4 F8 T7 C3 Cz C4 T8 P7 P3 Pz					
Signal source level O Raw signal O Assembled signal O Filtered signal Presets << select to load preset >> Load signal selection from loaded preset To raw mode Source level To raw mode Source level Signal source level Signal Signal	P4 P8 O1 O2 FCz Select all Select none Invert					

Signal selection tab allows you to change selection or e.g. set up decomposition of the whole signal page by page. Settings in other tabs correspond to the parameters of decomposition that will be written to the configuration file. Their meaning is explained in the next chapter.

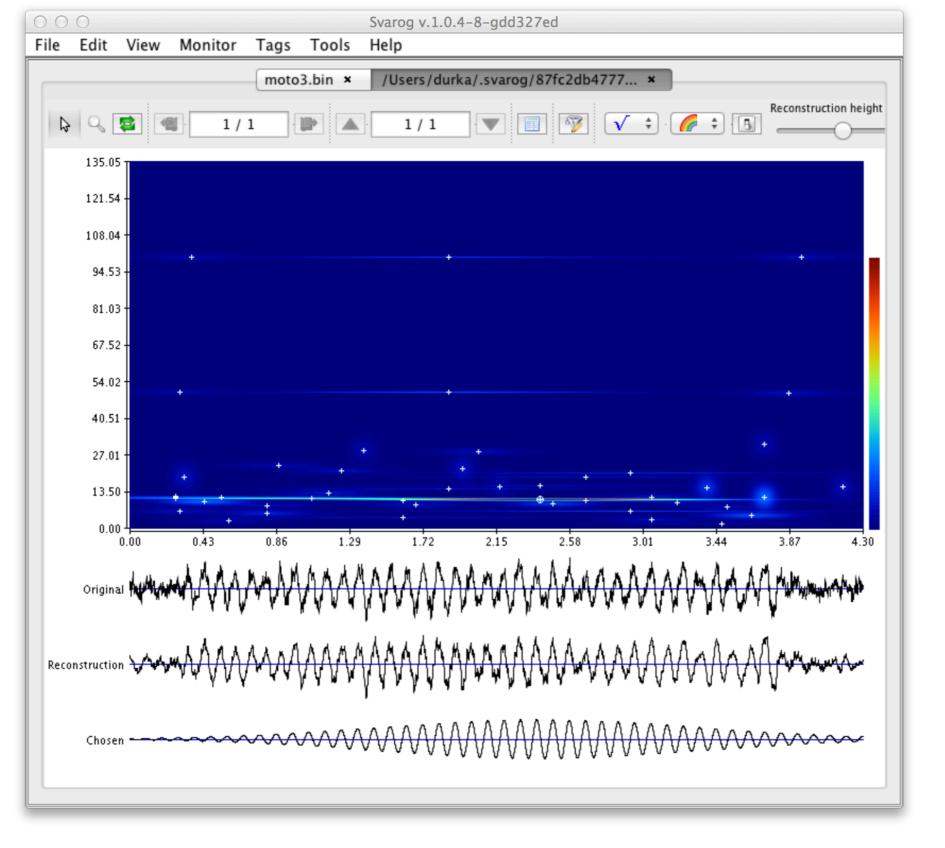
After pressing Ok we get the window showing the progress of the task

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Execution time depends strongly on selected settings and length of the signal being decomposed. It can be quite long for multichannel decompositions. If a single channel was selected (as in the picture above) with default settings, and the sampling frequency was not extremely high (that is, there are no more than few thousand points in the selected epoch) decomposition should not take more than a minute on average PC.

You can move this window aside and continue working with Svarog in the meantime.

After the task completes, Get result button will appear at the bottom of the window. Pressing it opens a dialog where you can choose Open in the viewer and/or Save to disk. The former option opens a new tab in Svarog and diplays the timefrequency map of the signal's energy density, computed from the decomposition:



Below the map the program displays the original signal, its reconstruction from all the atoms from the (just computed) decomposition, and the reconstruction computed from the selected atoms. You can select atoms by clicking the crosses in their centers. After clicking the magnifying glass in the top left, you can zoom selected area of the map.

Detailed settings and structure of the mp5 config file

Parameters of the decomposition, entered in subsequent tabs of the MP Decomposition configuration window, are written to a configuration file, which is in turn passed by Svarog to the mp5 binary. Mp5 config file consists of lines, which can be divided into three groups:

```
\alpha ,
```

- Comments.
- Obligatory settings.
- Additional settings.

Each line, which is not a comment, is treated as a setting for the program. Each setting consists of its name and assigned value. If any of the obligatory settings is missing, mp5 will return error. If the user does not set the additional settings, mp5 sets the default values. Lines with commands and comments can occur within the config file in any order.

Comments in the config file

The user can add his own comments to the configuration file by placing a # sign in front of the line. There are two type of comments:

• lines starting with # are treated as comments and will exist only in the config file, but will be neglected by mp5, for example:

Moly set the numberOfChannels to 5

• lines starting with ## are also treated as comments but they will be copied into the output file (decomposition book), e.g.:

channels selected above motor cortex decomposed with very small dictionary

Although in general the results are written to the binary file, the comments are saved as text and can be viewed by standard text editor.

Input settings — obligatory settings

Signal (time series) for mp5 decomposition must be stored in a binary file as a 4-byte float numbers.

In case of multivariate (multichannel) recordings the values should be multiplexed, that is: slch1, slch2, ..., slchK, s2ch1, s2ch2, ..., s2chK where sxchy is the X-th sample at channel Y, K is the number of channels

nameOfDataFile

Full path to the input file.

nameOfOutputDirectory

The path to the output directory, where the file with results should be saved. The default is the same as location of the input file.

writingMode

- CREATE create a new file for writing (saving results of decomposition).
- APPEND append the results to an already existing file or create a new file if there is no file to append to

Names of the output files are generated automatically; the name of the file with the data is appended with: _smp.b in case of a single channel mp algorithm (SMP), or _mmp.b in case of one of the multichannel mp algorithms (MMP1, MMP2, MMP3). The decomposition book includes also the original analyzed epoch.

numberOfChannels

Number of channels in the input file. Positive integer, range: 1 - 65535.

selectedChannels

In either of the algorithms—monochannel or multivariate—one does not have to use all the channels present in the data file. For example, using SMP we may decompose only few channels, and for MMP we may need to neglect non-EEG channels. This option lists the channels which we want to analyze. That is, to analyze all the channels from a 20-channel datafile config must contain:

numberOfChannels 20 selectedChannels 1-20

Channels are numbered starting from 1. Selection is written either as comma-separated list 1,2,3 or as ranges 1-3. Both can be combined in one line, e.g.:

```
selectedChannels 1, 3, 5, 7-11, 19
```

type: positive integer, range: 1 - 65535

numberOfSamplesInEpoch

When decomposing susequent epochs of a long signal, this option determines the length of the epoch measured as number

of samples. Argument's value: positive integer. Range: 1 - 2147483647

selectedEpochs

Chooses which epochs (of the length defined by numberOfSamplesInEpoch) will be analysed. First epoch in the file is number 1. Written as comma-separated list with ranges as in selectedChannels.

typeOfDictionary

- OCTAVE_FIXED functions distributed in a way to optimally cover the ranges of parameters reasonable for a given signal in a way that the distance between any two neighboring functions does not exceed the threshold given by a user (Kuś et al. 2013)
- OCTAVE_STOCH distribution as in OCTAVE_FIXED, plus a stochastic element to remove the possible bias (Durka et al. 2001). The dictionary with Gabors is created according to the first parameter energyError command, but in the next step a fraction of randomly selected functions are removed from dictionary. The number of atoms left is controlled by the second value passed to the EnergyError parameter.

energyError

This parameter regulates the density of the dictionary. That is, for the same energyError, and different sizes of the analyzed epoch, effective sizes of the dictionary will be larger for longer epochs, but accuracy of the decomposition should be equivalent (except for the border effects). Changing this parameter in the "Basic settings" tab automatically computes the amount of RAM necessary to store the dictionary that will be prepared for decomposition.

The energyError parameter has two values:

- the first one is a threshold for a distance between two nearest atom in dictionary, this value should be in range (0 1)
- the second is value percentage of atoms left after stochastic dictionary reduction; this value is omitted by mp5, when typeOfDictionary is set to OCTAVE_FIXED

For example, the configuration of commands:

typeOfDictionary OCTAVE_FIXED
energyError 0.3 30.0

results in generation of dictionary in which the maximal distance between nearest Gabors does not exceed value 0.3.

With another configuration:

typeOfDictionary OCTAVE_STOCH
energyError 0.3 40.0

the distribution of the atoms in dictionary is performed in such way that the maximal distance between nearest Gabors does not exceed value 0.3. After generation of the dictionary, the number of Gabors will be reduced to 40% of their initial amount (determined by the threshold 0.3) by random selection.

type of the first parameter (energyError): float number range of the first parameter: $(0.0 \ 1.0)$

type of the second parameter (stochastic reduction): float number range of the second parameter: $(0.0\ 100.0>$

randomSeed

If provided, this value will be used for the seed of the random generator used in creations a stochastic dictionary. Otherwise, the seed will be generated from the computer's clock.

For example:

randomSeed 23432

The mp5 will used this number as a seed for random generator. The seed is the same for any stochastic reinitialization of

the dictionary (see the next command: reinitDictionary).

Second example:

randomSeed auto

The mp5 will used the time form the computer's clock as a seed for therandom generator. The seed is different for each reinitialization of the dictionary.

Type: string "auto" or positive integer number range in case of integer: 0 - 2147483647

reinitDictionary

When using the default stochastic (randomized) dictionary, the randomization is called reinitialization of the dictionary. It can be performed:

- REINIT_AT_ALL at each of the analyzed epochs, works only for SMP.
- REINIT_IN_CHANNEL_DOMAIN exactly the same dictionary will be used to decompose all epochs within a channel. Before the analysis of another channel, parameters will be reinitialized, i.e. randomized again. This setting can be applied only in the case of separate decomposition of channels using SMP option.
- REINIT_IN_OFFSET_DOMAIN dictionary will be reinitialized before decomposing each subsequent epoch.
- NO_REINIT_AT_ALL the same dictionary for decompositions in all channels and all epochs.

maximalNumberOfIterations

Maximum number of iterations equals the largest allowed numbers of waveforms fitted to the analyzed epoch. This command together with the next energyPercent control the number of waveforms used for the approximation of the analyzed epoch, which equals the number of algorithms iteration performed before the stopping criterion is fulfilled. The iterations stop either after reaching this number, or after explaining the percentage of signal's energy set in the energyPercent parameter, whatever comes first. That is, if we want to force the decomposition to contain exactly maximalNumberOfIterations iterations, we should set energyPercent to 100. It does not influence the accuracy of the parameters of the waveforms fitted to the signal, in the sense that the 5 waveforms fitted to the signal in 5 iterations run will be just as good as the first 5 fitted in the 100 iterations run.

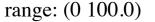
Time of computations is proportional to the number of algoritms iterations (except for the first, longer iteration), and obviously with more iterations we explain larger fraction of signal's energy; OTOH, too many iterations usually make no sense. Unfortunately, the meaning of "too many" depends on the properties of the signal and the aim of decomposition. For further considerations consult e.g. (Durka2007a) and (Durka2007b).

Type: positive integer range: <1 65535>

energyPercent

Algorithm stops after explaining energyPercent % of the energy of the analyzed epoch, that is, when the sum of energies of functions fitted to the signals reaches energyPercent/100 of the signal's energy, unless maximalNumberOfIterations was reached first.

Type: positive float



MP

There are significant differences between the 'classical' MP computed for the univariate time series, and the variety of possible versions of the multichannel MP. Choice of the appropriate flavor of multivariate algorithm is controlled by the MP variable, which can be assigned one of the following values. For equations and formulae see (Durka 2007a) and (Kuś et al. 2013)

- SMP monochannel MP (separate decomposition of single channels)
- MMP1 multichannel algorithm maximizing in each iteration the sum of energies explained in all channels
- MMP2 suboptimal, faster version of MMP1 maximizing in each iteration the sum of products of the waveform with

- all the channels. In theory may degrade in the case when opposite polarities are present across channels
- MMP3 as MMP1, but allowing different phases in each channel.

scaleToPeriodFactor

All Gabor functions, for which the number of periods of the sine, given by this parameter, exceeds the half-width of the Gaussian, will be removed from the dictionary. It regulates removal of non-physical cases. For example, 1 Hz oscillation which lasts 100 milliseconds mathematically represents a well defined waveform. However, the very notion of frequency is dubious in such case. Also, the actul peak-to-peak amplitude of such function may significantly differ from the doubled amplitude of the Gaussian envelope. See (Durka2004)

Type: positive float range: (0.0 3.402823466 E + 38>

samplingFrequency

Floating-point positive number. Decimal dot is obligatory, that is "128.0" and not "128". This parameter has no influence on the decomposition -- value is copied into the decomposition book file and can be later used for scaling the display of results.

pointsPerMicrovolt

Conversion rate from the actual values stored in the datafile to physical units, for EEG microvolts $[\mu V]$, for MEG femtotesla [fT]. This parameter has no influence on the decomposition -- value is copied into the decomposition book file and can be later used for scaling the display of results.

Decimal dot is obligatory. For example,

pointsPerMicrovolt 20.0

in EEG file says that the stored vaules must be multiplied by 20 to get microvolts.

Type: positive float range: (0.0 3.402823466 E + 38>

Input parameters — additional settings

normType

The experimental feature of mp5 program, do not change it. The default value is set to L2.

diracInDictionary

Generate dictionary including Dirac Function. The possible argument of command is YES or NO string.

gaussInDictionary

Generate dictionary including Gauss Function. The possible argument of command is YES or NO string.

sinCosInDictionary

Generate dictionary including Harmonic function. The possible argument of command is YES or NO string.

gaborInDictionary

Generate dictionary including Gabor functions. The possible argument of command is YES or NO string.

progressBar

Plot text bar representing the progress of calculation. The possible argument of command is YES or NO string.

Example configuration file

# OBLIGATORY PARAMETERS					
nameOfDataFile	test.dat				
nameOfOutputDirectory	•/				
writingMode	CREATE				
samplingFrequency	128.0				
numberOfChannels	24				
selectedChannels	1-19				
numberOfSamplesInEpoch	179				
selectedEpochs	1-31				
typeOfDictionary	OCTAVE_FIXED				
energyError	0.3 50.0				
randomSeed	auto				
reinitDictionary	NO_REINIT_AT_ALL				
maximalNumberOfIterations	100				
energyPercent	95.0				
MP	SMP				
scaleToPeriodFactor	1.0				
pointsPerMicrovolt	1.0				
# ADDITIONAL PARAMETERS	T 0				
normType	L2				
diracInDictionary	YES				
gaussInDictionary	YES				
sinCosInDictionary	YES				
gaborInDictionary	YES				
progressBar	ON				

References

(Mallat1993) Stéphane Mallat and Zhifeng Zhang Matching pursuit with time-frequency dictionaries. *IEEE Transactions on Signal Processing* 1993, 41:3397-3415

(Durka2007a) Piotr J. Durka <u>Matching Pursuit and Unification in EEG analysis</u>, Artech House 2007, ISBN 978-1-58053-304-1

(Durka2007b) Piotr J. Durka <u>Matching Pursuit</u>, Scholarpedia, p. 20910, 2007.

(Kuś et al.2013) Rafał Kuś, Piotr T. Różański and Piotr J. Durka Multivariate matching pursuit in optimal Gabor dictionaries: mathematical foundations and software for EEG/MEG (submitted to *Biomedical Engineering Online*)

(Durka et al. 2001) Piotr J. Durka, D. Ircha and K. J. Blinowska Stochastic time-frequency dictionaries for Matching Pursuit, *IEEE Transactions on Signal Processing*, vol. 49, No. 3, pp. 507-510, March 2001.

(Durka2004) Adaptive time-frequency parametrization of epileptic EEG spikes P.J. Durka, Physical Review E, vol. 69,

