How to use

```
[matrixProfile, profileIndex, motifIndex, discordIndex] =
interactiveMatrixProfile(data, subLen);
```

• Input

- data: input time series
- subLen: subsequence length

Output

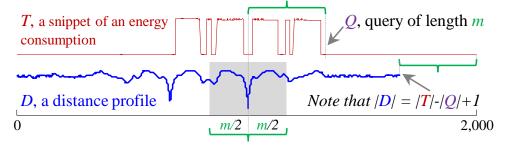
- matrixProfile: the approximated matrix profile when stopped or complete
- profileIndex: the approximated matrix profile index when stopped or complete
- motifIndex: the detected motif index when stopped or complete (3x2 cell)

1 st motif	1 st motif's NN within certain radius
2 nd motif	2 nd motif's NN within certain radius
3 rd motif	3 rd motif's NN within certain radius

• discordIndex: vector contains the discord's index when stop or complete

Exclusion zone

Remember we have talk about exclusion zone in the paper with the following graph.



It can be used to avoid matching the query with subsequence within certain time difference (e.g., if we set it to 10 sec, any two event happens in +-10 sec will not be recognized as motif set). The size of exclusion zone can be change if you look at line 10 in the source code. Currently, it is set to half of the subsequence length as shown in the figure below.

