Burst Variance Analysis: Help & Documentation

Kristofer Gryte¹

¹Department of Condensed Matter Physics, Clarendon Laboratory, University of Oxford, Oxford, United Kingdom

October 21, 2011

Contents

Jverview	1
Burst Variance Analysis	2
Installation	2
Input Arguments	3
Algorithm	4
Visualization	6
BVA Contour Plot	6
Efficiency Histogram	10
Standard Deviation Histogram	11
Concluding Remarks	12
Acknowledgments	13
Code	16
BVA Script	16
	23
BVA Analysis	26
Data Filtering	45

CONTENTS	CONTEN	NTS
Monte Carlo Simulation		54 50

Overview

Burst variance analysis (BVA) attempts to detect dynamic fluctuations in Förster Resonance Energy Transfer (FRET) as individual molecules emitting donor and acceptor fluorescence diffuse through a confocal volume. Motivations for this method reside in the attempt to distinguish between static and dynamic heterogeneity and distribution broadening in confocal histograms. Existing methods proved unable to adequately address the static-dynamic distinction, thus necessitating a burst-by-burst analysis [1].

As detailed in Torella et al, BVA employs a statistical criterion to compare observed experimental standard deviations to simulated standard deviations generated according to theoretical prediction. Observed systems exceeding the strict confidence bounds are considered 'dynamic', where dynamics are assumed to arise due to state interconversion for a sample species. We note that BVA is not conclusive, however, in ascertaining whether observed dynamics arise due to conformational fluctuations, alone. Depending on experimental conditions and setup, artifacts originating from fluorescent dye quenching, dye isomerization, or optical misalignment can obfuscate underlying conformational dynamics, the presence or lack thereof.* On the other hand, what BVA can validate is distribution broadening due to species dynamics, whatever those 'dynamics' might be, as opposed to static heterogeneity, where sample species are in accordance with theoretical prediction. Hence, we urge that BVA be used as one of a suite of tools and experimental controls designed to effectively eliminate artifacts and determine system heterogeneity.

What follows is an overview of the BVA software package and its implementation. The creator of this software has endeavored to the best of his ability to ensure that the functions are well-documented and error-free. Should the user happen to be confused or receive any error messages, s/he is asked to consult the HELP documentation provided as well as the individual HELP sections of the functions provided herein. And should any programmatic mistakes be found, the user is asked to make a detailed note of the problem and report this to the author. Any feedback (comments, criticisms, or questions) is certainly welcome. The author hopes that this package accomplishes its aims and wishes the user the best in unraveling the mystery of single-molecule behavior. Enjoy!

^{*}NOTE: efforts are currently underway within the Kapanidis laboratory to better address this issue and more clearly delineate conformational contributions.

Burst Variance Analysis

Installation

MATLAB® is the programming language for the BVA package. To install BVA,

- download the package either from the MATLAB® Central File Exchange or the laboratory webpage;
- save and unzip the *.zip file to a local directory;
- either add the path to the BVA package directory to startup.m (e.g., addpath(C:\...\BVA)), set the path from the MATLAB® command window (File → Set Path... → Add Folder), or navigate to the BVA folder within MATLAB® to make this the current directory;*
- once the top-level folder for BVA is accessible, the following functions may be run from the command-line:
 - InitalizeAxes.m: configure figure axes.
 - Confocal2ColorALExMultDataBVA_Analysis.m: BVA analysis.
 - PlotResults.m: plot BVA results.
- to run the demo script BurstVarianceAnalysis.m, navigate to the BVA folder within MATLAB® to make this the current directory (sample data is contained in the file MyArrivalMatrix.mat) and type BurstVarianceAnalysis at the command-line;
- edit the demo script accordingly to analyze user data.

The BVA package is OS independent. At the present time, users need a STATS toolbox license to generate random numbers drawn from a binomial distribution. The author apologizes for any inconvenience this may cause.

^{*}NOTE: only the top-level folder of the BVA package needs to be included in the path, as sub-folder access is controlled internally.

Input Arguments

BVA attempts to answer the question as to whether behavior consistent with 'dynamic' fluctuations is present for a sample species. Analysis is strictly dependent on two inputs: (i) a 'colored' photon *arrival matrix* and (ii) the specification of a *window* parameter. Additional input may be provided, such as filtering parameters (e.g, photon thresholds) and confidence interval adjustment (to either strengthen or weaken the barrier for fluctuations to be classified as dynamic). Default parameter values are built-in for all inputs except the data to be analyzed as well as filters/thresholds.* A full list of possible inputs is provided below:

- **ARRIVALMATRIX**: input data on which to perform BVA. The input matrix should be Nx3 or 3xN, where *N* equals the number of photons and with the following as vector inputs:
 - 1. Photon Arrival Times;
 - Detection Channel (DexDem = 0; DexAem = 2; AexAem = 1; AexDem = 3);^{†‡}
 - 3. Burst Classification (those photons belonging to a burst should have an index corresponding to a unique burst ID; background photons should be indexed as NaN)§

The composition of the data matrix should be as follows:

$$ARRIVALMATRIX = \left[ArrivalTimes \vdots Color ID \vdots BurstNumber\right]$$

- **WINDOW**: size of the sliding window (number of photons) over which FRET will be calculated. Default: 5.
- CLUSTERBINS: number of bins over which to cluster bursts characterized by their mean FRET value. Mean standard deviations are calculated across bursts belonging to a particular bin and compared

^{*}See Torella *et al* for an analysis on the effect of parameter values on dynamics detection. The default values are the same as those described elsewhere [1].

^{*}Key: D = donor; A = acceptor; ex = excitation; em = emission

[‡]NOTE: we assume Alternating Laser Excitation (ALEx) spectroscopy [2], although single laser excitation of the donor is compatible.

[§]NOTE: burst classification presumes a burst search has already been performed. The included software package does not include burst search functionality. Needed information includes the START and END time for each burst. For information regarding the burst search method, consult [4].

against a test statistic to determine significance (and here, the presence of dynamics). Default: 20.

- **CLUSTERMINWINDOWS**: threshold for the minimum number of total windows within a cluster from all bursts of the cluster analyzed (i.e., we need to ensure statistical significance). Default: 50.
- **CONFIDENCEINTERVAL**: $(1-\alpha)$ to determine statistical significance of cluster results. Default: 0.999^* .
- **FILTERS**: input structure providing minimum and maximum thresholds for the following data streams[†]:
 - DURATION: the temporal length of a detected burst [seconds]
 - LENGTH: the number of photons within a detected burst [photons]
 - DEXDEM: the number of photons detected in the donor channel upon donor excitation.
 - DEXAEM: the number of photons detected in the acceptor channel upon donor excitation.
 - AEXAEM: the number of photons detected in the acceptor channel upon acceptor excitation.
 - SUMPHOTONS: the number of photons detected in the donor & acceptor channels upon donor excitation.
 - TOTALPHOTONS: the number of photons detected in the donor & acceptor channels upon donor and acceptor excitation. (This excludes photons detected in donor channel upon acceptor excitation: AexDem.)
 - EFFICIENCY: the FRET efficiency.
 - STOICHIOMETRY: the relative fluorophore brightness.

Algorithm

The implementation of BVA is relatively straightforward:

^{*}NOTE: the strict α value stems from a Bonferroni correction for multiple-hypothesis testing, which is dependent, here, on CLUSTERBINS. See [3] for its justification.

[†]NOTE: once again, we assume ALEx; for single-laser excitation, ALEx applicable fields may remain empty.

size: Nx3; N = total photons

- 1. for each burst, bin the photons by arrival time into non-overlapping bins (bin size is defined by WINDOW).
- 2. calculate the FRET (E_i^*) for each bin.
- 3. calculate the standard deviation of E^* across all bins.
- 4. for all bursts with similar mean FRET $\{\mu^{E^*} : a \leq \mu^{E^*} < b\}$, calculate the cluster standard deviation across all bins for all satisfying bursts.
- perform a hypothesis test to determine if the observed standard deviation exceeds confidence bounds for static behavior, thus being indicative of dynamics.

Pseudo-code is provided below:

Input: ARRIVALMATRIX

```
Input: WINDOW
Output: DATA
Var: C = CLUSTERBINS; \alpha = CI; W = WINDOW
Filter bursts;
Partition E^* \to \mu_{\{C\}}^{E^*} : [0:1/C:1]
(C_i = \text{cluster index});
for k := 1 to N
     • Extract photons for burst k from ARRIVALMATRIX
     (L_k = \text{total photons});
     \circ Divide photons by arrival time into M_k non-overlapping windows
     (M_k = L_k/W);
     for j := 1 to M_k
         \circ Calculate E_j^* = F_j^A/L;
                                                                              F^A = \sum_{i=1}^{W} (w_i = 2)
     \begin{array}{l} \circ \text{ Calculate } \mu_k^{E^*}; \\ \circ \text{ Calculate } \sigma_k^{E^*}; \\ \circ \text{ Determine } \mu_k^{E^*} \ \in \forall C_i; \end{array} 
end
for i := 1 to C
```

```
 \circ \forall \text{ bursts satisfying } \mu_k^{E^*} \in C_i \text{, calculate } \mu_{C_i}^{\sigma^{E^*}}; \\ \circ \text{ Calculate shot-noise prediction: } P(\sigma_{SN}^{E^*}) \\ f(p,W,T); p = \mu_{C_i}^{E^*}; T = \Sigma \ (M \ \forall \ \mu_k^{E^*} \in C_i); \\ \circ \text{ Conduct multiple-hypothesis test } (\alpha) \text{ on } P(\sigma_{SN}^{E^*}) \text{ for upper bound: } UB_i; \\ \circ \text{ Compare } \mu_{C_i}^{\sigma^{E^*}} \text{ to } UB_i \text{ for significance; } \\ \text{end} \\ DATA \leftarrow \text{RESULTS;} \\ \text{Visualize } DATA; \\ \end{cases}
```

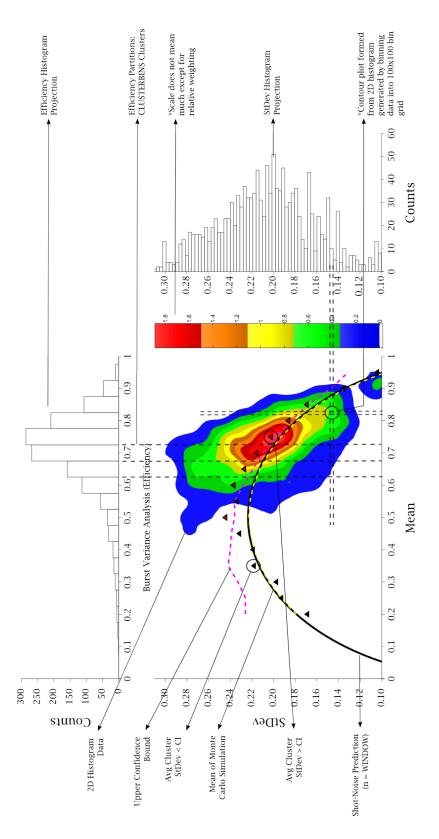
Visualization

BVA visualization is shown in Figure 1. The chosen format for graphical representation is an exploded plot having three components: (i) smoothed 2D histogram contour plot, (ii) projection of abscissa data (clustered FRET) in 1D histogram format, and (iii) projection of ordinate data (StDev FRET) in 1D histogram format.

BVA Contour Plot

BVA data (burst FRET mean versus standard deviation) is binned into a 100x100 bin grid, where the 2D bin size is determined by the minimum and maximum values along the respective dimensions. Obviously, this

Figure 1 (following page): Burst Variance Analysis (BVA) visualization. By plotting mean FRET versus standard deviation (StDev), a smoothed 2D histogram gives rise to the contour plot displayed. Red contours indicate areas of highest data density, whereas white represents zero density. The black line is the shot-noise prediction based on WINDOW. The dotted yellow line corresponds to the Monte Carlo shot-noise simulation based on experimental statistics. The Monte Carlo simulation and shot-noise prediction should agree. The pink line is the upper confidence bound for a static species. Triangles represent the mean values of the bursts grouped by partitioning the mean FRET axis. Black colored triangles are values falling below the upper confidence bound; whereas red colored triangles are values exceeding the upper bound and are indicative of dynamics. The colorbar displays the relative weighting of the colors used in the contour plot. The histograms are 1D projections along their respective axes. Further discussion is provided in the main text.



relative bin sizing should be remembered, particularly in comparing different data sets. Because the bin sizes are not fixed between samples, one cannot compare data sets absolutely—meaning data set 1 has more events at (x_1,y_1) than data set 2 has at the same location. What can be inferred is relative weighting within the respective data sets—meaning the relative weighting within data set 1 is similar in shape/distribution/et cetera to that in data set 2. The relative bin size approach is used for two reasons: (a) to allow more direct visual comparison between the relative distributions among data sets and (b) to accommodate the fact that the data distribution is not necessarily known beforehand. Whereas some prefer to think in bin sizes, others prefer to consider bin numbers. Here, we take a bin numbers approach, although the user is welcome to fix bin locations and size internally using the histon m function called in plotresults.m.*

Once the BVA is binned, we smooth the histogram using a spline smoothing function [5]. We have found a smoothing parameter of 20 yielded best visual results, balancing the trade-off between histogram noise and losing distinct histogram characteristics. This choice of default is dependent on data amount and quality. For those uncomfortable with a heuristic parameter, consult the smoothn.m documentation, which permits automatic parameter determination via generalized cross-validation. We do note, however, that consistency in smoothing parameter choice is more important than rigorous statistical validation, for, as long as the same smoothing parameter (chosen within reason) is applied across all samples (no analysis bias), any qualitative conclusions will remain valid.

The smoothed data yields the contour plot shown in the figure, where red represents high data density and white zero density. The colorbar values should not be considered as indicating absolute numbers of events, but be viewed in terms of relative data density.

Overlaid atop the contour plot are five data results from BVA:

1. a black parabolic curve representing the shot-noise prediction for static species, which is given by:

$$\sigma^{E^*} = \sqrt{\frac{E^*(1 - E^*)}{n}} \tag{1}$$

^{*}For source code, see Bruno Luong, "N-dimensional histogram", http://www.mathworks.com/matlabcentral/fileexchange/23897.

where n=WINDOW. Hence, in considering FRET alone, the shotnoise prediction provides a basis for visualizing the expected relation between μ^{E^*} and σ^{E^*} .

- a yellow curve showing the mean of the Monte Carlo simulated standard deviation distribution. This curve serves as a sanity check that the simulation produced reliable results in accordance with theory and should provide reassurance in the event that the upper confidence bound yields results which do not conform to the user's expectations.
- 3. a pink data series displaying the upper confidence bound generated from a multiple hypothesis test [3].
- 4. filled black triangles represent μ^{E^*} and σ^{E^*} for the group (cluster) of bursts residing between two partitions and which have values consistent with static species. The edges of the cluster bin, assuming CLUSTERBINS = 20, are located ± 0.025 of μ^{E^*} .
- 5. filled red triangles represent μ^{E^*} and σ^{E^*} for the group (cluster) of bursts residing between two partitions and which have values inconsistent with static species and are indicative of dynamics. The edges of the cluster bin are the same as for the black triangles.

A few notes regarding data characteristics:

• one feature of the contour plot is the increasing spread observed in 2D distribution as μ^{E^*} trends toward 0.5 (emphasis on breath of blue dispersion). While heterogeneity is expected among molecules, as well as increased standard deviation as we move toward mid-FRET values, we would not expect increased heterogeneity (standard deviation) in the distribution of standard deviations. We would expect the standard deviation of standard deviations to remain constant, thus leading to a more or less constant spread as we trend toward 0.5.

We address this on two fronts: (i) recall that each individual burst is plotted separately, and, for the data presented, we did not explicitly apply a filter or threshold on burst length or total photons. Accordingly, the blue area is a region of relatively sparse events, many of which are bursts of low photon numbers (i.e., few windows over which to compute standard deviation). We can thus explain the

larger size of the blue area trending toward lower standard deviation by noting that, depending on the timescale and characteristics of dynamics, we expect that two neighboring non-overlapping windows have a greater probability of sharing a similar μ^{E^*} value than two distant windows within the same burst. Hence, for bursts having only enough photons for, say, two or three windows (defined by WINDOW), we expect these bursts to exhibit lower standard deviations than 'normal'. In the limit of infinite windows per burst, the contour distribution would exhibit zero standard deviation, but be a trending line. In which case, for this kind of analysis, the distribution spread says something about the timescale of dynamics. The information contained in these bursts regarding interconversion rates provides insight similar to auto-correlation analysis. (ii) Secondly, if we remove the blue contour in its entirety, we notice that the width of the contour distribution remains relatively constant, which is in accordance with our expectation.

• another feature is that the triangles do not consistently follow the weight of the contour distribution. With reference to the above, we should not necessarily expect these values to align with the distribution density, but rather the mean of the distribution spread within the partition. Hence, if we consider the two partitions centered at $\mu^{E^*} = \{0.65, 0.70\}$, the standard deviation values are the means of the distributions within those partitions. These means do not necessarily equal the median values. Consequently, in the absence of the contour plot and assuming the interconversion of multiple states, the shape of the data series represented by triangles tells the user something about the timescale of dynamics.

Efficiency Histogram

Above the 2D histogram contour plot is a projection of the x-axis (cluster) data binned as a 1D histogram. We chose to bin the cluster data as opposed to the individual burst data to provide visual representation of the number of bursts summarized by the triangles in Figure 1. This information allows the user to better infer the data input for Monte Carlo shotnoise prediction and the associated upper confidence bound. Intuitively, the greater the support for a given cluster the more confident a user may be in determining the presence or absence of dynamics.

Standard Deviation Histogram

To the right of the 2D histogram contour plot is a projection of the y-axis data binned as a 1D histogram. In contrast to the x-axis projection, we chose to bin the individual burst data as opposed to the cluster data to provide visual representation of the underlying data structure supporting the contour plot in Figure 1. The 1D histogram serves as a reminder of the 100x100 bin grid and of the relative data density gradient represented by the colorbar. The cross-sections (areas under the curve) represented by the individual histogram bars along with knowledge of the shape of the distribution displayed in the x-axis histogram allow the user to infer approximately how many events support a given region of the contour plot.

Concluding Remarks

BVA can be an effective method to detect dynamics in single-molecule data. Used in conjunction with other analytical methods, such as probability distribution analysis (PDA) and correlation analysis, BVA provides a model-free method to hypothesis test for dynamics, helping eliminate those models, such as multiple static species, which yield similar distributions to that observed but are incompatible with fluctuations identified within individual bursts. We do note, however, that 'dynamics' is a broad term, including not only conformational fluctuations, but also photophysical effects resulting from quantum yield changes, dye sticking/stacking (restricted orientation), and other events which obfuscate biological behavior. Hence, from BVA alone, one cannot confidently attribute fluctuations solely to conformational dynamics. Appropriate experimental controls and optimization are needed to mitigate artifacts and conclusively explain distribution broadening. As such, BVA builds upon previous work investigating shot-noise limited distributions and provides a quick test for static versus dynamic heterogeneity as the origin of excess distribution width.

We hope this method and associated software prove beneficial in aiding single-molecule research, and we welcome any feedback (comments, criticisms, and questions) and input user's may have. Thank you for choosing to use BVA, and we look forward to your success in probing single-molecule behavior. All the best...

Acknowledgments

The author acknowledges the contributions of the following individuals who assisted in developing this software package, either in providing initial code or insights at various stages of development: Joseph Torella (Harvard University, Boston, USA), Johannes Hohlbein (University of Oxford, Oxford, UK), Yusdi Santoso (Boston Consulting Group, Indonesia), and many others who have tested and re-tested the code, providing feedback along the way. The author was supported by the Clarendon Fund (University of Oxford, Oxford, UK). The published work was supported by a European Commission Seventh Framework Programme (FP7/20072013) grant (HEALTH-F4-2008-201418, entitled READNA) and a Biotechnology and Biological Sciences Research Council grant (BB/H01795X/1) to Achillefs Kapanidis (University of Oxford, Oxford, UK).

REFERENCES REFERENCES

References

[1] Torella JP, Holden SJ, Santoso Y, Hohlbein J, Kapanidis AN (2011), "Identifying molecular dynamics in single-molecule FRET experiments with burst variance analysis", *Biophys J* 100(6):1568-77.

- [2] Kapanidis AN, Lee NK, Laurence TA, Doose S, Margeat E, Weiss S (2004) "Flourescence-aided molecule sorting: analysis of structure and interactions by alternating-laser excitation of single molecules", *PNAS* 101(24):8936-8941.
- [3] Abidi, H (2007), "Bonferroni and Sidak corrections for multiple comparisons", *Encyclopedia of Measurement and Statistics* Sage, Thousand Oaks (CA) 103-107.
- [4] Nir E, Michalet X, Hamadani KM, Laurence TA, Neuhauser D, Kovchegov Y, Weiss S (2006), "Shot-noise limited single-molecule FRET histograms: comparison between theory and experiments", *J Phys Chem B* 110(44):22103-22124.
- [5] Garcia, D (2009), "Robust smoothing of gridded data in one and higher dimensions with missing values", Comput Stat Data An 54:1167-1178. MATLAB code available at http://www.mathworks.com/matlabcentral/fileexchange/25634

List of Figures	List	of	Fig	gu	res
------------------------	------	----	-----	----	-----

1 Figure: Burst Variance Analysis 6

Code

BVA Script

```
2 %% Burst Variance Analysis:
  % Script:
      This MATLAB script allows the user to perform Burst Variance Analysis
      (BVA) and output this information to the user workspace and a figure.
      Details concerning the BVA method and its application may be found in
      the following publication:
  응
  응
          Torella, JP, Holden, SJ, Santoso, Y, Hohlbein, J, Kapanidis, AN
          (2011). Identifying molecular dyanmics in single-molecule FRET
11
          experiments with burst variance analysis. Biophys J.
12
          100(6):1568-77.
  응
13
14
  응
      To run the script, set the Current Directory to the folder containing
15
      the BVA package and run the script BURSTVARIANCEANALYSIS.M either from
  응
16
      the editor or from the command-line using the following command:
17
  응
  응
      ' BurstVarianceAnalysis '.
18
19
  응
  9
      The test data is real experimental data. The input data to be analyzed
20
      must meet two requirements:
  응
21
          [1] Individual photons are sorted by their arrival times and are
23
  응
          sorted according to a ALEx (see Kapanidis) color identification
24
  읒
          scheme: (i) upon donor excitation (Dex): donor emission (Dem) -->
25
          0, acceptor emission (Aem) --> 2; (ii) upon acceptor excitation
26
          (Aex): donor emission (Dem) --> 3, acceptor emission (Aem) --> 1.
27
          Thus, [DexDem, DexAem, AexAem, AexDem] --> [0, 2, 1, 3].
28
30
          [2] A BURST SEARCH must have already been performed. Consult early
  응
          work out of the Siedel laboratory for this methodology. The two
31
          most important outputs of the burst search are the start and end
 9
32
          times of the bursts. Such information can then be used to label
  응
          each photon as either belonging to a burst or not, and, if
34
  응
          belonging, to which burst it belongs.
35
  응
  응
      Accordingly, the input data should be of the following format:
38
          DATA = [ Arrival Times
                                  Photon ID (0, 2, 1, 3)
  응
                                                         Burst ID]
39
40
  응
  응
      As an example:
          DATA = [ 0.40, 0, NaN; ...
42
                   0.51, 1, NaN; ...
  응
43
```

```
0.52, 1,
44
                                    1; . . .
  2
                       0.53, 0,
45
                                    1; . . .
  응
                       0.54, 2,
                                    1; . . .
                       0.55, 0,
                                    1; . . .
                       0.62, 3, NaN; ...
48
49
50
51
                      51.70, 2, NaN; ...
52
                      51.74, 0, 100;...
  응
53
                      51.75, 2, 100; ...
  응
                      51.76, 2, 100; ...
55
                      51.77, 1, 100;...
  9
56
  용
                      51.78, 1, 100;...
57
                      52.81, 2, NaN];
  응
59
```

In the 'Testing' section, an example demonstrates how an Arrival Matrix containing only arrival times and photon IDs can be transformed to accommodate burst search input, as well as highlights some considerations regarding the data (e.g., long bursts being split into two, resulting in two adjacent bursts sharing the same photon, and the need to increasing end times by a fractional amount to ensure appropriate binning [NOTE: this, of course, can be avoided using a binning method other than HISTC.M]).

Following the 'Testing' section, find the 'Initial Parameters' section, which provides a starting point for input parameters. The BVA parameters are typical values, as detailed in Torella, et al (2011); whereas, the filter parameters will vary depending on the experimental setup and conditions. For the example data included, the filter values are found appropriate.

And finally, in the 'BVA' section, the figure generation, analysis, and plotting is broken down into 4 steps (functions), with example usage included.

For additional questions and concerns regarding the individual functions themselves, please consult each function's HELP. Importantly, more detail concerning the nature and structure of the input and output parameters is provided. Further information concerning BVA interpretation may be found in the documentation provided: HELP.pdf .

Lastly, if any bugs or breaks are found in the code, please contact the author of this package detailing the problem and provide any additional evidence of the error/problem, which includes, but is not limited to, 88 error messages, figures, improper output data structure, et cetera. The author's contact information may be found below. 90 응

92 응 NOTES:

60

63

64

65 응

67

71

73

75

79 응

80

81

82 83

84 85 응

86

87

91

9

응 68

응 69 응

응 72 응

으 76 응

```
[1] Need STATS toolbox for generating random numbers drawn from a
93
           binomial distribution.
94
   응
95
   응
       Publications:
97
           [1] Torella, JP, Holden, SJ, Santoso, Y, Hohlbein, J, Kapanidis, AN
98
           (2011). Identifying molecular dyanmics in single-molecule FRET
99
100
           experiments with burst variance analysis. Biophys J. 100(6):1568-77.
   응
101
       NOTICE:
   응
102
           If this software is used in the analysis of data which is to be
103
  응
           presented in publication, please cite the above work and
104
   응
           acknowledge the authors of this software.
105
106
           For purposes of redistribution, use, and/or promotion of products
           derived, consult the included license agreement.
108
109
          Additionally, further license agreements are contained in
110
  용
111
  응
          specialized subfunctions included in the directory 'Other'. Use,
           redistribution, and derivative works are subject to any additional
           requirements imposed therein.
113
114
       Copyright (c) 2011. Kristofer Gryte. University of Oxford.
115
116
   응
117
  응
118
       Version: 1.0 (2011-08-04)
  응
120
   응
121
       Author Information:
122
   응
          Kristofer Gryte
  응
           Clarendon Laboratory
124
          University of Oxford
125
  9
126
  응
           Oxford, United Kingdom
           e-mail: k.gryte1 (at) physics.ox.ac.uk
127
128
129
130
   응
       TODO:
131
132
  9
133
134
  응
137 %% EOP
138
139
141 %% Testing:
```

```
142
   addpath(...
143
        'Other/lightspeed',...
144
        'Other/SplitVec');
145
146
   clear ArrivalMatrix StartTimes EndTimes
147
148
149
   load('MyArrivalMatrix.mat');
150
  % Check!!!! Before binning photons, need to ensure that no StartTimes
151
  % and EndTimes are equal (i.e., no burst ends when another begins), as
   % this creates problems when creating the edges (eliminates a non-burst
   % bin).
154
   AdjacentBursts = intersect_sorted(...
155
       StartTimes, ...
156
       EndTimes);
157
158
   if isempty(AdjacentBursts) == false
159
        % Adjacent bursts have been found!!!
160
        fprintf('%d bursts were found to be immediately adjacent.\n',...
161
            numel(AdjacentBursts));
162
163
        % Cycle through each Adjacent Burst:
164
        for ADJBURST = 1 : numel(AdjacentBursts)
165
166
            % Find the burst:
167
168
            AdjBurstIndex = find(...
                EndTimes == AdjacentBursts(ADJBURST),...
169
                1, ...
170
                'first');
171
            % Increase the arrival time of any photons which
173
            % arrived at the previous start time by a fractional amount:
174
175
            ArrivalMatrix(ArrivalMatrix(:,1) == EndTimes(AdjBurstIndex, 1), 1) ...
                ArrivalMatrix(ArrivalMatrix(:,1) == EndTimes(AdjBurstIndex, ...
176
                    1), 1)...
177
                1e-10; % should be less than NI card time resolution...
178
179
            % Increase the start time of the next burst by a fractional amount:
180
            EndTimes(AdjBurstIndex, 1) = ...
181
                EndTimes(AdjBurstIndex, 1)...
182
                + . . .
183
                1e-10;
184
185
       end % end FOR
186
187
188
```

```
end % end IF
190
191
   % Configure the ArrivalMatrix:
   Edges = union_sorted_rows(...
193
       StartTimes, ...
194
       EndTimes + 1e-10); % Bump the end times by a fractional amount so that ...
195
          last photon can be placed in appropriate burst bin.
196
  % Histogram the arrival times:
197
  [Counts, BinIndices] = histc(ArrivalMatrix(:,1), Edges);
199
   % Determine which bins to throw away: (e.g., all those bins containing
200
   % photons which do not belong to a burst)
201
   % Discard the even BinIndices, as these correspond to bins defined by
203
   % EdgeA = EndTime(1); EdgeB = StartTime(2). Hence, this bin (EdgeA,
204
  % EdgeB) corresponds to photons which do not belong to a burst (i.e., photons
  % occurring after the end of the previous burst and before the start of
  % the next burst):
  VALS1 = 0:2:max(BinIndices); % Note: we include 0 to account for photons ...
      not falling into any defined bin.
   TF = ismember_sorted(BinIndices, VALS1);
209
210
211
212 % Remove all photons not belonging to a burst, thus creating a 'burst
  % arrival matrix':
  BinIndices(TF == true) = NaN; %[];
214
215
  % Tack on the BinIndices (i.e., Burst Indices) to the Arrival Matrix:
216
  ArrivalMatrix(:,3) = BinIndices;
217
218
  clear BinIndices TF VALS1 Counts Edges
219
220
221
  222
223 %% Initial Parameters:
224
225 \text{ WINDOW} = 5;
  CLUSTERBINS = 20;
226
227 CLUSTERMINWINDOWS = 50;
  CONFIDENCEINTERVAL = 0.999;
228
229
   % ----- %
230
   % Filters:
231
   232
233
234 % DexDem:
235 FILTERS.DexDem.Min = [];
```

```
236 FILTERS.DexDem.Max = [];
237
  % DexAem:
238
  FILTERS.DexAem.Min = [];
240 FILTERS.DexAem.Max = [];
241
242 % AexAem:
243 FILTERS.AexAem.Min = [30];
244 FILTERS.AexAem.Max = [];
245
246 % Sum Photons: DexDem + DexAem
247 FILTERS.SumPhotons.Min = [];
248 FILTERS.SumPhotons.Max = [];
249
  % Total Photons: DexDem + DexAem + AexAem
  FILTERS.TotalPhotons.Min = [];
252 FILTERS. Total Photons. Max = [];
253
254 % Efficiency:
255 FILTERS.Efficiency.Min = [];
256 FILTERS.Efficiency.Max = [];
257
  % Stoichiometry:
259 FILTERS.Stoichiometry.Min = [0.45];
260 FILTERS.Stoichiometry.Max = [];
261
262 % Duration: [seconds]
263 FILTERS.Duration.Min = [];
264 FILTERS.Duration.Max = [];
265
   % Length: [photons]
266
  FILTERS.Length.Min = [];
267
  FILTERS.Length.Max = [];
268
269
270
271
  272
273
   응응 BVA:
274
275
   % [1] Create a figure:
  handles.figure = figure;
276
277
  set (handles.figure, ...
278
       'Position', [520, 196, 900, 700]); % [x-pos, y-pos, width, height] % ...
279
          NOTE: the specified values are appropriate for the INITIALIZEAXES \dots
          function to follow. Should the figure window be too large or ...
          small, subsequent re-adjustments in the INITIALIZEAXES function ...
          may be needed.
280
```

```
281 % [2] Initialize the figure and axes:
282 handles = InitializeAxes(handles);
283
  % [3] Perform BVA:
  [DATA] = Confocal2ColorALExSolnMultDataBVA_Analysis(...
285
      ArrivalMatrix,...
286
      'window', WINDOW,...
287
      'clusterbins', CLUSTERBINS,...
288
289
      'clusterminwindows', CLUSTERMINWINDOWS,...
      'confidenceinterval', CONFIDENCEINTERVAL,...
290
      'filters', FILTERS);
291
292
293 % [4] Plot the results:
294 handles = PlotResults(DATA, WINDOW, handles);
296
298 %% EOS
```

CODE Initialize Axes

Initialize Axes

```
1 function handles = InitializeAxes(handles)
2 % INITIALIZEAXES
      Initialize axes within a figure window according to defined
      specifications. Three subplots are generated: [i] 2D histogram subplot
      and [ii] 2 1D histogram suplots along the upper and right edges of the
      2D histogram subplot. Orientations of the 1D histogram subplots are
      such that they are the projections of the data along the abscissa and
      ordinate axes, respectively.
      INPUTS:
10
  응
         HANDLES: structure with the following fields:
12 %
             FIGURE: handle to the figure window. Suggested figure size:
              [520, 196, 900, 700].
13 %
14 %
15 %
16 %
      OUTPUTS:
17 %
          HANDLES: structure with the following fields:
               SUBPLOT: nested structure with the following fields:
18
                   BVA: nested structure with the following fields:
19
                       AXES: handles to the 2D Histogram axes (subplot).
20 응
               SUBPLOT: nested structure with the following fields:
21 응
                  HISTOGRAM: nested structure with the following fields:
                       MEAN: nested structure with the following fields:
24 %
                          AXES: handle to the axes into which the histogram
25 %
                             will be plotted.
                       STDEV: nested structure with the following fields:
27
                          AXES: handle to the axes into which the histogram
28 응
                             will be plotted.
29 %
30 %
31 %
32 %
      DEPENDENCIES:
33 %
          (none)
34
  응
35
      EXAMPLE USAGE:
  응
  응
        handles = InitializeAxes(handles);
  응
39
      History:
40
          [1] Kristofer Gryte. University of Oxford (2011).
41
      Copyright (c) 2011. Kristofer Gryte. University of Oxford.
43 %
44 %
45 %
```

CODE Initialize Axes

```
46
     Version: 1.0 (2011-08-02)
 응
47
48
 응
     Author Information:
50
51 %
       Kristofer Gryte
        Clarendon Laboratory
52
        University of Oxford
       Oxford, United Kingdom
54
       e-mail: k.gryte1 (at) physics.ox.ac.uk
 응
55
56 %
57 %
     TODO:
58 %
59
61
63 %% EOP
67 %% Initialization:
69 % Activate figure window:
70 figure(handles.figure);
71
72 % Configure the figure colormap:
73 load('BVAcolormap.mat', 'BVAcolormap');
75 set(handles.figure,...
     'Colormap', BVAcolormap);
76
77
80 %% Initialize Subplots:
81
82 % Mean Histogram Subplot:
83 handles.Subplot.Histogram.Mean.Axes = subplot(3,1,1);
85 % Standard Deviation Histogram Subplot:
86 handles.Subplot.Histogram.StDev.Axes = subplot(3,1,2);
87
88 % BVA Subplot:
89 handles.Subplot.BVA.Axes = subplot(3,1,3);
91
94 %% Manipulate Subplots:
```

CODE Initialize Axes

```
95
% Get the current units:
97 OldUnits = get(handles.Subplot.Histogram.Mean.Axes, 'Units');
  % Set Units: % Note: Change to characters...as the number of pixels is
100 % relative to user monitor...thus, standardize...
  set([handles.Subplot.Histogram.Mean.Axes,...
      handles.Subplot.Histogram.StDev.Axes,...
103
      handles.Subplot.BVA.Axes], ...
       'Units', 'pixels');
104
105
106 % Set Mean Histogram Position:
107 set(handles.Subplot.Histogram.Mean.Axes,...
       'Position', [150 500 350 150]);
108
  % Set StDev Histogram Position:
  set (handles.Subplot.Histogram.StDev.Axes, ...
111
       'Position', [600 95 150 350]);
112
113
114 % Set BVA Subplot Position:
set (handles.Subplot.BVA.Axes,...
       'Position', [150 95 350 350]);
116
118 % Return units to 'characters':
119 set([handles.Subplot.Histogram.Mean.Axes,...
      handles.Subplot.Histogram.StDev.Axes,...
120
      handles.Subplot.BVA.Axes],...
       'Units', OldUnits); % Maybe Normalize?
122
123
124
126 %% EOF
```

BVA Analysis

```
1 function [DATA] = Confocal2ColorALExSolnMultDataBVA_Analysis(...
      varargin)
  % CONFOCAL2COLORALEXSOLNMULTDATABVA ANALYSIS
      Burst Variance Analysis (BVA) of confocal (2 color) solution
      alternating laser excitation (ALEx) data.
5
  응
  응
      INPUTS:
          ARRIVALMATRIX: input data on which to perform Burst Variance
          Analysis. The input matrix should be Nx3 or 3xN, with the following
          as vector inputs: (1) Photon Arrival Times; (2) Detection Channel
10
11
           (DexDem = 0; DexAem = 2; AexAem = 1; AexDem = 3); (3) Burst
          Classification (those photons belonging to a burst should have a
12
          corresponding index to a unique burst ID; background photons should
  9
13
 응
          be indexed as NaN): ARRIVALMATRIX = ArrivalTimes ID
15
  응
          WINDOW: size of the sliding window over which FRET will be
16
          calculated. Default: 5.
17
          CLUSTERBINS: number of bins over which to cluster bursts
19
          characterized by their mean FRET value. Mean standard deviation
20
          will be calculated across bursts belonging to a particular bin and
21
 용
 용
          compared against a test statistic to determine significance (and
22
23
          here, the presence of dynamics). Default: 20.
24
          CLUSTERMINWINDOWS: threshold for the minimum number of total windows
25
26
          within a cluster from all bursts of the cluster to be analyzed
           (i.e., we need to make sure we have enough data to be worth our
27
  응
          while). Default: 50.
28
29
  응
  응
          CONFIDENCEINTERVAL: (1-alpha) to determine statistical signifance
          of cluster results. Default: 0.999.
31 %
32 %
          FILTERS: nested structure providing minimum and maximum thresholds
33
          for the following data streams:
34
              DURATION: the temporal length of a detected burst [seconds]
35
36
  9
              LENGTH: the number of photons within a detected burst [photons]
  응
              DEXDEM: the number of photons detected in the donor channel
                   upon donor excitation.
              DEXAEM: the number of photons detected in the acceptor channel
39
                   upon donor excitation.
40
              AEXAEM: the number of photons detected in the acceptor channel
41
                  upon acceptor excitation.
43
              SUMPHOTONS: the number of photons detected in the donor &
                   acceptor channels upon donor excitation.
44 %
              TOTALPHOTONS: the number of photons detected in the donor &
```

```
acceptor channels upon donor and acceptor excitation.
46
                   (excluding photons detected in donor channel upon acceptor
47
                   excitation: AexDem)
               EFFICIENCY: the FRET efficiency.
               STOICHIOMETRY: the relative fluorophore brightness.
50
51
      OUTPUTS:
52
  응
          DATA: structure with the following data streams for each detected
53
          burst: (NOTE: data streams are filtered according to input FILTERS)
54
               DURATION: the temporal length of a detected burst [seconds]
  응
55
               LENGTH: the number of photons within a detected burst [photons]
  응
               DEXDEM: the number of photons detected in the donor channel
57
  응
                   upon donor excitation.
58
               DEXAEM: the number of photons detected in the acceptor channel
59
                   upon donor excitation.
               AEXAEM: the number of photons detected in the acceptor channel
61
                   upon acceptor excitation.
62
  9
               SUMPHOTONS: the number of photons detected in the donor &
63
  응
                   acceptor channels upon donor excitation.
               TOTALPHOTONS: the number of photons detected in the donor &
65
                   acceptor channels upon donor and acceptor excitation.
66
                   (excluding photons detected in donor channel upon acceptor
67
                   excitation: AexDem)
               EFFICIENCY: the FRET efficiency.
69
               STOICHIOMETRY: the relative fluorophore brightness.
  응
70
               RESULTS: an Nx2 array, where N is the number of bursts (after
71
  응
                   filtering). The first column corresponds to the mean ...
      Efficiency
  응
                   value for a burst and the second to the burst's Efficiency
73
  응
                   standard deviation. Array: E(efficiency) sqrt(V(efficiency))
74
               CLUSTERS: nested structure with the following fields:
75
                   CENTERS: vector containing the centers of the individual
76
  응
                      clusters.
77
  응
                   STDEVDISTRMEAN: the Monte Carlo simulated standard
                      deviation prediction.
                   STATISTICS: an Mx2 array, where M is the number of clusters
80
                       ('slices' along the efficiency axis, into which bursts are
81
                       grouped). The first column corresponds to the mean
82
                       Efficiency value for the cluster, and the second to the
                       standard deviation of the burst's within that cluster.
84
                   WINDOWMEANS: cellular array where each entry contains the
  응
85
                       the Efficiency values for all sliding windows for bursts
  응
                       belonging to a cluster.
87
                   DISTANCEFROMUPPERCI: a vector of the distance between the
88
                       mean standard deviations of each cluster from the upper
  9
89
                       confidence bound. This expresses something about the \dots
  응
      extent
                       of the 'dynamics'.
  2
91
92 응
               CONFIDENCEINTERVAL: nested structure with the following fields:
```

```
UPPER: the upper confidence bound generated from the Monte
93
   응
                        Carlo simulation and multiple hypothesis test.
94
   응
                    LOWER: the lower confidence bound generated from the Monte
                       Carlo simulation and the multiple hypothesis test.
97
       DEPENDENCIES:
   응
98
           SPLITVEC.M
99
100
            INTERSECT_SORTED.M (Lightspeed)
           UNION_SORTED_ROWS.M (Lightspeed)
101
           ISMEMBER_SORTED.M (Lightspeed)
   응
102
103
  응
           SETDIFF_SORTED.M (Lightspeed)
           FILTERDATA.M (private)
104
   응
           MONTECARLOPREDICTION.M (private; requires STATS toolbox)
105
106
       EXAMPLE USAGE:
108
            Confocal2ColorALExSolnMultDataBVA_Analysis (ArrivalMatrix);
109
110
111
  응
            Confocal2ColorALExSolnMultDataBVA_Analysis(ArrivalMatrix, 'window',
           5);
113
           Confocal2ColorALExSolnMultDataBVA_Analysis(ArrivalMatrix, 'window',
114
            5, 'clusterbins', 20, 'clusterminwindows', 50);
115
116
   응
117
   응
       History:
118
   응
            [1] Joseph Torella, Yusdi Santoso. University of Oxford (2009).
            [2] Johannes Hohlbein. University of Oxford (2010).
120
            [3] Kristofer Gryte. University of Oxford (2011).
   응
121
122
       Publications:
            [1] Torella, JP, Holden, SJ, Santoso, Y, Hohlbein, J, Kapanidis, AN
124
            (2011). Identifying molecular dyanmics in single-molecule FRET
125
126
   응
            experiments with burst variance analysis. Biophys J. 100(6):1568-77.
127
       NOTICE:
128
            If this software is used in the analysis of data which is to be
129
130
            presented in publication, please cite the above work and
            acknowledge the authors of this software.
131
132
       Copyright (c) 2011. Kristofer Gryte. University of Oxford.
   9
133
134
   응
135
   9
   응
136
       Version: 1.0 (2011-07-29)
   응
137
138
139
       Author Information:
   응
140
          Kristofer Gryte
141 %
```

```
Clarendon Laboratory
142 %
          University of Oxford
143
          Oxford, United Kingdom
          e-mail: k.gryte1 (at) physics.ox.ac.uk
146
147
  9
  응
      TODO:
148
          [1] Provide appropriate checks on filter values (e.g., ensure
          numeric, positive, etc.)
150
          [2] Make number of Monte Carlo simulations an input variable?
  응
151
152
  응
  응
153
  응
      See also
154
155
156
157
  158
159 %% EOP
160
161
162
  164
  %% StartUp/CleanUp:
165
166 StartUp();
167
168 C = onCleanup(@() CleanUp());
169
170
  171
172 %% Initialization:
173
  응 ---- 응
174
175 % Establish default values:
  응 ---- 응
  WINDOW = 5; % size of the sliding window (non-overlapping) over which FRET ...
      will be calculated; hence, for every 5 photons, we calculate FRET
  CLUSTERBINS = 20; % number of bins over which we will cluster bursts ...
     characterized by their mean FRET value
  CLUSTERMINWINDOWS = 50; % threshold for the minimum number of total ...
     windows within a cluster over all bursts for the cluster to be ...
      analyzed (i.e., we need to make sure we have enough data to be worth ...
      our while)
180
181 CONFIDENCEINTERVAL = 0.999; % (1-alpha) CI level to determine if observed ...
      variance is statistically significant
  NUMMONTECARLOSAMPLES = 500; % Number of Monte Carlo simulations to run for ...
     CI bound generation
183
```

```
FILTERS = struct(...
                         struct('Min', [], 'Max', []),...
185
        'DexDem',
        'DexAem',
                         struct('Min', [], 'Max', []),...
186
        'AexAem',
                         struct('Min', [], 'Max', []),...
187
                         struct('Min', [], 'Max', []),...
        'AexDem',
188
                         struct('Min', [], 'Max', []),...
        'Efficiency',
189
        'Stoichiometry', struct('Min', [], 'Max', []),...
190
                        struct('Min', [], 'Max', []),...
        'SumPhotons',
191
        'TotalPhotons', struct('Min', [], 'Max', []),...
192
        'Duration',
                         struct('Min', [], 'Max', []),...
193
                         struct('Min', [], 'Max', []));
194
        'Length',
195
   DATA = [];
196
197
   응 ---- 응
198
199
   % Parse input arguments:
200
201
202
   ArrivalMatrix = varargin{1};
203
   % Check!!!
204
   if ¬isnumeric(ArrivalMatrix)
205
206
       % error out!
       errordlg(['ERROR:invalid input argument for BVA function. Please input ...
207
            'a numeric array for the initial argument.'],...
208
            'ERROR:invalid input argument');
209
       return;
210
   end % end IF
211
212
   % Perform check for data input size and ensure dimensionality is correct
213
   % (column matrix)
214
   if size(ArrivalMatrix,1) < size(ArrivalMatrix, 2) % More columns than rows
215
       % Take the tranpose:
216
       ArrivalMatrix = ArrivalMatrix';
217
218
   elseif size (ArrivalMatrix, 2) \neq 3 % too many or too few columns
219
220
       % Error out!
221
       errordlg(['ERROR:invalid input argument for BVA function. Data input ',...
222
            'should be either (Nx3) or (3xN).'],...
223
            'ERROR: invalid input argument');
224
225
       return;
226
   end % end IF/ELSEIF
227
   % Check number of input arguments:
229
  if (mod(nargin, 2) - 1) \neq 0 \% odd number of input arguments
230
       % Throw a fit and error out!
231
```

```
232
        errordlg(['ERROR:invalid number of input arguments for BVA function. ',...
233
            'Please enter parameter-value pairs in addition to the data ...
234
                input.'],...
            'ERROR: invalid number of input arguments');
235
        return;
236
237
238
   end % end IF
239
   % Loop through and assign parameter-value pairs:
240
   for NumArg = 2 : 2 : nargin
241
242
        switch lower(char(varargin{NumArg}))
243
244
            case {'win', 'window', 'winsize'}
245
                % Update the WINDOW size:
246
                temp = varargin{NumArg + 1};
247
248
249
                % Check!
                if ¬isnumeric(temp)
                                         temp < 1
250
                     % Error out:
251
                     answer = questdlg(...
252
                          ['NOTICE: An invalid value was provided for the WINDOW ...
253
                             paramter. ',...
                          'Please supply an integer value greater than 1. ',...
254
                          'Would you like to continue with a default WINDOW = 5, ...
255
                             ',...
                          'or return to the command-line and supply an ...
256
                             appropriate input value?'],...
                          'WARNING:invalid input value',...
257
                          'Continue', 'Cancel',...
258
                          'Cancel');
259
260
261
                     % Parse the answer:
                     switch lower(answer)
262
                         case 'continue'
263
                              % Move along with default WINDOW value
264
                         case 'cancel'
265
                              % Abort!
266
267
                              return;
                     end % end SWITCH
268
269
                else
270
                     % Assign over the input WINDOW size:
271
                     WINDOW = temp;
272
                end % end IF/ELSE
273
274
            case {'clusterbins', 'bins', 'numbins', 'binnumber', 'binnum'}
275
276
```

```
% Update the number of cluster bins:
277
                temp = varargin{NumArg + 1};
278
279
                % Check!
                if ¬isnumeric(temp)
                                        temp < 1
281
                     % Error out:
282
                     answer = questdlg(...
283
284
                         ['An invalid value was provided for the CLUSTERBINS ...
                             paramter. ',...
                         'Please supply an integer value greater than 1. ',...
285
                         'Would you like to continue with a default CLUSTERBINS ...
286
                             = 20, ',...
                         'or return to the command-line and supply an ...
287
                             appropriate input value?'],...
                         'WARNING: invalid input value',...
                         'Continue', 'Cancel',...
289
                         'Cancel');
290
291
292
                     % Parse the answer:
                     switch lower(answer)
293
                         case 'continue'
294
                             % Move along with default CLUSTERBINS value
295
                         case 'cancel'
296
297
                             % Abort!
                             return;
298
                     end % end SWITCH
299
300
                else
301
                     % Assign over the input CLUSTERBINS value:
302
                     CLUSTERBINS = temp;
303
                end % end IF/ELSE
304
305
            case {'clusterminwindows', 'minwindows'}
306
307
                % Update the minimum number of windows within a cluster:
308
                temp = varargin{NumArg + 1};
309
310
311
                % Check!
                if ¬isnumeric(temp)
                                        temp \leq 1
312
313
                     % Error out:
                     answer = questdlg(...
314
                         ['An invalid value was provided for the ...
315
                             CLUSTERMINWNDOWS paramter. ',...
                         'Please supply an integer value greater than 1. ',...
316
                         'Would you like to continue with a default ...
317
                             CLUSTERMINWINDOWS = 50, ',...
                         'or return to the command-line and supply an ...
318
                             appropriate input value?'],...
                         'WARNING: invalid input value',...
319
```

```
'Continue', 'Cancel',...
320
                          'Cancel');
321
322
                     % Parse the answer:
323
                     switch lower(answer)
324
                         case 'continue'
325
                             % Move along with default CLUSTERMINWINDOWS value
326
                         case 'cancel'
327
328
                              % Abort!
                              return;
329
                     end % end SWITCH
330
331
332
                else
                     % Assign over the input CLUSTERMINWINDOWS:
333
                     CLUSTERMINWINDOWS = temp;
334
                end % end IF/ELSE
335
336
            case {'ci', 'confidenceinterval', 'conf', 'confidence', ...
337
                'confinterval'}
338
                % Update the confidence interval:
339
                temp = varargin{NumArg + 1};
340
341
342
                % Check!
                if ¬isnumeric(temp)
                                        (temp \ge 1 temp \le 0)
343
                    % Error out:
344
345
                     answer = questdlg(...
                          ['An invalid value was provided for the ...
346
                             CONFIDENCEINTERVAL paramter. ',...
                          'Please supply a numeric value greater than 0 and less ...
347
                             than 1. ',...
                          'Would you like to continue with a default ...
348
                             CONFIDENCEINTERVAL = 0.999, ',...
349
                          'or return to the command-line and supply an ...
                             appropriate input value?'],...
                          'WARNING: invalid input value',...
350
                          'Continue', 'Cancel',...
351
                          'Cancel');
352
353
                     % Parse the answer:
354
                     switch lower(answer)
355
356
                         case 'continue'
                              % Move along with default CONFIDENCEINTERVAL value
357
                         case 'cancel'
358
                              % Abort!
359
360
                              return;
                     end % end SWITCH
361
362
                else
363
```

```
% Assign over the input CONFIDENCEINTERVAL:
364
                     CONFIDENCEINTERVAL = temp;
365
                end % end IF/ELSE
366
367
            case {'filters', 'filter', 'thresh', 'thresholds', 'threshold', ...
368
                'filt'}
369
                % Update the FILTERS:
370
                temp = varargin{NumArg + 1};
371
372
                % Check!
373
                if ¬isstruct(temp)
374
                     % Error out:
375
                     answer = questdlg(...
376
                          ['An invalid value was provided for the FILTERS ...
377
                             paramter. ',...
                          'Please supply an appropriate structure containing ...
378
                             relevant fields. ',...
379
                          'Would you like to continue with a default FILTERS ...
                             (none), ',...
                          'or return to the command-line and supply an ...
380
                             appropriate input?'],...
                          'WARNING: invalid input value',...
381
                          'Continue', 'Cancel',...
382
                          'Cancel');
383
384
                     % Parse the answer:
385
                     switch lower(answer)
386
                         case 'continue'
387
                              % Move along with default FILTERS value
388
                         case 'cancel'
                              % Abort!
390
                              return;
391
                     end % end SWITCH
392
393
394
                     % Assign over the input FILTERS:
395
                     FILTERS = temp;
396
                end % end IF/ELSE
397
398
            otherwise
399
                % Throw a notice that the input argument was not recognized:
400
                msqbox(sprintf(...
401
                     ['NOTICE:Unrecognized input argument: %s. \n',...
402
                          'Parameter-value pair ignored']),...
403
                     char(varargin{NumArg}));
404
405
       end % end SWITCH
406
407
```

```
end % end FOR
409
410
  응 ---- 응
  % Calculate remaining parameters:
  PHOTONCUTOFF = floor(WINDOW/2); % Remove the first few photons of a burst ...
      (allows analyzed data to be centered within burst)
414
  CLUSTERWIDTH = (1-0)/CLUSTERBINS; % How wide are our bins in which we are ...
415
      clustering bursts (UNITS: none --> FRET efficiency); (0,1) defines the ...
      range of our FRET values.
416
  417
418
  419
  %% Filtering:
420
421
  % Filter the data:
422
  [DATA, REMOVEBURSTS] = FilterData(ArrivalMatrix, FILTERS); % this is a ...
      function!!!!
424
425
  427
428 %% Burst Variance Analysis:
429
430 % Start the timer...
431 tic
432
  % Launch the waitbar:
433
  h.Waitbar = waitbar(...
435
       'Commencing BVA...please be patient');
436
437
  % ----- %
438
  %% Winnow the Data:
439
440
  % Replace all the NaNs:
  ArrivalMatrix(isnan(ArrivalMatrix(:,3)),3) = 0;
442
443
  % Return the assigned IDs of the various runs: % recall that non-burst
444
  % photons are labeled with a '0'!!!!
  FLAGVALS = SplitVec(...
      ArrivalMatrix(:,3),...ArrivalMatrix(¬isnan(ArrivalMatrix(:,3)),3),.......
447
          % exclude any NaNs
       'equal',... % find all those runs of the same index (burst ID)
       'firstval'); % return just the first index marking a run.
449
450
451 % Extract some information about the bursts from the ArrivalMatrix:
```

```
STARTINDICES = SplitVec(...
452
       ArrivalMatrix(:,3),... % exclude any NaNs
453
       'equal',... % find all those runs of the same index (burst ID)
454
       'first'); % return just the first index marking a run.
455
456
   ENDINDICES = SplitVec(...
457
       ArrivalMatrix(:,3),... % exclude any NaNs
458
459
        'equal',... % find all those runs of the same index (burst ID)
        'last'); % return just the last index completing a run.
460
461
  STARTINDICES (FLAGVALS == 0) = []; % NOTE:
462
   ENDINDICES(FLAGVALS == 0) = [];
463
464
   % Get the start times and end times for each burst:
465
   STARTTIMES = ArrivalMatrix(STARTINDICES, 1);
466
467
   ENDTIMES = ArrivalMatrix(ENDINDICES, 1);
468
  % Grab the total number of bursts:
469
470
  TOTALBURSTS = numel(STARTINDICES);
  % Create an index array:
472
  BURSTINDICES(:,1) = 1 : TOTALBURSTS; % column vector
473
   % Check!!!! Before binning photons, need to ensure that no StartTimes
475
  % and EndTimes are equal (i.e., no burst ends when another begins), as
  % this creates problems when creating the edges (eliminates a non-burst
477
  % bin).
   AdjacentBursts = intersect_sorted(...
479
       STARTTIMES, ...
480
481
       ENDTIMES);
482
   if isempty(AdjacentBursts) == false
483
       % Adjacent bursts have been found!!!
484
       fprintf('%d bursts were found to be immediately adjacent.\n',...
485
           numel(AdjacentBursts));
486
487
       % Cycle through each Adjacent Burst:
488
       for ADJBURST = 1 : numel(AdjacentBursts)
489
490
           % Find the burst:
491
           AdjBurstIndex = find(...
492
                STARTTIMES == AdjacentBursts(ADJBURST),...
493
494
                1, ...
                'first');
495
496
           % Increase the arrival time of any photons which
           % arrived at the previous start time by a fractional amount:
498
           ArrivalMatrix(ArrivalMatrix(:,1) == STARTTIMES(AdjBurstIndex, 1), ...
499
               1) = ...
```

```
ArrivalMatrix(ArrivalMatrix(:,1) == STARTTIMES(AdjBurstIndex, ...
500
                   1), 1)...
501
                +...
                1e-10; % should be less than NI card time resolution...
502
503
            % Increase the start time of the next burst by a fractional amount:
504
            STARTTIMES (AdjBurstIndex, 1) = ...
505
506
                STARTTIMES (AdjBurstIndex, 1) ...
                +...
507
                1e-10;
508
509
       end % end FOR
510
511
512
   end % end IF
513
514
515
  % Create Edges:
516
517
   Edges = union_sorted_rows(...
       STARTTIMES (:, 1), ...
518
       ENDTIMES(:,1) + 1e-10);
519
520
   % Histogram the arrival times:
   [Counts, BinIndices] = histc(ArrivalMatrix(:,1), Edges);
522
523
  % Determine which bins to throw away: (e.g., all those bins containing
524
525
  % photons which do not belong to a burst)
526
  % Discard the even BinIndices, as these correspond to bins defined by
527
   % EdgeA = EndTime(1); EdgeB = StartTime(2). Hence, this bin (EdgeA,
528
   % EdgeB) corresponds to photons which do not belong to a burst (i.e., photons
  % occurring after the end of the previous burst and before the start of
  % the next burst):
531
  VALS1 = 0:2:max(BinIndices); % Note: we include 0 to account for photons ...
      not falling into any defined bin.
  TF = ismember_sorted(BinIndices, VALS1);
533
534
535
   % Remove all photons not belonging to a burst, thus creating a 'burst
536
   % arrival matrix':
537
  ArrivalMatrix(TF == true,:) = [];
538
  BinIndices(TF == true) = [];
540
541
542
   %% Burst Removal:
544
  % Remove the filtered bursts:
545
546 TF = ismember_sorted((BinIndices + 1) ./ 2, REMOVEBURSTS);
```

```
547
  ArrivalMatrix(TF == true, :) = [];
548
  BinIndices(TF == true, :) = [];
551
  % Now, we need to rid ourselves of all photons originating from red
552
  % excitation: DexDem = 0; DexAem = 2; AexAem = 1; AexDem = 3; => all
  % photons labeled 1 or 3... (NOTE: this also removes acceptor-only
  % bursts!)
555
556 LogicalIndexArray = (ArrivalMatrix(:,2) == 1 ArrivalMatrix(:,2) == 3);
  VALS2 = unique(BinIndices(LogicalIndexArray, :)); % This gives all burst ...
      indices which contain acceptor excitation photons
558
  ArrivalMatrix(LogicalIndexArray,:) = []; % All acceptor excitation photons ...
559
      removed.
  BinIndices(LogicalIndexArray,:) = []; % Note: this array will be important ...
      now, as all photons can be grouped by their unique identification with ...
      a Bin...
  % NOTE: now VALS and unique (BinIndices) do not necessarily
  % correspond!!!
563
  % Determine which, if any, bursts have been removed from further
   % analysis:
566
  REMOVEBURSTS = [REMOVEBURSTS; (setdiff_sorted(VALS2, unique(BinIndices)) + ...
      1) ./ 2];
567
568
  % REMOVEBURSTS now contains all bursts which either did not meet the
   % filtering criteria above or were acceptor-only...
569
570
  % Frequently, bursts have values which are not numbers; find them:
571
  NaNBURSTS = find(isnan(DATA.Efficiency(:,1)));
572
573
  REMOVEBURSTS = [REMOVEBURSTS; NanbursTS];
574
575
576
577
578
579
580
   % Clustering: cluster the bursts according to mean Efficiency values:
581
582
  % Define the cluster bounds: % NOTE: this creates an extra bin!!!
  CLUSTEREDGES = -CLUSTERWIDTH/2:CLUSTERWIDTH:1+CLUSTERWIDTH/2;
584
  CLUSTEREDGES = CLUSTEREDGES';
585
586
   % Each burst is histogrammed into a cluster bin:
   [Counts, CLUSTERBINIDX] = histc(...
588
       DATA.Efficiency(:,1),...
589
       CLUSTEREDGES);
590
```

```
591
592
593
594
595
   % Split the Arrival Matrix into Bursts:
596
   IndexCell = SplitVec(...
597
       BinIndices, ...
       'equal',... % Group all elements which are equal
599
       'loc'); % Return the locations of these elements in a cellular array
600
601
  % Each cellular element of IndexCell corresponds to a burst; within each
602
   % cell element, the individual photons are assigned a location
603
   % corresponding to their index in the (filtered) ArrivalMatrix.
604
606
607
  %% BVA:
608
609
  waitbar(0,...
610
       h.Waitbar,...
611
       'Performing sliding window analysis...');
612
614 % Initialize a BurstVarianceAnalysis array:
615 DATA.Results = nan(TOTALBURSTS, 2);
616
617
  % Loop through the bursts and perform BVA:
  COUNTER = 0;
618
  CLUSTERSTATS = cell(CLUSTERBINS+1, 1);
619
620
   TRACKER = 0;
621
   for BURST = 1 : TOTALBURSTS
622
623
624
       % Check!!!
       if ismember(BURST, REMOVEBURSTS) % Determine if BURST is blacklisted...
625
            % Blacklisted burst found! Move along to next burst...
626
            continue;
627
628
       else
            % Update our counter for our non-blacklisted bursts...
629
            COUNTER = COUNTER + 1;
630
       end % end IF
631
632
       % Grab the current (non-blacklisted) BURST indices:
633
       INDICES = IndexCell{COUNTER};
634
635
       % Remove a 'cutoff' number of photons:
       if numel(INDICES) < PHOTONCUTOFF</pre>
637
            % Go to next iteration...
638
            REMOVEBURSTS (end+1,1) = BURST;
639
```

```
TRACKER = TRACKER + 1;
640
           continue;
641
642
       else
           INDICES(1:PHOTONCUTOFF) = [];
643
       end % end IF/ELSE
644
645
646
647
       % Calculate total number of photons:
       TOTALPHOTONS = numel(INDICES);
648
649
       % Check!!!
650
       if TOTALPHOTONS < WINDOW
651
           % Go to next iteration, as not enough photons in 'burst'...
652
           REMOVEBURSTS (end+1,1) = BURST;
653
           TRACKER = TRACKER + 1;
654
           continue;
655
       end % end IF
656
657
658
659
       % We must apply a window to the burst photons; this window is
660
       % non-overlapping, and thus, in most cases, the number of windows
661
       % within a burst is not an integer, but rather, a remainder of
662
       % photons cannot be used. Let us discard these photons, so we can
663
       % reshape our photon array:
664
       RemPhotons = rem(TOTALPHOTONS, WINDOW);
665
666
       % We have some leftover photons; remove them...
667
       INDICES(end-RemPhotons+1:end) = [];
668
669
670
       % Reshape the photons from the arrival matrix into an array of
671
       % non-overlapping windows: (NOTE: photons within each window are
672
       % placed along the columns, and the window number within the burst
673
       % is placed along the rows)
674
       PhotonArray = transpose(reshape(ArrivalMatrix(INDICES, 2), WINDOW, ...
675
           [])); % No apriori knowledge of how many windows within a burst
676
       % Calculate the Efficiency value (FRET) within each window: (NOTE:
677
       % we exploit the fact that DexDem photons are labeled 0):
678
       EfficiencyVector = sum((PhotonArray == 2), 2) ./ WINDOW; % where ...
679
           WINDOW = sum of DexDem + DexAem and the numerator is the number of ...
           DexAem photons
680
       % Calculate the standard deviation of the efficiency values:
681
       EfficiencyStDev = std(EfficiencyVector);
683
       % Place the mean and stdev of the efficiency in our BVA array:
684
       DATA.Results(BURST, :) = [mean(EfficiencyVector), EfficiencyStDev];
685
```

```
686
687
       % Place the efficiency vector with the burst's respective cluster:
688
       WhichCluster = CLUSTERBINIDX(BURST);
689
690
       % Check!!!
691
       if WhichCluster == (CLUSTERBINS+2)
692
693
            % This accounts for values which are precisely equal to the
            % last edge. We place this data in the last bin.
694
            WhichCluster = WhichCluster - 1;
695
       end % end IF/ELSE
696
697
       CLUSTERSTATS{WhichCluster} = [CLUSTERSTATS{WhichCluster}; ...
698
           EfficiencyVector];
699
700
       % Update our waitbar:
701
       waitbar(BURST/TOTALBURSTS, h.Waitbar);
702
703
704
   end % end FOR
705
706
   fprintf(['%d bursts did not have sufficient photons ',...
707
        'to perform a sliding window calculation.\n\n'],...
708
       TRACKER);
709
710
   % Tidy-up our Cluster Statistics:
  RemoveClusters = cellfun(@isempty, CLUSTERSTATS);
712
   CLUSTERSTATS(RemoveClusters) = [];
   CLUSTEREDGES(RemoveClusters) = [];
714
715
   RemoveClusters = cell2mat(cellfun(@(x) (numel(x) < CLUSTERMINWINDOWS),...
716
       CLUSTERSTATS, ...
717
       'UniformOutput', false));
718
  CLUSTERSTATS(RemoveClusters) = [];
719
   CLUSTEREDGES(RemoveClusters) = [];
720
721
722
  IDs = setdiff(BURSTINDICES, REMOVEBURSTS); % Retain only 'good' bursts
723
724
725 DATA.Results = DATA.Results(IDs,:);
726
727 % Remove any data in which the mean efficiency is either 0 or 1: (why?
728 % these 'events' tend to arise when having low photon statistics and will
  % bunch up at the corner of the plots. Because of discrete photon
   % statistics, these will occur often and affect the relatively weighting of
   % the data elsewhere which has higher photon statistics and is more
732 % interesting (usually). Hence, we throw away this data, if present, and
733 % only keep that data which has more 'realistic' mean values.)
```

```
DATA.Results(DATA.Results(:,1) == 0 DATA.Results(:,1) == 1, :) = [];
735
  % Remove any data in which the standard deviation is zero: (why? Having a
736
  % single STD is not realistic, as fluctuations should be present simply to
  % to photon statistics. This also might be indicative of insufficient
  % number of WINDOWS allowing for STD calculation. Once again, with low
  % photon statistics, this is possible, and can bias the weighting of your
  % 2D histogram.)
742 DATA.Results(DATA.Results(:,2) == 0, :) = [];
743
744 % Remove all NaN rows:
DATA.Results(isnan(DATA.Results(:,1)), :) = [];
746
747
  % Tidy-up:
  clear ArrivalMatrix
750
  clear BinIndices
751
752
                        %% Cluster Statistics:
753
754
  waitbar(0,...
755
756
       h.Waitbar,...
       'Analyzing cluster statistics...');
757
758
759
  % ----- %
  % \ [ \ ] Group bursts with similar mean values and calculate population
761
  % standard deviation:
  ClusterCenters = CLUSTEREDGES(1:end-1) + CLUSTERWIDTH/2;
764
765
  % Initialize some variables:
766 NumClusters = numel(CLUSTERSTATS);
767 HighCIvec = zeros(NumClusters, 1);
768 LowCIvec = zeros(NumClusters, 1);
  StDevDistrMean = zeros(NumClusters, 1);
770
771
  ClusterStats = nan(NumClusters, 2);
772
773
  % Loop through each 'slice' (or 'cluster' of data) from the 2D Histogram
  % and calculate the statistics:
774
  for CLUSTER = 1 : NumClusters
775
776
       ClusterStats(CLUSTER, 2) = std(CLUSTERSTATS{CLUSTER});
777
       ClusterStats(CLUSTER,1) = mean(CLUSTERSTATS{CLUSTER});
778
       % Calculate StDev Distribution:
780
       Edges = 0 : (1-0)/1000 : 0.6; % why 0.6? Vertical upper bound (could ...
781
          be something else, but for practical purposes, we make this realistic)
```

```
NumWindows = numel(CLUSTERSTATS{CLUSTER});
782
       StDevDistr = MonteCarloPrediction(...
783
           Edges, ...
784
           NumWindows,... % number of windows
           WINDOW,... % number of photons within a window
786
           ClusterCenters(CLUSTER),... % probability of energy transfer
787
           NUMMONTECARLOSAMPLES); % number of samples generated by Monte ...
788
               Carlo simulation
789
       CumDistr = cumsum(StDevDistr);
790
791
       INV = 1 - CumDistr;
       StDevDistrMean(CLUSTER) = sum(StDevDistr .* Edges); % What is this? ...
792
           This should correspond to the theoretical shot noise prediction ...
           curve based on the number of photons specified by WINDOW
793
       %-- CI --
794
       %-- ADJUST CI BY THE MULT HYPOTH TEST --
795
       absvec = abs(CumDistr - CONFIDENCEINTERVAL);
796
797
       [M, closest] = min(absvec);
798
       absvecLOW = abs(INV - CONFIDENCEINTERVAL);
799
       closestLOW = find(absvecLOW == min(absvecLOW),...
800
801
           1, 'last');
802
       % Why are we calculating the CI off an index? The index tells us
803
       % something of about the number of false positives.
804
       HighCIvec(CLUSTER,:) = (closest-1)*.001; % 0.001 is 99.9% Confidence ...
       LowCIvec(CLUSTER,:) = (closestLOW-1)*.001;
806
807
       % Update our waitbar:
808
       waitbar(CLUSTER / NumClusters,...
809
           h.Waitbar);
810
811
812
813 end % end FOR
814
815 % Assign over to application data:
816 DATA.Clusters.Centers = ClusterCenters;
817 DATA.ConfidenceInterval.Upper = HighCIvec;
818 DATA.ConfidenceInterval.Lower = LowCIvec;
819 DATA.Clusters.StDevDistrMean = StDevDistrMean;
820 DATA.Clusters.Statistics = ClusterStats;
821 DATA.Clusters.WindowMeans = CLUSTERSTATS;
  DATA.Clusters.DistanceFromUpperCI = ...
822
       ClusterStats(:,2) - HighCIvec;
823
824
825
```

```
%% Tidy-up:
828
829
830
  % Delete waitbar:
831
  delete(h.Waitbar);
832
833
835
  TOTALTIME = toc
836
837
838
839
840
  return;
842
  843
 %% EOF
844
845
846
847
  848
  %% StartUp:
850
  function StartUp()
851
  응
852
  응
853
854
  addpath(...
855
      'Other/lightspeed',...
856
      'Other/SplitVec');
857
858
859
860
  861
 %% CleanUp:
862
  function CleanUp()
863
864
865
866
867
  rmpath(...
      'Other/lightspeed',...
869
      'Other/SplitVec');
870
```

Data Filtering

```
1 function [DATA, REMOVEBURSTS] = FilterData(...
      ArrivalMatrix, FILTERS)
  % FILTERDATA
      Filter a photon arrival matrix using supplied filter parameters.
5
  응
      INPUTS:
  응
          ARRIVALMATRIX: input data on which to perform Burst Variance
          Analysis. The input matrix should be Nx3 or 3xN, with the following
          as vector inputs: (1) Photon Arrival Times; (2) Detection Channel
          (DexDem = 0; DexAem = 2; AexAem = 1; AexDem = 3); (3) Burst
10
          Classification (those photons belonging to a burst should have a
          corresponding index to a unique burst ID; background photons should
12
          be indexed as NaN): ARRIVALMATRIX = ArrivalTimes
  9
                                                             ID
                                                                   BurstNumber
13
14 %
          FILTERS: nested structure providing minimum and maximum thresholds
15
          for the following data streams:
16
               DURATION: the temporal length of a detected burst [seconds]
17
               LENGTH: the number of photons within a detected burst [photons]
               DEXDEM: the number of photons detected in the donor channel
19
                   upon donor excitation.
20
               DEXAEM: the number of photons detected in the acceptor channel
21 응
22 %
                   upon donor excitation.
               AEXAEM: the number of photons detected in the acceptor channel
23
                   upon acceptor excitation.
24
               SUMPHOTONS: the number of photons detected in the donor &
25
26
                   acceptor channels upon donor excitation.
               TOTALPHOTONS: the number of photons detected in the donor &
27
  9
                   acceptor channels upon donor and acceptor excitation.
28
                   (excluding photons detected in donor channel upon acceptor
29
 응
  %
                   excitation: AexDem)
               EFFICIENCY: the FRET efficiency.
31
               STOICHIOMETRY: the relative fluorophore brightness.
  응
32
  응
33
      OUTPUTS:
34
  0
          DATA: structure with the following data streams for each detected
35
          burst:
  9
36
  응
               DURATION: the temporal length of a detected burst [seconds]
               LENGTH: the number of photons within a detected burst [photons]
               DEXDEM: the number of photons detected in the donor channel
39
                   upon donor excitation.
40
               DEXAEM: the number of photons detected in the acceptor channel
41
                   upon donor excitation.
43
              AEXAEM: the number of photons detected in the acceptor channel
44 %
                   upon acceptor excitation.
45 %
               SUMPHOTONS: the number of photons detected in the donor &
```

```
acceptor channels upon donor excitation.
46
             TOTALPHOTONS: the number of photons detected in the donor &
47
                acceptor channels upon donor and acceptor excitation.
                 (excluding photons detected in donor channel upon acceptor
                 excitation: AexDem)
50
51 %
             EFFICIENCY: the FRET efficiency.
             STOICHIOMETRY: the relative fluorophore brightness.
52 %
         REMOVEBURSTS: a vector providing indices for the bursts which
54
         failed filter criteria.
55 %
56 %
57 %
     DEPENDENCIES:
58 %
        SPLITVEC.M
59 %
  응
      EXAMPLE USAGE:
61
         [DATA, REMOVEBURSTS] = FilterData(ArrivalMatrix, FILTERS);
62
63
 9
 응
     History:
65
       [1] Kristofer Gryte. University of Oxford (2011).
66
67
      Copyright (c) 2011. Kristofer Gryte. University of Oxford.
69
 9
70
71 %
     Version: 1.0 (2011-07-30)
72 응
73 %
74 응
 응
      Author Information:
75
        Kristofer Gryte
76
         Clarendon Laboratory
 응
77
78 %
        University of Oxford
 은
        Oxford, United Kingdom
        e-mail: k.gryte1 (at) physics.ox.ac.uk
81 %
82 %
     TODO:
83
       [1] Add checks for input variables.
84
85
86 %
89 %% EOP
 %% Initialization:
94
```

```
REMOVEBURSTS = [];
95
96
  Apply = true; % FLAG to determine if apply filters or not. THIS IS NEEDED, ...
      FOR NOW, EVEN IF NO FILTERS ARE APPLIED!!!!
98
   %% Assemble Single Data Matrix:
  % Replace all the NaNs:
102
  ArrivalMatrix(isnan(ArrivalMatrix(:,3)),3) = 0;
103
  % Return the assigned IDs of the various runs: % recall that non-burst
105
  % photons are labeled with a '0'!!!!
106
   FLAGVALS = SplitVec(...
107
       ArrivalMatrix(:,3),...ArrivalMatrix(¬isnan(ArrivalMatrix(:,3)),3),.......
          % exclude any NaNs
       'equal',... % find all those runs of the same index (burst ID)
109
       'firstval'); % return just the first index marking a run.
110
111
112
   응 --- 응
113
   % [1] Start Times:
114
   STARTINDICES = SplitVec(...
       ArrivalMatrix(:,3),...ArrivalMatrix(¬isnan(ArrivalMatrix(:,3)),3),........
116
          % exclude any NaNs
       'equal',... % find all those runs of the same index (burst ID)
117
118
       'first'); % return just the first index marking a run.
119
  STARTINDICES (FLAGVALS == 0) = []; % NOTE:
120
121
  DATA.StartTimes = ArrivalMatrix(STARTINDICES, 1);
122
123
   응 --- 응
124
   % [2] End Times:
125
   ENDINDICES = SplitVec(...
126
       ArrivalMatrix(:,3),...ArrivalMatrix(¬isnan(ArrivalMatrix(:,3)),3),.......
127
           % exclude any NaNs
       'equal',... % find all those runs of the same index (burst ID)
128
       'last'); % return just the last index completing a run.
130
131 ENDINDICES (FLAGVALS == 0) = []; % NOTE:
  DATA.EndTimes = ArrivalMatrix(ENDINDICES, 1);
133
134
  응 --- 응
135
  % [3] Duration:
  DATA.Duration = DATA.EndTimes - DATA.StartTimes;
137
138
139 % --- %
```

```
140 % [4] Length:
  DATA.Length = ENDINDICES - STARTINDICES;
142
  응 --- 응
  BURSTCELL(:,1) = SplitVec(...
144
       ArrivalMatrix(:, 3),...ArrivalMatrix(¬isnan(ArrivalMatrix(:,3)), 2),...
145
       'equal',...
146
       'loc'); % this is a cellular array!
148
  BURSTCELL (FLAGVALS == 0) = []; % NOTE:
149
150
151 % ----- %
152 % Free up some memory:
  % clear ArrivalMatrix
   응 _____ 응
155
   % [5] DexDem: % DexDem = 0
156
  DATA.DexDem(:,1) = cellfun(@(x) sum(ArrivalMatrix(x,2) == 0), BURSTCELL, ...
      'UniformOutput', true);
158
  % [6] DexAem: % DexAem = 2
160 DATA.DexAem(:,1) = cellfun(@(x) sum(ArrivalMatrix(x,2) == 2), BURSTCELL, ...
      'UniformOutput', true);
161
162 % [7] AexAem: % AexAem = 1
163 DATA.AexAem(:,1) = cellfun(@(x) sum(ArrivalMatrix(x,2) == 1), BURSTCELL, ...
      'UniformOutput', true);
164
  용 --- 왕
165
  % [8] Sum Photons: DexDem + DexAem
  DATA.SumPhotons = DATA.DexDem + DATA.DexAem;
168
  응 --- 응
169
  % [9] Total Photons: DexDem + DexAem + AexAem
171 DATA.TotalPhotons = DATA.SumPhotons + DATA.AexAem;
172
  응 --- 응
173
  % [10] Efficiency:
  DATA.Efficiency = DATA.DexAem ./ (DATA.SumPhotons);
176
177 % --- %
178 % [11] Stoichiometry:
  DATA.Stoichiometry = DATA.SumPhotons ./ (DATA.TotalPhotons);
180
183 %% Threshold Data:
184
185 if Apply == true
```

```
186
       187
       % Min Duration:
188
       if ¬isempty(FILTERS.Duration.Min)
189
190
           IDs = find(DATA.Duration < FILTERS.Duration.Min); % Column vector</pre>
191
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
192
193
           DATA.Duration(DATA.Duration < FILTERS.Duration.Min,:) = NaN;
194
195
       end
196
197
       198
       % Max Duration:
199
       if ¬isempty(FILTERS.Duration.Max)
200
201
           IDs = find(DATA.Duration > FILTERS.Duration.Max); % Column vector
202
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
203
204
           DATA.Duration(DATA.Duration > FILTERS.Duration.Max,:) = NaN;
205
206
       end
207
208
209
       $$$$$$$$$$$$$$$$$$$$$$$$$$
210
       % Min Length:
211
212
       if ¬isempty(FILTERS.Length.Min)
213
           IDs = find(DATA.Length < FILTERS.Length.Min); % Column vector</pre>
214
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
215
216
           DATA.Length (DATA.Length < FILTERS.Length.Min,:) = NaN;
217
218
219
       end
220
       221
       % Max Length:
222
223
       if ¬isempty(FILTERS.Length.Max)
224
           IDs = find(DATA.Length > FILTERS.Length.Max); % Column vector
225
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
226
227
           DATA.Length (DATA.Length > FILTERS.Length.Max,:) = NaN;
228
229
       end
230
232
233
       234
```

```
% Min DexcDem:
235
       if ¬isempty(FILTERS.DexDem.Min)
236
237
            IDs = find(DATA.DexDem < FILTERS.DexDem.Min); % Column vector</pre>
238
            REMOVEBURSTS = [REMOVEBURSTS; IDs];
239
240
            DATA.DexDem(DATA.DexDem < FILTERS.DexDem.Min,:) = NaN;</pre>
241
242
       end
243
244
       88888888888888888888888888888888
245
        % Max DexcDem:
246
       if ¬isempty(FILTERS.DexDem.Max)
247
248
            IDs = find(DATA.DexDem > FILTERS.DexDem.Max); % Column vector
            REMOVEBURSTS = [REMOVEBURSTS; IDs];
250
251
            DATA.DexDem(DATA.DexDem > FILTERS.DexDem.Max,:) = NaN;
252
253
       end
254
255
       $$$$$$$$$$$$$$$$$$$$$$$$$$
256
        % Min DexcAem:
257
        if ¬isempty(FILTERS.DexAem.Min)
258
259
            IDs = find(DATA.DexAem < FILTERS.DexAem.Min); % Column vector</pre>
260
261
            REMOVEBURSTS = [REMOVEBURSTS; IDs];
262
            DATA.DexAem(DATA.DexAem < FILTERS.DexAem.Min,:) = NaN;</pre>
263
264
       end
265
266
       267
       % Max DexcAem:
268
       if ¬isempty(FILTERS.DexAem.Max)
269
270
            IDs = find(DATA.DexAem > FILTERS.DexAem.Max); % Column vector
271
272
            REMOVEBURSTS = [REMOVEBURSTS; IDs];
273
            DATA.DexAem(DATA.DexAem > FILTERS.DexAem.Max,:) = NaN;
274
275
276
       end
277
       $$$$$$$$$$$$$$$$$$$$$$$$$$$$$
278
        % Min AexcAem:
279
       if ¬isempty(FILTERS.AexAem.Min)
281
            IDs = find(DATA.AexAem < FILTERS.AexAem.Min); % Column vector</pre>
282
            REMOVEBURSTS = [REMOVEBURSTS; IDs];
283
```

```
284
           DATA.AexAem(DATA.AexAem < FILTERS.AexAem.Min,:) = NaN;
285
286
       end
287
288
       289
       % Max AexcAem:
290
291
       if ¬isempty(FILTERS.AexAem.Max)
292
           IDs = find(DATA.AexAem > FILTERS.AexAem.Max); % Column vector
293
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
294
295
           DATA.AexAem(DATA.AexAem > FILTERS.AexAem.Max,:) = NaN;
296
297
       end
298
299
       300
       % Min Sum DD+DA:
301
302
       if ¬isempty(FILTERS.SumPhotons.Min)
303
           IDs = find(DATA.SumPhotons < FILTERS.SumPhotons.Min); % Column vector</pre>
304
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
305
           DATA.SumPhotons(DATA.SumPhotons < FILTERS.SumPhotons.Min,:) = NaN;
307
308
       end
309
310
       311
       % Max Sum DD+DA:
312
       if ¬isempty(FILTERS.SumPhotons.Max)
313
314
           IDs = find(DATA.SumPhotons > FILTERS.SumPhotons.Max); % Column vector
315
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
316
317
           DATA.SumPhotons(DATA.SumPhotons > FILTERS.SumPhotons.Max,:) = NaN;
318
319
       end
320
321
       % Min Total Signal:
323
       if ¬isempty(FILTERS.TotalPhotons.Min)
324
325
           IDs = find(DATA.TotalPhotons < FILTERS.TotalPhotons.Min); % Column ...</pre>
326
              vector
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
327
           DATA. Total Photons (DATA. Total Photons < FILTERS. Total Photons. Min,:) ...
329
              = NaN;
330
```

```
end
331
332
       $$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$
333
       % Max Total Signal:
334
       if ¬isempty(FILTERS.TotalPhotons.Max)
335
336
           IDs = find(DATA.TotalPhotons > FILTERS.TotalPhotons.Max); % Column ...
337
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
338
339
           DATA.TotalPhotons(DATA.TotalPhotons > FILTERS.TotalPhotons.Max,:) ...
340
              = NaN;
341
       end
342
       344
       % Min Efficiency:
345
       if ¬isempty(FILTERS.Efficiency.Min)
346
347
           IDs = find(DATA.Efficiency < FILTERS.Efficiency.Min); % Column vector
348
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
349
350
           DATA.Efficiency (DATA.Efficiency < FILTERS.Efficiency.Min,:) = NaN;
351
352
       end
353
354
       355
       % Max Efficiency:
356
       if ¬isempty(FILTERS.Efficiency.Max)
357
358
           IDs = find(DATA.Efficiency > FILTERS.Efficiency.Max); % Column vector
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
360
361
362
           DATA.Efficiency (DATA.Efficiency > FILTERS.Efficiency.Max,:) = NaN;
363
       end
364
365
       366
       % Min Stoichiometry:
367
       if ¬isempty(FILTERS.Stoichiometry.Min)
368
369
           IDs = find(DATA.Stoichiometry < FILTERS.Stoichiometry.Min); % ...</pre>
370
              Column vector
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
371
372
           DATA. Stoichiometry (DATA. Stoichiometry < ...
              FILTERS.Stoichiometry.Min,:) = NaN;
374
375
       end
```

```
376
       $$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$$
377
       % Max Stoichiometry:
378
       if ¬isempty(FILTERS.Stoichiometry.Max)
379
380
           IDs = find(DATA.Stoichiometry > FILTERS.Stoichiometry.Max); % ...
381
              Column vector
           REMOVEBURSTS = [REMOVEBURSTS; IDs];
383
           DATA.Stoichiometry (DATA.Stoichiometry > ...
384
              FILTERS.Stoichiometry.Max,:) = NaN;
385
       end
386
387
  end % end IF
388
389
390
  % Retain only the unique indices:
391
  REMOVEBURSTS = unique(REMOVEBURSTS);
393
394
395
397 % FOR NOW...
398 return;
399
400
401
403 %% EOF
```

Monte Carlo Simulation

```
1 function STDDISTRIBUTION = MonteCarloPrediction(EDGES, EVENTS,...
      TRIALS, PROBABILITY, ITERATIONS)
3 % MONTECARLOPREDICTION
      Use Monte Carlo methods to generate a distribution of standard
      deviation (std) values observed from a binomial process (Bernoulli
      Trial) with a specified probability of success.
  응
      INPUTS:
         EDGES: a vector defining the bin edges into which the simulated
          standard deviations will be histogrammed.
10
12 %
          EVENTS: the number of events (collections of trials) to be
          sampled from the distribution.
13 %
14 %
          TRIALS: the number of trials.
15 %
16 %
17 %
          PROBABILITY: the probability of success. For example, a series of
          coin flips is a collection of events characterized by Bernoulli
18
          trials with probability 0.5.
19
20 응
          ITERATIONS: the number of times to repeat each collection of events
21 응
          so as to arrive at an average value for the calculated standard
          deviations. Default: 500.
23 %
24 %
25 %
26
  응
      OUTPUTS:
          STDDISTRIBUTION: a probability distribution of the standard
27
          deviation with support defined by EDGES. Output is a vector.
 9
28
29 %
 응
      DEPENDENCIES:
         BINORND.M (STATS toolbox)
31 %
32 %
  응
33
      EXAMPLE USAGE:
34
         MonteCarloPrediction([0:0.0001:1], 5, 100, 0.5);
  응
35
36
  응
  응
          MonteCarloPrediction([0:0.0001:1], 5, 100, 0.5, 500);
  응
  용
39
      History:
  용
40
          [1] Joseph Torella, Yusdi Santoso. University of Oxford (2009).
  응
41
          [2] Kristofer Gryte. University of Oxford (2011).
43 %
      Copyright (c) 2011. Kristofer Gryte. University of Oxford.
44 %
45 %
```

```
46
  응
47
     Version: 1.0 (2011-07-30)
  응
50
     Author Information:
  9
51
        Kristofer Gryte
52
53
         Clarendon Laboratory
         University of Oxford
54
        Oxford, United Kingdom
55 %
         e-mail: k.gryte1 (at) physics.ox.ac.uk
56 %
57 %
58 %
      TODO:
59
      [1] Generate a function independent of the STATS toolbox.
61
62
      ______
63
 9
 응
     See also BINORND
65 %
66
%% EOP
69
70
71
72
73
76 %% Check Input Arguments:
77
78 % Check!
79 if nargin ≤ 4
     ITERATIONS = 500; % default value
80
81 end % end IF
82
83 % Check EDGES:
 if size(EDGES,1) \neq 1 && size(EDGES,2) \neq 1
84
85
     % Throw an error:
      errordlg(['ERROR:invalid input argument. Please input a vector for the ...
86
        ',...
         'EDGES argument.'],...
87
         'ERROR: invalid input argument');
88
89
      return;
91 end % end IF
92
93 % Check EVENTS, TRIALS, PROBABILITY, and ITERATIONS
```

```
temp = [EVENTS, TRIALS, ITERATIONS];
   if sum((\neg isinteger(temp)) + (temp>0)) \neq 6
       % Throw an error:
96
       errordlg(['ERROR:invalid input argument. Please input positive ...
          integers for the ',...
           'EVENTS, TRIALS, and ITERATIONS arguments.'],...
98
           'ERROR: invalid input argument');
100
       return;
101
  end % end IF
102
103
   if (PROBABILITY < 0
                          PROBABILITY > 1)
104
       % Throw an error:
105
       errordlg(['ERROR:invalid input argument. Please input a probability ...
106
          between 0 and 1 ',...
           'for the PROBABILITY argument.'],...
107
           'ERROR: invalid input argument');
108
109
       return;
110
  end % end IF
112
113
114
115
117 %% Initialization:
119
  응 --- 응
  % Parameters:
120
  Runs = 3; % used in the event of a large number of requested standard ...
      deviations, in which case, for memory purposes, we need to break the ...
      work into smaller chunks.
122
123
124
  %% Generate Monte Carlo Prediction:
126
   if EVENTS*ITERATIONS ≤ 10000 % 10,000 ? A magical number discovered ...
128
      heuristically.
129
       % Generate the matrix of numbers drawn from a binomial distribution
130
       % with total trials equal to TRIALS and probability equal to
131
       % PROBABILITY:
132
       MyMatrix = binornd(...
133
           TRIALS, ...
134
           PROBABILITY, ...
135
          EVENTS, ...
136
           ITERATIONS); % this is a matrix of size: EVENTS x ITERATIONS
137
```

```
138
       % Calculate the standard deviation along each column
139
       StdMatrix = std(MyMatrix ./ TRIALS);
140
141
   elseif EVENTS*ITERATIONS > 10000
142
143
       % With many iterations, we run into the problem of very large matrices
144
145
       % and memory issues. Let us break it down more manageably:
146
147
       RemainingSamples = ITERATIONS;
148
149
       % Pre-allocate our array:
150
       StdMatrix = nan(1, ITERATIONS);
151
       % Initialize an index counter:
153
       Counter = 1;
154
155
156
       for k = 1: Runs
157
            % Determine the number of samples to run:
158
            Samples = floor(RemainingSamples / (Runs-k+1));
159
160
            % Generate the matrix of numbers drawn from a binomial distribution
161
            % with total trials equal to TRIALS and probability equal to
162
            % PROBABILITY:
163
164
            MyMatrix = binornd(...
                TRIALS, ...
165
                PROBABILITY, ...
166
                EVENTS, ...
167
                Samples);
168
169
            % Calculate the standard deviation along each column and append to
170
171
            % our standard deviation matrix:
            StdMatrix(Counter : (Counter+Samples-1)) = std(MyMatrix ./ TRIALS);
172
173
            % Calculate the remaining number of samples:
174
175
            RemainingSamples = RemainingSamples - Samples;
176
            % Update our counter:
177
            Counter = Counter + Samples;
178
179
180
       end % end FOR
181
182
   end % end IF/ELSEIF
184
185
  % Histogram the standard deviations where the bin edges are defined by
```

Visualization

```
1 function [DATA, handles] = PlotResults(DATA, WINDOW, handles)
2 % PLOTRESULTS
      Plots the DATA results generated by Burst Variance Analysis into the
      axes supplied by HANDLES. Configuration and aesthetics of the axes is
      currently controlled internally. NOTE: the BVA data is smoothed from a
      discretized histogram presentation and transformed to a continuous
      contour. The reference for this smoothing is " Garcia D, Robust
      smoothing of gridded data in one and higher dimensions with missing
      values. Computational Statistics & Data Analysis, 2010. "
10
      INPUTS:
          DATA: structure with the following data streams for each detected
12
          burst: (NOTE: data streams are filtered according to input FILTERS)
 9
13
14 %
              DURATION: the temporal length of a detected burst [seconds]
              LENGTH: the number of photons within a detected burst [photons]
15
              DEXDEM: the number of photons detected in the donor channel
16
17
                   upon donor excitation.
              DEXAEM: the number of photons detected in the acceptor channel
                  upon donor excitation.
19
              AEXAEM: the number of photons detected in the acceptor channel
20
                  upon acceptor excitation.
21 응
22 %
              SUMPHOTONS: the number of photons detected in the donor &
                  acceptor channels upon donor excitation.
23
              TOTALPHOTONS: the number of photons detected in the donor &
24
                   acceptor channels upon donor and acceptor excitation.
25
26
                   (excluding photons detected in donor channel upon acceptor
                   excitation: AexDem)
27
              EFFICIENCY: the FRET efficiency.
 용
28
              STOICHIOMETRY: the relative fluorophore brightness.
29
  %
              RESULTS: an Nx2 array, where N is the number of bursts (after
                   filtering). The first column corresponds to the mean ...
31 %
      Efficiency
  응
                  value for a burst and the second to the burst's Efficiency
32
                   standard deviation. Array: E(efficiency) sqrt(V(efficiency))
33
              CLUSTERS: nested structure with the following fields:
34
                  CENTERS: vector containing the centers of the individual
35
 9
  응
                      clusters.
                   STDEVDISTRMEAN: the Monte Carlo simulated standard
38
                      deviation prediction.
                   STATISTICS: an Mx2 array, where M is the number of clusters
39
                       ('slices' along the efficiency axis, into which bursts are
40
                       grouped). The first column corresponds to the mean
42 %
                       Efficiency value for the cluster, and the second to the
                       standard deviation of the burst's within that cluster.
43 %
              CONFIDENCEINTERVAL: nested structure with the following fields:
```

```
UPPER: the upper confidence bound generated from the Monte
45
                      Carlo simulation and multiple hypothesis test.
46
                   LOWER: the lower confidence bound generated from the Monte
47
                      Carlo simulation and the multiple hypothesis test.
49
          WINDOW: size of the sliding window over which FRET will be
50
          calculated. Default: 5.
51
          HANDLES: structure with the following fields:
53
               FIGURE: handle to the figure window.
54
               SUBPLOT: nested structure with the following fields:
55
  응
                   BVA: nested structure with the following fields:
56
  9
                       AXES: handles to the 2D Histogram axes (subplot).
57
               SUBPLOT: nested structure with the following fields:
58
                   HISTOGRAM: nested structure with the following fields:
                       MEAN: nested structure with the following fields:
60
                          AXES: handle to the axes into which the histogram
61
                             will be plotted.
62
                       STDEV: nested structure with the following fields:
                          AXES: handle to the axes into which the histogram
                             will be plotted.
65
66
68
      OUTPUTS:
  응
69
          DATA: input structure with the following added data streams:
  응
70
  응
               HISTOGRAMS: nested structure with the following fields:
                   COUNTS: the counts per bin for the 2D histogram.
72
                   EDGES: the edges used to define the histogram bins.
  응
73
                   BINCENTERS: the locations of the center of each bin.
74
                   SMOOTHED: the histogram data after application of a
                      smoothing filter.
76
  응
77
          HANDLES: input structure with the following added fields:
               PLOTS: nested structure with the following fields:
                   CONTOUR: handle to the counter plot created from the 2D
80
                       histogram
81
                   SHOTNOISEPREDICTION: handle to the shot noise prediction
82
                      based on the number of photons specified in WINDOW.
84
                   CONFIDENCEINTERVAL: nested structure with the following
                      fields:
  응
85
                       UPPER: handle to the upper confidence interval
  응
                          generated via a Monte Carlo simulation. Cluster STD
87
                          values above this limit are indicative of dynamics.
88
                       MEAN: handle to the average of upper and lower
89
                          confidence bounds. This should correspond to the
                          SHOTNOISEPREDICTION curve.
91
 응
                   STATIC: STD clusters which fall below the upper confidence
92
                      interval are considered 'static'. This is the handle to
93 응
```

```
these data points.
94
  응
                    DYNAMICS: STD clusters which are above the upper confidence
95
                       interval are considered 'dynamic'. This is the handle to
   응
                       these data points.
                    UNKNOWN: STD clusters which fall below the lower confidence
98
                       interval are categorized as 'unknown'. The causes of this
                       behavior are not well understood.
100
                SUBPLOT: nested structure with the following fields:
                    HISTOGRAM: nested structure with the following fields:
102
                        MEAN: nested structure with the following fields:
   응
103
                            PLOTS: nested structure with the following fields:
104
  응
                                BAR: handle to the Mean bar graph data.
105
106 %
                        STDEV: nested structure with the following fields:
107 %
                            PLOTS: nested structure with the following fields:
                                 BAR: handles to the StDev bar graph data.
                COLORBAR: handle to the colobar corresponding to the 2D
109
110
                   histogram data.
111
   9
112
  응
113
       DEPENDENCIES:
114
           SMOOTHN.M
115
           HISTCN.M
117
   응
118
       EXAMPLE USAGE:
119
   응
   응
           [DATA, handles] = PlotResults(DATA, WINDOW, handles);
   응
121
   응
122
       History:
123
           [1] Kristofer Gryte. University of Oxford (2011).
124
   응
125
       Copyright (c) 2011. Kristofer Gryte. University of Oxford.
126
   9
127
  응
128
129
       Version: 1.0 (2011-08-02)
130
131
132
133
   응
       Author Information:
          Kristofer Gryte
134 %
135
  응
           Clarendon Laboratory
           University of Oxford
136
           Oxford, United Kingdom
137
           e-mail: k.gryte1 (at) physics.ox.ac.uk
138
139
140
141 %
       TODO:
       [1] Allow axes configurations to be an input structure.
142 %
```

```
[2] Provide checks on input arguments.
          [3] Figure out why 'histon' appears to reset the path(s).
144
145
147
  148
  %% EOP
149
151
  152
  %% StartUp/CleanUp:
153
154
  StartUp();
155
156
  C = onCleanup(@() CleanUp());
157
158
159
  160
  %% Initialization:
161
  응 --- 응
163
  % Check!!!
164
  if nargin \neq 3
      % Throw an error:
166
      errordlg(['ERROR:not enough input arguments. Please provide the ...
167
         following ',...
          'input arguments: DATA, WINDOW, HANDLES. See the HELP for the more ...
168
             ١, . . .
          'information.'],...
169
          'ERROR: not enough input arguments');
170
      return;
  end % end IF
172
173
  174
  % Default Configurations:
175
176
  Config.Histogram2D.Smoothing = 20; % Smoothing parameter
177
178
  Config.ShotNoise.Prediction.OrdinateVals = 0:0.01:1; % Efficiency sampling....
  Config.ShotNoise.Prediction.StDev = sqrt(...
180
       (Config.ShotNoise.Prediction.OrdinateVals .* ...
181
         (1-Config.ShotNoise.Prediction.OrdinateVals))...
      /...
182
      WINDOW);
183
184
  Config.Axes.XLabel.FontSize = 14;
  Config.Axes.XLabel.String = 'Mean';
186
187
  Config.Axes.YLabel.FontSize = 14;
```

```
Config.Axes.YLabel.String = 'StDev';
189
190
   Config.Axes.Title.FontSize = 14;
191
   Config.Axes.Title.String = 'Burst Variance Analysis (Efficiency)';
192
193
   Config.Axes.TickDir = 'out';
194
   Config.Axes.Box = 'off';
195
196
   Config.Figure.BackgroundColor = 'w';
197
198
   Config.Legend.Box = 'off';
199
   Config.Legend.Location = 'south';
200
201
202
   Config.Histogram.Mean.Axes.YLabel.FontSize = 14;
203
   Config.Histogram.Mean.Axes.YLabel.String = 'Counts';
204
205
   Config.Histogram.Mean.Axes.Box = 'off';
206
207
   Config.Histogram.Mean.Axes.TickDir = 'out';
208
209
   Config.Histogram.StDev.Axes.XLabel.FontSize = 14;
210
211
   Config.Histogram.StDev.Axes.XLabel.String = 'Counts';
212
   Config.Histogram.StDev.Axes.Box = 'off';
213
   Config.Histogram.StDev.Axes.TickDir = 'out';
214
215
   216
217
218
   응 _____ 응
219
   %% Plot:
220
221
  % Activate our figure:
222
  figure (handles.figure);
223
224
   % Activate our BVA Axes:
225
226
   subplot (handles.Subplot.BVA.Axes);
   % Get the histogram counts: (NOTE: default is 100 Bins)
228
   [DATA.Histogram.Counts, DATA.Histogram.Edges, DATA.Histogram.BinCenters] = ...
229
      histcn(...
       DATA.Results);
230
231
   StartUp(); %%% NOTE!!! For some reason, 'histc'/'histcn' appears to reset ...
232
      the path(s)
233
  % 'Smooth' the histogram:
234
235 DATA.Histogram.Smoothed = smoothn(...
```

```
DATA. Histogram. Counts, ...
236
       Config.Histogram2D.Smoothing);
237
238
   % [1] Create the contour plot of the burst standard deviations:
240
   [ContourMatrix, handles.Plots.Contour] = contourf(...
241
242
       DATA.Histogram.BinCenters{1},...
243
       DATA.Histogram.BinCenters{2},...
       DATA.Histogram.Smoothed(:,:,1)',...
244
       10, ...
245
       'EdgeColor', 'none', ...
246
       'tag', 'Per Burst StDev');
247
248
   hold(handles.Subplot.BVA.Axes, 'on');
249
   % ----- %
251
   % [2] Display the shot noise prediction:
252
  handles.Plots.ShotNoisePrediction = plot(...
253
254
       Config.ShotNoise.Prediction.OrdinateVals,...
       Config.ShotNoise.Prediction.StDev,...
255
       'k-',...
256
       'LineWidth', 3.0,...
257
       'tag', 'Shot Noise Prediction');
258
259
   % ----- %
260
   % [3] Plot the confidence bounds:
261
262
   handles.Plots.ConfidenceInterval.Upper = plot(...
       DATA.Clusters.Centers,...
263
       DATA.ConfidenceInterval.Upper,...
264
265
       'm-.',...
       'LineWidth', 2.5,...
266
       'tag', 'Upper CI');
267
268
  handles.Plots.ConfidenceInterval.Mean = plot(...
       DATA.Clusters.Centers(:, 1),...
270
       DATA.Clusters.StDevDistrMean(:, 1),...
271
       'y--', ...
272
       'LineWidth', 2.0,...
273
       'MarkerSize', 10,...
274
       'tag', 'Mean MC Prediction');
275
276
  % ----- %
277
  % [4] Determine where the clustered populations reside relative to the
278
   % confidence bounds:
279
  LogID = (DATA.Clusters.Statistics(:,2) \leq DATA.ConfidenceInterval.Upper...
280
281
       DATA.Clusters.Statistics(:,2) > DATA.ConfidenceInterval.Lower);
282
283
284 handles.Plots.Static = plot(...
```

```
DATA.Clusters.Centers(LogID, 1),...
285
        DATA.Clusters.Statistics(LogID, 2),...
286
        'k^',...
287
        'LineWidth', 1.5,...
288
        'MarkerFaceColor', 'k', ...
289
        'MarkerSize', 8,...
290
        'tag', 'Static');
291
292
293
   LogID = (DATA.Clusters.Statistics(:,2) > DATA.ConfidenceInterval.Upper);
294
   handles.Plots.Dynamics = plot(...
295
        DATA.Clusters.Centers(LogID, 1),...
296
        DATA.Clusters.Statistics(LogID, 2),...
297
        'k^',...
298
        'LineWidth', 1.5,...
        'MarkerFaceColor', 'r', ...
300
        'MarkerSize', 8,...
301
        'tag', 'Dynamics');
302
303
   LogID = (DATA.Clusters.Statistics(:,2) < DATA.ConfidenceInterval.Lower);</pre>
304
   handles.Plots.Unknown = plot(...
305
        DATA.Clusters.Centers(LogID, 1),...
306
307
        DATA.Clusters.Statistics(LogID, 2),...
        'k^',...
308
        'LineWidth', 1.5,...
309
        'MarkerFaceColor', 'g', ...
310
311
        'MarkerSize', 8,...
        'tag', 'Unknown');
312
313
314
315
   hold(handles.Subplot.BVA.Axes, 'off');
316
317
318
   % [6] Compute Histograms:
319
320
   DATA.Histogram.Edges\{1,1\} = -0.025 : 0.05 : 1.025; % Comment out to retain ...
321
       3D hist edges
322
   Counts1 = histc(DATA.Results(:,1),...
323
       DATA.Histogram.Edges{1,1});
324
325
   handles.Subplot.Histogram.Mean.Plots.Bar = ...
326
        bar (handles. Subplot. Histogram. Mean. Axes, ...
327
            DATA.Histogram.Edges{1,1}, Counts1, 'histc');
328
   Counts2 = histc(DATA.Results(:,2),...
330
       DATA.Histogram.Edges{1,2});
331
332
```

```
handles.Subplot.Histogram.StDev.Plots.Bar = ...
       barh (handles.Subplot.Histogram.StDev.Axes, ...
334
            DATA.Histogram.Edges{1,2}, Counts2, 'histc');
335
336
   % ----- %
337
   %% Adjust Axes:
338
339
340
   % BVA:
   xlim(handles.Subplot.BVA.Axes,...
341
        [min(Config.ShotNoise.Prediction.OrdinateVals),...
342
       max(Config.ShotNoise.Prediction.OrdinateVals)]);
343
344
   ylim (handles.Subplot.BVA.Axes,...
345
        [0, max(DATA.Results(:,2))]);
346
347
   set (handles.figure, ...
348
        'color', Config.Figure.BackgroundColor)
349
350
351
   ylabel(handles.Subplot.BVA.Axes,...
       Config.Axes.YLabel.String, ...
352
        'FontSize', Config.Axes.YLabel.FontSize);
353
354
355
   xlabel(handles.Subplot.BVA.Axes,...
       Config.Axes.XLabel.String, ...
356
        'FontSize', Config.Axes.XLabel.FontSize)
357
358
359
   title (handles.Subplot.BVA.Axes, ...
       Config.Axes.Title.String, ...
360
        'FontSize', Config.Axes.Title.FontSize);
361
362
   set (handles.Subplot.BVA.Axes, ...
363
        'TickDir', Config.Axes.TickDir,...
364
        'Box', Config.Axes.Box);
365
366
367
   % Histograms:
368
   xlim(handles.Subplot.Histogram.Mean.Axes,...
369
370
        [min(Config.ShotNoise.Prediction.OrdinateVals),...
       max(Config.ShotNoise.Prediction.OrdinateVals)]);
371
372
   ylabel(handles.Subplot.Histogram.Mean.Axes, ...
373
       Config. Histogram. Mean. Axes. YLabel. String, ...
374
        'FontSize', Config.Histogram.Mean.Axes.YLabel.FontSize);
375
376
   set (handles.Subplot.Histogram.Mean.Axes,...
377
        'Box', Config.Histogram.Mean.Axes.Box,...
378
        'TickDir', Config.Histogram.Mean.Axes.TickDir);
379
380
381
```

```
ylim(handles.Subplot.Histogram.StDev.Axes,...
       [0, max(DATA.Results(:,2))]);
383
384
  xlabel(handles.Subplot.Histogram.StDev.Axes,...
385
       Config. Histogram. StDev. Axes. XLabel. String, ...
386
       'FontSize', Config.Histogram.StDev.Axes.XLabel.FontSize);
387
388
389
   set (handles.Subplot.Histogram.StDev.Axes,...
       'Box', Config.Histogram.StDev.Axes.Box,...
390
       'TickDir', Config.Histogram.StDev.Axes.TickDir);
391
392
393
   % ----- %
394
   % Create a colorbar:
395
  APOS = get(handles.Subplot.BVA.Axes, 'position');
397
398
  handles.Colorbar = colorbar('peer', handles.Subplot.BVA.Axes);
399
400
  CPOS = get(handles.Colorbar, 'position');
401
402
  set (handles.Colorbar, ...
403
      'Position', [APOS(1)+APOS(3)+0.01, CPOS(2), CPOS(3), CPOS(4)]);
404
405
406
  % Configure the Legend:
407
  % handles.Plots.Current = [handles.Plots.Contour, ...
      handles.Plots.ShotNoisePrediction,...
        handles.Plots.ConfidenceInterval.Upper, ...
409
      handles.Plots.ConfidenceInterval.Lower,...
        handles.Plots.ConfidenceInterval.Mean,...
410
        handles.Plots.Static, handles.Plots.Dynamics, handles.Plots.Unknown];
411
412
  % handles.Legend = legend(...
413
       handles.Plots.Current,...
414
        get(handles.Plots.Current, 'tag'));
415
416
417
  % set (handles.Legend, ...
        'Box', DATA.BVA.Config.Legend.Box,...
418
         'Location', DATA.BVA.Config.Legend.Location);
419
420
421
422
424 %% EOF
425
426
427
```

```
429 %% StartUp:
430
  function StartUp()
431
432
433
434
435 % Add path(s):
  addpath(...
436
      'Other/histcn',...
437
      'Other/smoothn');
438
439
440
441
442
445 %% CleanUp:
446 function CleanUp()
447 %
449
450
  % Remove path(s):
451
452 rmpath(...
453
      'Other/histcn',...
      'Other/smoothn');
454
```