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;+
; NAME:
;     DEMO
;
; AUTHORS:
;     Cherrie Kong & Mark B. Cannell.
;     Dept. Physiology, University of Auckland, New Zealand.
;
; DESCRIPTION:
;     This routine demonstrates how to use the MFDA and accessory routines to detect Ca2+ sparks in experimental datasets.
;     This demonstration is intended for line-scanned laser-scanning confocal images, with x-t oriented to x-y
;     respectively and in TIFF format. Use of other formats require the use of the appropriate read routines.
;     The orientation of images is important as the supplied model is oriented in that direction.
;     A simple normalization of the data is carried out by DEMO_NORMALIZER which simply averages all lines as
;     representative of the background. The user is strongly urged to think about what is appropriate normalization!
;     In this demonstration the data is broken into 2048-line chunks. This is not needed for the MFDA per se but simply
;     because it is likely the background will change after so many line scans. If the background changes significantly,
;     a warning is printed -please do not simply ignore this warning -something was happening during the experiment...
;     (Normalization provided by the user usually involves selecting a non-spark region to be used to create the average array
;     required and users can supply their own normalized datasets and models in the MFDA by directly using the FINDOBJECT
;     routine below.) This routine WILL NOT re-enter with a revised model and repeat the fit
;     -it's only a demonstration of the calling procedures!
;
; CALLING SEQUENCE:
;     DEMO [,NORMDAT,MODEL,RESULTS,REFINED]
;
; INPUTS:
;     The user will be prompted for filename and model parameters.
;
; OUTPUTS:
;     All outputs are optional and allow the user to use these elements as required:
;
;         NORMDAT:      normalized data array
;         MODEL:        original model that the user created
;         RESULTS: array holding results of the MFDA. Each line gives 3 values:
;             [0]: p-value associated with detected location
;             [1]: x-ordinate of detected location
;             [2]: y-ordinate of detected location
;         REFINED: refined model based on curve-fitting to an average of the detected events
;
; COPYRIGHT:The IDL codes and algorithms in this program are to be used for
;           non-commerical purposes only. For commerical use, contact the authors.
;-
FORWARD_FUNCTION DEMO_READER,DEMO_CELLNONCELL,KERNELSIZE,DEMO_NORMALIZER
PRO DEMO,normdat,model,results,refined
!EXCEPT = 0      ;kill a bug in Windows implementation of IDL

;INTRODUCTION
PRINT,''
PRINT,':::MFDA DEMONSTRATION:::'
PRINT,''
PRINT,'This is a demonstration of the Matched Filter Detection Algorithm (MFDA) and is intended for '
PRINT,'line-scanned laser-scanning confocal images (x-t) in TIFF format. x-t orientation must '
PRINT,'correspond with x-y, respectively, as per standard definition. '
PRINT,'Proceed? Y/N ' & proceed = STRUPCASE(GET_KBRD()) ;ask user if they wish to proceed

IF (proceed EQ 'Y') THEN BEGIN
;READ IN DATA & DISPLAY
PRINT,'' & PRINT,':::STEP 1: READ IN DATA... '

filename = ''      ;data file
WHILE (filename EQ '') DO BEGIN    ;query user until appropriate filepath and name entered
    PRINT,'Please type in full TIFF filepath and name  (e.g. C:\images\data.tif)'
    READ,filename
    query = QUERY_TIFF(filename)      ;check file exists and is in TIFF format
    IF (query EQ 0) THEN BEGIN      ;if query unsuccessful
        filename = ''
        PRINT,'Bad filename. Please try again.'
    ENDIF ELSE BEGIN
        data = READ_TIFF(filename)      ;read in TIFF
        szda = SIZE(data);find size of DATA
        IF (szda[0] NE 2) THEN BEGIN   ;must be 2D array

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        PRINT,'Data is not a 2D array. Do you want to read another file?' & proceed = STRUPCASE(GET_KBRD())
        IF (proceed EQ 'Y') THEN filename = '' ELSE RETURN
    ENDIF ELSE BEGIN
        IF (szda[4] LT 10000) THEN BEGIN ;DATA must have > 10000 pixels
            PRINT,'Dataset is too small. Do you want to try again?' & proceed = STRUPCASE(GET_KBRD())
            IF (proceed EQ 'Y') THEN filename = '' ELSE RETURN
        ENDIF
    ENDELSE
    ENDELSE
ENDWHILE

IF (szda[1] LT 256) THEN zoom = 256/szda[1] ELSE zoom = 1 ;zoom up to show pixels
SLIDE_IMAGE,BYTSCl(REBIN(data,szda[1]*zoom,szda[2]*zoom,/SAMPLE)),XVISIBLE=256,YVISIBLE=256,TITLE='Data' ;display DATA in window

;OR PARAMETERS AND CREATE MODEL
PRINT,'' & PRINT,':::STEP 2: CREATE A MODEL...'

peak = 1. & PRINT,'What is its peak change in fluorescence (dF; e.g. 1)? ' & READ,peak
bkgd = 1. & PRINT,'What is its basal fluorescence (F0; e.g. 1.0)? ' & READ,bkgd
fwhm = 5. & PRINT,'What is its FWHM (e.g. 5 px)? ' & READ,fwhm
rise = 1.8 & PRINT,'What is its rise time (e.g. 1.8 px)? ' & READ,rise
decay = 28. & PRINT,'What is its decay time (e.g. 28 px)? ' & READ,decay

model = FLTARR(256,256) ;create MODEL array
params = [peak,fwhm,rise,decay,128,bkgd] ;model parameters with centered spark
MAKESPARK_FUNC,model,params ;create MODEL spark
WINDOW,xsize=256*zoom,ysize=256*zoom,title='Model' ;open new window

TVSCL,REBIN(model,256*zoom,256*zoom,/SAMPLE) ;display MODEL in new window with same zoom as DATA

;FOR MODEL OBJECT IN DATASET USING THE MFDA
PRINT,'' & PRINT,':::STEP 3: EMPLOY MATCHED FILTER DETECTION ALGORITHM (sigp = 0.001)...'
sigp = 0.001 ;assign value of SIGP
;if SZDA[2]>1024 lines, then DATA will be segmented for analysis
settings = KERNELSIZE(data,model) ;find overlap region of segments based on border region required for FINDOBJECT
y_bord = settings[1] + settings[3];overlap region stored in Y_BORD
normdat = FLTARR(szda[1],szda[2]) ;empty array to store normalized segments in DATA coordinates (for viewing purposes)

nseg = (szda[2] / 1024) > 1
print,nseg
linestodo = szda[2] / nseg

segment=0,nseg-1 DO BEGIN
    endl = (segment+1)*linestodo
    IF (segment GT 0) THEN start = segment * linestodo - 2 * y_bord ELSE start = 0
    datn = data[*,start:endl-1]
    datn = DEMO_NORMALIZER(datn) ;normalize segment
    szdatn = SIZE(datn) ;size of segment
    normdat[*,start:endl-1] = datn ;fill NORMDAT

;MFDA DETECTION
FINDOBJECT,datn,model,sigp,resultsn,detectn,settings ;MFDA
IF (segment EQ 0) THEN BEGIN ;create/fill detect and results arrays for first segment
    print,detectn
    detected = detectn
    results = resultsn
ENDIF ELSE BEGIN ;for second plus segments
    resultsn[2,*] += start ;adjust detected y-ordinates to full data grid from segment grid
    results = [[results],[resultsn]]
    detected += detectn
ENDELSE

;RESULTS
PRINT,STRING(13B),'Total number of detected events: ',detected
PRINT,'Print full results? Y/N ' & proceed = STRUPCASE(GET_KBRD())
IF (proceed EQ 'Y') THEN PRINT,' P-value | X-ordinate | Y-ordinate ',STRING(13B),results
PRINT,'Show detected locations on image? Y/N ' & proceed = STRUPCASE(GET_KBRD())
IF (proceed EQ 'Y') THEN BEGIN
    vrtbd = settings[3]-1 & hrtd = settings[2]-1 ;create array that shows border region of no detection
    fdata = normdat ;copy of normdat
    fdata[vrtbd-1:hrtd,vrtbd:szda[2]-vrtbd] = 0 ;left border

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fdata[szda[1]-hrtbd-1:szda[1]-hrtbd,vrtbd:szda[2]-vrtbd] = 0 ;right border
fdata[hrtbd:szda[1]-hrtbd,vrtbd-1:vrtbd] = 0 ;top border
fdata[hrtbd:szda[1]-hrtbd,szda[2]-vrtbd-1:szda[2]-vrtbd] = 0 ;bottom border
SLIDE_IMAGE,BYTSCL(fdata),SLIDE_WINDOW=resim,XVISIBLE=256,YVISIBLE=256,TITLE='MFDA: Detected Locations'
WSET,resim ;display NORMDAT with detected events
FOR i=0,detected-1 DO ARROW,results[1,i],results[2,i]-10,results[1,i],results[2,i],THICK=2
ENDIF

;ANALYZE RESULTS BY FITTING AVERAGE
IF (detected GT 0) THEN BEGIN ;only give this option if there are detected events!
    PRINT,'' & PRINT,::::(Optional) Step 4: parameter fitting of the average detected event...
    PRINT,'Fit detected events by least-squared method to normalized data? Y/N ' & proceed = STRUPCASE(GET_KBRD())
ENDIF

IF (proceed EQ 'Y') THEN BEGIN
    initp = [params[0],params[1],params[3]] ;initial parameters for fit based on parameters given for the search object (model)
    ANALYZESPART,normdat,results,detected,initp,parmfit,refined,chi ;least-squares fit routine
    IF (chi EQ 0) THEN BEGIN ;if fit was successful then print parameters
        PRINT,'Least-squares fit was successful. The fitted parameters are: '
        PRINT,'dF: ',parmfit[0]
        PRINT,'F0 (basal fluorescence): ',parmfit[6]
        PRINT,'FWHM: ',parmfit[1],' px'
        PRINT,'rise time constant: ',parmfit[2],' px'
        PRINT,'decay time constant: ',parmfit[3],' px'
        WINDOW,xs=512,ys=512,title='Refined Model'
        TVSCL,refined[(szda[1]/2-256)>0:(szda[1]/2+255)<(szda[1]-1),(szda[2]/2-256)>0:(szda[2]/2+255)<(szda[2]-1)]
    ENDIF ELSE PRINT,'Least-squares fit was unsuccessful. Try changing the initial parameters.' ;if fit not successful
ENDIF
ENDIF
ENDIF ELSE PRINT,'No selected: MFDA demonstration not initiated.'
PRINT,''
PRINT,::::END OF MFDA DEMONSTRATION:::'
END

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;+
; NAME:      DEMO_NORMALIZER
; PURPOSE: This routine is used by DEMO and aids the user in normalizing a line-scanned
;
;           laser-scanning confocal image prior      to its use in the MFDA. It subtracts the mean of the data and then
;           scales the data. For many experiments this is not the best estimate of background or way to normalize the
;           data, the user should modify this routine as appropriate.
; INPUT: data:          2D data array
; OUTPUT:  data:          normalized 2D data array
; COPYRIGHT: The IDL6.3 codes and algorithms in this program are to be used for
;           non-commercial purposes only. For commercial use, contact the authors.
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FUNCTION DEMO_NORMALIZER,data
    szda = SIZE(data)
    normdat = data
    offset = MIN(data) - 1.0
    normdat = data - FLOAT(offset) ;estimate offset of image
                                    ;remove offset
    avgline = REBIN(normdat,szda[1],1)
    avg = FLOAT(REBIN(avgline,szda[1],szda[2])) ;average of all lines
    normdat = normdat/FLOAT(avg) ;resize average array to data size for division
                                ;normalize
    RETURN,normdat
END
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;+
; NAME: FINDOBJECT
; PURPOSE: This routine computes the MFDA: it detects objects against a model object defined by
;           the user. It returns a -log(P-value) map, P-values and coordinates
;           of local P-value minima for detected Ca2+ spark events.
; INPUTS:  data:   2D array containing search space, usually normalized data
;           model:  object to be detected, usually a synthetic Ca2+ spark
;           sigp:   significant P-value level
; OUTPUTS: results: array holding output of MFDA, with length of second dimension equal to number
;           of detected events (unless no sparks are detected)
;
;           [0]: pvalue
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;
; [1]: x-ordinate
; [2]: y-ordinate
; detected:number of detected events
; settings:settings calculated for MFDA
; [0]: shar[0]
; [1]: shar[1]
; [2]: area[0]
; [3]: area[1]
; COPYRIGHT:The IDL6.3 codes and algorithms in this program are to be used for
; non-commerical purposes only. For commercial use, contact the authors.
;-

FORWARD_FUNCTION FFTCORREL,RSTOP_CALC,CORREL_FIND,STATMAP,NEXTSP,KERNELSIZE
PRO FINDOBJECT,data,model,sigp,results,detected,settings
!EXCEPT = 0          ;kill a bug in Windows implementation of IDL
ON_ERROR,1

;NON-STATIONARY BEHAVIOR CHECK & WARNING
DEMO_NONSTATWARNING,data

;CREATE INTERMEDIATE & FINAL RESULTS ARRAYS FOR MFDA
szda = SIZE(data)           ;dimensions of DATA
smo = SIZE(model);dimensions of MODEL
results = FLTARR(3)

;FFT CORRELATION & FIND RSTOP (GIVEN USER-DEFINED SIGP)
correl = FFTCORREL(data,model)      ;initial cross-correlation to find
rstop = RSTOP_CALC(data,model,correl) ;calculate rstop

;FIND APPROPRIATE KERNEL SIZE
settings = KERNELSIZE(data,model)
shar = settings[0:1]
area = settings[2:3]

;SPEARMAN RANK CORRELATION
detected = 0      ;initiate detection count
WHILE (MAX(correl) GT rstop) DO BEGIN          ;begin detection
    pre = CORREL_FIND(correl,area,shar)        ;find tentative spark locations
    res = STATMAP(data,model,area,shar,sigp,pre) ;non-parametric correlation
    IF (res[0] LE sigp) THEN BEGIN              ;if P-value < SIGP
        IF (detected EQ 0) THEN results = res ELSE results = [[results],[res]] ;fill results array
        detected += 1      ;increase detection count
    ENDIF          ;empty array for output in stand-alone program
    correl = NEXTSP(correl,model,pre,area)      ;remove detected event from FFT correlation array
ENDWHILE
END

;+
; NAME:          DEMO_NONSTATWARNING
; PURPOSE:       This routine is used by DEMO and warns the user of
;                 non-stationary behavior that may affect MFDA performance. The routine averages
;                 line-scan data in the x-dimension to produce the time-dependent average signal.
;                 This signal is then reversed in time and compared to itself and R computed. A cut-off
;                 of 0.2 is used to detect weak trends in the data. YMMV.
; INPUT: data    : 2D data array
; OUTPUT: status :      0 = non-stationary behavior not detected
;                  1 = non-stationary behavior detected and warning is printed
; COPYRIGHT:The IDL6.3 codes and algorithms in this program are to be used for
; non-commerical purposes only. For commercial use, contact the authors.
;-

PRO DEMO_NONSTATWARNING,data,status
szda = SIZE(data)
timeavg = REFORM(REBIN(FLOAT(data),1,szda[2]))
result = CORRELATE(timeavg,REVERSE(timeavg))
IF (ABS(result) GT 0.2) THEN BEGIN
    PRINT,'Warning: Non-stationary behavior detected!'
    status = 1
ENDIF ELSE status = 0
END

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;+
; NAME: FFTCORREL
; PURPOSE: This routine correlates two datasets by FFT
; INPUTS:    imA:    data to be convolved
;           imB:    second dataset or 'PSF' (centred and enlarged)
; KEYWORDS: AUTO:    computes auto-correlation of A
; REFERENCE: This IDL6.3 routine has been modified from CONVOLVE, originally written by
;           Frank Varosi, NASA/GSFC 1992.
;-

FUNCTION FFTCORREL,imA,imB,AUTO=auto
  A = imA - MEAN(imA)          ;mean subtraction to relate to Pearson's
  sza = SIZE(A)                ;dimensions of A
  sc = sza/2
  na = N_ELEMENTS(A)
  aFT = FFT(A,-1) ;FFT of A
  IF KEYWORD_SET(auto) THEN BEGIN
    acor = SHIFT(na*REAL_PART(FFT(aFT*CONJ(aFT),1)),sc[1],sc[2]) ;autocorrelation of A
    normsiga = (na-1) * VARIANCE(A)
    acor = acor / FLOAT(normsiga)          ;normalization to relate to Pearson's
    RETURN, acor
  ENDIF ELSE BEGIN
    szb = SIZE(imB)
    B = imB - MEAN(imB)          ;mean subtraction to relate to Pearson's
    psf = (sc - szb/2) > 0      ;center B in new array
    s = (szb/2 - sc) > 0        ;scaling of B
    s2 = (s + sza-1) < (szb-1)
    bFT = CONJ(A)*0
    bFT[psf[1],psf[2]] = B[s[1]:s2[1],s[2]:s2[2]]
    bFT = FFT(bFT,-1,/OVERWRITE)
    cor = na * REAL_PART(FFT(aFT * CONJ(bFT),1)) ;cross-correlation
    sc = sc + (sza MOD 2)
    cor = SHIFT(cor,sc[1],sc[2]) ;return to correct position
    normsig = (na-1) * STDEV(A) * STDEV(B)
    cor = cor / FLOAT(normsig)    ;normalization to relate to Pearson's
    RETURN,cor
  ENDELSE
END

```

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;+
; NAME: RSTOP_CALC
; PURPOSE: This routine calculates the appropriate value of RSTOP by correlating scrambled data with
;           the model to find an upper-bound for correlation values obtained with noise of the same
;           energy as the normalized data itself.
; INPUTS:  data: 2D array holding dataset
;           model: 2D array holding model object
; OUTPUTS: correl: 2D array holding the correlation between data and model (given by FFTCORREL)
; COPYRIGHT: rstop: the appropriate value for RSTOP
;           The IDL6.3 codes and algorithms in this program are to be used for
;           non-commercial purposes only. For commercial use, contact the authors.
;-


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FUNCTION RSTOP_CALC,data,model,correl
  szda = SIZE(data);find size of data array
  noise_est = REFORM(data(SORT(RANDOMU(seed,szda[4]))),szda[1],szda[2])
  cor_noise = FFTCORREL(noise_est,model) ;cross-correlate noise with model
  RETURN,MEAN(cor_noise) + 6.0 * STDEV(cor_noise) ;six standard deviations from mean correlation
END

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;+
; NAME: KERNELSIZE
; PURPOSE: This routine finds the appropriate kernel size to use in STATMAP
; INPUTS:  data: 2D array holding dataset
;           model: 2D array holding model object
; OUTPUTS: shar: half of shift area for centre of model object
;           [0]: x-dimension
;           [1]: y-dimension
;
```

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; area:      2-element array defining kernel size for Spearman Rank Correlation
;           [0]: half WINA in x-dimension
;           [1]: half WINA in y-dimension
; COPYRIGHT:The IDL6.3 codes and algorithms in this program are to be used for
; non-commerical purposes only. For commerical use, contact the authors.
;-

FUNCTION KERNELSIZE,data,model
  szda = SIZE(data);find size of data
  smo = SIZE(model);find size of model
  shar = [5,5]          ;half shift area to test for most likely spark location
  peakloc = ARRAY_INDICES(model,WHERE(model EQ MAX(model))) ;find location of peak
  halfflu = ((MAX(model)-MIN(model))/2.+MIN(model)) ;half-max fluorescence value
  fwhm = N_ELEMENTS(WHERE(model[*],peakloc[1] GT halfflu));find FWHM
  fdhm = N_ELEMENTS(WHERE(model[peakloc[0],*] GT halfflu));find full-duration half maximum (FDHM)
  area = [1.5*fwhm,1.5*fdhm]           ;kernel size is 3 times FWHM in x and 3 times FWHD in y
  ulimx = ROUND((szda[1] - 2 * shar[0]) / 8 - 0.5);kernel cannot be larger than one quarter data size (minus shift region)
  ulimy = ROUND((szda[2] - 2 * shar[1]) / 8 - 0.5)
  llimx = 5           ;kernel should not be smaller than that which gives enough pixels for
  llimy = 5           ;Student's t distribution approximation (Spearman Rank Correlation test)
  IF (area[0] GE ulimx) THEN area[0] = ulimx - 1
  IF (area[0] LE llimx) THEN area[0] = llimx
  IF (area[1] GE ulimy) THEN area[1] = ulimy - 1
  IF (area[1] LE llimy) THEN area[1] = llimx
  RETURN,[shar,area]
END

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;+
; NAME:      CORREL_FIND
; PURPOSE:   This routine finds the first maximum cross-correlation coefficient and
;             returns its x,y coordinates on the original data grid
; INPUTS:    correl: 2D array holding R of data cross-correlated with model
;             area: defines kernel size for Spearman Rank Correlation
;                   [0]: half WINA in x-dimension
;                   [1]: half WINA in y-dimension
;             shar: half of shift area for centre of model object
;                   [0]: x-dimension
;                   [1]: y-dimension
; OUTPUTS:   pre:     tentative location for detected Ca2+ spark
;                   [0]: tentative x-ordinate
;                   [1]: tentative y-ordinate
; COPYRIGHT:The IDL6.3 codes and algorithms in this program are to be used for
; non-commerical purposes only. For commerical use, contact the authors.
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FUNCTION CORREL_FIND,correl,area,shar
  sco = SIZE(correl) ;dimensions of CORREL
  mask = BYTARR(sco[1],sco[2]) ;create mask to allow edge-space for STATMAP
  mask[shar[0]+area[0]:sco[1]-shar[0]-area[0]-1,shar[1]+area[1]:sco[2]-shar[1]-area[1]-1] = 1
  correl = mask * correl ;apply mask
  RETURN,ARRAY_INDICES(correl,WHERE(correl EQ MAX(correl))) ;find first maximum R value location
END

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;+
; NAME:  STATMAP
; PURPOSE: This routine cross-correlates regions of interest with a model object
;           using the Spearman Rho Rank method. It returns a map showing the
;           - log ( P-value ) of the regions and the final locations of detected events.
; INPUTS:  data: 2D array containing search space, usually normalized data
;           model: 2D array containing model object, usually a synthetic CLSM spark
;           area: 2-element array defining kernel size for Spearman Rank Correlation
;                   [0]: half WINA in x-dimension
;                   [1]: half WINA in y-dimension
;           shar: half of shift area for centre of model object
;                   [0]: x-dimension
;                   [1]: y-dimension
;           pre:   tentative location for detected Ca2+ spark
;                   [0]: tentative x-ordinate
;                   [1]: tentative y-ordinate
;
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; OUTPUTS:    xy:          final location for detected Ca2+ spark
;                      [0]: final x-ordinate
;                      [1]: final y-ordinate
; COPYRIGHT:The IDL6.3 codes and algorithms in this program are to be used for
; non-commerical purposes only. For commercial use, contact the authors.
;-

FORWARD_FUNCTION MINP
FUNCTION STATMAP,data,model,area,shar,sigp,pre
  ssp = SIZE(model);define model kernel, ARC
  arc = model[ssp[1]/2-area[0]:ssp[1]/2+area[0]-1,ssp[2]/2-area[1]:ssp[2]/2+area[1]-1]
  ind = WHERE(arc GT MIN(model))      ;use only pixels with intensity above basal fluorescence
  arc = arc[ind]
  pmap = FILTARR(2*(area[0]+shar[0]),2*(area[1]+shar[1]),2);array to hold individual rho- and P-value maps
  pmap[*,*,1] = 1. ;default P-value in map is 1.0
FOR xim = pre[0]-shar[0],pre[0]+shar[0]-1 DO BEGIN ;shift in X over area (defined by SHAR) around PRE
  FOR yim = pre[1]-shar[1],pre[1]+shar[1]-1 DO BEGIN ;shift in Y over area (defined by SHAR) around PRE
    aoi = data[xim-area[0]:xim+area[0]-1,yim-area[1]:yim+area[1]-1];define AOI in DATA
    aoi = aoi[ind];use IND pixels only
    pmap[xim-pre[0]+shar[0]+area[0],yim-pre[1]+shar[1]+area[1],*] = R_CORRELATE(aoi,arc) ;Spearman Rank Correlation
  ENDFOR
ENDFOR
RETURN, MINP(area,shar,pmap,pre) ;find XY by locating minimum P-value
END

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;+
; NAME: MINP
; PURPOSE: This routine returns the coordinates of the center of
;           the detected object as determined by the Spearman Rank Rho Correlation,
;           used in STATMAP.
; INPUTS:   area:    2-element array defining kernel size for Spearman Rank Correlation
;                      [0]: half WINA in x-dimension
;                      [1]: half WINA in y-dimension
;   shar:    half of shift area for centre of model object
;                      [0]: x-dimension
;                      [1]: y-dimension
;   pmap:   P-value map for individual object
;   pre:     tentative location for detected Ca2+ spark
;                      [0]: tentative x-ordinate
;                      [1]: tentative y-ordinate
; OUTPUT:   res:    [pvalue,x,y], where
;                      pvalue = minimum P-value in pmap
;                      x = final x-ordinate
;                      y = final y-ordinate
; COPYRIGHT:The IDL6.3 codes and algorithms in this program are to be used for
; non-commerical purposes only. For commercial use, contact the authors.
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FUNCTION MINP,area,shar,pmap,pre
  pvalue = MIN(pmap[*,*,1]);minimum value in PMAP[*,*,1] is the pvalue
  locxy = ARRAY_INDICES(pmap[*,*,1],WHERE(pmap[*,*,1] EQ pvalue));locate pvalue
IF (N_ELEMENTS(locxy) GT 2) THEN BEGIN ;if more than one pixel contains minimum take center
  locxy[0,0] = MEAN(locxy[0,*])
  locxy[1,0] = MEAN(locxy[1,*])
ENDIF
locxy[0,0] = pre[0] - area[0] - shar[0] + locxy[0,0];convert coordinates from PMAP grid to DATA grid
locxy[1,0] = pre[1] - area[1] - shar[1] + locxy[1,0]
RETURN,[pvalue,locxy[*,0]] ;return pvalue and coordinates
END

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;+
; NAME: NEXTSP
; PURPOSE: This routine removes a found object in the initial correlation
;           array to prepare it for subsequent object searches. Removal uses
;           the model object auto-correlation.
; INPUTS:   correl: initial 2D FFT correlation array
;   model:   2D array holding model object
;   xy:       initial location for detected Ca2+ spark

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;+                                     [0]: final x-ordinate
;+                                     [1]: final y-ordinate
; area:   2-element array defining kernel size for Spearman Rank Correlation
;+                                     [0]: half WINA in x-dimension
;+                                     [1]: half WINA in y-dimension
; OUTPUTS:    correl: amended 2D FFT correlation array
; COPYRIGHT:The IDL6.3 codes and algorithms in this program are to be used for
;+                           non-commerical purposes only. For commercial use, contact the authors.
;-

FUNCTION NEXTSP,correl,model,xy,area
  ssp = SIZE(model);dimensions of MODEL
  sco = SIZE(correl) ;dimensions of CORREL
  spmask = SMOOTH(FFTCORREL(model,/AUTO),2*area[0],/EDGE) > 0
;note autocorrelate model as term for subtraction and blur to account for differences in detected objects
  spmask = spmask * MAX(correl) / MAX(spmask) ;scale amplitude to cross-correlation
IF (sco[1] LT ssp[1]) THEN spmask = spmask[ssp[1]/2-sco[1]/2:ssp[1]/2-sco[1]/2+sco[1]-1,*] ;if x-dimension of model is > data
IF (sco[2] LT ssp[2]) THEN spmask = spmask[*,:ssp[2]/2-sco[2]/2:ssp[2]/2-sco[2]/2+sco[2]-1] ;if y-dimension of model is > data
  mask = FILTARR(sco[1],sco[2]);create MASK array to size of CORREL
  ssp = SIZE(spmask);dimensions of SPMASK
  mask[sco[1]/2-ssp[1]/2:sco[1]/2-ssp[1]/2+sco[1]-1,sco[2]/2-ssp[2]/2:sco[2]/2+ssp[2]-ssp[2]/2-1]=spmask ;fill MASK
  mask = SHIFT(mask,xy[0]-sco[1]/2-1,xy[1]-sco[2]/2-1);position correctly
  RETURN,correl - mask ;remove detected event
END

```

```

;+
; NAME: MAKESPARK_FUNC
; PURPOSE: This routine generates a synthetic CLSM Ca2+ spark in a 2D array
; INPUT: arr: 2D floating point array to hold Ca2+ spark
; p: 7-element floating point array holding initial spark parameters
;+                                     [0]: df
;+                                     [1]: FWHM (px)
;+                                     [2]: tau rise (px)
;+                                     [3]: tau decay (px)
;+                                     [4]: model center (x-ordinate), <= array x-dimensions
;+                                     [5]: model center (y-ordinate), <= array y-dimensions
;+                                     [6]: F0
; OUTPUT: arr: 2D floating point array holding Ca2+ spark
; start: (optional) start time (px) of spark time course
; tpeak: (optional) time to peak (px)
; thalf: (optional) half-time to decay (px)
; COPYRIGHT:The IDL6.3 codes and algorithms in this program are to be used for
;+                           non-commerical purposes only. For commercial use, contact the authors.
;-

```

```

PRO MAKESPARK_FUNC,arr,p,start,tpeak,thalf
!EXCEPT=0
sz = SIZE(arr) ;dimensions of ARR
y = FINDGEN(sz[2]) - P[5];generate time axis with shift
;amplitude(t) function
  ampt = EXP(-EXP(-y/P[2])-y/P[3]);sigmoidal rising phase and exponential decay phase
  ampt = ampt * P[0] / EXP(P[2]*( ALOG(P[2]/P[3])-1.)/P[3]);scale amplitude
  start = WHERE(ampt GE 1E-20) & start = start[0];find start of rise
  maxp = -P[2]*ALOG(P[2]/P[3]) + P[5];find location of peak
  endhalf = WHERE(ampt GE (P[0]/2.));find time of half decay
  tpeak = maxp - start;find time to peak
  thalf = endhalf[N_ELEMENTS(endhalf)-1] - maxp;find time of half decay
;sigma(t) function
  sigma = P[1] / 2.355;convert FWHM to sigma
  sigmat = ((y-start+P[5])^2+0.0001)^0.25;sigma increases with positive and negative square root of time
  sigmat = sigmat * sigma / sigmat[maxp];scale sigma at peak
  ;fluorescence(x,t) function
  FOR i =0,sz[1]-1 DO arr[i,*] = ampt * EXP(-(i-P[4])^2/(2.*sigmat^2)) + P[6];create spark
END

```

```

;+
; NAME: MAKESPARK
; PURPOSE: This routine utilizes the MAKESPARK_FUNC routine to generate a
;+                           synthetic CLSM Ca2+ spark. This adapts MAKESPARK_FUNC for its use in
;
```

```

;
; CURVEFIT (a 1D non-linear least squares fitting routine within the IDL
; package), also providing analytical partial derivatives.
;
; INPUT:      x:           1D independent variable, where
;             [0]: size of data array in x
;             [1]: size of data array in y
;             [2:x[0]+1]: independent variable x
;             [x[0]+2:*]: independent variable y
;             p:           7-element floating point array holding initial spark parameters
;             [0]: df
;             [1]: FWHM (px)
;             [2]: tau rise (px)
;             [3]: tau decay (px)
;             [4]: model center (x-ordinate)
;             [5]: model center (y-ordinate)
;             [6]: F0
; OUTPUT:     z:           1D data array
;             pder: (optional) 2D array of partial derivatives
; COPYRIGHT:The IDL6.3 codes and algorithms in this program are to be used for
; non-commerical purposes only. For commerical use, contact the authors.
;-

PRO MAKESPARK,x,p,z,pder
  nx = LONG(x[0]) & ny = LONG(x[1])
  n = nx * ny
  z = FLTARR(nx,ny)
  ;data size in x- and y-dimensions
  ;no. elements in dataset
  ;create empty array for spark

MAKESPARK_FUNC,z,p,start
;call spark routine
z = REFORM(z,n,/OVERWRITE)
;2D to 1D spark
;partial derivatives
IF (N_PARAMS() GE 4) THEN BEGIN
  xp = (x[2:nx+1]) # REPLICATE(1,ny)
  ;x independent variables
  yp = REPLICATE(1,nx) # (x[nx+2:])
  ;y independent variables
  a = P[0] / EXP( (P[2]/P[3]) * ( ALOG(P[2]/P[3])-1. ) )
  b = EXP(-(yp - P[5])/P[2])
  c = (yp-P[5])/P[3]
  d1 = P[2] * ALOG(P[3]/P[2]) + P[5] - start
  d2 = (xp-P[4])/P[1]
  d = 0.5 * d2^2 * (2.355*(d1^2+0.0001)^0.25/((yp-start)^2+0.0001)^0.25)^2
  e = EXP(-b-c-d)
  pder[*,*] = a*e / P[0]
  pder[*,*] = a*e*2.*d/P[1]
  pder[*,*] = -a*e*((ALOG(P[2]/P[3])-1.)/P[3]+(1/P[3]))+((yp-P[5])*b/P[2]^2+0.5*((d2*2.355)^2*d1* $ 
  (ALOG(P[3]/P[2])-1.)/((d1^2+0.0001)^0.5*((yp-start)^2+0.0001)^0.5) ) )
  pder[*,*] = -a*e*((-P[2]*(ALOG(P[2]/P[3])-1.)/P[3]^2 - P[2]/P[3]^2)-(c/P[3]-0.5*(d2*2.355)^2*d1* $ 
  P[2]/(P[3]*((d1^2+0.0001)^0.5*((yp-start)^2+0.0001)^0.5)))
  pder[*,*] = a*e*(xp-P[4])/((P[1]*((yp-start)^2+0.0001)^0.25/(2.355*((P[2]*ALOG(P[3]/P[2])+P[5]-start)^2+0.0001)^0.25))^2
  pder[*,*] = a*((-b/P[2])+1/P[3]-d)*e
  pder[*,*] = 1.
ENDIF
END

;+
; NAME:      ANALYZESPARK
; PURPOSE:   This routine fits parameters of Ca2+ spark using defined model/function by calling CURVEFIT and estimating
;            initial parameters with those used for finding sparks in FINDOBJECT and with locations
;            output from FINDOBJECT.
; INPUT: data:    data array
;             mfdares: array holding output of MFDA, with length of second dimension equal to number
;                       of detected events (unless no sparks are detected)
;             [0]: pvalue
;             [1]: x-ordinate
;             [2]: y-ordinate
;             p:           initial parameters for fit
;             [0]: df
;             [1]: FWHM at peak (px)
;             [2]: decay time constant (px)
; OUTPUT:    parmfilt: fitted parameters of a spark
;             [0]: df
;             [1]: FWHM at peak (px)
;             [2]: rise time constant (px)

```

```

;
; [3]: decay time constant (px)
; [4]: spark x-ordinate
; [5]: spark y-ordinate
; [6]: F0
;
; refined: optional output of 2D refined model array with fitted parameters
; chi:      status of curvefit, 0=success, 1=failed as Chi-square increased without bounds, 2=failed to converge in max. iterations
; COPYRIGHT: The IDL6.3 codes and algorithms in this program are to be used for non-commercial purposes only. For commercial use, contact the authors.
;-

PRO ANALYZESPARK,data,mfdares,detected,p,parmfilt,refined,chi

;ORGANISE DATA FOR CURVE-FITTING
szda = SIZE(data)          ;DATA dimensions
p = ABS(p)                 ;ensure no negative values
nx = (4. * FIX(P[1])) < (szda[2]/4 - 1)    ;fit includes data within 4 times expected FWHM
ny = (4. * FIX(P[2]) + 20) < (szda[2]/4 - 1)   ;fit includes data 20 px prior to detected location to 4 times expected half decay
n = nx * ny                ;size of dataset to be fitted
savg = FLTARR(nx,ny)
count = 0
FOR i=0,detected-1 DO BEGIN
    temp_x = mfdares[1,i] & temp_y = mfdares[2,i]
    IF (temp_x GT nx) AND (temp_y GT ny) AND (temp_x LT (szda[1]-nx)) AND (temp_y LT (szda[2]-ny)) THEN BEGIN ;events too close to edge
        temp = SHIFT(data,szda[1]/2-mfdares[1,i],20-mfdares[2,i]);centering detected objects in savg
        IF (count EQ 0) THEN savg = temp[szda[1]/2-nx/2:szda[1]/2+nx-nx/2-1,0:ny-1] ELSE $
            savg = [[[savg],[[temp[szda[1]/2-nx/2:szda[1]/2+nx-nx/2-1,0:ny-1]]]] ;cut out region of interest as defined by nx and ny
        count += 1
    ENDIF
ENDFOR
savg = REBIN(savg,nx,ny,1)      ;signal average all detected objects

;CURVE-FITTING
x = FINDGEN(nx) & y = FINDGEN(ny)          ;generate independent variables
parm = [P[0],P[1],0.1*P[2],P[2],nx/2,20,MEAN(data)] ;initial parameter estimates
result = CURVEFIT([nx,ny,x,y],REFORM(savg,n),weight,parm,FUNCTION_NAME="MAKESPARK",STATUS=chi) ;fitting with no weighting
parmfilt = [parm[0],parm[1],parm[2],parm[3],szda[1]/2,szda[2]/2,parm[6]]      ;fitted parameters in parmfilt

IF (parm[0] LT 0) OR (parm[1] LT 0) OR (parm[2] LT 0) OR (parm[3] LT 0) OR (parm[4] LT 0) OR (parm[5] LT 0) OR (parm[6] LT 0) THEN BEGIN
    chi = 1                         ;negative numbers as unsuccessful fit
    refined = INTARR(szda[1],szda[2])
ENDIF ELSE BEGIN
;GENERATE REFINED MODEL
    refined = FLTARR(szda[1],szda[2]) ;create array for refined model
    MAKESPARK_FUNC,refined,parmfilt    ;generate model and print time course characteristics
ENDELSE
END

;+
; NAME: MAKEGAUSS2D_FUNC
; PURPOSE: This routine generates a 2D Gaussian in a 2D array
; INPUT:    array: 2D floating point array to hold 2D Gaussian
;           p:       6-element floating point array holding initial 2D Gaussian parameters
;                   [0]: df
;                   [1]: FWHM in x (px)
;                   [2]: FWHM in y (px)
;                   [3]: model center (x-ordinate), =< array x-dimensions
;                   [4]: model center (y-ordinate), =< array y-dimensions
;                   [5]: F0
; OUTPUT:   array: 2D floating point array holding Ca2+ spark
; COPYRIGHT: The IDL6.3 codes and algorithms in this program are to be used for
;           non-commercial purposes only. For commercial use, contact the authors.
;-

PRO MAKEGAUSS2D_FUNC,array,p
sz = SIZE(array)          ;find dimensions of array
nx = LONG(sz[1])          ;x-dimension
ny = LONG(sz[2])          ;y-dimension
MAKEGAUSS2D,[nx,ny,FINDGEN(nx),FINDGEN(ny)],p,z    ;call 2D Gaussian function (requires 1D index)
array = REFORM(z,nx,ny)    ;reform array to 2D
END

```

```

;+
; NAME: MAKEGAUSS2D
; PURPOSE: This routine utilises the MAKEGAUSS2D_FUNC routine to generate a
;           2D Gaussian. This provides analytical partial derivatives and can be used with
;           CURVEFIT (a 1D non-linear least squares fitting routine within the IDL
;           package).
; INPUT:    x:          1D independent variable, where
;           [0]: size of data array in x
;           [1]: size of data array in y
;           [2:x[0]+1]: independent variable x
;           [x[0]+2:*]: independent variable y
;           p:          6-element floating point array holding initial 2D Gaussian parameters
;           [0]: df
;           [1]: FWHM in x (px)
;           [2]: FWHM in y (px)
;           [3]: model center (x-ordinate)
;           [4]: model center (y-ordinate)
;           [5]: F0
; OUTPUT:   z:          1D data array
;           pder: optional 2D array of partial derivatives
; REFERENCE: Modified from IDL6.3 GAUSS2DFIT routine.
;-

PRO    MAKEGAUSS2D,x,p,z,pder
nx = LONG(X[0]) & ny = LONG(X[1])                                ;x- and y-dimensions
xp = (X[2:nx+1]-P[3]) # REPLICATE(1.0/P[1],ny)                ;expand X values
yp = REPLICATE(1.0/P[2],nx) # (x[nx+2:]-P[4])                  ;expand Y values
n = nx * ny                                                       ;total number of elements
u = REFORM(EXP(-0.5 * (xp^2 + yp^2)),n)                      ;make exp() term 1D
z = P[5] + P[0] * u                                            ;partial derivatives
IF (N_PARAMS() GE 4) THEN BEGIN
  s = 0.0 & c = 1.0
  pder = FLTARR(n,N_ELEMENTS(P))
  pder[*,0] = u
  u = P[0] * u
  pder[*,1] = u * xp^2 / P[1]
  pder[*,2] = u * yp^2 / P[2]
  pder[*,3] = u * (c/P[1] * xp + s/P[2] * yp)
  pder[*,4] = u * (-s/P[1] * xp + c/P[2] * yp)
  pder[*,5] = 1.0
ENDIF
END

;+
; NAME: ANALYZEGAUSS
; PURPOSE: This routine fits parameters of a 2D Gaussian using defined model/function by calling CURVEFIT and estimating
;           initial parameters with those used for finding sparks in FINDOBJECT and with locations
;           output from FINDOBJECT.
; INPUT: data:  data array
;           mfdares: array holding output of MFDA, with length of second dimension equal to number
;                     of detected events (unless no sparks are detected)
;           [0]: pvalue
;           [1]: x-ordinate
;           [2]: y-ordinate
;           p:          initial parameters for fit
;           [0]: df
;           [1]: FWHM in x (px)
;           [2]: FWHM in y (px)
; OUTPUT:  parmfat: fitted parameters of a spark
;           [0]: df
;           [1]: FWHM in x (px)
;           [2]: FWHM in y (px)
;           [3]: object x-ordinate
;           [4]: object y-ordinate
;           [5]: F0
;           refined: optional output of 2D refined model array with fitted parameters
;           chi:      status of curvefit, 0=success, 1=failed as Chi-square increased without bounds, 2=failed to converge in max. iterations
; COPYRIGHT: The IDL6.3 codes and algorithms in this program are to be used for non-commercial purposes only. For commercial use, contact the authors.
;-

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PRO ANALYZEGAUSS,data,mfdares,detected,p,parmfit,refined,chi

;ORGANISE DATA FOR CURVE-FITTING
szda = SIZE(data) ;data dimensions
p = ABS(p) ;ensure no negative values
nx = ROUND(6. * P[1] + 0.5) < (szda[1]/4 - 1) ;fit area includes 6 times estimated FWHMx
ny = ROUND(6. * P[2] + 0.5) < (szda[2]/4 - 1);fit area includes 6 times estimated FWHMy
n = nx * ny ;total number of elements
count = 0
savg = FLTARR(nx,ny)
FOR i=0,detected-1 DO BEGIN
    temp_x = mfdares[1,i] & temp_y = mfdares[2,i]
    IF (temp_x GT nx) AND (temp_y GT ny) AND (temp_x LT (szda[1]-nx)) AND (temp_y LT (szda[2]-ny)) THEN BEGIN ;events too close to edge
        temp = SHIFT(data,szda[1]/2-mfdares[1,i],szda[2]/2-mfdares[2,i]);centering detected objects in savg
        IF (count EQ 0) THEN savg = temp[szda[1]/2-nx/2:szda[1]/2+nx-nx/2-1,szda[2]/2-ny/2:szda[2]/2+ny-ny/2-1] ELSE $
            savg = [[savg],[[temp[szda[1]/2-nx/2:szda[1]/2+nx-nx/2-1,szda[2]/2-ny/2:szda[2]/2+ny-ny/2-1]]]]
        count += 1
    ENDIF
ENDFOR
savg = REBIN(savg,nx,ny,1) ;signal average all detected objects

;CURVE-FITTING
x = FINDGEN(nx) & y = FINDGEN(ny) ;generate independent variables
parm = [P[0],P[1],P[2],nx/2,ny/2,MEAN(data)] ;initial parameter estimates
result = CURVEFIT([nx,ny,x,y],REFORM(savg,n),weight,parm,function_name="MAKEGAUSS2D",STATUS=chi) ;fitting with no weighting
parmfit = [parm[0],parm[1],parm[2],szda[1]/2,szda[2]/2,parm[5]] ;fitted parameters in parmfit

IF (parm[0] LT 0) OR (parm[1] LT 0) OR (parm[2] LT 0) OR (parm[3] LT 0) OR (parm[4] LT 0) OR (parm[5] LT 0) THEN BEGIN
    chi = 1 ;negative parameters as unsuccessful fit
    refined = INTARR(szda[1],szda[2])
ENDIF ELSE BEGIN
;GENERATE REFINED MODEL
    refined = FLTARR(szda[1],szda[2]) ;create array for refined model
    MAKEGAUSS2D_FUNC,refined,parmfit ;generate model and print time course characteristics
ENDELSE
END

```