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crose the image display righte it open, the crose the gar righter.
function segmentationautomation(varargin)
% SEGMENTATIONAUTOMATION Automated identification of contact
% zones
8
% Syntax
%
% CONTACTAUTOMATION prompts the user to select a folder with Imaris
% data to process. A new Imaris instance is opened to load and process
% the data.
%
% CONTACTAUTOMATION(strFolder) processes Imaris files in the folder
% indicated by the fully qualified path string strFolder. If the folder
% indicated by the fully qualified path string strionder. If the folder with a indicated by strFolder does not exist, the function errors and exits
% CONTACTAUTOMATION(strEolder imarigID) connects to the Imaria
% CONTACTAUTOMATION(strFolder, imarisID) connects to the Imaris
<pre>% instance specified by imarisID. The imarisID input can be zero or % positive-valued integer. If no running instance of Imaris has an ID</pre>
> nogifive-valued integer of no running ingtance of Imaria had an TD
% equal to imarisID, a new instance will be created with the specified
\$ equal to imarisID, a new instance will be created with the specified $$$ ID.
% equal to imarisID, a new instance will be created with the specified

Parse the optional inputs.

```
segmentationautomationParser = inputParser;
segmentationautomationParser.addOptional('xFolder', '', @(arg)exist('arg', 'di
segmentationautomationParser.addOptional('xImarisID', '', ...
    @(arg)...
    (isnumeric(arg) && rem(arg, 1) == 0) || ...
    (~isnan(str2double(arg)) && rem(str2double(arg), 1) == 0))
parse(segmentationautomationParser, varargin{:})

xFolder = segmentationautomationParser.Results.xFolder;
xImarisID = segmentationautomationParser.Results.xImarisID;
```

Create the figure.

```
desktopPos = get(0, 'MonitorPositions');
quiWidth = 552;
guiHeight = 361;
figPos = [...
    (desktopPos(1, 3) - guiWidth)/2, ...
    (desktopPos(1, 4) - guiHeight)/2, ...
    guiWidth, ...
    guiHeight];
guiSegmentation = figure(...
    'Color', 'k', ...
    'CloseRequestFcn', {@closecontactautomationfcn}, ...
    'MenuBar', 'None', ...
    'Name', 'Contact segmentation', ...
    'NumberTitle', 'Off', ...
    'Position', figPos, ...
    'Resize', 'Off', ...
    'Tag', 'guiSegmentation');
% Create the open and process buttons.
uicontrol(...
    'Background', 'k', ...
    'Callback', {@pushfolderopen, guiSegmentation}, ...
    'FontSize', 12, ...
    'Foreground', 'w', ...
    'Position', [20 324 75 24], ...
    'String', 'Folder', ...
    'Style', 'pushbutton', ...
    'Tag', 'pushOpen', ...
    'TooltipString', 'Open a folder with Imaris data');
uicontrol(...
    'Background', 'k', ...
    'Callback', {@pushfolderprocess, quiSeqmentation}, ...
    'FontSize', 12, ...
    'Foreground', 'w', ...
```

```
'Position', [457 324 75 24], ...
    'String', 'Process', ...
    'Style', 'pushbutton', ...
    'Tag', 'pushProcess', ...
    'TooltipString', 'Process all Imaris files in the folder');
% Create a uicontrol to display the selected folder.
textFolder = uicontrol(...
    'Background', 'k', ...
    'FontSize', 10, ...
    'Foreground', 'w', ...
    'HorizontalAlignment', 'Left', ...
    'Position', [115 320 322 24], ...
    'String', '', ...
    'Style', 'text', ...
    'Tag', 'textFolder');
% Create the display axes.
axesContours = axes(...
    'Units', 'Pixels', ...
    'Color', 'k', ...
    'Parent', guiSegmentation, ...
    'Position', [20 40 512 256], ...
    'XColor', 'k', ...
    'YColor', 'k');
% Setup the status bar.
hStatus = statusbar(guiSegmentation, xFolder);
hStatus.CornerGrip.setVisible(false)
hStatus.ProgressBar.setForeground(java.awt.Color.black)
hStatus.ProgressBar.setString('')
hStatus.ProgressBar.setStringPainted(true)
```

Process

Folder

Store the inputs.

```
setappdata(guiSegmentation, 'xFolder', xFolder)
setappdata(guiSegmentation, 'xImarisID', xImarisID)
```

Nested function to select a folder

```
function pushfolderopen(varargin)

% PUSHOPEN Select a folder with data to process
%
%
```

Get the folder to process.

```
xFolder = getappdata(guiSegmentation, 'xFolder');
xFolder = uigetdir(xFolder);
setappdata(guiSegmentation, 'xFolder', xFolder)
```

Update the folder text box.

```
slashIdxs = strfind(xFolder, filesep);
folderEnd = xFolder(slashIdxs(end - 1) + 1:end);
set(textFolder, 'String', ['...\' folderEnd])
end % pushfolderopen
```

Nested function to process the Imaris files

```
function pushfolderprocess(varargin)

% PUSHFOLDERPROCESS Process the Imaris files in the selected folder.
%
%
```

Get all the Imaris files in the folder.

```
xFolder = getappdata(guiSegmentation, 'xFolder');
xFiles = dir(fullfile(xFolder, '*.ims'));
if isequal(xFolder, 0) || isempty(xFiles)
    return
end % if
fStartTime = zeros(length(xFiles), 1);
```

Connect to Imaris.

```
xImarisID = getappdata(guiSegmentation, 'xImarisID');
if ~isempty(xImarisID)
    xImarisApp = xtconnectimaris(xImarisID);
else
    [xImarisApp, xImarisID] = xtconnectimaris;
    setappdata(guiSegmentation, 'xImarisID', xImarisID)
end % if
```

Setup the progress bar.

```
hStatus.ProgressBar.setVisible(true)
hStatus.ProgressBar.setMaximum(length(xFiles))
completionEstimate = 'N/A';
```

Segment the contacts in all the files.

```
for f = 1:length(xFiles)
```

Record the start time for processing.

```
fStartTime(f) = datenum(datetime('now'));
```

Load the data file into the connected Imaris instance.

```
xImarisApp.FileOpen(fullfile(xFolder, xFiles(f).name), 'reader="Imaris
```

Get the data set properties.

```
[~, fFileName, ~] = fileparts(char(xImarisApp.GetCurrentFileName));
hStatus.setText(['Processing: ' fFileName ' | Estimated finish: ' comp
% Get the data set dimensions.
xDataSet = xImarisApp.GetDataSet;

xSize = xDataSet.GetSizeX;

ySize = xDataSet.GetSizeY;
yMax = xDataSet.GetExtendMaxY;

cSize = xDataSet.GetSizeC;
tSize = xDataSet.GetSizeT;
```

Identify the channels. (1-based indices)

```
channelNames = cell(cSize, 1);
for c = 1:cSize
    channelNames{c} = char(xDataSet.GetChannelName(c - 1));
end % for c

qdotChannel = find(~cellfun(@isempty, ...
    regexp(channelNames, '^(Q(D|d)ot)', 'Match', 'Once')));
contourChannel = find(~cellfun(@isempty, ...
    regexp(channelNames, '(C|c)ontours', 'Match', 'Once')));
```

Use the first contour channel. If there is no contour channel, continue.

```
if length(contourChannel) > 1;
    contourChannel = contourChannel(1);
elseif isempty(contourChannel)
    continue
end % if
```

Get the movie images.

Crop the images to the cell path.

```
contourProjection = sum(uint8(contourImage), 3) > 0;
projectionRgnProps = regionprops(contourProjection, 'BoundingBox');
rectPos = [projectionRgnProps.BoundingBox];

cropXs(1) = max([1, floor(rectPos(1)) - 10]);
cropXs(2) = cropXs(1) + floor(rectPos(3)) + 19;
cropXs(2) = min([xSize, cropXs(2) + (rem(diff(cropXs), 2) - 1)]);

cropYs(1) = max([1, floor(rectPos(2)) - 10]);
cropYs(2) = cropYs(1) + floor(rectPos(4)) + 19;
cropYs(2) = min([ySize, cropYs(2) + (rem(diff(cropYs), 2) - 1)]);

qdotCropped = qdotImage(cropYs(1):cropYs(2), cropXs(1):cropXs(2), :);
% Normalize the images to the intensity range in the cell region.
qdotRange = [min(qdotCropped(:)) max(qdotCropped(:))];
qdotNormalized = (qdotImage - qdotRange(1))/(qdotRange(2) - qdotRange())
```

Expand the contour slightly.

```
contourExpanded = false(size(contourImage));
for t = 1:tSize
    contourDistance = 0.1*bwdist(contourImage(:, :, t));
    contourExpanded(:, :, t) = contourDistance < 0.25;
end % for t</pre>
```

Create a filtering kernel to smooth the data.

```
cKernel = fspecial('gaussian', [7 7], 1);
```

Detect regional minima and contour the holes.

Contacts segmentation results are stored as the output of regionprops.

```
cellContacts = cell(1, tSize);

for t = 1:tSize;
    % If the cell is not bound in the frame, continue.
    tContour = contourExpanded(:, :, t);
    if ~any(tContour(:))
```

```
continue
end % if
% Filter and detect maxima.
tQDot = qdotNormalized(:, :, t);
tRgnMin = imregionalmin(imfilter(tQDot, cKernel, 'Symmetric')) & .
    tContour & ...
    tQDot < median(tQDot(:));
tContacts = regionprops(tRgnMin, 'Area', 'Centroid', 'PixelIdxList
% Dilate and contour the seed points.
for s = 1:length(tContacts)
    % Create the input mask.
    sPhi = false(size(tQDot));
    sPhi(floor(tContacts(s).Centroid(2)), floor(tContacts(s).Centr
    sPhi = imdilate(sPhi, strel('disk', 1));
    % Contour.
    sPhi = activecontour(qdotNormalized(:, :, t), sPhi, 5, 'Chan-V
        'ContractionBias', 0, ...
        'SmoothFactor', 0);
    % Get the region properties.
    sContact = regionprops(sPhi, 'Area', 'Centroid', 'PixelIdxList
    if ~isempty(sContact)
        if length(sContact) == 1 && sContact.Area > 5
            tContacts(s) = deal(sContact);
        else % The region has fragmented into sub-regions. Keep th
            idxPrimaryRqn = find([sContact.Area] == max([sContact.
            sContact = sContact(idxPrimaryRqn);
            if sContact.Area > 5
                tContacts(s) = deal(sContact);
            end % if
        end % if
    end % if
end % for s
% Remove the isolated minima.
tContacts = tContacts([tContacts.Area] > 5);
% Merge overlapping regions. Use 50% overlap as a merge critera.
for s = 1:length(tContacts)
    % Get the segmentation data.
    sProps = tContacts(s);
    % If we've merged this region already, move on.
    if isempty(sProps.Centroid)
        continue
   end % if
    % Create the list of regions to test. Do not search the
    % region itself and do not search regions that have
```

```
% been merged with a lower index region.
pIdxs = 1:length(tContacts);
pIdxs(s) = nan;
pIdxs(arrayfun(@(s)isempty(s.Centroid), tContacts)) = nan;
pIdxs(isnan(pIdxs)) = [];
% Check each target for overlap.
for p = pIdxs
    % Get the overlapping pixels.
    pProps = tContacts(p);
    idxOverlap = ismember(sProps.PixelIdxList, pProps.PixelIdx
    if ~any(idxOverlap(:))
        continue
    else
        % Quantify the overlap.
        oMask = false(size(sPhi));
        oMask(sProps.PixelIdxList(idxOverlap)) = 1;
        oProps = regionprops(oMask, 'Area');
        oArea = sum([oProps.Area]);
        % Determine whether to merge.
        if oArea/sProps.Area >= 0.5 || oArea/pProps.Area >= 0.
            % Create the merged region.
            cMask = false(size(sPhi));
            cMask([sProps.PixelIdxList; pProps.PixelIdxList])
            sProps = regionprops(cMask, 'Area', 'Centroid', 'P
            % Delete the centroid data for the region
            % we merged in, so we skip over it when we
            % reach it in the loop.
            if s < p
                % The p-index region is now merged into the
                % s-index region. Remove the centroid value fo
                % and update s.
                tContacts(p).Centroid = [];
                tContacts(s) = sProps;
            else
                % The s-index region is now merged into
                % the p-index region. Remove the
                % centroid value for s and update p.
                % Break out of the p loop, as s is now
                % invalid.
                tContacts(p) = sProps;
                tContacts(s).Centroid = [];
                break
            end % if
        end % if
    end %if
end % for p
```

```
% Assign Spot IDs to match Imaris and add to contact cell
    % array.
    rgnCount = sum(cellfun(@length, cellContacts));
    iSpotIDs = num2cell(rgnCount:rgnCount + length(tContacts) - 1);
    [tContacts.ID] = deal(iSpotIDs{:});
    cellContacts{t} = tContacts;
    % Create the all-contacts mask.
    tMask = false(size(tQDot));
    tMask(vertcat(tContacts.PixelIdxList)) = true;
    % Display the slice.
    cla(axesContours)
    imshow(ind2rgb(gray2ind(tQDot, 256), colorramp('o', 256, 'k')), ...
        'Parent', axesContours)
    % imshow(tQDot, [], 'Parent', axesContours)
    set(axesContours, 'NextPlot', 'Add')
    % Draw the cell border.
    contour(axesContours, contourExpanded(:, :, t), [1 1], 'Color', [0
    % Draw the new regions on the preview image.
    contour(axesContours, tMask > 0, [1 1], 'g')
    title(axesContours, ['Time point ' num2str(t) ' of ' num2str(tSize
        'Color', 'w')
    drawnow
end % for t
% Save the contact data.
save(fullfile(xFolder, ['Contacts ' fFileName, '.mat']), 'cellContacts
```

% Remove the regions that are now merged into other regions.
mergedRgnIdxs = arrayfun(@(s)isempty(s.Centroid), tContacts);

Transfer the contacts to Imaris as spots.

end % for s

tContacts(mergedRgnIdxs) = [];

Allocate arrays for the spots.

```
end % if
    tSpotCount = length(tContacts);
    rEnd = rStart + tSpotCount - 1;
   posXYImaris = 0.1*(vertcat(tContacts.Centroid));
   posXYImaris(:, 2) = yMax - posXYImaris(:, 2);
   posSpots(rStart:rEnd, 1:2) = posXYImaris;
   posSpots(rStart:rEnd, 3) = 0.5;
    timeSpots(rStart:rEnd) = t - 1;
    radiiSpots(rStart:rEnd) = 0.1*sqrt([tContacts.Area]/pi);
    rStart = rStart + tSpotCount;
end % for t
% Create and place the spots object.
xSpots = xImarisApp.GetFactory.CreateSpots;
xSpots.Set(posSpots, timeSpots, radiiSpots);
xSpots.SetName('Contacts')
xSpots.SetColorRGBA(rgbtripleto24bit([0 0.5 1]))
xImarisApp.GetSurpassScene.AddChild(xSpots, -1)
```

Save the processed file.

```
xImarisApp.FileSave(fullfile(xFolder, xFiles(f).name), 'writer="Imaris
```

Update the status bar and estimate the time required to finish.

```
timeElapsed = datenum(datetime('now')) - fStartTime(1);
completionEstimate = datestr(datenum(datetime('now')) + ...
        timeElapsed/f*(length(xFiles) - f), 'HH:MM PM');

hStatus.setText(['Processing: ' fFileName ' | Estimated finish: ' comp
hStatus.ProgressBar.setValue(f)
end % for f
```

Reset the status bar.

```
hStatus.setText('Finished processing folder')
hStatus.ProgressBar.setValue(0)
hStatus.ProgressBar.setVisible(0)
end % pushfolderprocess
end % contactsegmentationautomation

function closecontactautomationfcn(guiContactSegmentation, ~)
% CLOSECONTACTAUTOMATIONFCN Close the GUI and figure window
```

00 00

Close the image display figure if open, the close the gui figure.

```
figContours = getappdata(guiContactSegmentation, 'figContours');
if ishandle(figContours)
        delete(figContours)
end % if

delete(guiContactSegmentation)
end % closecontactautomationfcn
```

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