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/*
 * This code assumes that you have a folder of images stacks grouped in three, where image 1 is
 * a cell fill and images 2 and 3 are punctae that can be mapped to the cell fill. This code
 * requires FIJI and the plugins FeatureJ by Eric Meijering and the 3D Suite by Thomas Boudier.
 * In order for the code to run, this file ("Synapse and DendriteAnalysis.ijm") must be saved in
 * your FIJI "macros" folder. In addition the files "Segment_Structures.ijm",
 * "Distance_Transform_Soma.ijm", "Analyze_Skeleton.ijm", "Distance_Analysis.ijm", and
 * "Threshold.ijm" must be in the FIJI "macros" folder as well.
 *
 *
 * In general, aspects of this code were optimzied for our microscope setup (ie., file naming,
 * pixel size, etc). However, the code can be easily modified to match any file format. So if you
 * are interested in using the code, please contact me (Nick Galati) at nick.galati80@gmail.com
 * or go through GitHub at https://github.com/nickgalati80. I'd be glad to help get the code
 * running on any image format and to help you sort through the data.
 *
 * Thanks!
 */

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/*
 * Synapse_and_Dendrite_Analysis.ijm
 *
 * Save the following code as "Synapse_and_Dendrite_Analysis.ijm" to your macros folder.
 * Open this code in the Script editor and press run to start the macro.
 */

setBatchMode(true);
setForegroundColor(255,255,255);
run("Set Measurements...", "area mean standard modal min centroid center perimeter bounding fit
shape feret's integrated median skewness kurtosis area_fraction stack display redirect=None
decimal=3");
file=File.openDialog("Choose a Neuron File");
open(file);
rawFile=File.getName(file);
rawFile2=replace(rawFile, "_w1488", "_");
rawFile3=replace(rawFile2, ".TIF", "");
file2=replace(file, "_w1488", "_w2568");
file3=replace(file, "_w1488", "_w3647");
directory=File.getParent(file)+"\\ "+rawFile3;
print(directory);
open(file2);
open(file3);

selectImage(1);
rename("Neuron");
selectImage(2);
rename("VGAT");
selectImage(3);
rename("VGlut1");

//Cell Fill Segmentation
selectImage("Neuron");
runMacro("Segment_Structures.ijm", "Neuron"+"\\t"+"Neuron");
//Puncta 1 Segmentation
selectImage("VGAT");
runMacro("Segment_Structures.ijm", "Puncta"+"\\t"+"VGAT");
//Puncta 2 Segmentation
selectImage("VGlut1");
runMacro("Segment_Structures.ijm", "Puncta"+"\\t"+"VGlut1");
//
selectImage("Segmented_Soma");
runMacro("Distance_Transform_Soma.ijm");

selectImage("Segmented_Neuron");
runMacro("Analyze_Skeleton.ijm");

runMacro("Distance_Analysis.ijm",directory);
setBatchMode("exit and display");

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/*
 * Segment_Structures.ijm
 *
 * Save the following code as "Segment_Structures.ijm" to your macros folder
 */
argument=split(getArgument, "\t");
sliceStart=1;
sliceEnd=nSlices;
sliceRange=toString(sliceStart)+"-"+toString(sliceEnd);
run("FeatureJ Laplacian", "compute smoothing=1.0");
rename("Mean_Lap");
selectImage("Mean_Lap");
run("Invert", "stack");
run("Z Project...", "start=1 stop=&sliceEnd projection=[Max Intensity]");
getStatistics(area, mean, min, max, std, histo);
run("Measure");
close();
mode=getResult("Mode");
selectWindow("Results");
run("Close");
selectImage("Mean_Lap");
if(argument[0]=="Puncta"){
    setMinAndMax(mean, mean+(5*std));
}
if(argument[0]=="Neuron"){
    setMinAndMax(mode+(0.1*std), mode+(0.1*std));
}
run("8-bit");
if(argument[0]=="Neuron"){
    setMinAndMax(0,0);
    run("Apply LUT", "stack");
    run("Find Connected Regions", "allow_diagonal display_one_image display_results regions_for_values_over=100 minimum_number_of_points=1 stop_after=-1");
    rename("Segmented_Neuron");
    selectImage("Mean_Lap");
    close();
    sizeArray=newArray(nResults);
    for(a=0; a<nResults; a++){
        sizeArray[a]=getResult("Points In Region", a);
    }
    Array.getStatistics(sizeArray, min, neuronmax, mean, std);
    for(a=0; a<nResults; a++){
        value=sizeArray[a];
        if(value==neuronmax){
            thresholdMin=a+1;
            thresholdMax=a+1;
        }
    }
    selectImage("Segmented_Neuron");
    for(a=0; a<nSlices; a++){
        setSlice(a+1);
        setThreshold(thresholdMin, thresholdMax);
        run("Create Selection");
        if(selectionType>=0){
            run("Fill", "slice");
            run("Clear Outside", "slice");
        }
        if(selectionType===-1){
            run("Select All");
            run("Clear", "slice");
        }
        run("Select None");
        resetThreshold();
    }
    selectWindow("Results");
    run("Close");

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runMacro("Threshold.ijm");
run("Grays");
}
///
if(argument[0]=="Neuron"){
selectImage("Neuron");
run("Duplicate...", "title=Soma duplicate range=&sliceRange");
run("Enhance Contrast...", "saturated=0.4 use");
run("8-bit");
run("Gaussian Blur...", "sigma=10 stack");
run("Z Project...", "start=1 stop=&sliceEnd projection=[Sum Slices]");
run("8-bit");
setAutoThreshold("RenyiEntropy dark");
getThreshold(minSoma,maxSoma);
close();
selectImage("Soma");
setThreshold(minSoma,maxSoma);
for(a=0; a<nSlices; a++){
selectImage("Soma");
setSlice(a+1);
run("Create Selection");
if(selectionType>=0){
run("Enlarge...", "enlarge=-2.5");
run("Fill", "slice");
run("Clear Outside", "slice");
}
if(selectionType==1){
run("Select All");
run("Clear", "slice");
}
run("Select None");
}
run("Find Connected Regions", "allow_diagonal display_one_image display_results
regions_for_values_over=100 minimum_number_of_points=1 stop_after=-1");
rename("Segmented_Soma");
selectImage("Soma");
close();
sizeArray=newArray(nResults);
for(a=0; a<nResults; a++){
sizeArray[a]=getResult("Points In Region", a);
}
Array.getStatistics(sizeArray, min,neuronmax,mean,std);
for(a=0; a<nResults; a++){
value=sizeArray[a];
if(value==neuronmax){
thresholdMin=a+1;
thresholdMax=a+1;
}
}
selectImage("Segmented_Soma");
for(a=0; a<nSlices; a++){
setSlice(a+1);
setThreshold(thresholdMin, thresholdMax);
run("Create Selection");
if(selectionType>=0){
run("Fill", "slice");
run("Clear Outside", "slice");
}
if(selectionType==1){
run("Select All");
run("Clear", "slice");
}
run("Select None");
resetThreshold();
}
selectWindow("Results");
run("Close");
runMacro("Threshold.ijm");
run("Grays");

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run("8-bit");
}
if(argument[0]=="Puncta"){
run("3D Fast Filters","filter=MaximumLocal radius_x_pix=2.0 radius_y_pix=2.0
radius_z_pix=1.0 Nb_cpus=4");
runMacro("Threshold.ijm");
run("3D Spot Segmentation", "seeds_threshold=0 local_background=1 radius_0=3 radius_1=3
radius_2=4 weighth=0.50 radius_max=10 sd_value=.95 local_threshold=[Constant]
seg_spot=Classical watershed volume_min=1 volume_max=5 seeds=3D_MaximumLocal spots=Mean_Lap
radius_for_seeds=2 output=[Label Image]");
runMacro("Threshold.ijm");
rename("Segmented_"+argument[1]);
selectImage("3D_MaximumLocal");
close();
selectImage("Mean_Lap");
close();
}

/*
 * Distance_Transform_Soma.ijm
 *
 * Save the following code as "Distance_Transform_Soma.ijm" to your macros folder
 */
run("3D Manager");
Ext.Manager3D_AddImage();
Ext.Manager3D_MassCenter3D(0, cmx,cmy,cmz);
Ext.Manager3D_Reset();
run("Replace value", "pattern=0 replacement=1");
setSlice(cmz);
makePoint(cmx,cmy);
setForegroundColor(0,0,0);
run("Draw", "slice");
setForegroundColor(255,255,255);
run("Exact Euclidean Distance Transform (3D)");
rename("Distance_Map");
for(a=0; a<nImages; a++){
selectImage(a+1);
run("Specify...", "width=450 height=450 x=&cmx y=&cmy slice=&cmz oval centered");
run("Clear Outside", "stack");
run("Select None");
}
selectImage("Segmented_Soma");
run("Replace value", "pattern=1 replacement=0");

/*
 * Analyze_Skeleton.ijm
 *
 * Save the following code as "Analyze_Skeleton.ijm" to your macros folder
 */
selectImage("Segmented_Neuron");
sliceStart=1;
sliceEnd=nSlices;
sliceRange=toString(sliceStart)+"-"+toString(sliceEnd);
run("Duplicate...", "title=DistanceSkeleton duplicate range=&sliceRange");
distanceSkeleton=getImageID();
run("Duplicate...", "title=Skeleton duplicate range=&sliceRange");
skeleton=getImageID();
selectWindow("Skeleton");
run("Skeletonize (2D/3D)");
selectWindow("Distance");
run("Replace value", "pattern=0 replacement=1");
run("Replace value", "pattern=255 replacement=3");
selectWindow("Skeleton");

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run("Replace value", "pattern=255 replacement=3");
imageCalculator("Subtract create stack", "Distance", "Skeleton");
rename("DistanceMap");
run("3D Distance Map", "threshold=0");
rename("Diameter");
for(a=0; a<nSlices; a++){
    selectImage("Segmented_Neuron");
    setSlice(a+1);
    setThreshold(1,255);
    run("Create Selection");
    if(selectionType>=0){
        roiManager("Add");
        selectImage("Diameter");
        setSlice(a+1);
        roiManager("Select",0);
        run("Clear Outside", "slice");
        roiManager("Select", 0);
        roiManager("Delete");
    }
    if(selectionType===-1){
        selectImage("Diameter");
        setSlice(a+1);
        run("Select All");
        run("Clear", "slice");
    }
    run("Select None");
}
selectImage("Segmented_Neuron");
run("Select None");
resetThreshold();
selectImage("Distance");
close();
selectImage("DistanceMap");
close();
selectImage("Skeleton");
rename("Segmented_Skeleton");
runMacro("Threshold.ijm");
selectImage("Diameter");
run("Gaussian Blur...", "sigma=10 stack");

/*
 * Distance_Analysis.ijm
 *
 * Save the following code as "Distance_Analysis.ijm" to your macros folder
 */

argument=getArgument();
run("3D Manager");
nameArray=newArray("Neuron", "Soma", "Skeleton", "VGAT_TFX", "VGAT_NonTFX", "VGlut1_TFX",
"VGlut1_NonTFX");
Ext.Manager3D_MonoSelect();
//
selectImage("Segmented_Neuron");
Ext.Manager3D_AddImage();
Ext.Manager3D_Select(0);
Ext.Manager3D_Rename("Neuron Object");
Ext.Manager3D_Save(argument+"Neuron"+" .zip");
Ext.Manager3D_Measure3D(0, "Vol", totalNeuronVolume);
Ext.Manager3D_Reset();
//
selectImage("Segmented_Skeleton");
Ext.Manager3D_AddImage();
Ext.Manager3D_Select(0);

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Ext.Manager3D_Rename("Skeleton Object");
Ext.Manager3D_Save(argument+"Skeleton"+" .zip");
Ext.Manager3D_Measure3D(0, "Vol", totalNeuronLength);
selectImage("Diameter");
Ext.Manager3D_Quantif3D(0, "Mean", averageNeuronDiameter);
Ext.Manager3D_Reset();

//
selectImage("Segmented_VGAT");
Ext.Manager3D_AddImage();
Ext.Manager3D_Select(0);
Ext.Manager3D_Rename("VGAT Object");
Ext.Manager3D_Save(argument+"VGAT_Total"+" .zip");
Ext.Manager3D_Reset();
//
selectImage("Segmented_VGlut1");
Ext.Manager3D_AddImage();
Ext.Manager3D_Select(0);
Ext.Manager3D_Rename("VGlut1 Object");
Ext.Manager3D_Save(argument+"VGlut1_Total"+" .zip");
Ext.Manager3D_Reset();
////////////////////////////////////
selectImage("Segmented_VGAT");
Ext.Manager3D_Segment(1,255);
close;
Ext.Manager3D_AddImage();
close;
Ext.Manager3D_Count(objects);
Ext.Manager3D_MultiSelect();
selectImage("Segmented_Neuron");
for(a=0; a<objects; a++){
    Ext.Manager3D_Quantif3D(a, "Max", value);
    if(value>0){
        Ext.Manager3D_Select(a);
    }
}
Ext.Manager3D_Save(argument+"VGAT_TFX"+" .zip");
newImage("VGAT_TFX", "8-bit black", 512, 512, 13);
Ext.Manager3D_FillStack(255,255,255);
Ext.Manager3D_Delete();
Ext.Manager3D_MultiSelect();
Ext.Manager3D_SelectAll();
Ext.Manager3D_Save(argument+"VGAT_NonTFX"+" .zip");
newImage("VGAT_NonTFX", "8-bit black", 512, 512, 13);
Ext.Manager3D_FillStack(255,255,255);
Ext.Manager3D_DeselectAll();
Ext.Manager3D_Reset();
////////////////////////////////////
////////////////////////////////////
selectImage("Segmented_VGlut1");
Ext.Manager3D_Segment(1,255);
close;
Ext.Manager3D_AddImage();
close;
Ext.Manager3D_Count(objects);
Ext.Manager3D_MultiSelect();
selectImage("Segmented_Neuron");
for(a=0; a<objects; a++){
    Ext.Manager3D_Quantif3D(a, "Max", value);
    if(value>0){
        Ext.Manager3D_Select(a);
    }
}
Ext.Manager3D_Save(argument+"VGlut1_TFX"+" .zip");
newImage("VGlut1_TFX", "8-bit black", 512, 512, 13);
Ext.Manager3D_FillStack(255,255,255);
Ext.Manager3D_Delete();
Ext.Manager3D_MultiSelect();
Ext.Manager3D_SelectAll();

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Ext.Manager3D_Save(argument+"VGlut1_NonTFX"+" .zip");
newImage("VGlut1_NonTFX", "8-bit black", 512, 512, 13);
Ext.Manager3D_FillStack(255,255,255);
Ext.Manager3D_DeselectAll();
Ext.Manager3D_Reset();
////////////////////////////////////

////////////////////////////////////

newImage("Cross_Synapses", "8-bit black", 512, 512, 13);
titleArray=newArray("VGAT", "VGAT", "VGlut1", "VGlut1");
secondTitleArray=newArray("VGlut1_TFX", "VGlut1_NonTFX", "VGAT_TFX", "VGAT_NonTFX");
binArray=newArray("binOne", "binTwo", "binThree", "binFour", "binFive", "binSix", "binSeven", "binEight",
, "binNine");
selectionArray=newArray(36);
totalObjectsArray=newArray(4);
counter=0;
for(a=0; a<4; a++){
    zipFile=argument+nameArray[a+3]+".zip";
    Ext.Manager3D_Load(zipFile);
    Ext.Manager3D_Count(objects);
    totalObjectsArray[a]=objects;
    selectImage(secondTitleArray[a]);
    Ext.Manager3D_AddImage();
    masterArray=newArray(objects*8);
    bin1Array=newArray();
    bin2Array=newArray();
    bin3Array=newArray();
    bin4Array=newArray();
    bin5Array=newArray();
    bin6Array=newArray();
    bin7Array=newArray();
    bin8Array=newArray();
    bin9Array=newArray();
    newImage(nameArray[a+3]+"_Total", "32-bit black", 8, objects, 1);
    for(b=0; b<objects; b++){
        tempArray=newArray(8);
        selectImage(8);
        Ext.Manager3D_Quantif3D(b, "Mean", somadistance);
        if(somadistance<=25){bin=1;}
        if(somadistance>25 && somadistance<=50){bin=2;}
        if(somadistance>50 && somadistance<=75){bin=3;}
        if(somadistance>75 && somadistance<=100){bin=4;}
        if(somadistance>100 && somadistance<=125){bin=5;}
        if(somadistance>125 && somadistance<=150){bin=6;}
        if(somadistance>150 && somadistance<=175){bin=7;}
        if(somadistance>175 && somadistance<=200){bin=8;}
        if(somadistance>200){bin=9;}
        selectImage(titleArray[a]);
        Ext.Manager3D_Quantif3D(b, "Max", samemax);
        Ext.Manager3D_Closest(b, "cc", sameclosest);
        Ext.Manager3D_Dist2(b, sameclosest, "c1b2", closest);
        selectImage(10);
        Ext.Manager3D_Quantif3D(b, "Max", dendritediameter);
        Ext.Manager3D_Dist2(b, objects, "c1b2", otherdistance);
        if(otherdistance<1){
            selectImage("Cross_Synapses");
            Ext.Manager3D_Select(b);
            fillColor=(a+10)*10;
            Ext.Manager3D_FillStack(fillColor, fillColor, fillColor);
            cross=1;
            notcross=0;
        }
        if(otherdistance>=1){
            cross=0;
            notcross=1;
        }
        index=b*8;

        masterArray[index+0]=bin; masterArray[index+1]=somadistance; masterArray[index+2]=samemax; m

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asterArray[index+3]=closest;masterArray[index+4]=dendritediameter;masterArray[index+5]=otherdistance;masterArray[index+6]=cross;masterArray[index+7]=notcross;
selectImage(nameArray[a+3]+"_Total");

setPixel(0,b,bin);setPixel(1,b,somadistance);setPixel(2,b,samemax);setPixel(3,b,closest);
setPixel(4,b,dendritediameter);setPixel(5,b,otherdistance);setPixel(6,b,cross);setPixel(7,b,notcross);
}
for(b=0; b<objects; b++){
  showStatus("HI"+a);
  showProgress(b/objects);
  index=b*8;
  bin=masterArray[index];
  slice=Array.slice(masterArray,index, index+8);
  if(bin==1){bin1Array=Array.concat(bin1Array, slice);}
  if(bin==2){bin2Array=Array.concat(bin2Array, slice);}
  if(bin==3){bin3Array=Array.concat(bin3Array, slice);}
  if(bin==4){bin4Array=Array.concat(bin4Array, slice);}
  if(bin==5){bin5Array=Array.concat(bin5Array, slice);}
  if(bin==6){bin6Array=Array.concat(bin6Array, slice);}
  if(bin==7){bin7Array=Array.concat(bin7Array, slice);}
  if(bin==8){bin8Array=Array.concat(bin8Array, slice);}
  if(bin==9){bin9Array=Array.concat(bin9Array, slice);}
}
length1=bin1Array.length/8;
length2=bin2Array.length/8;
length3=bin3Array.length/8;
length4=bin4Array.length/8;
length5=bin5Array.length/8;
length6=bin6Array.length/8;
length7=bin7Array.length/8;
length8=bin8Array.length/8;
length9=bin9Array.length/8;
heightArray=newArray(9);

heightArray[0]=length1;heightArray[1]=length2;heightArray[2]=length3;heightArray[3]=length4;heightArray[4]=length5;heightArray[5]=length6;heightArray[6]=length7;heightArray[7]=length8;heightArray[8]=length9;
if(length1==0){
  heightArray[0]=1;
  bin1Array=newArray(8);

  bin1Array[0]=1;bin1Array[1]=0;bin1Array[2]=0;bin1Array[3]=0;bin1Array[4]=0;bin1Array[5]=0;bin1Array[6]=0;bin1Array[7]=0;
}

for(b=0; b<9; b++){
  height=heightArray[b];

  binArray=newArray("binOne","binTwo","binThree","binFour","binFive","binSix","binSeven","binEight","binNine");
  newImage(nameArray[a+3]+"_"+binArray[b], "32-bit black",8,height,1);
}
for(b=0; b<9; b++){
  selectImage(nameArray[a+3]+"_"+binArray[b]);

  if(b==0)array=bin1Array;if(b==1)array=bin2Array;if(b==2)array=bin3Array;if(b==3)array=bin4Array;if(b==4)array=bin5Array;if(b==5)array=bin6Array;if(b==6)array=bin7Array;if(b==7)array=bin8Array;if(b==8)array=bin9Array;
  selectionArray[counter]=array.length/8;
  counter++;
  for(c=0; c<array.length/8; c++){
    for(d=0; d<8; d++){
      index=c*8;
      setPixel(d,c,array[index+d]);
    }
  }
}
}
Ext.Manager3D_Reset();

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}
run("Images to Stack", "method=[Copy (top-left)] name=Binned_Results title=bin use");
run("Images to Stack", "method=[Copy (top-left)] name=Total_Results title=_Total use");
////////////////////////////////////////
selectImage("Binned_Results");
averageBinnedResults=newArray(288);
counter=0;
counterTwo=0;
for(b=0; b<36; b++){
    slice=b+1;
    setSlice(slice);
    height=selectionArray[counter];
    counter++;
    for(c=0; c<8; c++){
        run("Specify...", "width=1 height=&height x=&c y=0 slice=&slice");
        getStatistics(area,mean,min,max,std,histo);
        if(c<=5){value=mean;}
        else{value=round(mean*area);}
        averageBinnedResults[counterTwo]=value;
        run("Select None");
        counterTwo++;
    }
}
selectImage("Total_Results");
averageTotalResults=newArray(32);
counter=0;
for(b=0; b<4; b++){
    slice=b+1;
    setSlice(slice);
    height=totalObjectsArray[b];
    for(c=0; c<8; c++){
        run("Specify...", "width=1 height=&height x=&c y=0 slice=&slice");
        getStatistics(area,mean,min,max,std,histo);
        if(c<=5){value=mean;}
        else{value=round(mean*area);}
        averageTotalResults[counter]=value;
        run("Select None");
        counter++;
    }
}
}
////
selectImage("Cross_Synapses");
Ext.Manager3D_AddImage();
Ext.Manager3D_Select(0);
Ext.Manager3D_Save(argument+"CrossSyn_VGAT_TFX.zip");
Ext.Manager3D_Select(1);
Ext.Manager3D_Save(argument+"CrossSyn_VGAT_NonTFX.zip");
Ext.Manager3D_Select(2);
Ext.Manager3D_Save(argument+"CrossSyn_VGluT1_TFX.zip");
Ext.Manager3D_Select(3);
Ext.Manager3D_Save(argument+"CrossSyn_VGluT1_NonTFX.zip");
Ext.Manager3D_Reset();
//
selectImage("Distance_Map");
run("Z Project...", "start=1 stop=13 projection=[Min Intensity]");
rename("CenterPoint");
setThreshold(0, 0);
run("Create Selection");
roiManager("Add");
roiManager("Split");
selectImage("CenterPoint");
close();
selectImage("Segmented_Skeleton");
run("Z Project...", "start=1 stop=13 projection=[Max Intensity]");
rename("Sholl");
roiManager("Select", 1);
run("Measure");
xCenter=getResult("BX", 0);

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yCenter=getResult("BY", 0);
selectWindow("Results");
run("Close");
selectImage("Sholl");
makePoint(xCenter,yCenter);
run("ShollAnalysis ", "starting=1 ending=250 radius=25 radius=5 span=Median");
selectWindow("Sholl Analysis for Sholl");
Plot.getValues(distance, crossings);
Array.getStatistics(crossings,min,max,mean,std);
totalCrossings=floor(mean*crossings.length);
selectWindow("Sholl");
close();
selectWindow("Sholl Analysis for Sholl");
close();
roiManager("Deselect");
roiManager("Delete");
//
selectImage("Segmented_Neuron");
run("Replace value", "pattern=255 replacement=1");
selectImage("Segmented_Skeleton");
run("Replace value", "pattern=255 replacement=1");
imageCalculator("Multiply create 32-bit stack", "Segmented_Neuron","Distance_Map");
rename("Distance_Volume");
imageCalculator("Multiply create 32-bit stack", "Segmented_Skeleton","Distance_Map");
rename("Distance_Skeleton");
//
binnedVolumeArray=newArray(9);
for(a=0; a<9; a++){
    selectImage("Distance_Volume");
    Ext.Manager3D_Count(startObjects);
    lowThreshold=a*25+1;
    highThreshold=lowThreshold+25;
    Ext.Manager3D_Segment(lowThreshold, highThreshold);
    close();
    Ext.Manager3D_AddImage();
    close();
    Ext.Manager3D_Count(endObjects);
    if((endObjects-startObjects)>1){
        Ext.Manager3D_MultiSelect();
        Ext.Manager3D_SelectFor(startObjects, endObjects, 1);
        Ext.Manager3D_Merge();
        Ext.Manager3D_Count(measureObject);
        Ext.Manager3D_Measure3D(measureObject-1, "Vol", volume);
        Ext.Manager3D_DeselectAll();
    }
    Ext.Manager3D_Count(measureObject);
    Ext.Manager3D_Measure3D(measureObject-1, "Vol", volume);
    binnedVolumeArray[a]=volume;
}
binnedLengthArray=newArray(9);
binnedDiameterArray=newArray(9);
for(a=0; a<9; a++){
    selectImage("Distance_Skeleton");
    Ext.Manager3D_Count(startObjects);
    lowThreshold=a*25+1;
    highThreshold=lowThreshold+25;
    Ext.Manager3D_Segment(lowThreshold, highThreshold);
    close();
    Ext.Manager3D_AddImage();
    close();
    Ext.Manager3D_Count(endObjects);
    if((endObjects-startObjects)>1){
        Ext.Manager3D_MultiSelect();
        Ext.Manager3D_SelectFor(startObjects, endObjects, 1);
        Ext.Manager3D_Merge();
        Ext.Manager3D_DeselectAll();
    }
    Ext.Manager3D_Count(measureObject);
    Ext.Manager3D_Measure3D(measureObject-1, "Vol", volume);
}

```

```

selectImage("Diameter");
Ext.Manager3D_Quantif3D(measureObject-1, "Mean", diameter);
binnedLengthArray[a]=volume;
binnedDiameterArray[a]=diameter;
}
Ext.Manager3D_Save(argument+"BinnedMorphology"+" .zip");
Ext.Manager3D_DeselectAll();
Ext.Manager3D_MultiSelect();
Ext.Manager3D_SelectFor(9,17,1);
Ext.Manager3D_Merge();
selectImage("Diameter");
Ext.Manager3D_Quantif3D(9,"Mean",totalAverageDiameter);
Ext.Manager3D_Reset();
///
counter=0;
counterTwo=0;
for(a=0; a<4; a++){
    thirdTitleArray=newArray("VGAT_TFX", "VGAT_NonTFX", "VGlut1_TFX", "VGlut1_NonTFX");
    for(b=0; b<9; b++){
        setResult("Label", b, "Bin-"+averageBinnedResults[counter]);
        setResult(thirdTitleArray[a]+"_"+"Bin", b, averageBinnedResults[counter]);
        setResult(thirdTitleArray[a]+"_"+"Distance From Soma", b,
            averageBinnedResults[counter+1]);
        setResult(thirdTitleArray[a]+"_"+"Intensity", b, averageBinnedResults[counter+2]);
        setResult(thirdTitleArray[a]+"_"+"Closest Same", b, averageBinnedResults[counter+3]);
        setResult(thirdTitleArray[a]+"_"+"Dend. Diameter @ Synapse", b,
            averageBinnedResults[counter+4]);
        setResult(thirdTitleArray[a]+"_"+"Closest Other", b, averageBinnedResults[counter+5]);
        setResult(thirdTitleArray[a]+"_"+"Total Cross", b, averageBinnedResults[counter+6]);
        setResult(thirdTitleArray[a]+"_"+"Total Non Cross", b, averageBinnedResults[counter+7]);
        setResult(thirdTitleArray[a]+"_"+"Total", b,
            averageBinnedResults[counter+6]+averageBinnedResults[counter+7]);
        setResult(thirdTitleArray[a]+"_"+"Volume", b, binnedVolumeArray[b]);
        setResult(thirdTitleArray[a]+"_"+"Length", b, binnedLengthArray[b]);
        setResult(thirdTitleArray[a]+"_"+"Dendrite Diameter", b, binnedDiameterArray[b]);
        setResult(thirdTitleArray[a]+"_"+"Crossings", b, crossings[b]);
        setResult(thirdTitleArray[a]+"_"+"Cross Syn./Length", b,
            averageBinnedResults[counter+6]/binnedLengthArray[b]);
        setResult(thirdTitleArray[a]+"_"+"Cross Syn./Volume", b,
            averageBinnedResults[counter+6]/binnedVolumeArray[b]);
        setResult(thirdTitleArray[a]+"_"+"Non Cross Syn./Length", b,
            averageBinnedResults[counter+7]/binnedLengthArray[b]);
        setResult(thirdTitleArray[a]+"_"+"Non Cross Syn./Volume", b,
            averageBinnedResults[counter+7]/binnedVolumeArray[b]);
        setResult(thirdTitleArray[a]+"_"+"Total Syn./Length", b,
            (averageBinnedResults[counter+6]+averageBinnedResults[counter+7])/binnedLengthArray[b]);
        setResult(thirdTitleArray[a]+"_"+"Total Syn./Volume", b,
            (averageBinnedResults[counter+6]+averageBinnedResults[counter+7])/binnedVolumeArray[b]);

        counter=counter+8;
    }
    setResult("Label", 9, "Unbinned Average");
    setResult(thirdTitleArray[a]+"_"+"Distance From Soma", 9, averageTotalResults[counterTwo+1]);
    setResult(thirdTitleArray[a]+"_"+"Intensity", 9, averageTotalResults[counterTwo+2]);
    setResult(thirdTitleArray[a]+"_"+"Closest Same", 9, averageTotalResults[counterTwo+3]);
    setResult(thirdTitleArray[a]+"_"+"Dend. Diameter @ Synapse", 9,
        averageTotalResults[counterTwo+4]);
    setResult(thirdTitleArray[a]+"_"+"Closest Other", 9, averageTotalResults[counterTwo+5]);
    setResult(thirdTitleArray[a]+"_"+"Total Cross", 9, averageTotalResults[counterTwo+6]);
    setResult(thirdTitleArray[a]+"_"+"Total Non Cross", 9, averageTotalResults[counterTwo+7]);
    setResult(thirdTitleArray[a]+"_"+"Total", 9,
        averageTotalResults[counterTwo+6]+averageTotalResults[counterTwo+7]);
    setResult(thirdTitleArray[a]+"_"+"Volume", 9, totalNeuronVolume);
    setResult(thirdTitleArray[a]+"_"+"Length", 9, totalNeuronLength);
    setResult(thirdTitleArray[a]+"_"+"Dendrite Diameter", 9, totalAverageDiameter);
    setResult(thirdTitleArray[a]+"_"+"Crossings", 9, totalCrossings);
    setResult(thirdTitleArray[a]+"_"+"Cross Syn./Length", 9,
        averageTotalResults[counterTwo+6]/totalNeuronLength);

```

```

setResult(thirdTitleArray[a]+"_"+"Cross Syn./Volume", 9,
averageTotalResults[counterTwo+6]/totalNeuronVolume);
setResult(thirdTitleArray[a]+"_"+"Non Cross Syn./Length", 9,
averageTotalResults[counterTwo+7]/totalNeuronLength);
setResult(thirdTitleArray[a]+"_"+"Non Cross Syn./Volume", 9,
averageTotalResults[counterTwo+7]/totalNeuronVolume);
setResult(thirdTitleArray[a]+"_"+"Total Syn./Length", 9,
(averageTotalResults[counterTwo+6]+averageTotalResults[counterTwo+7])/totalNeuronLength);
setResult(thirdTitleArray[a]+"_"+"Total Syn./Volume", 9,
(averageTotalResults[counterTwo+6]+averageTotalResults[counterTwo+7])/totalNeuronVolume);
counterTwo=counterTwo+8;
updateResults();
selectWindow("Results");
saveAs("Results", argument+"Binned_"+thirdTitleArray[a]+".xls");
selectWindow("Results");
run("Close");
}
selectImage("Total_Results");
saveAs("TIFF", argument+"_Total.tif");
selectImage("Binned_Results");
saveAs("TIFF", argument+"_Binned.tif");

```