

Algorithm for Finding Granuloma Clusters

Because the number of granulomas is typically in the dozens, this algorithm as designed doesn't require the terms to have memory pre-allocation.

First, we make a list of ID's of all coupled intersections of individual granulomas.

Ex. *IndivIntersectList*=[1,2 ; 1,3 ; 1,4 ; 2,3 ; 2,7 ; 6,10]

means the following:

IndivGran(1) intersects with *IndivGran*(2), *IndivGran*(3), and *IndivGran*(4)

IndivGran(2) intersects with *IndivGran*(1), *IndivGran*(3), and *IndivGran*(7)

IndivGran(3) intersects with *IndivGran*(1) and *IndivGran*(2)

IndivGran(4) intersects with *IndivGran*(1)

IndivGran(6) intersects with *IndivGran*(10)

IndivGran(7) intersects with *IndivGran*(2)

and the rest of the individual granulomas don't intersect any others.

Algorithm:

IndivIntersectList: list of ID's of pairs of intersecting individual granulomas

LenIntersectList: number of pairs in *IndivIntersectList*

TotIndivGrans: total number of individual granulomas

IndivGranObject: data structure of all individual granulomas

ti: time

PositionX(*ti*), *PositionY*(*ti*), *PositionZ*(*ti*): x-,y-,z-coordinates of center of granuloma at time *ti*

Radius(*ti*): radius of individual granuloma at time *ti*

IndivIntersectList=[]

LenIntersectList=0

if *TotIndivGrans*>1

for *ID1*=1 to *ID1*=*TotIndivGrans*-1

x_ID1=*IndivGranObject*(*ID1*).*PositionX*(*ti*)

y_ID1=*IndivGranObject*(*ID1*).*PositionY*(*ti*)

z_ID1=*IndivGranObject*(*ID1*).*PositionZ*(*ti*)

Radius_ID1=*IndivGranObject*(*ID1*).*Radius*(*ti*)

for *ID2*=*ID1*+1 to *ID2*=*TotIndivGrans*

x_ID2=*IndivGranObject*(*ID2*).*PositionX*(*ti*)

y_ID2=*IndivGranObject*(*ID2*).*PositionY*(*ti*)

z_ID2=*IndivGranObject*(*ID2*).*PositionZ*(*ti*)

Radius_ID2=*IndivGranObject*(*ID2*).*Radius*(*ti*)

TempDist=sqrt((*x_ID1*-*x_ID2*)² + (*y_ID1*-*y_ID2*)² + (*z_ID1*-*z_ID2*)²)

if *TempDist*<=(*Radius_ID1*+*Radius_ID2*)

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        LenIntersectList=LenIntersectList+1
        IndivIntersectList(LenIntersectList,1)=ID_1
        IndivIntersectList(LenIntersectList,2)=ID_2
    end "if" loop
end "for" loop
end "for" loop
end "if" loop

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Next, we build a list of individual granuloma ID's that make up all granuloma clusters.

Ex. with the example above, *IndivIntersectList*=[1,2 ; 1,3 ; 1,4 ; 2,3 ; 2,7 ; 6,10] with

TotIndivGrans=10

we have *GranulomaClusterList*={ [1,2,3,4,7] , [6,10] , 5 , 8 , 9 }

which means that we have a total of 5 granuloma "clusters": a cluster made up by individual granulomas of ID's 1,2,3,4, and 7; a cluster made up by individual granulomas of ID's 6 and 10; a "cluster" made up by individual granulomas of ID 5; a "cluster" made up by individual granulomas of ID 8; and a "cluster" made up by individual granulomas of ID 9.

Algorithm:

GranulomaClusterList: the main output of the algorithm: a list of individual granuloma ID's that make up each granuloma cluster

LenGranulomaClusterList: the total number of granuloma clusters

IndivIntersectList: list of ID's of pairs of intersecting individual granulomas (from above)

LenIntersectList: number of pairs in *IndivIntersectList* (from above)

IndivIntersectListTEMP: begins a copy of *IndivIntersectList*, but empties as the algorithm progresses

LenIntersectList: number of pairs in *IndivIntersectListTEMP*

TotIndivGrans: total number of individual granulomas

IndivGranIDList: list of all ID's for individual granulomas

UnassignedIDList: list of all individual granuloma ID's that haven't been assigned to a granuloma cluster

LenUnassignedIDList: number of individual granuloma ID's that haven't been assigned to a granuloma cluster—it begins the same as *IndivGranIDList*, but empties as the algorithm progresses

AssignedIDList: list of all individual granuloma ID's that have been assigned to a granuloma cluster—each individual granuloma ID must belong to either *UnassignedIDList* or *AssignedIDList*, but not both

LenUnassignedIDList: number of individual granuloma ID's that have been assigned to a granuloma cluster

TempGranClusterList: a list of individual granuloma ID's that each iteration is dynamically constructed, added to *GranulomaClusterList*, then purged before repeating

LenTempGranClusterList: number of individual granuloma ID's in *GranulomaClusterList*

TempSearchList: a dynamically-changing list of individual granuloma ID's that is a part of the current cluster but still pending a move to *TempGranClusterList*—each iteration it begins at 1, then gets added to and one by one moves its elements to *TempGranClusterList* until it is empty

LenTempSearchList: number of individual granuloma ID's that are in *TempSearchList*

TempSearchID: a single individual granuloma ID from *TempSearchList* that is being checked against to see whether there are other individual granuloma ID's paired with *TempSearchID* in *IndivIntersectListTEMP*

RemoveList: a list of pairs of items: (1) the index for which pair of ID's *IndivIntersectListTEMP* that should be removed from *IndivIntersectListTEMP*, and (2) the 2nd ID associated with the index from (1)—each iteration it begins empty then grows dynamically

LenRemoveList: the number of pairs of items in *RemoveList*

RemoveList2:

LenRemoveList2:

TempValI:

AddToSearchList:

RemoveFromList:

IndivGranIDList=1 , 2 , ... , *TotIndivGrans*

if *LenIntersectList*==0

GranulomaClusterList=*IndivIDGranList*

LenGranulomaClusterList=*TotIndivGrans*

else

IndivIntersectListTEMP=*IndivIntersectList*; *LenIntersectListTEMP*=*LenIntersectList*

UnassignedIDList=*IndivIDGranList*; *LenUnassignedIDList*=*TotIndivGrans*

AssignedIDList=zeros(size(*UnassignedIDList*)); *LenAssignedIDList*=0

GranulomaClusterList=[]; *LenGranulomaClusterList*=0

while *LenUnassignedIDList*>0

LenTempGranClusterList=0

TempGranClusterList=[];

LenTempSearchList=1

TempSearchList(*LenTempSearchList*)=*UnassignedIDList*(1)

UnassignedIDList(1)=[];

LenUnassignedIDList=*LenUnassignedIDList*-1

AssignedIDList(*LenAssignedIDList*)=*TempSearchList*(*LenTempSearchList*);

LenAssignedIDList=*LenAssignedIDList*+1

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while LenTempSearchList>0
  TempSearchList=sort(TempSearchList)
  TempSearchID=TempSearchList(1)
  TempSearchList(1)=[ ]
  LenTempSearchList=LenTempSearchList-1
  LenGranulomaClusterList=LenGranulomaClusterList+1
  GranulomaClusterList(LenGranulomaClusterList)=TempSearchID

  if LenIntersectListTEMP>0
    RemoveIDList=[ ]; LenRemoveList=0

    for IntersectListIndex=1 to IntersectListIndex =LenIntersectListTEMP
      if IndividIntersectListTEMP(IntersectListIndex,1)==TempSearchID
        LenTempSearchList=LenTempSearchList+1

        TempSearchList(LenTempSearchList)=IndividIntersectListTEMP(IntersectListIndex,2)
        LenRemoveList=LenRemoveList+1
        RemoveList(LenRemoveList,1)= IntersectListIndex
        RemoveList(LenRemoveList,2)=TempSearchList(LenTempSearchList)
      end "if"
      if IndividIntersectListTEMP(IntersectListIndex,1)>TempSearchID
        break
      end "if"
    end "for"

    if LenRemoveList>0
      for RemoveListIndex=1 to RemoveListIndex=LenRemoveList
        for IntersectListIndex=1 to IntersectListIndex=LenIntersectListTEMP
          if
            IndividIntersectListTEMP(IntersectListIndex,2)==RemoveList(LenRemoveList,2)
            && IndividIntersectListTEMP(IntersectListIndex,1)~=TempSearchID
            AddToSearchList='yes'
            if LenTempSearchList>0
              for SearchListIndex=1 to SearchListIndex=LenTempSearchList
                if TempSearchList(SearchListIndex)==
                  IndividIntersectListTEMP(IntersectListIndex,1)
                  AddToSearchList='yes'
                  break
                end "if"
              end "for"
            end "if"
            if AddToSearchList=='yes'
              LenTempSearchList=LenTempSearchList+1
            end "if"
          end "if"
        end "for"
      end "for"

      TempSearchList(LenTempSearchList)=IndividIntersectListTEMP(IntersectListIndex,1)
    end "if"
  end "if"
end "while"

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        LenRemoveList=LenRemoveList+1
        RemoveList(LenRemoveList,1)=IntersectListIndex
        RemoveList(LenRemoveList,2)=TempSearchList(LenTempSearchList)
    end "if"
end "for"
end "for"
end "if"

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if LenRemoveList>0
    IndivIntersectListTEMP(RemoveList(~,1),~)=[ ]
    LenIntersectListTEMP=LenIntersectListTEMP-LenRemoveList
    RemoveList2=zeros(size(RemoveList))
    LenRemoveList2=0

    for RemoveListIndex=1 to RemoveListIndex=LenRemoveList
        TempVal=RemoveList(RemoveListIndex,2)
        for UnassignedListIndex=1 to UnassignedListIndex=LenUnassignedList
            if UnassignedList(UnassignedListIndex)==TempVal
                RemoveFromList='yes'
                if RemoveListIndex>1
                    for RemoveListIndex2=1 to RemoveListIndex2=RemoveListIndex-1
                        if RemoveList(RemoveListIndex2,2)==TempVal
                            RemoveFromList='no'
                            break
                        end "if"
                    end "for"
                end "if"
                if RemoveFromList=='yes'
                    LenRemoveList2=LenRemoveList2+1
                    RemoveList2(LenRemoveList2,1)=UnassignedListIndex
                    RemoveList2(LenRemoveList2,2)=TempVal
                    break
                end "if"
            end "if"
        end "for"
    end "for"

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if LenRemoveList2>0
    RemoveList2=RemoveList2(1:LenRemoveList2,~)
    UnassignedList(RemoveList2(~,1))=[ ]
    LenUnassignedList=LenUnassignedList-LenRemoveList2

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AssignedList(LenAssignedList+1:LenAssignedList+LenRemoveList2)=Transpose(RemoveList2(
~,2))

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    LenAssignedList= LenAssignedList+LenRemoveList2
end "if"

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end “if”

end “if”

end “while”

LenGranulomaClusterList=LenGranulomaClusterList+1

GranulomaClusterList{LenGranulomaClusterList}=IndivIntersectListTEMP

end “while”

end “if”