Agglomerative Hierarchical Clustering The C++ implmentation for AHC with OpenMP

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Chapter 1

Agglomerative Hierarachical Clustering (AHC)

The program includes basically two aspects of AHC

- AHC of three linkages (will generate cluster result information)
 - Single linkage
 - Complete linkage
 - Average linkage
- The hierarchical L method to find optimal number of clusters (only generate optimal cluster number)
 - It is a global search of knee point along the clusters

Number of clusters as input

The program supports two kinds of input for number of clusters

- Direct input after the query information > Input cluster number among [0, 1000] for norm X:
- · Read the cluster numbers from a txt file
 - The txt file is called 'cluster_number' in the /dataset/ folder
 - The 'cluster number' has the following format
 - 0:10 // for similarity measure 0, the input of cluster number is 10 1:10 // for similarity measure 1, the input of cluster number is 10 2:10 4:10 12:10 13:10 14:10 15:10 17:10
 - for better batch processing especially in our experiment when the code is automatically run on the server

Chapter 2

Class Index

2.1 Class List

Here are the classes, structs, unions and interfaces with brief descriptions:

AHC	 																						7	ĺ
DataSet .	 											 					 						22	
DistNode	 											 					 						23	
Ensemble																							24	

4 Class Index

Chapter 3

File Index

3.1 File List

Here is a list of all files with brief descriptions:

AHC.cpp																								27
AHC.h																								27
main.cpp																								27
Predefined	.cp)																						28
Predefined	.h																							28

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Chapter 4

Class Documentation

4.1 AHC Class Reference

#include <AHC.h>

Collaboration diagram for AHC:

Public Member Functions

- AHC ()
- AHC (const int &argc, char **argv)
- ∼AHC ()
- void performClustering ()

Private Member Functions

- void extractFeatures (const std::vector < int > &storage, const std::vector < std::vector < int > > &neighbor ← Vec, const Eigen::MatrixXf ¢roid)
- void setDataset (const int &argc, char **argv)
- const float getDistAtNodes (const vector< int > &firstList, const vector< int > &secondList, const int &Linkage)
- void hierarchicalMerging (std::unordered_map< int, Ensemble > &node_map, std::vector< DistNode > &d↔
 NodeVec, std::vector< Ensemble > &nodeVec)
- void setLabel (const std::vector< Ensemble > &nodeVec, vector< vector< int > > &neighborVec, vector< int > &storage, Eigen::MatrixXf ¢roid)
- string getLinkageStr ()
- void getEntropyRatio (const std::vector< int > &storage, float &EntropyRatio)
- string getNormStr ()
- string getEntropyStr (const float &EntropyRatio)
- void setValue_merge (std::vector < DistNode > &dNodeVec, std::unordered_map < int, Ensemble > &node ← _map)
- void setValue (std::vector < DistNode > &dNodeVec, std::unordered map < int, Ensemble > &node map)
- void performClustering_by_norm ()

Private Attributes

- MetricPreparation object
- int normOption
- bool isPBF
- std::vector< int > group
- std::vector < string > activityList
- std::vector< string > timeList
- DataSet ds
- int numberOfClusters
- · int initializationOption
- int linkageOption
- bool IMethod
- bool readCluster
- bool isPathlines
- std::vector< float > curveValue [4]

4.1.1 Detailed Description

Definition at line 36 of file AHC.h.

4.1.2 Constructor & Destructor Documentation

```
4.1.2.1 AHC::AHC()
```

Definition at line 12 of file AHC.cpp.

```
13 {
14
15 }
```

4.1.2.2 AHC::AHC (const int & argc, char ** argv)

Definition at line 26 of file AHC.cpp.

4.1.2.3 AHC:: \sim AHC()

Definition at line 38 of file AHC.cpp.

```
39 {
40          deleteDistanceMatrix(ds.dataMatrix.rows());
41 }
```

4.1.3 Member Function Documentation

4.1.3.1 void AHC::extractFeatures (const std::vector< int > & storage, const std::vector< std::vector< int > > & neighborVec, const Eigen::MatrixXf & centroid) [private]

Definition at line 473 of file AHC.cpp.

```
475 {
        const int& Row = ds.dataMatrix.rows();
476
477
       const int& Column = ds.dataMatrix.cols();
478
479
        std::cout << "Final group number information: " << std::endl;</pre>
480
        for (int i = 0; i < storage.size(); ++i)</pre>
481
482
            std::cout << storage[i] << " ";
483
484
       std::cout << std::endl;
485
486
        /* record labeling information */
487
       // IOHandler::generateGroups(neighborVec);
488
489
       IOHandler::printClusters(ds.dataVec, group, storage, "norm"+to_string(
      normOption), ds.fullName, ds.dimension);
490
491
        struct timeval start, end;
492
       double timeTemp;
493
        /\star compute the centroid coordinates of each clustered group \star/
494
495
496
       gettimeofday(&start, NULL);
497
498
       vector<vector<float> > closest(numberOfClusters);
499
       vector<vector<float> > furthest(numberOfClusters);
500
        /* extract the closest and furthest streamlines to centroid */
501
502 #pragma omp parallel for schedule(static) num_threads(8)
       for (int i=0;i<numberOfClusters;++i)</pre>
503
504
505
            float minDist = FLT_MAX;
506
            float maxDist = -10;
            int minIndex = -1, maxIndex = -1;
507
            const std::vector<int>& groupRow = neighborVec[i];
508
509
           const Eigen::VectorXf& eachCentroid = centroid.row(i);
510
            for (int j = 0; j < groupRow.size(); ++j)
511
512
                float distance = getDisimilarity(eachCentroid, ds.dataMatrix, groupRow[j],
     normOption, object);
513
                if (minDist>distance)
514
                {
515
                    minDist = distance;
516
                    minIndex = groupRow[j];
517
518
                if (maxDist<distance)
519
                {
520
                    maxDist = distance;
521
                    maxIndex = groupRow[j];
522
523
            closest[i] = ds.dataVec[minIndex];
524
            furthest[i] = ds.dataVec[maxIndex];
525
526
527
528
        // re-assign centroid coordinates to the vector<vector<float>>
529
       std::vector<std::vector<float> > center_vec(numberOfClusters, vector<float>(Column));
532
533
            for (int j = 0; j < Column; ++j)
534
535
                center_vec[i][j] = centroid(i,j);
536
537
538
539
        // calculate the normalized entropy
540
        float EntropyRatio;
541
       getEntropyRatio(storage, EntropyRatio);
542
        std::cout << "Entropy ratio is: " << EntropyRatio << std::endl;</pre>
543
544
545
        // record the time for feature extraction
546
       gettimeofday(&end, NULL);
```

```
547
548
      activityList.push_back("Feature extraction for norm "+to_string(
normOption)+ " takes: ");
549
550
        timeList.push_back(to_string(timeTemp)+" s");
551
552
          / calculate the normalized validity measurement
553
        ValidityMeasurement vm;
554
         vm.computeValue(normOption, ds.dataMatrix, group, object,
      isPBF);
555
        activityList.push_back("AHC Validity measure is: ");
556
         stringstream fc_ss;
557
        fc_ss << vm.f_c;
558
        timeList.push_back(fc_ss.str());
559
560
         // calculate the silhouette, the Gamma statistics and DB index
561
         gettimeofday(&start, NULL);
562
        Silhouette sil;
563
         sil.computeValue(normOption, ds.dataMatrix, ds.
      dataMatrix.rows(),ds.dataMatrix.cols(),
564
                            group, object, numberOfClusters,
       isPBF):
565
         gettimeofday(&end, NULL);
        566
567
        activityList.push_back("Silhouette calculation for norm " +to_string(
568
      normOption) + " takes: ");
569
        timeList.push_back(to_string(timeTemp)+" s");
570
571
        std::cout << "Finishing extracting features!" << std::endl;</pre>
572
573
        stringstream ss;
574
        ss << "norm_" << normOption;
575
        /* measure closest and furthest rotation */
std::vector<float> closestRotation, furthestRotation;
576
577
        const float& closestAverage = getRotation(closest, closestRotation);
578
579
        const float& furthestAverage = getRotation(furthest, furthestRotation);
580
581
         // record the linkage type, norm option and normalized entropy
        string linkage = getLinkageStr();
string normStr = getNormStr();
582
583
584
         string entropyStr = getEntropyStr(EntropyRatio);
585
        // create the .vtk for streamline labels and cluster representatives
IOHandler::printFeature(ds.dataName+"_AHC_"+linkage+"_closest_"+ss.str()+".vtk", closest, sil
586
587
       .sCluster,
        closestRotation, ds.dimension);
IOHandler::printFeature(ds.dataName+"_AHC_"+linkage+"_furthest_"+ss.str()+".vtk", furthest,
588
589
      sil.sCluster,
         furthestRotation, ds.dimension);
IOHandler::printFeature(ds.dataName+"_AHC_"+linkage+"_centroid_"+ss.str()+".vtk", center_vec,
590
591
        sil.sCluster,ds.dimension);
592
        IOHandler::printToFull(ds.dataVec, sil.sData, "AHC_SValueLine_"+ss.str(),
593
      ds.fullName, ds.dimension);
   IOHandler::printToFull(ds.dataVec, group, sil.sCluster, "AHC_SValueCluster_"+ss.str(),
594
      ds.fullName, ds.dimension);
595
596
         // generate README for evaluation metrics
         activityList.push_back("numCluster is: ");
597
598
        timeList.push_back(to_string(numberOfClusters));
599
600
         activityList.push_back("Average Silhouette is: ");
601
         timeList.push_back(to_string(sil.sAverage));
602
603
         activityList.push_back("Average rotation of closest is: ");
604
         timeList.push_back(to_string(closestAverage));
605
606
         activityList.push_back("Average rotation of furthest is: ");
607
         timeList.push_back(to_string(furthestAverage));
608
         IOHandler::generateReadme(activityList,timeList);
IOHandler::writeReadme("Linkage: "+linkage+", "+"norm option is "+normStr+", ");
609
610
         IOHandler::writeGroupSize(storage);
611
612
         /* print entropy value for the clustering algorithm */
IOHandler::writeReadme(EntropyRatio, sil, "For norm "+to_string(normOption));
613
614
615
         IOHandler::writeReadme(closestAverage, furthestAverage);
616
         //curveValue[0].push_back(sil.sAverage);
617
         //curveValue[1].push_back(sil.gammaStatistic);
//curveValue[2].push_back(sil.dbIndex);
618
619
620
         //curveValue[3].push_back(vm.f_c);
621 }
```

4.1.3.2 const float AHC::getDistAtNodes (const vector < int > & firstList, const vector < int > & secondList, const int & Linkage) [private]

Definition at line 722 of file AHC.cpp.

```
723 {
        const int& m = firstList.size();
const int& n = secondList.size();
724
725
726
        assert (m!=0):
727
        assert(n!=0);
728
        /* 0: single linkage, min(x_i,y_j)
729
         * 1: complete linkdage, max(x_i,y_j)
730
        * 2: average linkage, sum/x_i/y_j
731
732
        float result, value;
733
        switch (Linkage)
734
735
        case 0: //single linkage
736
            {
737
                result = FLT_MAX;
             #pragma omp parallel for reduction(min:result) num_threads(8)
738
                 for (int i=0; i < m; ++i)</pre>
739
740
741
                     for (int j=0; j<n; ++j)</pre>
742
743
                         if(distanceMatrix)
                             value = distanceMatrix[firstList[i]][secondList[j]];
744
745
                         else
746
                              value = getDisimilarity(ds.dataMatrix, firstList[i], secondList[j],
      normOption, object);
747
                         result = std::min(result, value);
748
749
                 }
750
751
            break;
752
753
        case 1: //complete linkage
754
755
                result = -1.0;
            756
757
758
759
                     for(int j=0;j<n;++j)</pre>
760
761
                         if(distanceMatrix)
                             value = distanceMatrix[firstList[i]][secondList[j]];
762
763
                         else
764
                             value = getDisimilarity(ds.dataMatrix, firstList[i], secondList[j],
      normOption, object);
765
                         result = std::max(result, value);
766
767
                }
768
769
            break;
770
771
        case 2: // average linkage
772
773
                result = 0;
             #pragma omp parallel for reduction(+:result) num_threads(8)
774
775
                 for (int i=0; i < m; ++i)</pre>
776
777
                     for(int j=0;j<n;++j)</pre>
778
779
                         if (distanceMatrix)
780
                             value = distanceMatrix[firstList[i]][secondList[j]];
781
                         else
782
                              value = getDisimilarity(ds.dataMatrix, firstList[i], secondList[j],
      normOption, object);
783
                         result+=value;
784
                     }
785
                result/=m*n;
786
787
788
            break;
789
            ault: // no linkage option, should print "Error" information and exit the program std::cout << "error!" << std::endl;
790
791
792
            exit(1):
793
        return result;
795 }
```

4.1.3.3 void AHC::getEntropyRatio (const std::vector < int > & storage, float & EntropyRatio) [private]

Definition at line 829 of file AHC.cpp.

```
830 {
831     EntropyRatio = 0;
832     const int& Row = ds.dataMatrix.rows();
833     for (int i = 0; i < storage.size(); ++i)
834     {
835          float ratio = float(storage[i])/float(Row);
836          EntropyRatio-=ratio*log2f(ratio);
837     }
838     EntropyRatio/=log2f(storage.size());
839 }</pre>
```

4.1.3.4 string AHC::getEntropyStr (const float & EntropyRatio) [private]

Definition at line 860 of file AHC.cpp.

4.1.3.5 string AHC::getLinkageStr() [private]

Definition at line 802 of file AHC.cpp.

```
803 {
804
         string result;
switch(linkageOption)
805
806
807
         case 0:
             result = "single";
808
809
             break;
810
811
         case 1:
812
            result = "complete";
813
             break;
814
815
         case 2:
             result = "average";
816
817
             break;
818
         return result;
820 }
```

4.1.3.6 string AHC::getNormStr() [private]

Definition at line 846 of file AHC.cpp.

```
847 {
848 stringstream ss;
849 ss << normOption;
850 return ss.str();
851 }
```

4.1.3.7 void AHC::hierarchicalMerging (std::unordered_map< int, Ensemble > & node_map, std::vector< DistNode > & dNodeVec, std::vector< Ensemble > & nodeVec) [private]

Definition at line 270 of file AHC.cpp.

```
272 {
273
        std::map<int, float> dist map;
274
275
        /\star would store distance matrix instead because it would save massive time \star/
276
        struct timeval start, end;
277
        double timeTemp;
278
        gettimeofday(&start, NULL);
279
280
        const int Row = ds.dataMatrix.rows();
281
282
        DistNode poped;
283
        /* find node-pair with minimal distance */
284
        float minDist = FLT_MAX;
int target = -1;
285
286
287
        for (int i = 0; i < dNodeVec.size(); ++i)</pre>
288
289
            if (dNodeVec[i].distance<minDist)</pre>
290
291
                target = i;
292
                minDist = dNodeVec[i].distance;
293
294
295
        // find which distNode is to be popped
296
297
        poped = dNodeVec[target];
298
299
        int index = Row, currentNumber;
300
        301
302
            if (lMethod) // if the l-method is enabled, record the number of clusters and merged
       distance
303
            {
304
                dist_map.insert(std::make_pair(node_map.size(), poped.distance));
305
306
            //create new node merged and input it into hash unordered_map
307
            vector<int> first = (node_map[poped.first]).element;
            vector<int> second = (node_map[poped.second]).element;
308
309
310
            /\star index would be starting from Row \star/
311
            Ensemble newNode(index);
312
            newNode.element = first;
313
            newNode.element.insert(newNode.element.end(), second.begin(), second.end());
314
            node_map.insert(make_pair(index, newNode));
315
316
            //delete two original nodes
317
            node_map.erase(poped.first);
318
            node_map.erase(poped.second);
319
320
            /* the difficulty lies how to update the min-heap with linkage
321
             * This would take 2NlogN.
322
             * Copy all node-pairs that are not relevant to merged nodes to new vec.
323
             * For relevant, would update the mutual distance by linkage
324
325
326
            /* how many clusters exist */
327
            currentNumber = node_map.size();
328
329
            target = -1, minDist = FLT_MAX;
330
331
            // create new distNode vector
332
            std::vector<DistNode> tempVec(currentNumber*(currentNumber-1)/2);
333
            // update and find the minimal distance for next merging
334
335
            int current = 0, i_first, i_second;
336
            for (int i=0;i<dNodeVec.size();++i)</pre>
337
338
                \verb|i_first=dNodeVec[i].first|, | \verb|i_second=dNodeVec[i].second|; \\
                /* not relevant, directly copied to new vec */
if(i_first!=poped.first&&i_first!=poped.second&&i_second!=poped.
339
340
      first&&i_second!=poped.second)
341
342
                     tempVec[current] = dNodeVec[i];
343
                     if(tempVec[current].distance<minDist)</pre>
344
345
                         target = current;
346
                         minDist = tempVec[current].distance;
347
```

```
348
                      ++current;
349
                 }
350
             }
351
352
             // merge two nodes and update the node-distance relative to these two nodes
353
             for (auto iter=node_map.begin();iter!=node_map.end();++iter)
354
355
                 if((*iter).first!=newNode.index)
356
                      tempVec[current].first = (*iter).first;
tempVec[current].second = newNode.index;
357
358
359
                      \texttt{tempVec[current].distance=} \\ \texttt{getDistAtNodes} \\ (\texttt{newNode.element,(*iter).second.} \\
      element, linkageOption);
360
                      if(tempVec[current].distance<minDist)</pre>
361
                          target = current;
minDist = tempVec[current].distance;
362
363
364
365
                      ++current;
366
                 }
367
            }
368
369
             if(target>=0 && tempVec.size()>=1)
370
371
                 poped = tempVec[target];
372
373
                 /\star judge whether current is assigned to right value \star/
374
                 assert(current==tempVec.size());
375
                 dNodeVec.clear();
376
                 dNodeVec = tempVec;
377
                 tempVec.clear();
378
                 ++index;
379
380
381
         }while(node_map.size()!=numberOfClusters); //merging happens whenever requested
       cluster is not met
382
383
        if (lMethod) // invoke the l-method to find the optimal number of clusters
384
385
             /\star perform L-method computation to detect optimal number of AHC \star/
386
             DetermClusterNum dcn;
387
             dcn.iterativeRefinement(dist_map);
             std::cout << "Otimal number of clusters by L-Method is " << dcn.getFinalNumOfClusters() <<
388
      std::endl;
389
            dcn.recordLMethodResult(normOption);
390
391
392
        else
               // otherwise, just perform the AHC clustering
393
394
             nodeVec=std::vector<Ensemble>(node map.size());
395
             int tag = 0;
396
             for(auto iter=node_map.begin();iter!=node_map.end();++iter)
397
                 nodeVec[tag++]=(*iter).second;
398
             gettimeofday(&end, NULL);
399
            timeTemp = ((end.tv_sec - start.tv_sec) * 1000000u + end.tv_usec - start.tv_usec) / 1.e6;
400
             activityList.push_back("Hirarchical clustering of norm "+to_string(
402
      normOption) + " for "+
403
                                      to_string(numberOfClusters) +" groups takes: ");
             timeList.push_back(to_string(timeTemp)+" s");
404
405
             /* task completed, would delete memory contents */
406
             dNodeVec.clear();
407
            node_map.clear();
408
             /st use alpha function to sort the group by its size in ascending order st/
409
             std::sort(nodeVec.begin(), nodeVec.end(), [](const Ensemble& e1, const
      Ensemble& e2)
410
            {return el.element.size() < e2.element.size() | (e1.element.size() == e2.element.size() &&</pre>
      e1.index<e2.index);});
411
412 }
```

4.1.3.8 void AHC::performClustering ()

Definition at line 165 of file AHC.cpp.

```
166 {
167    /* 0: Euclidean Norm
168    1: Fraction Distance Metric
169    2: piece-wise angle average
```

```
3: Bhattacharyya metric for rotation
171
             4: average rotation
172
            5: signed-angle intersection
173
             6: normal-direction multivariate distribution
            7: Bhattacharyya metric with angle to a fixed direction 8: Piece-wise angle average \times standard deviation 9: normal-direction multivariate un-normalized distribution
174
175
176
177
             10: x*y/|x||y| borrowed from machine learning
178
            11: cosine similarity
179
            12: Mean-of-closest point distance (MCP)
180
             13: Hausdorff distance min_max(x_i,y_i)
             14: Signature-based measure from http://ieeexplore.ieee.org/stamp/stamp.jsp?tp=&arnumber=6231627
181
             15: Procrustes distance take from http://ieeexplore.ieee.org/stamp/stamp.jsp?tp=&arnumber=6787131
182
             16: entropy-based distance metric taken from http://vis.cs.ucdavis.edu/papers/pg2011paper.pdf
183
184
             17: time-series MCP distance from https://www.sciencedirect.com/science/article/pii/
      S0097849318300128
185
                 for pathlines only
186
187
188
        // read the input number for different similarity measures from the file "cluster_number"
        std::unordered_map<int,int> clusterMap;
189
190
        if(readCluster)
191
            IOHandler::readClusteringNumber(clusterMap, "cluster number");
192
193
194
        // std::vector<int> cluster_array;
195
196
        //for(int i=2; i<=100; ++i)
197
             //cluster_array.push_back(i);
198
             for (normOption=0; normOption<=17; ++normOption)</pre>
199
200
201
                 if(isPathlines) // for pathlines, will consider d_T (17)
202
203
                     if(normOption!=0 && normOption!=1 &&
      normOption!=2 && normOption!=4 && normOption!=12
                        && normOption!=13 && normOption!=14 &&
204
      normOption!=15 && normOption!=17)
205
                         continue:
206
207
                 else
                       // for streamlines, will not consider d_T (17)
208
                 {
                     if (normOption!=0 && normOption!=1 &&
209
      normOption!=2 && normOption!=4 && normOption!=12
                     && normOption!=13 && normOption!=14 &&
210
      normOption!=15)
211
                         continue:
212
213
                                                                                        ----" << std::endl;
                 std::cout << "----
                 std::cout << "norm " << normOption << " starts....." << std::endl;
214
215
                 timeList.clear();
216
                 activityList.clear();
217
218
                 /\star L-method is not performed. It's a normal AHC procedure \star/
219
                 if(!lMethod)
220
                 {
221
                     const int& Row = ds.dataMatrix.rows();
222
                     if(readCluster)
223
224
                         numberOfClusters = clusterMap[normOption];
225
                     }
226
                     else
227
                     {
                         std::cout << "-----
228
                                                                           ----" << std::endl;
                         std::cout << "Input cluster number among [0, " << Row << "] for norm " <<
229
      normOption << ": ";</pre>
230
                         std::cin >> numberOfClusters;
231
                         assert(numberOfClusters>0 && numberOfClusters<Row);
232
233
                     //numberOfClusters = i;
234
                     assert(numberOfClusters>0 && numberOfClusters<Row);</pre>
235
236
                 /\star perform L-method for detecting optimal num of clusters \star/
                 else if(lMethod)
237
238
                 {
239
                     numberOfClusters = 1:
240
241
                 // perform clustering by given input of norm option
242
                 performClustering_by_norm();
243
        //}
244
245
246
        /*std::ofstream curve("../dataset/curveValue.txt", ios::out);
247
        for (int i=0; i<4; ++i)
248
249
             for(int j=0; j<curveValue[0].size(); ++j)</pre>
250
```

4.1.3.9 void AHC::performClustering_by_norm() [private]

Definition at line 53 of file AHC.cpp.

```
55
        /\star very hard to decide whether needed to perform such pre-processing, but still create a
56
        *\ \texttt{MetricPreparation object in case some pre-calculation for similarity measures can be ready}
57
        * before the pairwise distance matrix calculation
58
59
        object = MetricPreparation(ds.dataMatrix.rows(), ds.dataMatrix.cols());
        object.preprocessing(ds.dataMatrix, ds.dataMatrix.rows(),
60
       ds.dataMatrix.cols(), normOption);
61
62
        /\star would store distance matrix instead because it would save massive time \star/
        struct timeval start, end;
63
        double timeTemp;
64
65
       gettimeofday(&start, NULL);
66
       // check whether the file for distance matrix exists or not
std::ifstream distFile(("../dataset/"+to_string(normOption)).c_str(), ios::in);
if(distFile.fail()) // not exist, will calculate the distance matrix and store them in local files
67
68
69
70
            distFile.close();
72
            // calculate the distance matrix
73
            getDistanceMatrix(ds.dataMatrix, normOption, object);
            // store the distance matrix values in the local files
std::ofstream distFileOut(("../dataset/"+to_string(normOption)).c_str(), ios::out);
74
7.5
76
            for(int i=0;i<ds.dataMatrix.rows();++i)</pre>
77
78
                 for(int j=0; j<ds.dataMatrix.rows();++j)</pre>
79
                      distFileOut << distanceMatrix[i][j] << " ";</pre>
80
81
                 distFileOut << std::endl;
82
83
84
            distFileOut.close();
85
86
        else
                 // the file for distance matrix exists, then directly reads in the pair-wise values
87
88
            std::cout << "read distance matrix..." << std::endl;
89
            distanceMatrix = new float*[ds.dataMatrix.rows()];
91
        #pragma omp parallel for schedule(static) num_threads(8)
92
            for (int i = 0; i < ds.dataMatrix.rows(); ++i)</pre>
93
94
                 distanceMatrix[i] = new float[ds.dataMatrix.rows()];
95
            }
97
             // read the distance values from the .txt file
98
            int i=0, j;
99
            string line;
100
             stringstream ss:
101
             while (getline (distFile, line))
102
             {
103
                  j=0;
104
                  ss.str(line);
105
                  while(ss>>line)
106
107
                       if(i==j)
108
                           distanceMatrix[i][j]=0;
109
110
                           distanceMatrix[i][j] = std::atof(line.c_str());
111
                       ++j;
112
                  ++i;
113
                  ss.str("");
114
115
                  ss.clear();
116
117
              distFile.close();
118
119
120
         // record the time for distance matrix computation time
121
         gettimeofday(&end, NULL);
```

```
122
123
     activityList.push_back("Distance matrix computing for norm "+to_string(
normOption)+" takes: ");
124
125
       timeList.push_back(to_string(timeTemp)+" s");
126
127
        // create node-related parameters for AHC clustering
128
       std::unordered_map<int, Ensemble> node_map;
129
        std::vector<DistNode> dNodeVec;
130
       std::vector<Ensemble> nodeVec;
131
        /* set the ditNode vector */
132
133
       setValue_merge(dNodeVec, node_map);
134
135
        /\star perform hiarchical clustering where within each step would merge two nodes \star/
136
       hierarchicalMerging(node_map, dNodeVec, nodeVec);
137
        if(!lMethod) // perform the AHC clustering with lMethod not activated
138
139
140
            vector<vector<int>> neighborVec(numberOfClusters);
141
            // element size for all groups
142
            vector<int> storage(numberOfClusters);
143
144
            // geometric center
145
           Eigen::MatrixXf centroid = Eigen::MatrixXf::Zero(numberOfClusters,
     ds.dataMatrix.cols());
146
147
            // set label information
148
           setLabel(nodeVec, neighborVec, storage, centroid);
149
150
           nodeVec.clear();
151
152
            extractFeatures(storage, neighborVec, centroid);
153
154 }
```

4.1.3.10 void AHC::setDataset (const int & argc, char ** argv) [private]

Definition at line 634 of file AHC.cpp.

```
635 {
636
        if(argc!=3)
637
638
             std::cout << "Input argument should have 3!" << endl
                       << "./cluster inputFile_name(in dataset folder) "
<< "data_dimension(3)" << endl;</pre>
639
640
641
            exit(1);
642
643
        // get the attribute for data set
644
        ds.strName = string("../dataset/")+string(argv[1]);
        ds.dataName = string(argv[1]);
ds.dimension = atoi(argv[2]);
645
646
647
648
        /* get the bool tag for isPBF */
649
        std::cout << "It is a PBF dataset? 1.Yes, 0.No" << std::endl;
650
        int PBFjudgement;
651
        std::cin >> PBFjudgement;
        assert(PBFjudgement==1||PBFjudgement==0);
652
        isPBF = (PBFjudgement==1);
653
654
655
        /* get the bool tag for isPBF */
656
        std::cout << "It is a pathline dataset? 1.Yes, 0.No" << std::endl;</pre>
657
        std::cin >> PBFjudgement;
658
        assert(PBFjudgement==1||PBFjudgement==0);
        isPathlines = (PBFjudgement==1);
659
660
661
        // set up the sample option by user input and data set type (pathlines or streamlines)
662
        int sampleOption;
663
        if(isPathlines) // default direct-repeating for pathlines to match the time steps
            sampleOption = 1;
664
665
        else
               // streamline sample option can be versatile
666
667
            std::cout << "choose a sampling method for the dataset?" << std::endl</pre>
                       << "1.directly filling with last vertex; 2. uniform sampling." << std::endl;
668
669
             std::cin >> sampleOption;
670
            assert(sampleOption==1||sampleOption==2);
671
672
673
        // read the coordinates into the member variales
        IOHandler::readFile(ds.strName, ds.dataVec, ds.vertexCount,
```

```
ds.dimension,ds.maxElements);
675
676
        // get the path of full name and print the streamlines vtk
        ds.fullName = ds.strName+"_full.vtk";
IOHandler::printVTK(ds.fullName, ds.dataVec, ds.
677
678
      vertexCount, ds.dimension);
679
680
          perform sampling operation with user parameters
681
        if (sampleOption==1)
682
            IOHandler::expandArray(ds.dataMatrix,ds.dataVec,ds.
      dimension, ds.maxElements);
683
      else if(sampleOption==2)
            IOHandler::sampleArray(ds.dataMatrix, ds.dataVec, ds.
684
      dimension, ds.maxElements);
685
686
        // create the label index for each individual streamline/pathline
        group = std::vector<int>(ds.dataMatrix.rows());
687
688
689
        // whether to activate the L-method or not
        std::cout << "Perform L-method to detect optimal num of clusters? 0: No, 1: Yes! " << std::endl;
690
691
        std::cin >> lMethod;
692
        assert(lMethod==0 || lMethod==1);
        lMethod = (lMethod==1);
693
694
695
        // which linkage type to be selected
                                                ---" << std::endl;
696
        std::cout << "
697
        std::cout << "Input linkage option: 0.single linkage, 1.complete linkage, 2.average linkage" <</pre>
      std::endl;
698
        std::cin >> linkageOption;
        assert(linkageOption==0||linkageOption==1||linkageOption==2);
699
700
701
           lMethod is not activated, so will ask for number of cluster as input
702
703
704
            std::cout << "-----" << std::endl;
            std::cout << "Choose cluster number input method: 0.user input, 1.read from file: " << std::endl;
705
706
            int clusterInput;
            std::cin >> clusterInput;
708
            assert(clusterInput==0||clusterInput==1);
709
            readCluster = (clusterInput==1);
710
711 }
```

4.1.3.11 void AHC::setLabel (const std::vector< Ensemble > & nodeVec, vector< vector< int > & storage, Eigen::MatrixXf & centroid) [private]

Definition at line 426 of file AHC.cpp.

```
429 // group tag by increasing order
430
        int groupID = 0;
431
        // element list for each group
432
433
        vector<int> eachContainment;
434
435
        // find group id and neighboring vec
436
        for(auto iter = nodeVec.begin(); iter!=nodeVec.end();++iter)
437
438
            eachContainment = (*iter).element;
            neighborVec[groupID] = eachContainment;
439
        #pragma omp parallel num_threads(8)
441
442
            #pragma omp for nowait
443
                for(int i=0;i<eachContainment.size();++i)</pre>
444
445
                    // update the label index for each streamline candidates
446
                    group[eachContainment[i]] = groupID;
                #pragma omp critical
448
                    // update the centroid coordinates of the cluster
449
                    centroid.row(groupID) += ds.dataMatrix.row(eachContainment[i]);
450
                }
451
452
            storage[groupID] = (*iter).element.size();
453
            centroid.row(groupID)/=eachContainment.size();
454
            ++groupID;
455
            eachContainment.clear();
456
457 }
```

4.1.3.12 void AHC::setValue (std::vector < DistNode > & dNodeVec, std::unordered_map < int, Ensemble > & node_map) [private]

Definition at line 955 of file AHC.cpp.

```
956 {
957
        const int& Row = ds.dataMatrix.rows();
958
        dNodeVec = std::vector<DistNode>(Row*(Row-1)/2);
959
        int tag = 0;
960
        // record the node i, node j and their distance into the vector
        for (int i=0; i < Row-1; ++i)</pre>
961
962
963
            for (int j=i+1; j<Row; ++j)</pre>
964
965
                dNodeVec[tag].first = i;
966
                dNodeVec[tag].second = j;
967
                if (distanceMatrix)
                    dNodeVec[tag].distance = distanceMatrix[i][j];
968
969
970
                    dNodeVec[tag].distance = getDisimilarity(ds.dataMatrix, i, j,
      normOption, object);
971
                ++tag;
972
973
974
        assert(tag==dNodeVec.size());
975
        // record the index of the node
976
        for (int i=0; i < Row; ++i)
977
978
            node_map[i].element.push_back(i);
979
980 }
```

4.1.3.13 void AHC::setValue_merge (std::vector < DistNode > & dNodeVec, std::unordered_map < int, Ensemble > & node_map) [private]

Definition at line 874 of file AHC.cpp.

```
875 {
876
        const int& Row = ds.dataMatrix.rows();
877
878
         /* find the node of closest distance */
879
        std::vector<int> miniNode(Row);
880 #pragma omp parallel for schedule(static) num_threads(8)
881
         for (int i=0; i < Row; ++i)</pre>
882
883
             float miniDist = FLT_MAX, dist;
             int index = -1;
for(int j=0; j<Row; ++j)</pre>
884
885
886
             {
887
                 if(i==j)
888
                      continue;
889
                 if(distanceMatrix)
890
                     dist = distanceMatrix[i][j];
891
                 else
                     dist = getDisimilarity(ds.dataMatrix, i, j,
892
      normOption, object);
893
894
                 if(miniDist>dist)
895
                      miniDist=dist;
896
897
                      index=j;
898
899
900
             miniNode[i]=index;
901
902
903
        std::vector<bool> isIn(Row, false);
904
905
         // update the map for node
906
        int tag = 0;
        for (int i=0; i < Row; ++i)</pre>
907
908
909
             if(!isIn[i])
910
911
                 Ensemble en;
```

```
if (miniNode[miniNode[i]]==i)
914
                     en.element.push_back(i);
                     en.element.push_back(miniNode[i]);
915
916
                     isIn[i]=true;
                     isIn[miniNode[i]]=true;
917
918
                    node_map[tag] = en;
919
920
                else
921
922
                     en.element.push_back(i);
                     isIn[i]=true;
923
                    node_map[tag] = en;
924
925
926
                 ++tag;
927
            }
928
929
930
       // update the dNodeVec from the newly create nodes
931
        const int& mapSize = node_map.size();
932
        dNodeVec = std::vector<DistNode>(mapSize*(mapSize-1)/2);
933
934
        tag = 0;
935
        for(auto start = node_map.begin(); start!=node_map.end(); ++start)
936
937
            for(auto end = node_map.begin(); end!=node_map.end() && end!=start; ++end)
938
                dNodeVec[tag].first = start->first;
dNodeVec[tag].second = end->first;
939
940
                dNodeVec[tag].distance = getDistAtNodes(start->second.element,end->second.element
941
      , linkageOption);
942
                ++tag;
943
944
945
        assert(tag==dNodeVec.size());
946 }
```

4.1.4 Member Data Documentation

4.1.4.1 std::vector<**string**> **AHC::activityList** [private]

Definition at line 122 of file AHC.h.

4.1.4.2 std::vector<float> AHC::curveValue[4] [private]

Definition at line 167 of file AHC.h.

4.1.4.3 DataSet AHC::ds [private]

Definition at line 132 of file AHC.h.

4.1.4.4 std::vector<int> AHC::group [private]

Definition at line 117 of file AHC.h.

4.1.4.5 int AHC::initializationOption [private]

Definition at line 142 of file AHC.h.

```
4.1.4.6 bool AHC::isPathlines [private]
Definition at line 162 of file AHC.h.
4.1.4.7 bool AHC::isPBF [private]
Definition at line 112 of file AHC.h.
4.1.4.8 int AHC::linkageOption [private]
Definition at line 147 of file AHC.h.
4.1.4.9 bool AHC:: Method [private]
Definition at line 152 of file AHC.h.
4.1.4.10 int AHC::normOption [private]
Definition at line 107 of file AHC.h.
4.1.4.11 int AHC::numberOfClusters [private]
Definition at line 137 of file AHC.h.
4.1.4.12 MetricPreparation AHC::object [private]
Definition at line 102 of file AHC.h.
4.1.4.13 bool AHC::readCluster [private]
Definition at line 157 of file AHC.h.
4.1.4.14 std::vector<string> AHC::timeList [private]
Definition at line 127 of file AHC.h.
The documentation for this class was generated from the following files:
```

Generated by Doxygen

AHC.hAHC.cpp

4.2 DataSet Struct Reference

#include <Predefined.h>

Public Attributes

- vector< vector< float > > dataVec
- Eigen::MatrixXf dataMatrix
- int maxElements = -1
- int vertexCount = -1
- int dimension = -1
- string strName
- string fullName
- string dataName

4.2.1 Detailed Description

Definition at line 20 of file Predefined.h.

4.2.2 Member Data Documentation

4.2.2.1 Eigen::MatrixXf DataSet::dataMatrix

Definition at line 23 of file Predefined.h.

4.2.2.2 string DataSet::dataName

Definition at line 30 of file Predefined.h.

4.2.2.3 vector<vector<float> > DataSet::dataVec

Definition at line 22 of file Predefined.h.

4.2.2.4 int DataSet::dimension = -1

Definition at line 26 of file Predefined.h.

4.2.2.5 string DataSet::fullName

Definition at line 29 of file Predefined.h.

```
4.2.2.6 int DataSet::maxElements = -1
```

Definition at line 24 of file Predefined.h.

4.2.2.7 string DataSet::strName

Definition at line 28 of file Predefined.h.

4.2.2.8 int DataSet::vertexCount = -1

Definition at line 25 of file Predefined.h.

The documentation for this struct was generated from the following file:

· Predefined.h

4.3 DistNode Struct Reference

```
#include <Predefined.h>
```

Public Member Functions

- DistNode (const int &first, const int &second, const float &dist)
- DistNode ()

Public Attributes

```
• int first = -1
```

- int second = -1
- float distance = -1.0

4.3.1 Detailed Description

Definition at line 69 of file Predefined.h.

4.3.2 Constructor & Destructor Documentation

4.3.2.1 DistNode::DistNode (const int & first, const int & second, const float & dist) [inline]

Definition at line 74 of file Predefined.h.

```
74 :first(
    first), second(second), distance(dist)
75 {}
```

```
4.3.2.2 DistNode::DistNode( ) [inline]
```

Definition at line 77 of file Predefined.h.

```
78 {}
```

4.3.3 Member Data Documentation

```
4.3.3.1 float DistNode::distance = -1.0
```

Definition at line 72 of file Predefined.h.

```
4.3.3.2 int DistNode::first = -1
```

Definition at line 71 of file Predefined.h.

```
4.3.3.3 int DistNode::second = -1
```

Definition at line 71 of file Predefined.h.

The documentation for this struct was generated from the following file:

• Predefined.h

4.4 Ensemble Struct Reference

```
#include <Predefined.h>
```

Public Member Functions

- Ensemble (const int &index)
- Ensemble ()

Public Attributes

- int index = -1
- std::vector< int > element

4.4.1 Detailed Description

Definition at line 38 of file Predefined.h.

4.4.2 Constructor & Destructor Documentation

4.4.2.1 Ensemble::Ensemble (const int & index) [inline]

Definition at line 45 of file Predefined.h.

```
45 : index(index)
46 {}
```

4.4.2.2 Ensemble::Ensemble() [inline]

Definition at line 48 of file Predefined.h.

```
49 {}
```

4.4.3 Member Data Documentation

4.4.3.1 std::vector<int> Ensemble::element

Definition at line 43 of file Predefined.h.

4.4.3.2 int Ensemble::index = -1

Definition at line 40 of file Predefined.h.

The documentation for this struct was generated from the following file:

· Predefined.h

Chapter 5

File Documentation

5.1 AHC.cpp File Reference

```
#include "AHC.h"
Include dependency graph for AHC.cpp:
```

5.2 AHC.h File Reference

```
#include <unordered_set>
#include <unordered_map>
#include <map>
#include <string>
#include <algorithm>
#include "Predefined.h"
#include "ValidityMeasurement.h"
#include "DetermClusterNum.h"
Include dependency graph for AHC.h: This graph shows which files directly or indirectly include this file:
```

Classes

class AHC

5.3 main.cpp File Reference

```
#include "AHC.h"
Include dependency graph for main.cpp:
```

Functions

• int main (int argc, char **argv)

28 File Documentation

5.3.1 Function Documentation

```
5.3.1.1 int main ( int argc, char ** argv )
```

Definition at line 17 of file main.cpp.

```
18 {
19     AHC ahc(arge, argv);
20     ahc.performClustering();
21     return 0;
22 }
```

5.4 Predefined.cpp File Reference

```
#include "Predefined.h"
Include dependency graph for Predefined.cpp:
```

Functions

template < class T >
 void delete VecElements (std::vector < T > & original, const T & first, const T & second)

5.4.1 Function Documentation

5.4.1.1 template < class T > void deleteVecElements (std::vector < T > & original, const T & first, const T & second)

Definition at line 19 of file Predefined.cpp.

```
20 {
21
        std::size_t size = original.size();
        assert (size>2);
22
        vector<T> result(size-2);
        int tag = 0;
25
        for (int i=0; i < size; ++i)</pre>
26
             //meet with target elements, not copied
if(original[i]==first || original[i]==second)
2.7
28
                  continue;
             result[tag++]=original[i];
32
        assert(tag==size-2);
33
        original = result;
34 }
```

5.5 Predefined.h File Reference

```
#include "IOHandler.h"
#include "Initialization.h"
#include "Silhouette.h"
```

Include dependency graph for Predefined.h: This graph shows which files directly or indirectly include this file:

Classes

- struct DataSet
- struct Ensemble
- struct DistNode

Functions

template < class T > void deleteVecElements (std::vector < T > & origine, const T & first, const T & second)

5.5.1 Function Documentation

5.5.1.1 template < class T > void deleteVecElements (std::vector < T > & origine, const T & first, const T & second)

Definition at line 19 of file Predefined.cpp.

```
20 {
21
22
        std::size_t size = original.size();
        assert(size>2);
        vector<T> result(size-2);
int tag = 0;
for(int i=0;i<size;++i)</pre>
23
24
25
27
             //meet with target elements, not copied
             if(original[i] == first || original[i] == second)
28
2.9
             continue;
result[tag++]=original[i];
30
32
        assert(tag==size-2);
33
        original = result;
34 }
```

5.6 README.md File Reference

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