Instructions for data in S01_Example_PC

Description:

The data in S01_Example_PC is used for supporting the results "3.1 Overview of the method for recognizing SSE packing clusters" in this article, which gives the decomposition of a domain (d1atza_) into two SSE packing clusters (**Fig. 1B**).

File name	Description
d1atzaent	The PDB structure file for d1atza_
d1atza_PC1.pdb	The information and structure of SSE PC A
d1atza_PC2.pdb	The information structure of SSE PC B

The file "d1atza_PC1.pdb" not only gives the intact structure of SSE packing cluster A, but also presents the corresponding information of SSE packing cluster in the header of the file.

```
REMARK 111
            PDB Structure for a Packing Cluster.
REMARK 222
            Sites in Packing Cluster:
REMARK 222
            7,3,5,9,20,23,24,26,27,29,30,31,34,36,41,
REMARK 222
            43, 45, 47, 56, 52, 67, 71, 74, 110, 108, 112, 136, 138, 140, 160,
            164, 170, 171, 173, 174, 180, 181, 184, 15, 167, 12, 14, 81, 114,
REMARK 222
REMARK 333
            Packing Face 1: E 3,5,7,9,
REMARK 333
            Packing Face 2: H 20,23,24,26,27,29,30,31,
            Packing Face 3: E 41,43,45,47,
REMARK 333
            Packing Face 4: E 52,56,
REMARK 333
REMARK 333
            Packing Face 5: H 67,71,
REMARK 333
            Packing Face 6: E 108,110,112,
            Packing Face 7: E 136,138,140,
REMARK 333
REMARK 333
            Packing Face 8: H 170,171,173,174,
REMARK 444
            FaceInfor: SUM 8 H 3 E 5
         10 N
                  PRO A
                          2
                                  17.204
                                            6.905
                                                   41.225
                                                            1.00 19.54
ATOM
         11
             CA
                  PRO A
                          2
                                  17.879
                                            7.928
                                                   40.419
                                                            1.00 15.16
                                                                                  C
ATOM
                          2
                                  17.779
                                                                                  C
         12
             C
                  PRO A
                                            7.581
                                                   38.937
                                                            1.00 11.65
ATOM
ATOM
         13
             0
                  PRO A
                          2
                                  17.894
                                            6.414
                                                   38.554
                                                            1.00 10.50
                                                                                  0
ATOM
         14
             CB PRO A
                          2
                                  19.323
                                            7.844
                                                   40.903
                                                            1.00 16.80
                                                                                  C
                          2
         15
             \mathsf{C}\mathsf{G}
                 PRO A
                                            7.411
                                                   42.316
                                                                                  C
ATOM
                                  19.183
                                                            1.00 19.92
             CD
                 PRO A
                          2
                                  18.124
                                            6.341
                                                   42.225
                                                            1.00 19.12
                                                                                  C
ATOM
         16
                  LEU A
         17
                          3
                                  17.545
                                            8.596
ATOM
              N
                                                   38.118
                                                            1.00
                                                                  7.55
                                                                                  N
             CA LEU A
                          3
                                  17.414
                                            8.421
                                                                                  C
ATOM
         18
                                                   36.687
                                                            1.00
                                                                  6.70
         19
             C
                          3
                                  18.136
                                            9.532
                                                                                  C
ATOM
                  LEU A
                                                   35.945
                                                            1.00
                                                                  3.92
                  LEU A
                          3
                                                                                  0
         20
             0
                                  17.874
ATOM
                                           10.706
                                                   36.188
                                                            1.00
                                                                  3.73
ATOM
         21
             CB LEU A
                          3
                                  15.928
                                            8.457
                                                   36.276
                                                            1.00
                                                                  7.28
                                                                                  C
         22
             CG LEU A
                          3
                                  15.652
                                            8.527
                                                   34.761
                                                            1.00
                                                                  6.70
                                                                                  C
ATOM
                                                                                  C
                           3
                                  15.941
                                            7.179
ATOM
         23
             CD1 LEU A
                                                   34.085
                                                            1.00
                                                                  8.22
ATOM
         24
             CD2 LEU A
                                  14.206
                                            8.927
                                                   34.520
                                                            1.00
                                                                  8.60
```

REMARK 222 contains the residues involved in an SSE packing cluster. REMARK 333 contains the interfaces involved in an SSE packing cluster.