# The basic larcc module \*

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# 1 Basic representations

A few basic representation of topology are used in LARCC. They include some common sparse matrix representations: CSR (Compressed Sparse Row), CSC (Compressed Sparse Column), COO (Coordinate Representation), and BRC (Binary Row Compressed).

### 1.1 BRC (Binary Row Compressed)

We denote as BRC (Binary Row Compressed) the standard input representation of our LARCC framework. A BRC representation is an array of arrays of integers, with no requirement of equal length for the component arrays. The BRC format is used to represent a (normally sparse) binary matrix. Each component array corresponds to a matrix row, and contains the indices of columns that store a 1 value. No storage is used for 0 values.

**BRC format example** Let  $A = (a_{i,j} \in \{0,1\})$  be a binary matrix. The notation BRC(A) is used for the corresponding data structure.

$$A = \begin{pmatrix} 0,1,0,0,0,0,0,1,0,0 \\ 0,0,1,0,0,0,0,0,0,0 \\ 1,0,0,1,0,0,0,0,0,1 \\ 1,0,0,0,0,0,1,1,1,0,0 \\ 0,0,1,0,1,0,0,0,1,0 \\ 0,0,0,0,0,0,0,0,0,0,0 \\ 0,1,0,0,0,0,0,0,0,0,0 \\ 0,1,0,0,0,0,0,0,0,0,0 \\ 0,1,1,0,1,0,0,0,0,1,0 \\ 0,1,1,0,1,0,0,0,0,0,0 \end{pmatrix} \mapsto BRC(A) = \begin{bmatrix} [1,7], \\ [2], \\ [0,3,9], \\ [0,6], \\ [2,4,8], \\ [1,7,9], \\ [3,8], \\ [1,2,4]] \end{bmatrix}$$

#### 1.2 Format conversions

First we give the function triples2mat to make the transformation from the sparse matrix, given as a list of triples row, column, value (non-zero elements), to the scipy.sparse format corresponding to the shape parameter, set by default to "csr", that stands for Compressed Sparse Row, the normal matrix format of the LARCC framework.

```
⟨From list of triples to scipy.sparse 3a⟩ ≡

def triples2mat(triples,shape="csr"):
    n = len(triples)
    data = arange(n)
    ij = arange(2*n).reshape(2,n)
    for k,item in enumerate(triples):
        ij[0][k],ij[1][k],data[k] = item
    return scipy.sparse.coo_matrix((data, ij)).asformat(shape)
    ⋄
```

Macro referenced in 25.

The function brc2Coo transforms a BRC representation in a list of triples (row, column, 1) ordered by row.

Macro referenced in 25.

Two coordinate compressed sparse matrices coof and coof are created below, starting from the BRC representation FV and EV of the incidence of vertices on faces and edges, respectively, for a very simple plane triangulation.

```
⟨Test example of Brc to Coo transformation 3c⟩ ≡
    print "\n>>> brc2Coo"
    V = [[0, 0], [1, 0], [2, 0], [0, 1], [1, 1], [2, 1]]
    FV = [[0, 1, 3], [1, 2, 4], [1, 3, 4], [2, 4, 5]]
    EV = [[0,1], [0,3], [1,2], [1,3], [1,4], [2,4], [2,5], [3,4], [4,5]]
    cooFV = brc2Coo(FV)
    cooEV = brc2Coo(EV)
    assert cooFV == [[0,0,1], [0,1,1], [0,3,1], [1,1,1], [1,2,1], [1,4,1], [2,1,1],
    [2,3,1], [2,4,1], [3,2,1], [3,4,1], [3,5,1]]
    assert cooEV == [[0,0,1], [0,1,1], [1,0,1], [1,3,1], [2,1,1], [2,2,1], [3,1,1],
    [3,3,1], [4,1,1], [4,4,1], [5,2,1], [5,4,1], [6,2,1], [6,5,1], [7,3,1], [7,4,1],
    [8,4,1], [8,5,1]]
    ◊
```

Macro referenced in 26a.

Two CSR sparse matrices csrFV and csrEV are generated (by *scipy.sparse*) in the following example:

```
⟨Test example of Coo to Csr transformation 4b⟩ ≡
    csrFV = coo2Csr(cooFV)
    csrEV = coo2Csr(cooEV)
    print "\ncsr(FV) =\n", repr(csrFV)
    print "\ncsr(EV) =\n", repr(csrEV)
```

Macro referenced in 26a.

The *scipy* printout of the last two lines above is the following:

```
csr(FV) = <4x6 sparse matrix of type '<type 'numpy.int64'>'
  with 12 stored elements in Compressed Sparse Row format>
csr(EV) = <9x6 sparse matrix of type '<type 'numpy.int64'>'
  with 18 stored elements in Compressed Sparse Row format>
```

The transformation from BRC to CSR format is implemented slightly differently, according to the fact that the matrix dimension is either unknown (shape=(0,0)) or known.

```
⟨Brc to Csr transformation 4c⟩ ≡

def csrCreate(BRCmatrix,shape=(0,0)):
    triples = brc2Coo(BRCmatrix)
    if shape == (0,0):
        CSRmatrix = coo2Csr(triples)
    else:
        CSRmatrix = scipy.sparse.csr_matrix(shape)
        for i,j,v in triples: CSRmatrix[i,j] = v
    return CSRmatrix

⟩
```

Macro referenced in 25.

The conversion to CSR format of the characteristic matrix faces-vertices FV is given below for our simple example made by four triangle of a manifold 2D space, graphically shown in Figure 1a. The LAR representation with CSR matrices does not make difference between manifolds and non-manifolds, conversely than most modern solid modelling representation schemes, as shown by removing from FV the third triangle, giving the model in Figure 1b.

Macro referenced in 6d, 26a.

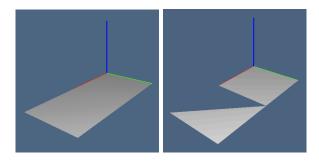


Figure 1: (a) Manifold two-dimensional space; (b) non-manifold space.

# 2 Matrix operations

As we know, the LAR representation of topology is based on CSR representation of sparse binary (and integer) matrices. Two Utility functions allow to query the number of rows and columns of a CSR matrix, independently from the low-level implementation (that in the following is provided by scipy.sparse).

Macro referenced in 25.

```
\langle Test examples of Query Matrix shape 6a \rangle \equiv
     print "\n>>> csrGetNumberOfRows"
     print "\ncsrGetNumberOfRows(csrFV) =", csrGetNumberOfRows(csrFV)
     print "\ncsrGetNumberOfRows(csrEV) =", csrGetNumberOfRows(csrEV)
     print "\n>>> csrGetNumberOfColumns"
     print "\ncsrGetNumberOfColumns(csrFV) =", csrGetNumberOfColumns(csrFV)
     print "\ncsrGetNumberOfColumns(csrEV) =", csrGetNumberOfColumns(csrEV)
Macro referenced in 26a.
\langle Sparse to dense matrix transformation 6b \rangle \equiv
     def csr2DenseMatrix(CSRm):
         nrows = csrGetNumberOfRows(CSRm)
          ncolumns = csrGetNumberOfColumns(CSRm)
          ScipyMat = zeros((nrows,ncolumns),int)
          C = CSRm.tocoo()
          for triple in zip(C.row,C.col,C.data):
              ScipyMat[triple[0],triple[1]] = triple[2]
          return ScipyMat
Macro referenced in 25.
\langle Test examples of Sparse to dense matrix transformation 6c \rangle \equiv
     print "\n>>> csr2DenseMatrix"
     print "\nFV =\n", csr2DenseMatrix(csrFV)
     print "\nEV =\n", csr2DenseMatrix(csrEV)
Macro referenced in 6d, 26a.
```

Characteristic matrices Let us compute and show in dense form the characteristic matrices of 2- and 1-cells of the simple manifold just defined. By running the file test/py/larcc/test08.py the reader will get the two matrices shown in Example 2

```
"test/py/larcc/test08.py" 6d ≡

import sys; sys.path.insert(0, 'lib/py/')

from larcc import *

⟨Test example of Brc to Csr transformation 5a⟩

⟨Test examples of Sparse to dense matrix transformation 6c⟩

⋄
```

**Example 1** (Dense Characteristic matrices). Let us notice that the two matrices below have the some numbers of columns (indexed by vertices of the cell decomposition). This very fact allows to multiply one matrix for the other transposed, and hence to compute the

matrix form of linear operators between the spaces of cells of various dimensions.

$$FV = \begin{bmatrix} \begin{bmatrix} 1 & 1 & 0 & 0 & 0 & 0 \\ & \begin{bmatrix} 1 & 1 & 0 & 1 & 0 & 0 \\ & \begin{bmatrix} 0 & 1 & 1 & 0 & 1 & 0 \\ & \begin{bmatrix} 0 & 1 & 1 & 0 & 1 & 0 \\ & \begin{bmatrix} 0 & 1 & 0 & 1 & 1 & 0 \end{bmatrix} \end{bmatrix} \end{bmatrix}$$

$$EV = \begin{bmatrix} \begin{bmatrix} 0 & 1 & 0 & 0 & 1 & 0 \\ & \begin{bmatrix} 0 & 1 & 0 & 0 & 1 & 0 \\ & \begin{bmatrix} 0 & 0 & 1 & 0 & 1 & 0 \end{bmatrix} \end{bmatrix}$$

$$\begin{bmatrix} \begin{bmatrix} 0 & 0 & 1 & 0 & 1 & 1 \\ & \begin{bmatrix} 0 & 0 & 0 & 1 & 1 & 1 \end{bmatrix} \end{bmatrix}$$

$$\begin{bmatrix} \begin{bmatrix} 0 & 0 & 0 & 1 & 1 & 1 \end{bmatrix}$$

Matrix product and transposition The following macro provides the IDE interface for the two main matrix operations required by LARCC, the binary product of compatible matrices and the unary transposition of matrices.

```
⟨ Matrix product and transposition 7⟩ ≡
    def matrixProduct(CSRm1,CSRm2):
        CSRm = CSRm1 * CSRm2
        return CSRm

def csrTranspose(CSRm):
        CSRm = CSRm.T
        return CSRm
```

Macro referenced in 25.

**Example 2** (Operators from edges to faces and vice-versa). As a general rule for operators between two spaces of chains of different dimensions supported by the same cellular complex, we use names made by two characters, whose first letter correspond to the target space, and whose second letter to the domain space. Hence FE must be read as the operator from edges to faces. Of course, since this use correspond to see the first letter as the space generated by rows, and the second letter as the space generated by columns. Notice that the element (i,j) of such matrices stores the number of vertices shared between the (row-)cell i and the

(column-)cell j.

```
FE = FV EV^{\top} = \begin{bmatrix} \begin{bmatrix} 2 & 1 & 1 & 0 & 1 & 0 \\ & 1 & 0 & 2 & 1 & 2 & 2 & 1 & 1 & 1 \\ & 1 & 1 & 1 & 2 & 2 & 1 & 0 & 2 & 1 \\ & \begin{bmatrix} 0 & 0 & 1 & 0 & 1 & 2 & 2 & 1 & 2 & 1 \end{bmatrix} \end{bmatrix} 
EF = EV FV^{\top} = \begin{bmatrix} \begin{bmatrix} 2 & 1 & 1 & 0 \\ & 1 & 2 & 1 & 1 \\ & 2 & 1 & 2 & 0 \end{bmatrix} 
\begin{bmatrix} 0 & 2 & 1 & 2 & 1 \\ & 1 & 2 & 2 & 1 \\ & 0 & 2 & 1 & 2 \end{bmatrix} 
\begin{bmatrix} 0 & 1 & 0 & 2 \\ & 1 & 2 & 1 \end{bmatrix} 
\begin{bmatrix} 0 & 1 & 1 & 2 \end{bmatrix}
```

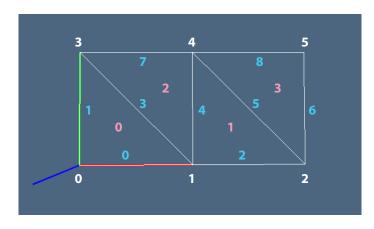


Figure 2: example caption

```
def csrBoundaryFilter(CSRm, facetLengths):
    maxs = [max(CSRm[k].data) for k in range(CSRm.shape[0])]
    inputShape = CSRm.shape
    coo = CSRm.tocoo()
    for k in range(len(coo.data)):
        if coo.data[k] ==maxs[coo.row[k]]: coo.data[k] = 1
        else: coo.data[k] = 0
    mtx = coo_matrix((coo.data, (coo.row, coo.col)), shape=inputShape)
    out = mtx.tocsr()
    return out
```

Macro referenced in 25.

```
\langle Test example of Matrix filtering to produce the boundary matrix 9a \rangle \equiv
     print "\n>>> csrBoundaryFilter"
     csrEF = matrixProduct(csrFV, csrTranspose(csrEV)).T
     facetLengths = [csrCell.getnnz() for csrCell in csrEV]
     CSRm = csrBoundaryFilter(csrEF, facetLengths).T
     print "\ncsrMaxFilter(csrFE) =\n", csr2DenseMatrix(CSRm)
Macro referenced in 26a.
\langle Matrix filtering via a generic predicate 9b \rangle \equiv
     def csrPredFilter(CSRm, pred):
         # can be done in parallel (by rows)
         coo = CSRm.tocoo()
         triples = [[row,col,val] for row,col,val
                   in zip(coo.row,coo.col,coo.data) if pred(val)]
         i, j, data = TRANS(triples)
         CSRm = scipy.sparse.coo_matrix((data,(i,j)),CSRm.shape).tocsr()
         return CSRm
     \Diamond
Macro referenced in 25.
\langle Test example of Matrix filtering via a generic predicate 9c\rangle \equiv
     print "\n>>> csrPredFilter"
     CSRm = csrPredFilter(matrixProduct(csrFV, csrTranspose(csrEV)).T, GE(2)).T
     print "\nccsrPredFilter(csrFE) =\n", csr2DenseMatrix(CSRm)
```

# 3 Topological operations

Macro referenced in 26a.

In this section we provide the matrix representation of operators to compute the more important and useful topological operations on cellular complexes, and/or the indexed relations they return. We start the section by giving a graphical tool used to test the developed software, concerning the graphical writing of the full set of indices of the cells of every dimension in a 3D cuboidal complex.

**Visualization of cell indices** As already outlined, the modelIndexing function return the hpc value assembling both the 1-skeletons of the cells of every dimensions, and the graphical output of their indices, located on the centroid of each cell, and displayed using colors and sizes depending on the rank of the cell.

```
from sysml import *
     def modelIndexing(shape):
        V, bases = larCuboids(shape,True)
        # bases = [[cell for cell in cellComplex if len(cell) == 2**k] for k in range(4)]
        color = [YELLOW, CYAN, GREEN, WHITE]
        nums = AA(range)(AA(len)(bases))
        hpcs = []
        for k in range(4):
           hpcs += [SKEL_1(STRUCT(MKPOLS((V,bases[k]))))]
           hpcs += [cellNumbering((V,bases[k]),hpcs[2*k])(nums[k],color[k],0.3+0.2*k)]
        return STRUCT(hpcs)
Macro defined by 9d, 10a.
Macro referenced in 25.
\langle Visualization of cell indices 10a \rangle \equiv
     """ Numbered visualization of a LAR model """
     def larModelNumbering(V,bases,submodel,numberScaling=1):
        color = [YELLOW, CYAN, GREEN, WHITE]
        nums = AA(range)(AA(len)(bases))
        hpcs = [submodel]
        for k in range(len(bases)):
            hpcs += [cellNumbering((V,bases[k]),submodel)
                      (nums[k],color[k],(0.3+0.2*k)*numberScaling)]
        return STRUCT(hpcs)
     \quad
Macro defined by 9d, 10a.
Macro referenced in 25.
```

**Drawing of oriented edges** The following function return the hpc of the drawing with arrows of the oriented 1-cells of a 2D cellular complex. Of course, each edge orientation is from second to first vertex, independently from the vertex indices. Therefore, the edge orientation can be reversed by swapping the vertex indices in the 1-cell definition.

```
⟨ Drawing of oriented edges 10b⟩ ≡
    """ Drawing of oriented edges (2D) """
    def mkSignedEdges (model):
        V,EV = model
        assert len(V[0])==2
        hpcs = []
        times = C(SCALARVECTPROD)
        for e0,e1 in EV:
            v0,v1 = V[e0], V[e1]
            vx,vy = DIFF([ v1, v0 ])
            nx,ny = [-vy, vx]
```

```
v2 = SUM([ v0, times(0.66)([vx,vy]) ])
v3 = SUM([ v0, times(0.6)([vx,vy]), times(0.06)([nx,ny]) ])
v4 = SUM([ v0, times(0.6)([vx,vy]), times(-0.06)([nx,ny]) ])
verts,cells = [v0,v1,v2,v3,v4],[[1,2],[3,4],[3,5]]
hpcs += [MKPOL([verts,cells,None])]
hpc = STRUCT(hpcs)
return hpc
```

Macro referenced in 25.

Example of oriented edge drawing An example of drawing of oriented edges is given in test/py/larcc/test11.py file, and in Figure 3, showing both the numbering of the cells and the arrows indicating the edge orientation.

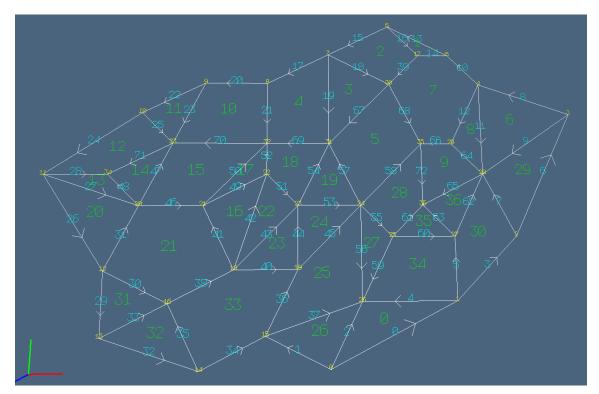


Figure 3: Example of numbered polytopal complex, including edge orientations.

```
"test/py/larcc/test11.py" 11 \equiv 
""" Example of oriented edge drawing """ 
import sys;sys.path.insert(0, 'lib/py/') 
from larcc import *
```

```
V = [[9,0],[13,2],[15,4],[17,8],[14,9],[13,10],[11,11],[9,10],[7,9],[5,9],[3,
8],[0,6],[2,3],[2,1],[5,0],[7,1],[4,2],[12,10],[6,3],[8,3],[3,5],[5,5],[7,6],
[8,5], [10,5], [11,4], [10,2], [13,4], [14,6], [13,7], [11,9], [9,7], [7,7], [4,7], [2,
6],[12,7],[12,5]]
FV = [[0,1,26],[5,6,17],[6,7,17,30],[7,30,31],[7,8,31,32],[24,30,31,35],[3,4,
28], [4,5,17,29,30,35], [4,28,29], [28,29,35,36], [8,9,32,33], [9,10,33], [11,10,
33,34],[11,20,34],[20,33,34],[20,21,32,33],[18,21,22],[21,22,32],[22,23,31,
32],[23,24,31],[11,12,20],[12,16,18,20,21],[18,22,23],[18,19,23],[19,23,24],
[15,19,24,26],[0,15,26],[24,25,26],[24,25,35,36],[2,3,28],[1,2,27,28],[12,13,
16], [13,14,16], [14,15,16,18,19], [1,25,26,27], [25,27,36], [36,27,28]]
VIEW(EXPLODE(1.2,1.2,1)(MKPOLS((V,FV))))
VV = AA(LIST)(range(len(V)))
_,EV = larFacets((V,FV+[range(16)]),dim=2,emptyCellNumber=1)
submodel = mkSignedEdges((V,EV))
VIEW(submodel)
VIEW(larModelNumbering(V, [VV, EV, FV], submodel, 3))
```

#### 3.1 Incidence and adjacency operators

Let us start by computing the more interesting subset of the binary relationships between the 4 decompositive and/or boundary entities of 3D cellular models. Therefore, in this case we denote with C, F, E, and V, the 3-cells and their faces, edges and vertices, respectively. The input is the full-fledged LAR representation provided by

$$CV := CSR(M_3) \tag{1}$$

$$FV := CSR(M_2) \tag{2}$$

$$EV := CSR(M_1) \tag{3}$$

$$VV := CSR(M_0) \tag{4}$$

Of course,  $CSR(M_0)$  coincides with the identity matrix of dimension |V| and can by excluded by further considerations. Some binary incidence and adjacency relations we are going to compute are:

$$CF := CV \times FV^t = CSR(M_3) \times CSR(M_2)^t$$
(5)

$$\mathtt{CE} := \mathtt{CV} \times \mathtt{EV}^t = \mathtt{CSR}(M_3) \times \mathtt{CSR}(M_1)^t \tag{6}$$

$$\mathtt{FE} := \mathtt{FV} \times \mathtt{EV}^t = \mathtt{CSR}(M_2) \times \mathtt{CSR}(M_1)^t \tag{7}$$

The other possible operators follow from a similer computational pattern.

The programming pattern for incidence computation A high-level function larIncidence useful to compute the LAR representation of the incidence matrix (operator) and the incidence relations is given in the script below.

```
⟨Some incidence operators 13a⟩ ≡
    """ Some incidence operators """
    def larIncidence(cells,facets):
        csrCellFacet = csrCellFaceIncidence(cells,facets)
        cooCellFacet = csrCellFacet.tocoo()
        larCellFacet = [[] for cell in range(len(cells))]
        for i,j,val in zip(cooCellFacet.row,cooCellFacet.col,cooCellFacet.data):
            if val == 1: larCellFacet[i] += [j]
            return larCellFacet

        ⟨Cell-Face incidence operator 13b⟩
        ⟨Cell-Edge incidence operator 13c⟩
        ⟨Face-Edge incidence operator 14a⟩
        ⟩

Macro referenced in 25.
```

Cell-Face incidence The csrCellFaceIncidence and larCellFace functions are given below, and exported to the larce module.

```
⟨ Cell-Face incidence operator 13b ⟩ ≡
    """ Cell-Face incidence operator """
    def csrCellFaceIncidence(CV,FV):
        return boundary(FV,CV)

def larCellFace(CV,FV):
        return larIncidence(CV,FV)
```

Cell-Edge incidence Analogously, the csrCellEdgeIncidence and larCellFace functions are given in the following script.

Macro referenced in 13a.

Macro referenced in 13a.

**Face-Edge incidence** Finally, the csrCellEdgeIncidence and larCellFace functions are provided below.

```
⟨ Face-Edge incidence operator 14a⟩ ≡
    """ Face-Edge incidence operator """
    def csrFaceEdgeIncidence(FV,EV):
        return boundary(EV,FV)

def larFaceEdge(FV,EV):
        return larIncidence(FV,EV)
```

Macro referenced in 13a.

**Example** The example below concerns a 3D cuboidal grid, by computing a full LAR stack of bases CV, FV, EV, VV, showing its fully numbered 3D model, and finally by computing some more useful binary relationships (CF, CE, FE), needed for example to compute the signed matrices of boundary operators.

```
"test/py/larcc/test10.py" 14b ==
    """ A mesh model and various incidence operators """
    import sys; sys.path.insert(0, 'lib/py/')
    from larcc import *
    from largrid import *

    shape = [2,2,2]
    V,(VV,EV,FV,CV) = larCuboids(shape,True)
    """

    CV = [cell for cell in cellComplex if len(cell)==8]
    FV = [cell for cell in cellComplex if len(cell)==4]
    EV = [cell for cell in cellComplex if len(cell)==2]
    VV = [cell for cell in cellComplex if len(cell)==2]
    VV = [cell for cell in cellComplex if len(cell)==1]
    """

    VIEW(modelIndexing(shape))

    CF = larCellFace(CV,FV)
    CE = larCellFace(CV,EV)
    FE = larCellFace(FV,EV)
```

#### 3.2 Boundary and coboundary operators

```
\langle From cells and facets to boundary operator 15a\rangle \equiv
     def boundary(cells,facets):
         csrCV = csrCreate(cells)
         csrFV = csrCreate(facets)
         csrFC = matrixProduct(csrFV, csrTranspose(csrCV))
         facetLengths = [csrCell.getnnz() for csrCell in csrCV]
         return csrBoundaryFilter(csrFC,facetLengths)
     def coboundary(cells,facets):
         Boundary = boundary(cells,facets)
         return csrTranspose(Boundary)
Macro referenced in 25.
\langle Test examples of From cells and facets to boundary operator 15b\rangle \equiv
     V = [[0.0, 0.0, 0.0], [1.0, 0.0, 0.0], [0.0, 1.0, 0.0], [1.0, 1.0, 0.0],
     [0.0, 0.0, 1.0], [1.0, 0.0, 1.0], [0.0, 1.0, 1.0], [1.0, 1.0, 1.0]]
     CV = [[0, 1, 2, 4], [1, 2, 4, 5], [2, 4, 5, 6], [1, 2, 3, 5], [2, 3, 5, 6],
     [3, 5, 6, 7]]
     FV = [[0, 1, 2], [0, 1, 4], [0, 2, 4], [1, 2, 3], [1, 2, 4], [1, 2, 5],
     [1, 3, 5], [1, 4, 5], [2, 3, 5], [2, 3, 6], [2, 4, 5], [2, 4, 6], [2, 5, 6],
     [3, 5, 6], [3, 5, 7], [3, 6, 7], [4, 5, 6], [5, 6, 7]]
     EV = [[0, 1], [0, 2], [0, 4], [1, 2], [1, 3], [1, 4], [1, 5], [2, 3], [2, 4],
     [2, 5], [2, 6], [3, 5], [3, 6], [3, 7], [4, 5], [4, 6], [5, 6], [5, 7],
     [6, 7]]
     print "\ncoboundary_2 =\n", csr2DenseMatrix(coboundary(CV,FV))
     print "\ncoboundary_1 =\n", csr2DenseMatrix(coboundary(FV,EV))
     print "\ncoboundary_0 =\n", csr2DenseMatrix(coboundary(EV,AA(LIST)(range(len(V)))))
```

Macro referenced in 26a.

```
\langle From cells and facets to boundary cells 16a\rangle \equiv
     def zeroChain(cells):
        pass
     def totalChain(cells):
        return csrCreate([[0] for cell in cells]) # ???? zero ??
     def boundaryCells(cells,facets):
        csrBoundaryMat = boundary(cells,facets)
        csrChain = totalChain(cells)
        csrBoundaryChain = matrixProduct(csrBoundaryMat, csrChain)
        for k,value in enumerate(csrBoundaryChain.data):
            if value % 2 == 0: csrBoundaryChain.data[k] = 0
        boundaryCells = [k for k,val in enumerate(csrBoundaryChain.data.tolist()) if val == 1]
        return boundaryCells
Macro referenced in 25.
\langle Test examples of From cells and facets to boundary cells 16b\rangle \equiv
     boundaryCells_2 = boundaryCells(CV,FV)
     boundaryCells_1 = boundaryCells([FV[k] for k in boundaryCells_2],EV)
     print "\nboundaryCells_2 =\n", boundaryCells_2
     print "\nboundaryCells_1 =\n", boundaryCells_1
     boundaryModel = (V,[FV[k] for k in boundaryCells_2])
     VIEW(EXPLODE(1.5,1.5,1.5) (MKPOLS(boundaryModel)))
```

Signed boundary matrix for simplicial complexes The computation of the *signed* boundary matrix starts with enumerating the non-zero elements of the mod two (unoriented) boundary matrix. In particular, the pairs variable contains all the pairs of incident ((d-1)-cell), corresponding to all the 1 elements in the binary boundary matrix. Of course, their number equates the product of the number of d-cell, times the number of (d-1)-facets on the boundary of each d-cell. For the case of a 3-simplicial complex CV, we have 4|CV| pairs elements. The actual goal of the function signedBoundary, in the macro

Macro referenced in 26a.

below, is to compute a sign for each of them.

The pairs values must be interpreted as (i, j) values in the incidence matrix FC (facets-cells), and hence as pairs of indices f and c into the characteristic matrices FV = CSR( $M_{d-1}$ ) and CV = CSR( $M_d$ ), respectively.

For each incidence pair f,c, the list vertLists contains the two lists of vertices associated to f and to c, called respectively the face and the coface. For each face, coface

pair (i.e. for each unit element in the unordered boundary matrix), the missingVertIndices list will contain the index of the coface vertex not contained in the incident face. Finally the  $\pm 1$  (signed) incidence coefficients are computed and stored in the faceSigns, and then located in their actual positions within the csrSignedBoundaryMat. The sign of the incidence coefficient associated to the pair (facet,cell), also called (face,coface) in the implementation below, is computed as the sign of  $(-1)^k$ , where k is the position index of the removed vertex in the facet  $\langle v_0, \ldots, v_{k-1}, v_{k+1}, \ldots, v_d \rangle$ . of the  $\langle v_0, \ldots, v_d \rangle$  cell.

```
(Signed boundary matrix for simplicial models 17) ≡
    def signedBoundary (CV,FV):
        # compute the set of pairs of indices to [boundary face,incident coface]
        coo = boundary(CV,FV).tocoo()
        pairs = [[coo.row[k],coo.col[k]] for k,val in enumerate(coo.data) if val != 0]

# compute the [face, coface] pair as vertex lists
        vertLists = [[FV[f], CV[c]] for f,c in pairs]

# compute the local (interior to the coface) indices of missing vertices
        def missingVert(face,coface): return list(set(coface).difference(face))[0]
        missingVertIndices = [c.index(missingVert(f,c)) for f,c in vertLists]

# signed incidence coefficients
        faceSigns = AA(C(POWER)(-1))(missingVertIndices)

# signed boundary matrix
        csrSignedBoundaryMat = csr_matrix( (faceSigns, TRANS(pairs)) )
        return csrSignedBoundaryMat
```

Computation of signed boundary cells Two simplices are said coherently oriented when their common facets have opposite orientations. If the boundary cells give a decomposition of the boundary of an orientable solid, that partitionates the embedding space in two subsets corresponding to the *interior* and the *exterior* of the solid, then the boundary cells can be coherently oriented. This task is performed by the function signedBoundaryCells below.

Macro referenced in 25.

The matrix of the signed boundary operator, with elements in  $\{-1,0,1\}$ , is computed in compressed sparse row (CSR) format, and stored in csrSignedBoundaryMat. In order to be able to return a list of signedBoundaryCells having a coherent orientation, we need to compute the coface of each boundary facet, i.e. the single d-cell having the facet on its boundary, and provide a coherent orientation to such chain of d-cells. The goal is obtained computing the sign of the determinant of the coface matrices, i.e. of square matrices having as rows the vertices of a coface, in normalised homogeneous coordinates.

The chain of boundary facets boundaryCells, obtained by multiplying the signed matrix of the boundary operator by the coordinate representation of the total *d*-chain, is coherently oriented by multiplication times the determinants of the cofaceMats.

The cofaceMats list is filled with the matrices having per row the position vectors of vertices of a coface, in normalized homogeneous coordinates. The list of signed face indices orientedBoundaryCells is returned by the function.

```
\langle Oriented boundary cells for simplicial models 18\rangle \equiv
     def signedBoundaryCells(verts,cells,facets):
        csrSignedBoundaryMat = signedBoundary(cells,facets)
        csrTotalChain = totalChain(cells)
        csrBoundaryChain = matrixProduct(csrSignedBoundaryMat, csrTotalChain)
        cooCells = csrBoundaryChain.tocoo()
        boundaryCells = []
        for k,v in enumerate(cooCells.data):
            if abs(v) == 1:
               boundaryCells += [int(cooCells.row[k] * cooCells.data[k])]
        boundaryCocells = []
        for k,v in enumerate(boundaryCells):
            boundaryCocells += list(csrSignedBoundaryMat[abs(v)].tocoo().col)
        boundaryCofaceMats = [[verts[v]+[1] for v in cells[c]] for c in boundaryCocells]
        boundaryCofaceSigns = AA(SIGN)(AA(np.linalg.det)(boundaryCofaceMats))
        def swap(mylist): return [mylist[1]]+[mylist[0]]+mylist[2:]
        orientedBoundaryCells = list(array(boundaryCells)*array(boundaryCofaceSigns))
        return orientedBoundaryCells
     \Diamond
Macro defined by 18, 20.
Macro referenced in 25.
Orienting polytopal cells
input: "cell" indices of a convex and solid polytopes and "V" vertices;
output: biggest "simplex" indices spanning the polytope.
m: number of cell vertices
d: dimension (number of coordinates) of cell vertices
```

d+1: number of simplex vertices

vcell : cell vertices

 ${\tt vsimplex} \, : \, {\rm simplex} \, \, {\rm vertices} \,$ 

Id: identity matrix

 ${\tt basis}$  : orthonormal spanning set of vectors  $e_k$ 

vector : position vector of a simplex vertex in translated coordinates

unUsedIndices: cell indices not moved to simplex

```
\langle Oriented boundary cells for simplicial models 20\rangle \equiv
     def pivotSimplices(V,CV,d=3):
        simplices = []
        for cell in CV:
           vcell = np.array([V[v] for v in cell])
           m, simplex = len(cell), []
           # translate the cell: for each k, vcell[k] -= vcell[0], and simplex[0] := cell[0]
           for k in range(m-1,-1,-1): vcell[k] = vcell[0]
           \# simplex = [0], basis = [], tensor = Id(d+1)
           simplex += [cel1[0]]
           basis = []
           tensor = np.array(IDNT(d))
           # look for most far cell vertex
           dists = [SUM([SQR(x) for x in v])**0.5 for v in vcell]
           maxDistIndex = max(enumerate(dists),key=lambda x: x[1])[0]
           vector = np.array([vcell[maxDistIndex]])
           # normalize vector
           den=(vector**2).sum(axis=-1) **0.5
           basis = [vector/den]
           simplex += [cell[maxDistIndex]]
           unUsedIndices = [h for h in cell if h not in simplex]
           # for k in \{2,d+1\}:
           for k in range(2,d+1):
              # update the orthonormal tensor
              e = basis[-1]
              tensor = tensor - np.dot(e.T, e)
              # compute the index h of a best vector
              # look for most far cell vertex
              dists = [SUM([SQR(x) for x in np.dot(tensor,v)])**0.5
              if h in unUsedIndices else 0.0
              for (h,v) in zip(cell,vcell)]
              # insert the best vector index h in output simplex
              maxDistIndex = max(enumerate(dists),key=lambda x: x[1])[0]
              vector = np.array([vcell[maxDistIndex]])
              # normalize vector
              den=(vector**2).sum(axis=-1) **0.5
              basis += [vector/den]
              simplex += [cell[maxDistIndex]]
              unUsedIndices = [h for h in cell if h not in simplex]
           simplices += [simplex]
        return simplices
     def simplexOrientations(V,simplices):
        vcells = [[V[v]+[1.0]] for v in simplex] for simplex in simplices]
        return [SIGN(np.linalg.det(vcell)) for vcell in vcells]
```

```
\langle Extraction of facets of a cell complex 22 \rangle \equiv
     def setup(model,dim):
         V, cells = model
         csr = csrCreate(cells)
         csrAdjSquareMat = larCellAdjacencies(csr)
         csrAdjSquareMat = csrPredFilter(csrAdjSquareMat, GE(dim)) # ? HOWTODO ?
         return V,cells,csr,csrAdjSquareMat
     def larFacets(model,dim=3,emptyCellNumber=0):
             Estraction of (d-1)-cellFacets from "model" := (V,d-cells)
             Return (V, (d-1)-cellFacets)
         V,cells,csr,csrAdjSquareMat = setup(model,dim)
         solidCellNumber = len(cells) - emptyCellNumber
         cellFacets = []
         # for each input cell i
         for i in range(len(cells)):
             adjCells = csrAdjSquareMat[i].tocoo()
             cell1 = csr[i].tocoo().col
             pairs = zip(adjCells.col,adjCells.data)
             for j,v in pairs:
                  if (i<j) and (i<solidCellNumber):</pre>
                      cell2 = csr[j].tocoo().col
                      cell = list(set(cell1).intersection(cell2))
                      cellFacets.append(sorted(cell))
         # sort and remove duplicates
         cellFacets = sorted(AA(list)(set(AA(tuple)(cellFacets))))
         return V, cellFacets
```

Macro referenced in 25.

```
\( \text{Test examples of Extraction of facets of a cell complex 23} \) \( \text{V} = [[0.,0.],[3.,0.],[0.,3.],[3.,3.],[1.,2.],[2.,2.],[1.,1.],[2.,1.]] \)
\( \text{FV} = [[0,1,6,7],[0,2,4,6],[4,5,6,7],[1,3,5,7],[2,3,4,5],[0,1,2,3]] \)
\( _,\text{EV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{VIEW}(\text{EXPLODE}(1.5,1.5,1.5)(\text{MKPOLS}((V,\text{EV})))) \)
\( \text{FV} = [[0,1,3],[1,2,4],[2,4,5],[3,4,6],[4,6,7],[5,7,8], # \text{full} \)
\( [1,3,4],[4,5,7], # \text{empty} \)
\( [0,1,2],[6,7,8],[0,3,6],[2,5,8] ] # \text{exterior} \)
\( _,\text{EV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{PV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{PV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{PV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{PV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{PV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{PV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{PV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{PV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{PV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{PV} = \text{larFacets}((V,\text{FV}),\dim=2) \)
\( \text{print "\nEV} = ",\text{EV} \)
\( \text{PV} = \text{larFacets}((V,\text{PV}),\dim=2) \)
\( \text{larFacets}((V
```

Macro referenced in 26a.

## 4 Exporting the library

#### 4.1 MIT licence

```
\langle The MIT Licence 24a\rangle \equiv """

The MIT License
```

Permission is hereby granted, free of charge, to any person obtaining a copy of this software and associated documentation files (the 'Software'), to deal in the Software without restriction, including without limitation the rights to use, copy, modify, merge, publish, distribute, sublicense, and/or sell copies of the Software, and to permit persons to whom the Software is furnished to do so, subject to the following conditions:

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 $\Diamond$ 

Macro referenced in 25.

Macro referenced in 25.

#### 4.2 Importing of modules or packages

```
⟨Importing of modules or packages 24b⟩ ≡
    from pyplasm import *
    import collections
    import scipy
    import numpy as np
    from scipy import zeros,arange,mat,amin,amax,array
    from scipy.sparse import vstack,hstack,csr_matrix,coo_matrix,lil_matrix,triu
    from lar2psm import *
```

## 4.3 Writing the library file

```
"lib/py/larcc.py" 25 \equiv
      # -*- coding: utf-8 -*-
      """ Basic LARCC library """
      ⟨The MIT Licence 24a⟩
      (Importing of modules or packages 24b)
      (From list of triples to scipy.sparse 3a)
      (Brc to Coo transformation 3b)
      (Coo to Csr transformation 4a)
      (Brc to Csr transformation 4c)
      Query Matrix shape 5b
      (Sparse to dense matrix transformation 6b)
      (Matrix product and transposition 7)
      (Matrix filtering to produce the boundary matrix 8)
      (Matrix filtering via a generic predicate 9b)
       From cells and facets to boundary operator 15a
       From cells and facets to boundary cells 16a
       Signed boundary matrix for simplicial models 17
       Oriented boundary cells for simplicial models 18, ... >
       Computation of cell adjacencies 21a
       Extraction of facets of a cell complex 22
      (Some incidence operators 13a)
      ⟨ Visualization of cell indices 9d, . . . ⟩
      (Numbered visualization of a LAR model?)
      ⟨ Drawing of oriented edges 10b⟩
      if __name__ == "__main__":
         (Test examples 26a)
```

#### 5 Unit tests

```
⟨Test examples 26a⟩ ≡
⟨Test example of Brc to Coo transformation 3c⟩
⟨Test example of Coo to Csr transformation 4b⟩
⟨Test example of Brc to Csr transformation 5a⟩
⟨Test examples of Query Matrix shape 6a⟩
⟨Test examples of Sparse to dense matrix transformation 6c⟩
⟨Test example of Matrix filtering to produce the boundary matrix 9a⟩
⟨Test example of Matrix filtering via a generic predicate 9c⟩
⟨Test examples of From cells and facets to boundary operator 15b⟩
⟨Test examples of From cells and facets to boundary cells 16b⟩
⟨Test examples of Computation of cell adjacencies 21b⟩
⟨Test examples of Extraction of facets of a cell complex 23⟩
```

Macro referenced in 25.

#### Comparing oriented and unoriented boundary

```
"test/py/larcc/test09.py" 26b ==

""" comparing oriented boundary and unoriented boundary extraction on a simple example """
import sys; sys.path.insert(0, 'lib/py/')
from largrid import *
from larcc import *

V,CV = larSimplexGrid1([1,1,1])
FV = larSimplexFacets(CV)

orientedBoundary = signedBoundaryCells(V,CV,FV)
def swap(mylist): return [mylist[1]]+[mylist[0]]+mylist[2:]
orientedBoundaryFV = [FV[-k] if k<0 else swap(FV[k]) for k in orientedBoundary]
VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS((V,orientedBoundaryFV))))

BF = boundaryCells(CV,FV)
boundaryCellsFV = [FV[k] for k in BF]
VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS((V,boundaryCellsFV))))</pre>
```

# A Appendix: Tutorials

#### A.1 Model generation, skeleton and boundary extraction

```
"test/py/larcc/test01.py" 27a \equiv
      import sys; sys.path.insert(0, 'lib/py/')
      from larcc import *
      from largrid import *
      (input of 2D topology and geometry data 27b)
      ⟨ characteristic matrices 27c ⟩
      (incidence matrix 27d)
      (boundary and coboundary operators 28a)
      (product of cell complexes 28b)
      (2-skeleton extraction 28c)
      (1-skeleton extraction 29a)
      (0-coboundary computation 29b)
      (1-coboundary computation 29c)
      (2-coboundary computation 30a)
      ⟨ boundary chain visualisation 30b⟩
\langle \text{ input of 2D topology and geometry data 27b} \rangle \equiv
      # input of geometry and topology
      V2 = [[4,10],[8,10],[14,10],[8,7],[14,7],[4,4],[8,4],[14,4]]
     EV = [[0,1],[1,2],[3,4],[5,6],[6,7],[0,5],[1,3],[2,4],[3,6],[4,7]]
      FV = [[0,1,3,5,6],[1,2,3,4],[3,4,6,7]]
Macro referenced in 27a.
\langle characteristic matrices 27c \rangle \equiv
      # characteristic matrices
      csrFV = csrCreate(FV)
      csrEV = csrCreate(EV)
      print "\nFV =\n", csr2DenseMatrix(csrFV)
      print "\nEV =\n", csr2DenseMatrix(csrEV)
Macro referenced in 27a.
\langle \text{ incidence matrix 27d} \rangle \equiv
     # product
      csrEF = matrixProduct(csrEV, csrTranspose(csrFV))
      print "\nEF =\n", csr2DenseMatrix(csrEF)
Macro referenced in 27a.
```

```
\langle boundary and coboundary operators 28a\rangle \equiv
     # boundary and coboundary operators
     facetLengths = [csrCell.getnnz() for csrCell in csrEV]
     boundary = csrBoundaryFilter(csrEF,facetLengths)
     coboundary_1 = csrTranspose(boundary)
     print "\ncoboundary_1 =\n", csr2DenseMatrix(coboundary_1)
Macro referenced in 27a.
\langle product of cell complexes 28b \rangle \equiv
     # product operator
     mod_2D = (V2, FV)
     V1, topol_0 = [[0.], [1.], [2.]], [[0], [1], [2]]
     topol_1 = [[0,1],[1,2]]
     mod_OD = (V1, topol_O)
     mod_1D = (V1, topol_1)
     V3,CV = larModelProduct([mod_2D,mod_1D])
     mod_3D = (V3,CV)
     VIEW(EXPLODE(1.2,1.2,1.2)(MKPOLS(mod_3D)))
     print "\nk_3 =", len(CV), "\n"
Macro referenced in 27a.
\langle 2-skeleton extraction 28c \rangle \equiv
     # 2-skeleton of the 3D product complex
     mod_2D_1 = (V2, EV)
     mod_3D_h2 = larModelProduct([mod_2D,mod_0D])
     mod_3D_v2 = larModelProduct([mod_2D_1,mod_1D])
     _{,FV_h} = mod_{3D_h2}
     _{,FV_v} = mod_{3D_v2}
     FV3 = FV_h + FV_v
     SK2 = (V3, FV3)
     VIEW(EXPLODE(1.2,1.2,1.2)(MKPOLS(SK2)))
     print "\nk_2 =", len(FV3), "\n"
```

Macro referenced in 27a.

```
\langle 1-skeleton extraction 29a\rangle \equiv
     # 1-skeleton of the 3D product complex
     mod_2D_0 = (V2,AA(LIST)(range(len(V2))))
     mod_3D_h1 = larModelProduct([mod_2D_1,mod_0D])
     mod_3D_v1 = larModelProduct([mod_2D_0,mod_1D])
     _{,EV_h} = mod_{3D_h1}
     \_, EV_v = mod_3D_v1
     EV3 = EV_h + EV_v
     SK1 = (V3, EV3)
     VIEW(EXPLODE(1.2,1.2,1.2)(MKPOLS(SK1)))
     print "\nk_1 =", len(EV3), "\n"
Macro referenced in 27a.
\langle 0-coboundary computation 29b \rangle \equiv
     # boundary and coboundary operators
     np.set_printoptions(threshold=sys.maxint)
     csrFV3 = csrCreate(FV3)
     csrEV3 = csrCreate(EV3)
     csrVE3 = csrTranspose(csrEV3)
     facetLengths = [csrCell.getnnz() for csrCell in csrEV3]
     boundary = csrBoundaryFilter(csrVE3,facetLengths)
     coboundary_0 = csrTranspose(boundary)
     print "\ncoboundary_0 =\n", csr2DenseMatrix(coboundary_0)
Macro referenced in 27a.
\langle 1-coboundary computation 29c \rangle \equiv
     csrEF3 = matrixProduct(csrEV3, csrTranspose(csrFV3))
     facetLengths = [csrCell.getnnz() for csrCell in csrFV3]
     boundary = csrBoundaryFilter(csrEF3,facetLengths)
     coboundary_1 = csrTranspose(boundary)
     print "\ncoboundary_1.T =\n", csr2DenseMatrix(coboundary_1.T)
     \Diamond
Macro referenced in 27a.
```

# A.2 Boundary of 3D simplicial grid

```
"test/py/larcc/test02.py" 30c ≡
import sys; sys.path.insert(0, 'lib/py/')
⟨boundary of 3D simplicial grid 31a⟩
⋄
```

```
\langle boundary of 3D simplicial grid 31a\rangle \equiv
     from simplexn import *
     from larcc import *
     V,CV = larSimplexGrid1([10,10,3])
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS((V,CV))))
     SK2 = (V,larSimplexFacets(CV))
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS(SK2)))
     _{,FV} = SK2
     SK1 = (V,larSimplexFacets(FV))
     _{,EV} = SK1
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS(SK1)))
     boundaryCells_2 = boundaryCells(CV,FV)
     boundary = (V,[FV[k] for k in boundaryCells_2])
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS(boundary)))
     print "\nboundaryCells_2 =\n", boundaryCells_2
     boundaryCells_2 = signedBoundaryCells(V,CV,FV)
     def swap(mylist): return [mylist[1]]+[mylist[0]]+mylist[2:]
     boundaryFV = [FV[-k] if k<0 else swap(FV[k]) for k in boundaryCells_2]
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS((V,boundaryFV))))
     print "\nboundaryCells_2 =\n", boundaryFV
```

Macro referenced in 30c.

#### A.3 Oriented boundary of a random simplicial complex

```
"test/py/larcc/test03.py" 31b ≡

⟨Importing external modules 31c⟩
⟨Generating and viewing a random 3D simplicial complex 32a⟩
⟨Computing and viewing its non-oriented boundary 32b⟩
⟨Computing and viewing its oriented boundary 32c⟩
⟨
Importing external modules 31c⟩ ≡

import sys; sys.path.insert(0, 'lib/py/')

from simplexn import *

from larcc import *

from scipy import *

from scipy.spatial import Delaunay
import numpy as np

⟨
```

Macro referenced in 31b.

```
\langle Generating and viewing a random 3D simplicial complex 32a\rangle \equiv
     verts = np.random.rand(10000, 3) # 1000 points in 3-d
     verts = [AA(lambda x: 2*x)(VECTDIFF([vert,[0.5,0.5,0.5]])) for vert in verts]
     verts = [vert for vert in verts if VECTNORM(vert) < 1.0]</pre>
     tetra = Delaunay(verts)
     cells = [cell for cell in tetra.vertices.tolist()
               if ((verts[cell[0]][2]<0) and (verts[cell[1]][2]<0)
                      and (verts[cel1[2]][2]<0) and (verts[cel1[3]][2]<0) ) ]
     V, CV = verts, cells
     VIEW(MKPOL([V,AA(AA(lambda k:k+1))(CV),[]]))
Macro referenced in 31b.
\langle Computing and viewing its non-oriented boundary 32b\rangle \equiv
     FV = larSimplexFacets(CV)
     VIEW(MKPOL([V,AA(AA(lambda k:k+1))(FV),[]]))
     boundaryCells_2 = boundaryCells(CV,FV)
     print "\nboundaryCells_2 =\n", boundaryCells_2
     bndry = (V,[FV[k] for k in boundaryCells_2])
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS(bndry)))
Macro referenced in 31b.
\langle Computing and viewing its oriented boundary 32c\rangle \equiv
     boundaryCells_2 = signedBoundaryCells(V,CV,FV)
     print "\nboundaryCells_2 =\n", boundaryCells_2
     def swap(mylist): return [mylist[1]]+[mylist[0]]+mylist[2:]
     boundaryFV = [FV[-k] if k<0 else swap(FV[k]) for k in boundaryCells_2]
     boundaryModel = (V,boundaryFV)
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS(boundaryModel)))
Macro referenced in 31b.
      Oriented boundary of a simplicial grid
"test/py/larcc/test04.py" 32d \equiv
     (Generate and view a 3D simplicial grid 33a)
     (Computing and viewing the 2-skeleton of simplicial grid 33b)
     ⟨ Computing and viewing the oriented boundary of simplicial grid 33c⟩
```

```
\langle Generate and view a 3D simplicial grid 33a\rangle \equiv
     import sys; sys.path.insert(0, 'lib/py/')
     from simplexn import *
     from larcc import *
     V,CV = larSimplexGrid1([4,4,4])
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS((V,CV))))
Macro referenced in 32d.
\langle Computing and viewing the 2-skeleton of simplicial grid 33b \rangle \equiv
     FV = larSimplexFacets(CV)
     EV = larSimplexFacets(FV)
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS((V,FV))))
Macro referenced in 32d.
\langle Computing and viewing the oriented boundary of simplicial grid 33c\rangle \equiv
     csrSignedBoundaryMat = signedBoundary (CV,FV)
     boundaryCells_2 = signedBoundaryCells(V,CV,FV)
     def swap(1): return [1[1],1[0],1[2]]
     boundaryFV = [FV[-k] if k<0 else swap(FV[k]) for k in boundaryCells_2]
     boundary = (V,boundaryFV)
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS(boundary)))
Macro referenced in 32d.
```

#### A.5 Skeletons and oriented boundary of a simplicial complex

```
"test/py/larcc/test05.py" 33d ≡
import sys; sys.path.insert(0, 'lib/py/')

⟨Skeletons computation and vilualisation 34a⟩
⟨Oriented boundary matrix visualization 34b⟩
⟨Computation of oriented boundary cells 34c⟩

⋄
```

```
\langle Skeletons computation and vilualisation 34a\rangle \equiv
     from simplexn import *
     from larcc import *
     V,FV = larSimplexGrid1([3,3])
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS((V,FV))))
     EV = larSimplexFacets(FV)
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS((V,EV))))
     VV = larSimplexFacets(EV)
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS((V,VV))))
Macro referenced in 33d.
\langle Oriented boundary matrix visualization 34b\rangle \equiv
     np.set_printoptions(threshold='nan')
     csrSignedBoundaryMat = signedBoundary (FV,EV)
     Z = csr2DenseMatrix(csrSignedBoundaryMat)
     print "\ncsrSignedBoundaryMat =\n", Z
     from pylab import *
     matshow(Z)
     show()
Macro referenced in 33d.
\langle Computation of oriented boundary cells 34c\rangle \equiv
     boundaryCells_1 = signedBoundaryCells(V,FV,EV)
     print "\nboundaryCells_1 =\n", boundaryCells_1
     def swap(mylist): return [mylist[1]]+[mylist[0]]+mylist[2:]
     boundaryEV = [EV[-k] if k<0 else swap(EV[k]) for k in boundaryCells_1]
     bndry = (V,boundaryEV)
     VIEW(EXPLODE(1.5,1.5,1.5)(MKPOLS(bndry)))
Macro referenced in 33d.
      Boundary of random 2D simplicial complex
"test/py/larcc/test06.py" 34d \equiv
     import sys; sys.path.insert(0, 'lib/py/')
     from simplexn import *
     from larcc import *
     from scipy.spatial import Delaunay
     ⟨ Test for quasi-equilateral triangles 35a⟩
      (Generation and selection of random triangles 35b)
     (Boundary computation and visualisation 36a)
```

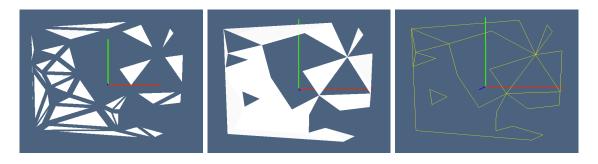


Figure 4: example caption

```
\langle Test for quasi-equilateral triangles 35a \rangle \equiv
     def quasiEquilateral(tria):
          a = VECTNORM(VECTDIFF(tria[0:2]))
          b = VECTNORM(VECTDIFF(tria[1:3]))
          c = VECTNORM(VECTDIFF([tria[0],tria[2]]))
         m = max(a,b,c)
          if m/a < 1.7 and m/b < 1.7 and m/c < 1.7: return True
          else: return False
Macro referenced in 34d.
\langle Generation and selection of random triangles 35b \rangle \equiv
     verts = np.random.rand(20,2)
     verts = (verts - [0.5, 0.5]) * 2
     triangles = Delaunay(verts)
     cells = [ cell for cell in triangles.vertices.tolist()
               if (not quasiEquilateral([verts[k] for k in cell])) ]
     V, FV = AA(list)(verts), cells
     EV = larSimplexFacets(FV)
     pols2D = MKPOLS((V,FV))
     VIEW(EXPLODE(1.5,1.5,1.5)(pols2D))
```

Macro referenced in 34d.

```
\langle Boundary computation and visualisation 36a\rangle \equiv
     boundaryCells_1 = signedBoundaryCells(V,FV,EV)
     print "\nboundaryCells_1 =\n", boundaryCells_1
     def swap(mylist): return [mylist[1]]+[mylist[0]]+mylist[2:]
     boundaryEV = [EV[-k] if k<0 else swap(EV[k]) for k in boundaryCells_1]
     bndry = (V,boundaryEV)
     VIEW(STRUCT(MKPOLS(bndry) + pols2D))
     VIEW(COLOR(RED)(STRUCT(MKPOLS(bndry))))
Macro referenced in 34d.
\langle Compute the topologically ordered chain of boundary vertices 36b\rangle \equiv
Macro never referenced.
\langle Decompose a permutation into cycles 36c\rangle \equiv
     def permutationOrbits(List):
        d = dict((i,int(x)) for i,x in enumerate(List))
        out = []
         while d:
            x = list(d)[0]
            orbit = []
            while x in d:
               orbit += [x],
               x = d.pop(x)
            out += [CAT(orbit)+orbit[0]]
        return out
     if __name__ == "__main__":
        print [2, 3, 4, 5, 6, 7, 0, 1]
        print permutationOrbits([2, 3, 4, 5, 6, 7, 0, 1])
        print [3,9,8,4,10,7,2,11,6,0,1,5]
        print permutationOrbits([3,9,8,4,10,7,2,11,6,0,1,5])
```

Macro never referenced.

## A.7 Assemblies of simplices and hypercubes

```
"test/py/larcc/test07.py" 37a ≡

import sys; sys.path.insert(0, 'lib/py/')

from simplexn import *

from larcc import *

from largrid import *

⟨Definition of 1-dimensional LAR models 37b⟩

⟨Assembly generation of squares and triangles 37c⟩

⟨Assembly generation of cubes and tetrahedra 38⟩

⋄
```

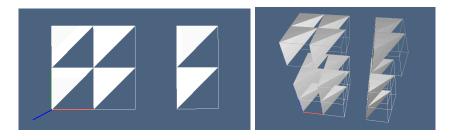


Figure 5: (a) Assemblies of squares and triangles; (b) assembly of cubes and tetrahedra.

Macro referenced in 37a.

Macro referenced in 37a.

# References

[CL13] CVD-Lab, *Linear algebraic representation*, Tech. Report 13-00, Roma Tre University, October 2013.