# Boundary operators on LAR $^\ast$

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#### Abstract

The various versions of boundary operators on Linear Algebraic Representation of cellular complexes are developed in this module, in order to maintain under focus their proper development, including the possible special cases.

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# 1 Introduction

In the current LarLib implementation, we have to distinguish between between dimension-independent, dimension-dependent, oriented and non-oriented operators. Therefore a code refactoring of LarLib—related to boundary/coboundary operators—started here, with the aim of both providing a precise mathematical definition within the LAR framework, and to simplify and generalise the implemented algorithms.

# 2 Implementation

We start this section by making a distinction between the (matrices of) boundary operators for the linear spaces  $C_k$  of chains over the field  $\mathbb{Z}_2 = \{0,1\}$  and over the field  $\mathbb{Z}$  of integer numbers. We call either non-oriented or oriented the corresponding boundary operators, respectively, since the matrix elements take values within the sets  $\{-1,0,+1\}$  or  $\{0,1\}$ , correspondingly. Of course, the associated matrices of coboundary operators are their transpose matrices.

## 2.1 Non-oriented operators

For several computations, the knowledge of the matrices of non-oriented boundary operators is sufficient. Therefore we will use such tool wherever possible, since its computation is much faster in term of computing time.

In the following we provide be binary operator matrices provided by two implementations, respectively named boundary and larUnsignedBoundary2. The first one works correctly only with convex cells; the second one works also with non-convex but path-connected cells.

#### 2.1.1 Dimension-independence

As we show in the following, in order to compute the non-oriented boundary operator  $\partial_d$ , it is sufficient to have knowledge of the  $M_d$  and  $M_{d-1}$  characteristic matrices of d-cells and their (d-1)-facets, at least in the case of cellular complexes with convex cells. Conversely, for more general non-convex but simply-connected cells, also the  $M_{d-2}$  matrix is needed.

Convex-cells The algorithm used is pretty easy to present. The compressed characteristic matrices of d-cells and (d-1)-cells, denoted as cells and facets, respectively, are first put in csr format as csrCV and csrFV. Then the incidence matrix csrFC in compressed sparse row format is computed by matrix product of the compressed characteristic matrices.

The element (i, j) of this matrix provides the number of vertices in the intersection of facet i and cell j, whereas the number of non-zero elements in each csrFV row gives the number of vertices of the facet represented by the row, and is stored in facetLengths.

The boundary function—to be used only with dimension-independent LAR convex cells—is written efficiently in the following script, by using only the standard functions and attributes of the scipy.sparse module.

The variable facetCoboundary stores in a list, for every facet (for h in range(m)) the list of cells in its *coboundary*, to be stored in the output csr\_matrix boundary matrix as column indices of elements with non-zero (i.e. 1) value.

Notice that both the computation of facetCoboundary contents, and the output of the compressed boundary matrix, are performed in the most efficient way—according to the internal design of the scipy's csr sparse data structure.

```
\langle \text{ convex-cells boundary operator } 3a \rangle \equiv
     """ convex-cells boundary operator --- best implementation """
     def larBoundary(cells,facets):
         lenV = max(max(CAT(cells)), max(CAT(facets)))+1
         csrCV = csrCreate(cells,lenV)
         csrFV = csrCreate(facets,lenV)
         csrFC = csrFV * csrCV.T
         facetLengths = [csrFacet.getnnz() for csrFacet in csrFV]
         m,n = csrFC.shape
         facetCoboundary = [[csrFC.indices[csrFC.indptr[h]+k]
              for k,v in enumerate(csrFC.data[csrFC.indptr[h]:csrFC.indptr[h+1]])
                  if v==facetLengths[h]] for h in range(m)]
         indptr = [0]+list(cumsum(AA(len)(facetCoboundary)))
         indices = CAT(facetCoboundary)
         data = [1]*len(indices)
         return csr_matrix((data,indices,indptr),shape=(m,n),dtype='b')
```

Macro referenced in 23b.

#### 2.2 Non-convex LAR cells

A more general larUnsignedBoundary2 operator is given in the following, aiming at compute the boundary matrix for general non-convex cellular decompositions, including multiply connected LAR models. Notice that in this case an input triple made by CV, FV, and EV is needed, where—more in general embedded in  $\mathbf{E}^d$ —they stand for the (binary compressed) characteristic matrices  $M_d$ ,  $M_{d-1}$ , and  $M_{d-2}$ .

#### Boundary operator from 3-chains to 2-chains

```
\langle path-connected-cells boundary operator 3b \rangle \equiv
```

```
""" path-connected-cells boundary operator """
     def larUnsignedBoundary2(CV,FV,EV):
         out = larBoundary(CV,FV)
         def csrRowSum(h):
             return sum(out.data[out.indptr[h]:out.indptr[h+1]])
         unreliable = [h for h in range(len(FV)) if csrRowSum(h) > 2]
         if unreliable != []:
             csrBBMat = larBoundary(FV,EV) * larBoundary(CV,FV)
             lenV = max(max(CAT(CV)), max(CAT(FV)), max(CAT(EV)))+1
             FE = larcc.crossRelationO(lenV,FV,EV)
             out = csrBoundaryFilter2(unreliable,out,csrBBMat,CV,FE)
         return out
     def boundary3(CV,FV,EV):
         out = larUnsignedBoundary2(CV,FV,EV)
         lenV = max(max(CAT(CV)), max(CAT(FV)), max(CAT(EV)))+1
         VV = AA(LIST)(range(lenV))
         csrBBMat = scipy.sparse.csc_matrix(larBoundary(FV,EV) * larUnsignedBoundary2(CV,FV,EV))
         def csrColCheck(h):
             return any([val for val in csrBBMat.data[csrBBMat.indptr[h]:csrBBMat.indptr[h+1]] if v
         unreliable = [h for h in range(len(CV)) if csrColCheck(h)]
         if unreliable != []:
             FE = larcc.crossRelationO(lenV,FV,EV)
             out = csrBoundaryFilter3(unreliable,out,csrBBMat,CV,FE)
         return out
Macro defined by 3b, 5a.
```

Boundary operator from 2-chains to 1-chains First the boundary operator for the convex case is computed within the out variable of csr\_matrix type. Then every out row (i.e. every (d-1)-facet of the d-complex) is tested for reliability, since every (d-1)-face can be shared by at most two d-cells in a d-complex. When this condition is not satisfied, deeper tests are needed to understand what row elements must be forced to value 1, since the (d-1)-face itself is a subset, but not actually a facet, of the corresponding d-cell.

Macro referenced in 23b.

In presence of some "unreliable" facets, the matrix csrBBMat of the operator  $\partial_{d-1} \circ \partial_d$  and the relation FE between faces of dimensions d-1 and d-2 are computed. Now, let us notice that the columns of csrBBMat report the number of incidences of the d-2 faces (as belonging to (d-1)-facets embedded on the boundary) and d-cells (that are associated to such matrix columns). Hence, in a regular (convex) d-complex, such numbers are always even, and in  $\mathbb{Z}_2$  arithmetic are reduced to zero, in order to satisfy the fundaments equation  $\partial \partial = 0$ .

Conversely, with non-convex LAR cells, some incidence numbers may get odd values, due to the non-strict coincidence between cell facets and vertex subsets. Therefore, for

"unreliable" h rows (facets) the csrBBMat columns tracked by ones in  $[\partial_d]$  are checked, looking for elements of (h, k) indices with value greater that 2.

```
\langle path-connected-cells boundary operator 5a \rangle \equiv
     """ path-connected-cells boundary operator """
     import larlib
     import larcc
     from larcc import *
     def csrBoundaryFilter2(unreliable,out,csrBBMat,cells,FE):
         for row in unreliable:
             for j in range(len(cells)):
                  if out[row,j] == 1:
                      cooCE = csrBBMat.T[j].tocoo()
                      flawedCells = [cooCE.col[k] for k,datum in enumerate(cooCE.data)
                      if all([facet in flawedCells for facet in FE[row]]):
                           out[row,j]=0
         return out
     def csrBoundaryFilter3(unreliable,out,csrBBMat,cells,FE):
         for col in unreliable:
              cooCE = csrBBMat.T[col].tocoo()
             flawedCells = [cooCE.col[k] for k,datum in enumerate(cooCE.data)
                          if datum>2]
             for j in range(out.shape[0]):
                  if out[j,col] == 1:
                      if all([facet in flawedCells for facet in FE[j]]):
                          out[j,col]=0
         return out
Macro defined by 3b, 5a.
Macro referenced in 23b.
\langle From cells and facets to boundary cells 5b\rangle \equiv
     def totalChain(cells):
         return csr_matrix(len(cells)*[[1]])
     def boundaryCells(cells,facets):
         csrBoundaryMat = larBoundary(cells,facets)
         csrChain = csr_matrix(totalChain(cells))
         csrBoundaryChain = csrBoundaryMat * csrChain
         out = [k for k,val in enumerate(csrBoundaryChain.data.tolist()) if val == 1]
         return out
```

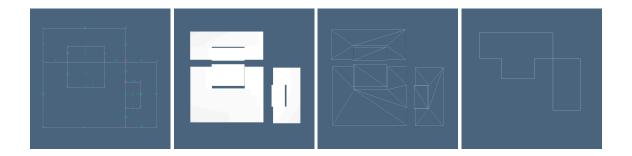


Figure 1: Non-convex LAR 2-complex with (two) 1-cells that are subsets of 2-cells without being their facets. Correctly disentangled by the larUnsignedBoundary2() function: (a) Indexing of 0-, 1-, and 2-cells; (b) exploded 2-cells; (c) triangulated and exploded 2-cells; (d) boundary of the 2-chain [1,1,1,1,1,0].

```
def boundary2Cells(cells,facets,faces):
    csrBoundaryMat = larUnsignedBoundary2(cells,facets,faces)
    csrChain = csr_matrix(totalChain(cells))
    csrBoundaryChain = csrBoundaryMat * csrChain
    out = [k for k,val in enumerate(csrBoundaryChain.data.tolist()) if val == 1]
    return out

def boundary3Cells(cells,facets,faces):
    csrBoundaryMat = boundary3(cells,facets,faces)
    csrChain = csr_matrix(totalChain(cells))
    csrBoundaryChain = csrBoundaryMat * csrChain
    out = [k for k,val in enumerate(csrBoundaryChain.data.tolist()) if val == 1]
    return out
```

#### 2.3 Correctness proof

Our goal is to get a constructive and, of course, correct representation of the matrix  $[\partial_3]$  starting only from  $M_1$ ,  $M_2$ , and  $M_3$ .

We have sufficient evidence here to support the correctness of our identification of the matrices of boundary operators as discussed in the previous section. Remember that  $M_1$ ,  $M_2$ , and  $M_3$  are the characteristic matrices of 1-cells, 2-cells and 3-cells as subsets of vertices, and that  $C_0, C_1, C_2, C_3$  are the linear spaces of 0-, 1-, 2-, and 3-chains, with coefficients in the field  $\mathbb{Z}_2 = \{0, 1\}$ .

In the following we give a dimension-independent proof, even our implementation is currently restricted to  $d \in \{1, 2, 3\}$ .

**Preamble** It is well known that a linear transformation  $T: V \to W$  between two linear spaces, with fixed bases of dimension n and m respectively, is represented uniquely by a matrix  $A \in \mathbb{R}^{m \times n}$ , that by columns contains the coordinate representations in W of the basis vectors of V.

**Theorem 1.** Consider the linear transformation  $\partial_d : C_d \to C_{d-1}$ . Having fixed the bases, made by singleton cells in  $C_d$  and  $C_{d-1}$ , the transformation  $\partial_d$  can be represented as a matrix product  $c \mapsto [\partial_d] c$ , where  $c \in C_d$  is the coordinate representation of a d-chain, and  $[\partial_d]$  is the  $m \times n$  binary matrix having by columns the coordinate representation in  $C_{d-1}$  of the boundary (d-1)-chains of cells in  $C_d$ .

*Proof.* The computation of the boundary operator matrix  $A_d = [\partial_d] \in \mathbb{Z}_2^{m \times n}$ , where m and n are the dimensions of the linear spaces  $C_d$  and  $C_{d-1}$ , respectively, is made in two steps, in the general case.

**First step** Consider the characteristic matrices  $M_d \in \mathbb{Z}_2^{m \times q}$  and  $M_{d-1} \in \mathbb{Z}_2^{n \times q}$  having as rows the images of characteristic functions of bases elements as subsets of vertices, where q is the number of vertices.

Then consider the product matrix  $M = M_{d-1}M_d^t = (m_{ij})$ , with values in the set  $\mathbb{N}$  of non-negative integers. Clearly  $m_{ij}$  will coincide with the number of vertices shared by cells  $c_i \in C_{d-1}$  and  $c_j \in C_d$ , i.e. with the cardinality of their intersection as discrete sets of vertices. The predicate

$$m_{ij} \equiv |c_i| \tag{1}$$

is a necessary condition for  $c_i$  to be a facet of  $c_j$ . Using the Iverson bracket notation, where [P] returns either 1 or 0 depending on the truth of predicate P, we can assign (to) the (i,j) element of our (tentative) boundary matrix the corresponding value:

$$A_d(i,j) := [m_{ij} \equiv |c_i|], \qquad 1 \le i \le m, \ 1 \le j \le n.$$
 (2)

Unfortunately, condition (1) is also sufficient for  $c_i$  being a facet of  $c_j$  only when both are convex cells. In other words, Equation (2) with  $A_d = [\partial_d]$  holds in full generality if and only if the cellular complex under consideration is made only by convex cells. With more general cells, the column j of the  $\partial_d$  matrix contains the coordinate representation in  $C_{d-1}$  of a possibly proper superset of  $\partial_d(c_j)$ .

**Second step** In order to reduce the columns of the approximate boundary matrix  $A_d$  to their exact value in  $\mathbb{Z}_2^m$ , with  $m = \dim C_{d-1}$ , we may compute an (approximate) matrix  $B \in \mathbb{N}^{p \times n}$  of the operator  $\partial_{d-1} \circ \partial_d : C_d \to C_{d-2}$ , with  $p = \dim C_{d-2}$ , by using the approximate representation  $A_d$  of the matrix  $[\partial_d]$ . Let us remark that the exact value of the latter is yet unknown at this point.

According to what asserted in the preamble, every column of this matrix should contain the coordinate representation (in  $C_{d-2}$ ) of the boundary of the boundary of a basis element in  $C_d$ , i.e. of a singleton d-chain.

Therefore, we will enforce the validity of the constraints  $\partial \partial = 0$ , by checking the values of columns in the product matrix  $B_d = A_{d-1}A_d \in \mathbb{N}^{p \times n}$ . In particular, every unit vector  $e_j$ , *i.e.*, the coordinate representation of a singleton chain  $c_j \in C_d$ , should be mapped by  $B_d$  to the zero vector in  $\mathbb{Z}_2^p$ :

$$e_j \mapsto B_d \, e_j = 0^p \in \mathbb{Z}_2^p, \qquad c_j \in C_d.$$
 (3)

Now, let consider the cells of LAR both as subsets of vertices and as cells of a cellular complex. Actually, in the LAR general case, there may be some (d-1)-facets of some d-cell that are subsets of other d-cells of which they are not faces. It is not difficult to provide some examples of this fact (see Figure ??).

When considering each column of B as the coordinate representation of the boundary (d-2)-chain of a d-cell, and noting that every d-cell must be orientable, and hence separating an interior space from an exterior space, we conclude that the number of occurrencies of each (d-2)-cell in a B column must be necessarily even. In particular, it must be either equal to 2 if the column (the d-cell) is locally manifold, or equal to some even number > 2 if the d-cell is locally non-manifold. But odd incidencies of (d-2)-cells along the d-cell boundary are not allowed.

Therefore, for each column  $B_j$   $(1 \le j \le n)$  we look for the subset of rows (i.e. boundary (d-2)-faces of  $c_j$ ) of odd value. If there are none, the column  $A_j$ , associated to  $c_j$ , is a correct representation of  $[\partial_d]_j$ . Otherways, we must look for the subsets of (d-1)-cells incorrectly considered facets of  $c_j$ .

Let us call  $R_j$  the subset of row indices corresponding to odd values in column  $B_j$ , and consider the subset of  $A_j$  rows with value  $a_{ij} = 1$ , i.e. the superset  $S_j$  of (d-1)-cells including the boundary (d-1)-chain of  $c_j$ . The redundant vertex subsets that are not boundary facets of  $c_j$  are easily discovered by looking at the (d-2)-boundary of each  $s \in S_j$ .

In particular, the (d-1)-cell  $s \in S_j$  is redundant or extraneous with respect to the  $c_j$  boundary if and only if  $\partial_{d-1}(s) \subseteq R_j$ , because that property has certainly introduced a spurious increment for each (d-2)-facet in the count of incidences of boundary facets.

At this point, we can finally compute the actual boundary matrix  $\partial_d$  for a general LAR cellular complex, using again the Iverson brackets:

$$[\partial_d]_{ij} = [(a_{ij} = 1) \land ((R_j = \emptyset) \lor (R_j \not\supseteq \partial_{d-1}(s_i)))] \qquad 1 \le i \le m, \ 1 \le j \le n$$

In words it sounds that the element (i, j) of the boundary matrix  $[\partial]_d$  equals that of the "approximate" matrix  $A_d$  if and only if either the redundant set  $R_j$  is empty, or if it does not contain the (d-1)-boundary chain of the i-th (d-1)-cell.

Just remember that the redundant set  $R_j$  contains the (d-2)-faces with odd incidencies on  $c_j \in C_d$ , computed via the "approximate" matrix  $B_d = A_{d-1} \circ A_d$ .

2.4 Oriented operators

#### 2.4.1 Oriented simplicial complexes

#### 2.4.2 Oriented LAR complexes

# 3 From relations to operators

The LAR approach to topology, implemented in the LarLib modules, allows the user to consider the topological relations of incidence and adjacency between faces of a cellular complex as *linear operators* between chains of cells of various dimensions.

The previous approach was to consider the incidence and adjaceny as set-theoretical relations, to be solved by using typical database tools. Conversely, according to the novel IT evolution towards big data and cloud-based storage, even for geometrical data, LarLib takes advantage of a conceptual framework based on linear-algebra and sparse matrices.

### 3.1 Classification of operators

In the standard solid modeling approach, mainly based on boundary representations, the standard topological operations concern the answers to queries, by reporting the subsets of boundary elements which are incident (different dimension) or adjacent (equal dimension) to assigned boundary elements. Boundary elements stand there for three type of boundary cells: aka  $faces\ F$ ,  $edges\ E$ , and  $vertices\ V$ . Nine binary relations may be considered, that are summarized in Table 1.

Table 1: Binary topological relations between boundary elements in boundary representations of solid models

	F	Е	V
F	FF	FE	FV
$\mathbf{E}$	EF	EE	EV
V	VF	VE	VV

Conversely, LAR models are normally based on cellular 3-complexes, so using four sets of cells, namely 3-cells C, 2-cells F, 1-cells E, and 0-cells V. The resulting tables of topological relations, and the associated linear operators, are shown in Tables 2a and 2b, respectively.

Table 2: Binary topological relations between cells of LAR decompositions of solid models and corresponding topological operators on  $\partial_{\circ}$  chains.

	С	F	Ε	V
C	CC	$\overline{\mathrm{CF}}$	CE	CV
F	FC	FF	$\overline{\text{FE}}$	FV
E	EC	$\mathbf{EF}$	EE	$\mathrm{EV}$
V	VC	$\overline{\mathrm{VF}}$	VE	VV

	С	F	E	V
С	$1_C^{ op} \circ 1_C$	$\partial_3$	$\partial_2 \circ \partial_3$	$1_C$
F	$\delta_2$	$1_F^\top \circ 1_F$	$\partial_2$	$1_F$
E	$\delta_2 \circ \delta_1$	$\delta_1$	$1_E^\top \circ 1_E$	$1_E$
V	$1_C^ op$	$1_F^\top$	$1_E^ op$	$1_V$

# 3.2 Topological relations

### 3.2.1 Adjacency relations

#### Adjacency relations examples

```
"test/py/boundary/test09.py" 10b =
    """ Adjacency relations examples """
    from larlib import *

    sys.path.insert(0, 'test/py/boundary/')
    from test07 import *

        (kfaces-to-kfaces relation 10a)
    print "\nCC =",CC
    print "\nFF =",FF
    print "\nEE =",EE,"\n"

        V,BF,BE = larUnsignedBoundary3(V,CV,FV,EV)([1,0])
        VIEW(STRUCT(MKTRIANGLES((V,[FV[h] for h in FF[-1]],EV),color=True)))
        VIEW(STRUCT(MKPOLS((V,[EV[h] for h in EE[-1]]))+[COLOR(RED)(MKPOLS((V,[EV[-1]]))[0])]))
```

#### 3.2.2 Incidence relations

```
⟨ mfaces-to-nfaces relations 11a⟩ ≡

""" mfaces-to-nfaces relations """

fcOp = larCells2Faces(CV,FV,EV)

CF = [fcOp([k]) for k in range(len(CV))]

FC = invertRelation(CF)

ecOp = larCells2Edges(CV,FV,EV)

CE = [ecOp([k]) for k in range(len(CV))]

EC = invertRelation(CE)

efOp = larFaces2Edges(FV,EV)

FE = [efOp([k]) for k in range(len(FV))]

EF = invertRelation(FE)

⋄

Macro referenced in 11b, 16a.
```

### Incidence relations examples

```
"test/py/boundary/test10.py" 11b =
    """ Incidence relations examples """
    from larlib import *

    sys.path.insert(0, 'test/py/boundary/')
    from test08 import *

    ⟨mfaces-to-nfaces relations 11a⟩
    print "\nFC =",FC
    print "\nEC =",EC
    print "\nEF =",EF,"\n"
    ⋄
```

### 3.3 Querying

The more important topological operations in a geometric system concern the answer to queries of the type: "what is the h-chain whose cells are (k-h)-incident to a given h-chain"?

An efficient answer is given by the three higher-level functions in this section, respectively denoted as larCells2Faces, larCells2Edges, and larFaces2Edges. Their first application, over the two or three necessary (compressed) characteristic matrices, returns the csr\_matrix of the topological operator, that can be so cached by the calling code. The second application, over the list of h-chain indices, returns the list of k-chain indices of the cells that share with them a (k-h)-face.

#### 3.3.1 Topological incidences

### Query from 3-chain to incident 2-chain

Macro referenced in 23b.

#### Query from 3-chain to incident 1-chain

Macro referenced in 23b.

Compute the signed 2-boundary matrix. Compute the signed 2-boundary matrix. The second definition extends the first one by considering LAR faces including holes. For this purpose the signed areas of all loops inside each lar face in FV are taken into account, and only the greates positive area and all the negative areas (except the most negative one) are considered.

```
⟨ Compute the signed 2-boundary matrix 12c⟩ ≡
    """ Compute the signed 2-boundary matrix """
    import triangulation,integr

def larSignedBoundary2(V,FV,EV):
```

```
efOp = larFaces2Edges(V,FV,EV)
   FE = [efOp([k]) for k in range(len(FV))]
   data,row,col = [],[],[]
   for f in range(len(FE)):
        cycle_data = triangulation.find_cycles(V,[EV[e] for e in FE[f]])
        vcycles = cycle_data['v_cycles']
        ecycles = cycle_data['e_cycles']
        pairs = zip(vcycles,ecycles)
        vertEdgeCycles = [zip(vcycle,[FE[f][e] for e in ecycle]) for vcycle,ecycle in pairs]
        vertEdgeCycles = sorted(vertEdgeCycles,key=len)[:-1]
        coefficients = [1 if v == EV[e][0] else -1 for cycle in vertEdgeCycles for (v,e) in cy
        ecycle = [e for cycle in vertEdgeCycles for (v,e) in cycle]
        data += coefficients
        row += ecycle
        col += [f]*len(ecycle)
        #print f,len(data),len(row),len(col)
    signedBoundary2 = coo_matrix((data,(row,col)), shape=(len(EV),len(FV)),dtype='b')
   return csr_matrix(signedBoundary2)
def larSignedBoundary2(V,FV,EV):
   efOp = larFaces2Edges(V,FV,EV)
   FE = [efOp([k]) for k in range(len(FV))]
   data,row,col = [],[],[]
   for f in range(len(FE)):
        Vcycles,Ecycles = triangulation.makeCycles((V,[EV[e] for e in FE[f]]))
        print "\nVcycles =",Vcycles
        print "Ecycles =",Ecycles
        Ecycles = [[FE[f][e] for e in cycle] for cycle in Ecycles]
        print "Ecycles =",Ecycles
        areas = integr.signedSurfIntegration((V,Vcycles,EV),signed=True)
        print "areas =",areas
        sortedAreas = sorted((area,k) for k, area in enumerate(areas))
        print "sortedAreas =",sortedAreas
        innerLoops = [zip(Vcycles[k], Ecycles[k]) for area,k in sortedAreas[1:] if area<0]
        print "innerLoops =",innerLoops
        outerLoop = [zip(Vcycles[sortedAreas[-1][1]],Ecycles[sortedAreas[-1][1]])]
        print "outerLoop =",outerLoop
        orientedFaceLoops = CAT(outerLoop+innerLoops)
        print "orientedFaceLoops =",orientedFaceLoops
        coefficients = [1 if v==EV[e][0] else -1 for v,e in orientedFaceLoops]
        print "coefficients =",coefficients
        ecycle = [e for v,e in orientedFaceLoops]
        print "ecycle =",ecycle
        data += coefficients
```

```
row += ecycle
col += [f]*len(ecycle)
    #print f,len(data),len(row),len(col)
signedBoundary2 = coo_matrix((data,(row,col)), shape=(len(EV),len(FV)),dtype='b')
return csr_matrix(signedBoundary2)
```

### Query from 2-chain to incident 1-chain

```
⟨ Query from 2-chain to incident 1-chain 14a⟩ ≡
    """ Query from 2-chain to incident 1-chain """

def larFaces2Edges(V,FV,EV):
    VV = AA(LIST)(range(len(V)))
    csrEF = larUnsignedBoundary2(FV,EV,VV)
    def larCells2Faces0(chain):
        chainCoords = csc_matrix((csrEF.shape[1],1),dtype='b')
        for k in chain: chainCoords[k,0] = 1
        out = csrEF * chainCoords
        return out.tocoo().row.tolist()
        return larCells2Faces0
```

Macro referenced in 23b.

#### 3.3.2 Topological adjacencies

#### kfaces-to-kfaces relations

```
\langle kfaces-to-kfaces relations """

def larCells2Cells(CV,FV,EV):
    csrMat = boundary3(CV,FV,EV)
    csrCC = csrMat.T * csrMat
    def larCells2Cells0(chain):
        chainCoords = csc_matrix((csrCC.shape[1],1),dtype='b')
        for k in chain: chainCoords[k,0] = 1
        out = csrCC * chainCoords
        return out.tocoo().row.tolist()
    return larCells2Cells0

def larFaces2Faces(FV,EV):
    lenV = max(CAT(FV)) + 1
    VV = AA(LIST)(range(lenV))
    csrMat = larUnsignedBoundary2(FV,EV,VV)
```

```
csrFF = csrMat.T * csrMat
   def larFaces2Faces0(chain):
        chainCoords = csc_matrix((csrFF.shape[1],1),dtype='b')
        for k in chain: chainCoords[k,0] = 1
        out = csrFF * chainCoords
        return out.tocoo().row.tolist()
   return larFaces2Faces0
def larEdges2Edges(EV,VV):
   lenV = len(VV)
   csrMat = larBoundary(EV,VV)
   csrEE = csrMat.T * csrMat
   def larFaces2Faces0(chain):
        chainCoords = csc_matrix((csrEE.shape[1],1),dtype='b')
        for k in chain: chainCoords[k,0] = 1
        out = csrEE * chainCoords
        return out.tocoo().row.tolist()
   return larFaces2Faces0
```

#### 3.4 Oriented operators

#### 3.4.1 Signed 2-boundary

**Signed 2-boundary** An efficient implementation of the sparse matrix of the signed 2-boundary operator  $\partial_2$  follows. It is computed using also the  $M_2$  and  $M_1$  characteristic matrices, or — better — their binary compressed representations FV and EV.

```
\langle \text{ Signed 2-boundary 15} \rangle \equiv
     """ Signed 2-boundary """
     def larSignedBoundary2(FV,EV):
         efOp = larFaces2Edges(FV,EV)
         FE = [efOp([k]) for k in range(len(FV))]
         data,row,col = [],[],[]
         for f in range(len(FE)):
              cycle_data = find_cycles(V,[EV[e] for e in FE[f]])
              coefficients = [cycle_data['ev_mapping'][k]['direction']
                    for k in range(len(cycle_data['ev_mapping']))]
              ecycle = FE[f]
             data += coefficients
             row += ecycle
              col += [f]*len(ecycle)
              #print f,len(data),len(row),len(col)
         signedBoundary2 = coo_matrix((data,(row,col)), shape=(len(EV),len(FV)),dtype='b')
         return csr_matrix(signedBoundary2)
```

 $\Diamond$ 

Macro never referenced.

#### Testing signed 2-boundary

## Transformation from chain coordinates to explicit chain data

```
⟨Transformation from chain coordinates to explicit chain data 16b⟩ ≡
    """ Transformation from chain coordinates to explicit chain data """
    def coords2chain(chainCoords):
        coo = coo_matrix(chainCoords)
        return [(e,val) for e,val in zip(coo.row,coo.data)]
        ◊
```

Macro referenced in 23b.

#### 3.5 Extraction of 3-cells from a 2-skeleton embedded in 3D

Need to characterize the two vertices of zero edge as "last" of previous edge in the chain, and "first" of following edge in the chain. Therefore the zero edge is oriented as from "last" to "first" and consequently, is "positive" iff "first" i, "last".

It is sufficient to characterize the "first" vertex of the edge following the zero edge. This one is the "last" of zero edge. the other vertex of zero edge is its "first". Zero edge is "positive" iff last(zero) ; first(zero).

Choose the next face on ordered coboundary of edge Choose the "next" face  $g_i$  on "ordered" coboundary of edge

```
⟨ Choose the next face on ordered coboundary of edge 16c⟩ ≡
   """ Choose the next face on ordered coboundary of edge """
   def adjFace(boundaryOperator,EV,EF_angle,faceChainOrientation):
        def adjFaceO(edge,orientation):
        if orientation > 0: edgeLoop = REVERSE(EF_angle[edge])
```

```
elif orientation < 0: edgeLoop = EF_angle[edge]
edgeLoop = edgeLoop + [edgeLoop[0]] # all positive indices

candidates = set([f for f,_ in faceChainOrientation]).intersection(edgeLoop)
if candidates != set([]):
    pivotFace = candidates.pop()
    if pivotFace in edgeLoop:
        pivotIndex = edgeLoop.index(pivotFace)
    else:
        pivotIndex = edgeLoop.index(-pivotFace)
    adjacentFace = edgeLoop[pivotIndex+1]
else: return None

theSign = boundaryOperator[edge,adjacentFace]
    return adjacentFace, -(theSign*orientation)
return adjFaceO</pre>
```

#### Choose the start facet in extracting the facet representation of a cell

 $\langle$  Choose the start facet in extracting the facet representation of a cell 17 $\rangle \equiv$ 

```
""" Choose the start facet in extracting the facet representation of a cell """
def chooseStartFace(FV,faceCounter):
   if faceCounter[0,0]==0: return (0,1)
   for f in range(len(FV)):
        if faceCounter[f,0]==1 and faceCounter[f,1]==0: return (f,-1)
        elif faceCounter[f,0]==0 and faceCounter[f,1]==1: return (f,1)
   for f in range(len(FV)):
        if faceCounter[f,0] == 0 and faceCounter[f,1] == 0: return (f,1)
   if sum(array(faceCounter))==2*len(FV): return (-1,999)
   else: print "ERROR: chooseStartFace"
def chooseStartFace(FV,faceCounter):
   for f in range(len(FV)):
        if faceCounter[f,0]==1 and faceCounter[f,1]==0: return (f,-1)
        elif faceCounter[f,0] == 0 and faceCounter[f,1] == 1: return (f,1)
   for f in range(len(FV)):
        if faceCounter[f,0] == 0 and faceCounter[f,1] == 0: return (f,1)
   if sum(array(faceCounter))==2*len(FV): return (-1,999)
   else: return (0,1)
```

Macro referenced in 23b.

#### Extract the signed representation of a basis element

```
\( \textract \the \textract \the \textract \textra
```

**Algoritm** The algorithm to compute the signed boundary matrix of a 3-complex is given below in pseudocode.

#### Return the signed boundary matrix of a 3-complex

```
\langle Return the signed boundary matrix of a 3-complex 18b\rangle \equiv
     """ Return the signed boundary matrix of a 3-complex """
     import boolean
     def larSignedBoundary3((V,FV,EV)):
         model = V,FV,EV
         faceCounter = zeros((len(FV),2),dtype='b')
         CF,m = [],len(FV)
         efOp = larFaces2Edges(V,FV,EV)
         FE = [efOp([k]) for k in range(len(FV))]
         EF_angle, _,_, = boolean.faceSlopeOrdering(model,FE)
         nonWorkedFaces,coboundary_2,cellNumber = set(range(m)),[],0
         boundaryOperator = larSignedBoundary2(V,FV,EV)
         FEbasis = signedBasis(boundaryOperator)
         row,col,data = [],[],[]
         longestrow,longestcol,longestdata,longestLength = [],[],[],0
         while True:
             startFace,orientation = chooseStartFace(FV,faceCounter)
             if startFace == -1: break
             nonWorkedFaces = nonWorkedFaces.difference({startFace})
             faceChainOrientation = {(startFace,orientation)}
             vect = csc_matrix((m,1),dtvpe='b')
             for face, orientation in faceChainOrientation:
                 vect[face] = orientation
```

# **Algorithm 1** Compute the signed $\partial_3$ matrix of a 3-complex, from 2-skeleton

```
1: function LARSIGNEDBOUNDARY3(LarModel)
         V, FV, EV \leftarrow \text{LarModel}
 2:
         [\partial_2] \equiv [\delta_1]^t \leftarrow \text{LARSIGNEDBOUNDARY2}(\text{LarModel})
 3:
         sort: EV \to FV \to \mathbb{R}^*: e \mapsto \delta_1(e) (ordered loops of signed 2-faces)
 4:
         [\partial_3] \equiv [\delta_2]^t \leftarrow [] (boundary 2-faces of 3-cells by row)
 5:
         S = \text{indices of 2-faces}
 6:
         while S \neq \emptyset (set of non-traversed 2-faces) do
 7:
              f \leftarrow \text{CHOOSE}(S) (first 2-face, reversing previous sign)
 8:
              F \leftarrow \{f\} (singleton 2-chain)
 9:
              while \partial_2(F) \neq \emptyset do (F non closed)
10:
                   E \leftarrow \partial_2(F) (1-cycle of signed edges)
11:
                   while E \neq \emptyset do
12:
                        for each e \in E do
13:
                            f_i \leftarrow next(f)(sort(\delta_1(e)))
14:
                            F \leftarrow F \cup \{f_i\}
15:
                            S \leftarrow S - \{f_i\} (non-traversed 2-faces)
16:
                            E \leftarrow E - \{e\}
17:
                        end for
18:
                        E \leftarrow \partial_2(F) (new oriented 1-cycle)
19:
                   end while
20:
              end while
21:
              [\partial_3] \leftarrow [\partial_3] + [F] (put F in new [\partial_3] column, i.e. new [\delta_2] row)
22:
         end while
23:
24:
         return [\partial_3] (operator's matrix)
25: end function
```

```
edgeCycleCoords = boundaryOperator * vect
             edgeCycle = coords2chain(edgeCycleCoords)
             while edgeCycle != []:
                 look4face = adjFace(boundaryOperator,EV,EF_angle,faceChainOrientation)
                 for edge,orientation in edgeCycle:
                     outPair = look4face(edge,orientation)
                     if outPair != None:
                         adjacentFace, orientation = outPair
                         faceChainOrientation = faceChainOrientation.union(
                              [(adjacentFace, orientation)])
                         nonWorkedFaces = nonWorkedFaces.difference([adjacentFace])
                 vect = csc_matrix((m,1),dtype='b')
                 for face, orientation in faceChainOrientation:
                     vect[face] = orientation
                 edgeCycleCoords = boundaryOperator * vect
                 edgeCycle = coords2chain(edgeCycleCoords)
                 #if edgeCycle!=[]: VIEW(STRUCT(MKPOLS((V,[EV[e] for e in TRANS(edgeCycle)[0]]))))
             for face, orientation in faceChainOrientation:
                 if orientation == 1: faceCounter[face,0]+=1
                 elif orientation == -1: faceCounter[face,1]+=1
             #VIEW(STRUCT(MKPOLS((V,[FV[f] for f in TRANS(faceChainOrientation)[0]]))))
             lastrow = [face for face,_ in faceChainOrientation]
             lastcol = [cellNumber for face, orientation in faceChainOrientation]
             lastdata = [orientation for _,orientation in faceChainOrientation]
             lastlength = len(lastrow)
             if lastlength >= longestLength:
                 lastrow,longestrow = longestrow,lastrow
                 lastcol,longestcol = longestcol,lastcol
                 lastdata,longestdata = longestdata,lastdata
                 lastlength,longestLength = longestLength,lastlength
             if lastlength != 0:
                 row += lastrow
                 col += lastcol
                 data += lastdata
                 CF += [lastrow]
                 cellNumber += 1
             print "\nfaceCounter =",faceCounter
         outMatrix = coo_matrix((data, (row,col)), shape=(m,cellNumber),dtype='b')
         signedBoundary = zip(longestrow,longestdata)
         return csr_matrix(outMatrix),CF,signedBoundary
Macro defined by 18b, 21.
Macro referenced in 23b.
```

#### Return the signed boundary matrix of a 3-complex

```
\langle Return the signed boundary matrix of a 3-complex 21 \rangle \equiv
     """ Return the signed boundary matrix of a 3-complex """
     import boolean
     def larSignedBoundary3((V,FV,EV)):
         model = V,FV,EV
         faceCounter = zeros((len(FV),2),dtype='b')
         CF,m = [],len(FV)
         efOp = larFaces2Edges(V,FV,EV)
         FE = [efOp([k]) \text{ for } k \text{ in } range(len(FV))]
         EF_angle, _,_, = boolean.faceSlopeOrdering(model,FE)
         nonWorkedFaces,coboundary_2,cellNumber = set(range(m)),[],0
         boundaryOperator = larSignedBoundary2(V,FV,EV)
         FEbasis = signedBasis(boundaryOperator)
         row,col,data = [],[],[]
         while True:
             startFace,orientation = chooseStartFace(FV,faceCounter)
             if startFace == -1: break
             nonWorkedFaces = nonWorkedFaces.difference({startFace})
             faceChainOrientation = {(startFace, orientation)}
             vect = csc_matrix((m,1),dtype='b')
             for face, orientation in faceChainOrientation:
                 vect[face] = orientation
             edgeCycleCoords = boundaryOperator * vect
             edgeCycle = coords2chain(edgeCycleCoords)
             while edgeCycle != []:
                 look4face = adjFace(boundaryOperator,EV,EF_angle,faceChainOrientation)
                 for edge, orientation in edgeCycle:
                      outPair = look4face(edge,orientation)
                      if outPair != None:
                          adjacentFace, orientation = outPair
                          faceChainOrientation = faceChainOrientation.union(
                              [(adjacentFace, orientation)])
                          nonWorkedFaces = nonWorkedFaces.difference([adjacentFace])
                 vect = csc_matrix((m,1),dtype='b')
                 for face, orientation in faceChainOrientation:
                      vect[face] = orientation
                 edgeCycleCoords = boundaryOperator * vect
                 edgeCycle = coords2chain(edgeCycleCoords)
                 #if edgeCycle!=[]: VIEW(STRUCT(MKPOLS((V,[EV[e] for e in TRANS(edgeCycle)[0]]))))
             row += [face for face,_ in faceChainOrientation]
             col += [cellNumber for face,orientation in faceChainOrientation]
             data += [orientation for _,orientation in faceChainOrientation]
             cellNumber += 1
             for face, orientation in faceChainOrientation:
```

```
if orientation == 1: faceCounter[face,0]+=1
    elif orientation == -1: faceCounter[face,1]+=1
    print "faceChainOrientation =",faceChainOrientation
    print "faceCounter =",faceCounter

CF += [[face for face,_ in faceChainOrientation]]
    print "CF =",CF
    print "\nfaceCounter =",faceCounter

outMatrix = coo_matrix((data, (row,col)), shape=(m,cellNumber),dtype='b')
    return csr_matrix(outMatrix),CF,faceCounter
```

Macro defined by 18b, 21. Macro referenced in 23b.

Macro referenced in 23b.

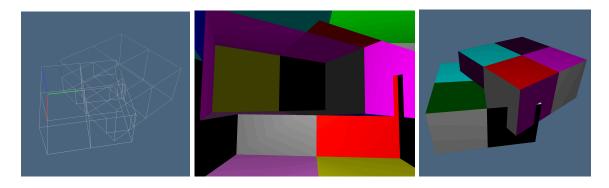


Figure 2: Arrangment of 3-complexes: (a) 1-skeleton of the fragmented (union) 3-complex; (b) view from the interior of the boundary 2-complex; (c) view from the exterior of the 3-complex. Notice that the colour patches correspond to the chain of 2-faces of the boundary of 3-complex.

#### Test the signed boundary matrix of a 3-complex

```
\( \text{Test the signed boundary matrix of a 3-complex 22} \) \( = \)
\( \""" \text{ Test the signed boundary matrix of a 3-complex """ } \)
\( \text{if } \__name__=="__main__": \)
\( \text{V,[VV,EV,FV,CV]} = \text{larCuboids([2,2,1],True)} \)
\( \text{cubeGrid} = \text{Struct([(V,FV,EV)],"cubeGrid")} \)
\( \text{cubeGrids} = \text{Struct(2*[cubeGrid,t(.5,.5,.5),r(0,0,PI/6)])} \)
\( \text{V,FV,EV} = \text{struct2Marshal(cubeGrids)} \)
\( \text{VIEW(EXPLODE(1.2,1.2,1.2)(BREP((V,FV,EV),color=False)))} \)
\( \text{\left}
\)
\( \text{\text{O}}
\)
\( \text{\text{o
```

## 3.6 Examples

#### A 3-cell with several holes

# 4 Exporting

```
"larlib/larlib/boundary.py" 23b \equiv
      """ boundary operators """
      from larlib import *
       (convex-cells boundary operator 3a)
       \langle \text{ path-connected-cells boundary operator } 3b, \dots \rangle
       \langle From cells and facets to boundary cells 5b\rangle
       (Marshalling a structure to a LAR cellular model 32)
        Compute the signed 2-boundary matrix 12c >
       Boundary of a 3-complex 33
        Query from 3-chain to incident 2-chain 12a
        Query from 3-chain to incident 1-chain 12b
        Query from 2-chain to incident 1-chain 14a
       \langle \text{ kfaces-to-kfaces relations } \frac{14b}{\rangle}
        Transformation from chain coordinates to explicit chain data 16b
        Choose the next face on ordered coboundary of edge 16c
        Choose the start facet in extracting the facet representation of a cell 17
       (Extract the signed representation of a basis element 18a)
       \langle Return the signed boundary matrix of a 3-complex 18b, ... \rangle
      \langle Test the signed boundary matrix of a 3-complex 22\rangle
```

# 5 Testing

#### 5.1 Non-oriented operators

Correct boundary extraction example The larBoundary() operator is applied here to a cellular 2-complex of convex cells, producing correct result. It is worth noting that the operator is dimension-independent, and must be applied to the *pair* of compressed characteristic matrices  $M_d$  and  $M_{d-1}$ , that — in list format — we call either CV,FV or FV,EV, depending on the dimension (either 3 or 2) of the embedding space.

```
"test/py/boundary/test01.py" 24a \equiv
     """ testing boundary operators (correct result) """
     from larlib import *
     filename = "test/svg/inters/boundarytest0.svg"
     lines = svg2lines(filename)
     VIEW(STRUCT(AA(POLYLINE)(lines)))
     V,FV,EV,polygons = larFromLines(lines)
     VV = AA(LIST)(range(len(V)))
     submodel = STRUCT(MKPOLS((V,EV)))
     VIEW(larModelNumbering(1,1,1)(V,[VV,EV,FV],submodel,0.2))
     VIEW(EXPLODE(1.2,1.2,1.2)(MKPOLS((V,[EV[e] for e in boundaryCells(FV,EV)],))))
     VIEW(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,FV,EV))))
     boundaryOp = larUnsignedBoundary2(FV,EV,VV)
     for k in range(1,len(FV)+1):
         faceChain = k*[1]
         BF = chain2BoundaryChain(boundaryOp)(faceChain)
         VIEW(STRUCT(MKPOLS((V,[EV[e] for e in BF]))))
     \Diamond
```

Wrong boundary extraction example The larBoundary() operator, applied to a cellular 2-complex wih some non-convex cells, produces incorrect results. In such cases a correct result may be produced only by chance (sometimes this happens). So, be careful to use it only when the precondition (of cell convexity) is everywhere verified. In order to get always a correct result, use the larUnsignedBoundary2 operator.

```
"test/py/boundary/test02.py" 24b =

""" testing boundary operators (wrong result) """

from larlib import *

filename = "test/svg/inters/boundarytest3.svg" # KO (MKTRIANGLES) with boundarytest3 !!!

#filename = "test/svg/inters/boundarytest4.svg"
```

```
lines = svg2lines(filename)
VIEW(STRUCT(AA(POLYLINE)(lines)))
V,FV,EV,polygons = larFromLines(lines)
VV = AA(LIST)(range(len(V)))
submodel = STRUCT(MKPOLS((V,EV)))
VIEW(larModelNumbering(1,1,1)(V,[VV,EV,FV],submodel,0.2))
boundaryOp = larUnsignedBoundary2(FV,EV,VV) # <<=====</pre>
#boundaryOp = larBoundary(FV,EV) # <<===== NB</pre>
BF = chain2BoundaryChain(boundaryOp)([1]*len(FV))
VIEW(EXPLODE(1.2,1.2,1.2)(MKPOLS((V,[EV[e] for e in BF]))))
VIEW(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,FV,EV),color=True)))
VIEW(SKEL_1(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,FV,EV)))))
for k in range(1,len(FV)+1):
    faceChain = k*[1]
    boundaryChain = chain2BoundaryChain(boundaryOp)(faceChain)
    VIEW(STRUCT(MKPOLS((V,[EV[e] for e in boundaryChain]))))
11 11 11
\Diamond
```

**Example** Comparison of two implementations of the  $\partial$  operator. Notice the difference between the penultimate rows. In particular, the penultimate row of the matrix generated by larBoundary(FV,EV) is plain wrong. It means that the edge  $e_{10}$  is shared by all the (three) 2-cells of the complex. Conversely, it is well known that, for a solid complex, i.e. a d-complex embedded in  $\mathbb{E}^d$ , every (d-1)-facet may be shared by no more than 2 d-cells. The resulting boundary of the total chain  $[f_0, f_1, f_2]$  codified in coordinates as [1, 1, 1], and shown in Figure 4d, is sonsequently incorrect.

```
In [1]: larBoundary(FV,EV).todense()
                                                      In [2]: larUnsignedBoundary2(FV,EV,VV).todense()
Out[1]:
                                                      Out[2]:
matrix([[0, 1, 0],
                                                      matrix([[0, 1, 0],
        [0, 0, 1],
                                                              [0, 0, 1],
        [1, 0, 1],
                                                               [1, 0, 1],
        [1, 0, 1],
                                                              [1, 0, 1],
        [0, 1, 1],
                                                               [0, 1, 1],
        [0, 1, 0],
                                                              [0, 1, 0],
        [1, 0, 1],
                                                               [1, 0, 1],
        [0, 0, 1],
                                                               [0, 0, 1],
        [0, 0, 1],
                                                              [0, 0, 1],
        [0, 1, 0],
                                                               [0, 1, 0],
        [1, 1, 1],
                                                               [1, 1, 0],
        [0, 1, 1]])
                                                               [0, 1, 1]])
```

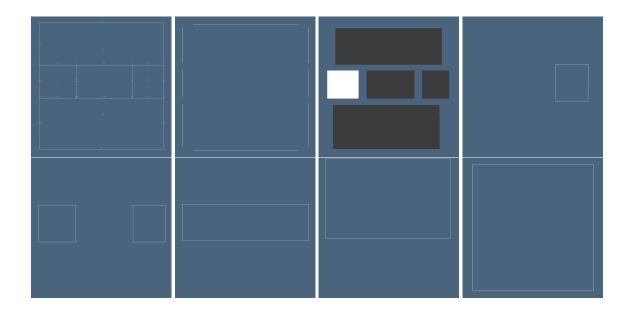


Figure 3: Convex-cell 2-complex. (a) Indexing of 0-,1-,and 2-cells; (b) exploded 2-boundary cells; (c) exploded 2-cells; (d) boundary of a singleton 2-chain; (e-h) boundaries of some 2-chains.

**3D non-convex LAR cells** In this example and in the next one we show the boundary computation of LAR models with non-contractible 3- and 2-cells.

```
"test/py/boundary/test03.py" 26 =
    """ 3D non-convex LAR cells """
    from larlib import *

V = [[0.25,0.25,0.0],[0.25,0.75,0.0],[0.75,0.75,0.0],[0.75,0.25,0.0],[1.0, 0.0,0.0],
    [0.0,0.0,0.0],[1.0,1.0,0.0],[0.0,1.0,0.0],[0.25,0.25,1.0],[0.25, 0.25,2.0],[0.25,0.75,
    2.0],[0.25,0.75,1.0],[0.25,0.75,-1.0],[0.25,0.25, -1.0],[0.75,0.75,-1.0],[0.75,0.25,
    -1.0],[0.75,0.25,1.0],[0.75,0.75,1.0], [1.0,0.0,1.0],[0.0,0.0,1.0],[1.0,1.0,1.0],
    [0.0,1.0,1.0],[0.75,0.75,2.0],[0.75,0.25,2.0]]

CV = [(0,1,2,3,4,5,6,7,8,11,16,17,18,19,20,21), (0,1,2,3,8,11,16,17),
    (0,1,2,3,12,13,14,15), (8,9,10,11,16,17,22,23)]

FV = [(2,3,16,17),(6,7,20,21),(12,13,14,15),(0,1,8,11),(1,2,11,17),(0,1,12,13),
    (4,6,18,20),(5,7,19,21),(0,3,13,15),(0,3,8,16),(0,1,2,3),
    (10,11,17,22),(2,3,14,15),(8,9,16,23),(8,11,16,17),
    (1,2,12,14),(16,17,22,23),(4,5,18,19),(8,9,10,11),(
    9,10,22,23),(0,1,2,3,4,5,6,7),(8, 11,16,17,18,19,20,21)]
```

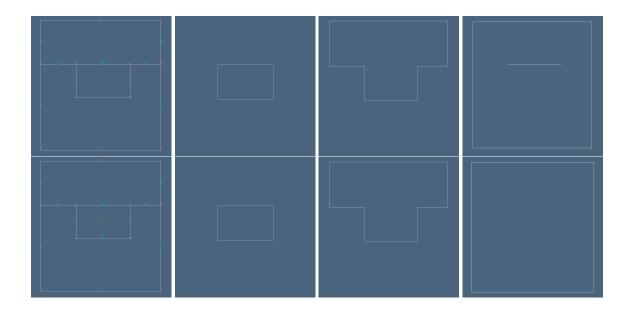


Figure 4: Non-working (i.e. *wrong*) example with boundary. (a) Indexing of 0-,1-,and 2-cells; (b) boundary of a singleton 2-chain; (c) exploded 2-cells; (d) boundary of a singleton 2-chain. Working (i.e. *exact*) example using larUnsignedBoundary2: (e-h) as above.

```
EV =[(3,15),(7,21),(10,11),(4,18),(12,13),(5,19),(8,9),(18,19),(22,23),(0,3),(1,11),
(16,17),(0,8),(6,7),(20,21),(3,16),(10,22),(18,20),(19,21),(1,2),(12,14),(4,5),(
8,11),(13,15),(16,23),(14,15),(11,17),(17,22),(2,14),(2,17),(0,1),(9,10),(8,16),
(4,6),(1,12),(5,7),(0,13),(9,23),(6,20),(2,3)]

VV = AA(LIST)(range(len(V)))
hpc = STRUCT(MKPOLS((V,EV)))
VIEW(larModelNumbering(1,1,1)(V,[VV,EV,FV,CV],hpc,0.6))

BF = boundary3Cells(CV,FV,EV)
VIEW(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,[FV[f] for f in BF],EV),color=True)))
```

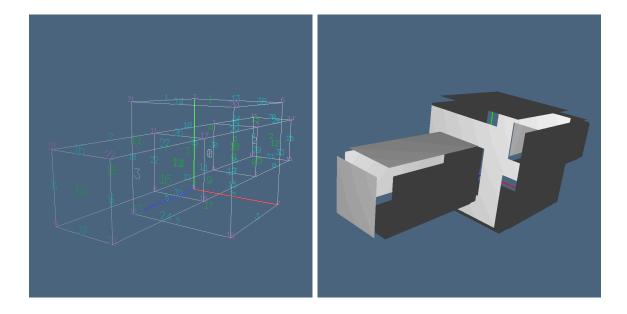


Figure 5: Non-convex 3-complex. (a) Indexing of 0-,1-,2- and 3-cells; (b) exploded 2-boundary cells. Notice that two faces are multiply-connected.

**3D non-convex LAR cells** In this example the 3D model is constructed partly in automated way, partly by hand. In particular, first we generate a structure of cuboidal complexes, then we transform it is a single complex using part of the computational pipeline being developed for the Boolean arrangments of complexes, so that all the included cells are mutually fragmented. Then the 3-cells are assembed as sets of 2-faces, giving the CF (cells-by-faces) variable. Finally this one is transformed automatically into CV (cells-by-vertices).

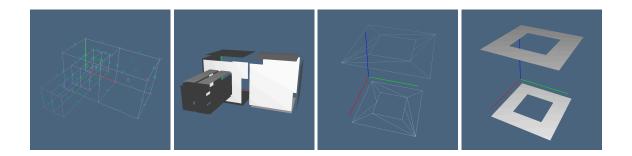


Figure 6: Non-convex 3-complex. (a) Indexing of 0-,1-,2- and 3-cells; (b) exploded 2-boundary cells —notice a drawing error on the back of the model—conversely, the data structures involved are correct, as shown by the two following pictures; (c) solid drawing of the 2-chain [FV[29],FV[30]]; (d) triangulation of the same 2-chain.

```
"test/py/boundary/test04.py" 29a ≡

""" 3D non-convex LAR cells """

from larlib import *

⟨Input of a cellular 3-complex 29b⟩

⟨Visualization of a 2-chain of a 3-complex 29c⟩

⟨Visualization of a 3-chain of a 3-complex 30a⟩

⋄
```

## Input of a cellular 3-complex

Macro referenced in 29a.

```
⟨Input of a cellular 3-complex 29b⟩ ≡
    """ Input of a cellular 3-complex """
    V,[VV,EV,FV,CV] = larCuboids([2,1,1],True)
    struct = Struct([(V,FV,EV),t(.25,.25,0),s(.25,.5,2),(V,FV,EV)])

    V,FV,EV = struct2Marshal(struct)
    CF = AA(sorted)([[20,12,21,5,19,6],[27,1,5,28,13,23],[12,14,25,17,10,4],
    [1,7,17,24,11,18],[30,29,26,16,8,22,10,11,4,18,24,25],[2,3,8,9,0,15]])
    CV = [list(set(CAT([FV[f] for f in faces]))) for faces in CF]

    VV = AA(LIST)(range(len(V)))
    hpc = STRUCT(MKPOLS((V,EV)))
    VIEW(larModelNumbering(1,1,1)(V,[VV,EV,FV,CV],hpc,0.6))
    ⋄
```

#### Visualization of a 2-chain of a 3-complex

```
\langle Visualization of a 2-chain of a 3-complex 29c \rangle \equiv
```

```
""" Visualization of the boundary 2-chain of a 3-complex """
     V,BF,BE = larUnsignedBoundary3(V,CV,FV,EV)(len(CV)*[1])
     VIEW(STRUCT(MKTRIANGLES((V,BF,EV),color=True)))
     VIEW(SKEL_1(STRUCT(MKTRIANGLES((V,BF,EV)))))
     boundaryEdges = chain2BoundaryChain(larUnsignedBoundary2(FV,EV,VV))
     edgeChain = boundaryEdges(29*[0]+[1]+[1])
     VIEW(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,FV[29:31],[EV[e] for e in edgeChain]),color=True)))
     VIEW(SKEL_1(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,FV[29:31],[EV[e] for e in edgeChain])))))
Macro referenced in 29a.
Visualization of a 3-chain of a 3-complex
\langle Visualization of a 3-chain of a 3-complex 30a \rangle \equiv
     """ Visualization of a 3-chain of a 3-complex """
     print "\n***** ECCOMI"
     V,BF,BE = larUnsignedBoundary3(V,CV,FV,EV)([0,0,0,0,1,1])
     VIEW(STRUCT(MKTRIANGLES((V,BF,BE))))
     VIEW(SKEL_1(STRUCT(MKTRIANGLES((V,BF,BE)) )))
Macro referenced in 29a.
"test/py/boundary/test05.py" 30b \equiv
     """ Boundary of a 3-complex """
     from larlib import *
     V, [VV,EV,FV,CV] = larCuboids([1,1,1],True)
     cube = Struct([ (V,FV,EV) ])
     assembly = Struct([cube, Struct([t(0,.5,0), r(PI/4,0,0), s(.5,.5,.5), cube])])
     V,FV,EV = struct2Marshal(assembly)
     VV = AA(LIST)(range(len(V)))
     hpc = STRUCT(MKPOLS((V,EV)))
     VIEW(larModelNumbering(1,1,1)(V,[VV,EV,FV],hpc,0.6))
     CF = [[1,2,3,4,6,7],[0,1,2,3,4,5,6,7,8,9,10,11]]
     CV = [list(set(CAT([FV[f] for f in faces]))) for faces in CF]
     V,BF,BE = larUnsignedBoundary3(V,CV,FV,EV)([0,1])
     VIEW(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,BF,BE),color=True)))
```

```
"test/py/boundary/test06.py" 31a \equiv
     """ Boundary of a 3-complex """
     from larlib import *
     V, [VV,EV,FV,CV] = larCuboids([1,1,1],True)
     cube = Struct([ (V,FV,EV) ])
     hole = Struct([t(0,.5,0), r(PI/4,0,0), s(.5,.5,.5), cube])
     assembly = Struct([ cube, hole, t(0,0,SQRT(0.5)), hole ])
     V,FV,EV = struct2Marshal(assembly) # WRONG: TODO: check ...
     VV = AA(LIST)(range(len(V)))
     hpc = STRUCT(MKPOLS((V,EV)))
     VIEW(larModelNumbering(1,1,1)(V,[[],[],FV],hpc,0.6))
     CF = [[4,5,7,16,17,19,20],[3,8,6,12,11,13],[0,1,10,20],[]]
     CV = [list(set(CAT([FV[f] for f in faces]))) for faces in CF]
     V,BF,BE = larUnsignedBoundary3(V,CV,FV,EV)([0,1,1,0])
     VIEW(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,BF,BE)))) # ERROR in MKTRIANGLES with non-manifold fa
     VIEW(EXPLODE(1.2,1.2,1.2)(MKFACES((V,BF,EV))))
     VIEW(SKEL_1(EXPLODE(1.2,1.2,1.2)(MKFACES((V,BF,EV)))))
"test/py/boundary/test07.py" 31b \equiv
     """ Boundary of a 3-complex """
     from larlib import *
     V,[VV,EV,FV,CV] = larCuboids([1,1,1],True)
     cube = Struct([ (V,FV,EV) ])
     hole = Struct([t(0,.5,0), r(PI/4,0,0), s(1,.5/SQRT(2),.5/SQRT(2)),cube])
     assembly = Struct([ cube, hole ])
     V,FV,EV = struct2Marshal(assembly) # WRONG: TODO: check ...
     VV = AA(LIST)(range(len(V)))
     hpc = STRUCT(MKPOLS((V,EV)))
     VIEW(larModelNumbering(1,1,1)(V,[VV,EV,FV],hpc,0.6))
     CF = [[1,3,6,7,12,11],[0,2,4,5,9,8]]
     CV = [list(set(CAT([FV[f] for f in faces]))) for faces in CF]
     V,BF,BE = larUnsignedBoundary3(V,CV,FV,EV)([1,0])
     VIEW(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,BF,EV),color=True)))
"test/py/boundary/test08.py" 31c \equiv
     """ Boundary of a 3-complex """
```

```
from larlib import *
V,[VV,EV,FV,CV] = larCuboids([1,1,1],True)
cube = Struct([ (V,FV,EV) ])
hole = Struct([t(0,.5,0), r(PI/4,0,0), s(1,.5/SQRT(2),.5/SQRT(2)), cube])
assembly = Struct([ cube, hole ])
assembly2 = Struct([ assembly, t(0,0,.5), s(0.5,1,1), hole ])
V,FV,EV = struct2Marshal(assembly2)
VV = AA(LIST)(range(len(V)))
hpc = STRUCT(MKPOLS((V,EV)))
VIEW(larModelNumbering(1,1,1)(V,[VV,EV,FV],hpc,0.7))
\#CF = [[1,3,6,14,17,18,19,0,4,9,11,15,16,2,5,7,8,10,13],[0,4,9,11,15,16],[2,5,7,8,10,13]]
CF = [[16,9,11,0,15,4],[10,12,7,2,17,15,9,5,18,8,16,3,6,14,19,4,1],[5,13,7,10,8,2]]
CV = [list(set(CAT([FV[f] for f in faces]))) for faces in CF]
n = len(CV)
for k in range(n):
   V,BF,BE = larUnsignedBoundary3(V,CV,FV,EV)(IDNT(n)[k])
   VIEW(STRUCT(MKTRIANGLES((V,BF,BE),color=True)))
   VIEW(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,BF,BE),color=True)))
   VIEW(SKEL_1(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,BF,BE)))))
V,BF,BE = larUnsignedBoundary3(V,CV,FV,EV)([1,1,1])
VIEW(SKEL_1(EXPLODE(1.2,1.2,1.2)(MKTRIANGLES((V,BF,BE)))))
```

#### A Utilities

Marshalling a structure to a LAR cellular model The function struct2Marshal transforms a Struct object, often used to define some assembly of simpler models, to a correctly defined LAR cellular model, i.e. to a cellular partition of the space, in other words a quasi-disjoint partition of the object into well-glued cells of suitable dimensions.

```
⟨Marshalling a structure to a LAR cellular model 32⟩ ≡

""" Marshalling a structure to a LAR cellular model """

import boolean,inters

def struct2Marshal(struct):
    W,FW,EW = struct2lar(struct)
    quadArray = [[W[v] for v in face] for face in FW]
    parts = boolean.boxBuckets3d(boolean.containmentBoxes(quadArray))
    Z,FZ,EZ = boolean.spacePartition(W,FW,EW, parts)
    V,FV,EV = inters.larSimplify((Z,FZ,EZ),radius=0.0001)
```

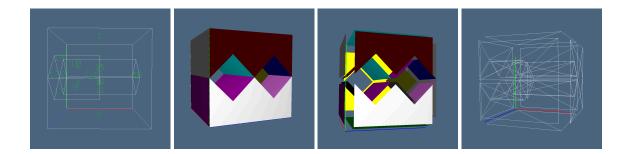


Figure 7: Decomposition of the unit 3-cube in a cellular 3-complex with three 3-cells. Two 3-cells are homeomorphic to the 3-ball, while the remaining one is homeomorphic to the 3-torus. Notice that one of 2-cells (as well one of 3-cells) are non-contractible and non-manifold, as well as non-convex: (a) Indexing of 2-faces of the 3-complex; (b) drawing of the (boundary of) non-convex 3-cell; (c) exploded drawing of the (boundary of) non-convex 3-cell; (d) triangulation of its 2-faces. The triangulation of LAR 2-faces is needed in order to draw them solidly.

```
return V,FV,EV
```

 $\Diamond$ 

### Boundary of a 3-complex

```
\langle Boundary of a 3-complex 33\rangle \equiv
     """ Boundary of a 3-complex """
     import larcc
     """ WHY wrong ???? TOCHECK !!
     def larUnsignedBoundary3(V,CV,FV,EV):
         VV = AA(LIST)(range(len(V)))
         operator3 = larcc.chain2BoundaryChain(boundary3(CV,FV,EV))
         operator2 = larcc.chain2BoundaryChain(larUnsignedBoundary2(FV,EV,VV))
         def larUnsignedBoundary30(chain):
             BF = operator3(chain)
             faceCoords = len(FV)*[0]
             for f in BF: faceCoords[f] = 1
             BE = operator2(faceCoords)
             return V,[FV[f] for f in BF],[EV[e] for e in BE]
         return larUnsignedBoundary30
     def larUnsignedBoundary3(V,CV,FV,EV):
         VV = AA(LIST)(range(len(V)))
         operator3 = larcc.chain2BoundaryChain(boundary3(CV,FV,EV))
```

```
operator2 = larcc.chain2BoundaryChain(larUnsignedBoundary2(FV,EV,VV))
def larUnsignedBoundary30(chain):
    BF = operator3(chain)
    BE = set()
    for f in BF:
        faceCoords = len(FV)*[0]
        faceCoords[f] = 1
        BE = BE.union(operator2(faceCoords))
    return V,[FV[f] for f in BF],[EV[e] for e in BE]
return larUnsignedBoundary30
```

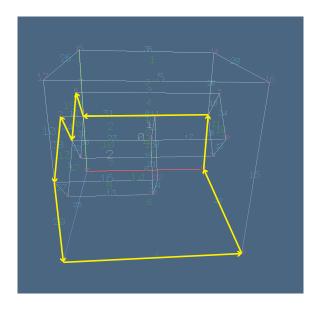


Figure 8: The oriented boundary 1-cycle of a partial 3-cell extraction from a 2-complex in 3D.

# References

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