

$\mathbb{Z}_2$ -Homology of 2-manifolds may be computed  
in  $O(n \log^*(n))$  time

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# Motivation

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- In general computing homology takes  $O(n^3)$  time.

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- In general computing homology takes  $O(n^3)$  time.
- For real problems in computer graphics, computer assisted proofs, electromagnetism where  $n$  could be  $> 10^6$  it is too much.

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- In general computing homology takes  $O(n^3)$  time.
- For real problems in computer graphics, computer assisted proofs, electromagnetism where  $n$  could be  $> 10^6$  it is too much.

To avoid the problem with complexity, two approach can be useful:

- Reduction algorithms - after reduction the input set is small enough to use standard algorithm.

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- For real problems in computer graphics, computer assisted proofs, electromagnetism where  $n$  could be  $> 10^6$  it is too much.

To avoid the problem with complexity, two approach can be useful:

- Reduction algorithms - after reduction the input set is small enough to use standard algorithm.
- For special input classes faster algorithms exist.

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- $R(A)$  - free module over  $R$  generated by  $A$

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- $R(A)$  - free module over  $R$  generated by  $A$
- $S$  be a finite set with a gradation  $S_q$

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- $R(A)$  - free module over  $R$  generated by  $A$
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- $\dim s = q$  iff  $s \in S_q$



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- $R(A)$  - free module over  $R$  generated by  $A$
- $S$  be a finite set with a gradation  $S_q$
- $\dim s = q$  iff  $s \in S_q$
- $\kappa : S \times S \rightarrow R$  s.t.

$$\kappa(s, t) \neq 0 \Rightarrow \dim s = \dim t + 1$$

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$(S, \kappa)$  is an  $S$ -complex if  $(R(S), \partial^\kappa)$  with  $\partial^\kappa : R(S) \rightarrow R(S)$  given by

$$\partial^\kappa(s) := \sum_{t \in S} \kappa(s, t)t$$

is a free chain complex with base  $S$

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We assume that the coding of  $S$  carries the information about  $\kappa$ .

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A *simplicial complex* consists of a collection  $\mathcal{S}$  of simplices. The simplicial complex  $\mathcal{S}$  has a natural gradation  $(\mathcal{S}_q)$ , where  $\mathcal{S}_q$  consists of simplices of dimension  $q$ .

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The simplicial complex  $\mathcal{S}$  has a natural gradation  $(\mathcal{S}_q)$ , where  $\mathcal{S}_q$  consists of simplices of dimension  $q$ .

Assume an ordering of  $\mathcal{S}_0$  is given and every simplex  $\sigma$  in  $\mathcal{S}$  is coded as  $[A_0, A_1, \dots, A_q]$ , where the vertices  $A_0, A_1, \dots, A_q$  are listed according to the prescribed ordering of  $\mathcal{S}_0$ .

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By putting

$$\kappa(\sigma, \tau) := \begin{cases} (-1)^i & \text{if } \sigma = [A_0, A_1, \dots, A_q] \\ & \text{and } \tau = [A_0, A_1, \dots, A_{i-1}, A_{i+1}, \dots, A_q] \\ 0 & \text{otherwise.} \end{cases}$$

we obtain an  $S$ -complex whose chain complex is the classical simplicial chain complex used in simplicial homology.

# Qubical complex

For every elementary cube  $Q$  and number  $j \in \{1, 2, \dots, d\}$  we define the  $j$ th *nondegeneracy number* of  $Q$  by

$$\nu(Q, j) := \begin{cases} \text{card} \{ i < j \mid \text{len } l_i = 1 \} & \text{if } \text{len } l_j = 1. \\ 0 & \text{otherwise.} \end{cases}$$

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A *cubical complex*  $\mathcal{C}$  in  $\mathbb{R}^d$  is a finite collection of elementary cubes in  $\mathbb{R}^d$  and the associated *cubical set* is the union of this collection. The cubical complex  $\mathcal{C}$  has a natural gradation  $(\mathcal{C}_q)_{q \in \mathbb{Z}}$ , where  $\mathcal{C}_q$  consists of elementary cubes of dimension  $q$ .

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$$\kappa(Q, P) := \begin{cases} (-1)^{\nu(Q, j)} & \text{if } Q = l_1 \times \dots \times l_j \times \dots \times l_d \\ & \text{and } P = l_1 \times \dots \times l_j^- \times \dots \times l_d \\ (-1)^{1+\nu(Q, j)} & \text{if } Q = l_1 \times \dots \times l_j \times \dots \times l_d \\ & \text{and } P = l_1 \times \dots \times l_j^+ \times \dots \times l_d \\ 0 & \text{otherwise.} \end{cases}$$

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An induction argument in  $d$  may be used to show that a cubical complex is an  $S$ -complex.

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$$\begin{aligned}\mathrm{bd}_S A &:= \{ t \in S \mid \kappa(s, t) \neq 0 \text{ for some } s \in A. \} \\ \mathrm{cbd}_S A &:= \{ s \in S \mid \kappa(s, t) \neq 0 \text{ for some } t \in A. \}\end{aligned}$$

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## Theorem

*If  $S' \subset S$  is such that for all  $t_1, t_2 \in S'$  and  $s \in S$*

$$s \in \mathrm{bd}_S t_1 \text{ and } t_2 \in \mathrm{bd}_S s \text{ implies } s \in S', \quad (1)$$

*then  $(S', \kappa')$  is an  $S$ -complex.*

- $S' \subset S$  satisfying (1) is *regular*

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*then  $(S', \kappa')$  is an S-complex.*

- $S' \subset S$  satisfying (1) is *regular*
- $S' \subset S$  is *closed* in  $S$  if  $\text{bd}_S S' \subset S'$
- $S' \subset S$  is *open* in  $S$  if  $S \setminus S'$  is closed in  $S$

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## Theorem

*If  $S' \subset S$  is closed in  $S$ , then  $S'$  and  $S \setminus S'$  are regular.*

## Theorem

*If  $S'$  is closed in  $S$ , then*

- (i)  $\partial^{\kappa'} = \partial^{\kappa}|_{R(S')}$
- (ii)  $R(S')$  is a subcomplex of  $R(S)$ .

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## Theorem (M.Mrozek, B.Batko - 2009)

*Assume  $S' \subset S$  is closed in  $S$ . Let  $S'' := S \setminus S'$  and  $\kappa'' := \kappa|_{S'' \times S''}$ . Then the inclusion and the projection*

$$\begin{aligned}\iota : (R(S'), \partial^{\kappa'}) &\rightarrow (R(S), \partial^{\kappa}) \\ \pi : (R(S), \partial^{\kappa}) &\rightarrow (R(S''), \partial^{\kappa''})\end{aligned}$$

*are chain maps. Moreover, we have the following short exact sequence*

$$0 \rightarrow R(S') \xrightarrow{\iota} R(S) \xrightarrow{\pi} R(S'') \rightarrow 0$$

*and the following long exact sequence of homology modules*

$$\dots H_q(R(S')) \xrightarrow{\iota_q} H_q(R(S)) \xrightarrow{\pi_q} H_q(R(S'')) \xrightarrow{\partial_k} H_{q-1}(R(S')) \dots$$

# Relative homology

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## Theorem

*If  $S'$  is closed in  $S$  then*

$$H(R(S'')) \cong H(R(S), R(S')).$$



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$T \subset S$  is a *nullset* of  $S$  if  $T$  is closed or open in  $S$  and  $H(R(T)) = 0$ .

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## Theorem

*If  $T$  is a nullset of  $S$  then  $H(R(S))$  and  $H(R(S \setminus T))$  are isomorphic.*

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## Theorem

*If  $T$  is a nullset of  $S$  then  $H(R(S))$  and  $H(R(S \setminus T))$  are isomorphic.*

- $(a, b)$  is a *reduction pair* if  $\kappa(a, b)$  is invertible
- $a$  is a *free face* of  $b$  in  $S$  if  $(a, b)$  is a reduction pair and  $\text{cbd}_S a = \{b\}$
- $b$  is a *free coface* of  $a$  in  $S$  if  $(a, b)$  is a reduction pair and  $\text{bd}_S b = \{a\}$

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- $b$  is a *free coface* of  $a$  in  $S$  if  $(a, b)$  is a reduction pair and  $\text{bd}_S b = \{a\}$

## Theorem

*Assume  $a, b \in S$ . If  $a$  is a free face of  $b$  or  $b$  is a free face of  $a$  then  $\{a, b\}$  is a nullset.*

# Coreduction algorithm [M.Mrozek, B.Batko - 2009]

```
Q := empty queue;
enqueue(Q,s);
while  $Q \neq \emptyset$  do
    s:=dequeue(Q);
    if  $\text{bd}_S s = \{t\}$  then
        remove(s);
        remove(t);
        foreach  $u \in \text{cbd}_S t$  do
            if  $u \notin Q$  then
                enqueue(Q, u);
            endif;
        else if  $\text{bd}_S s = \emptyset$  then
            foreach  $u \in \text{cbd}_S s$  do
                if  $u \notin Q$  then
                    enqueue(Q, u);
                endif;
            endif;
        endif;
    endwhile ;
```

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            if  $u \notin Q$  then
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            endif;
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                if  $u \notin Q$  then
                    enqueue(Q, u);
                endif;
            endif;
        endif;
    endwhile ;
```

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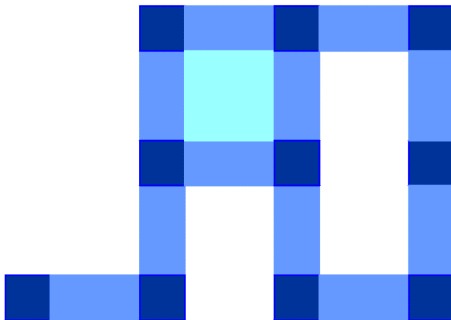
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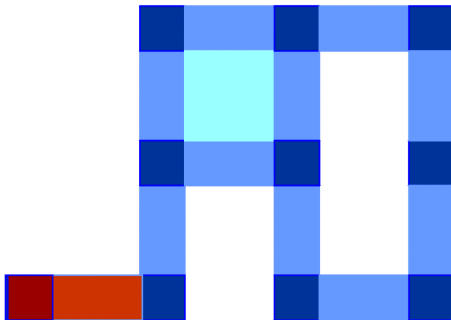
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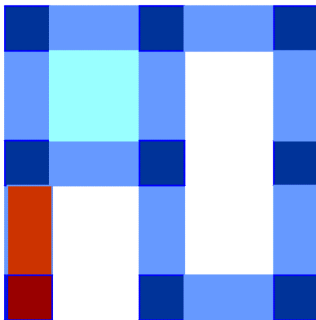
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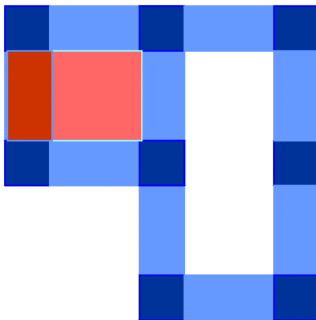
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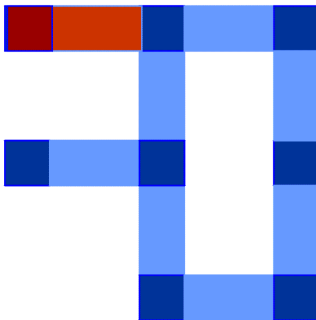
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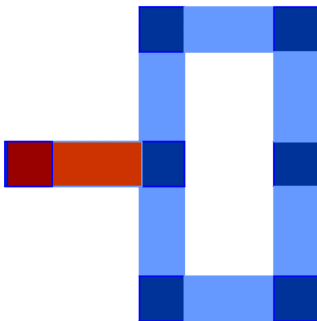
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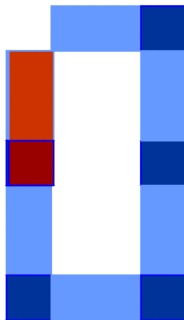
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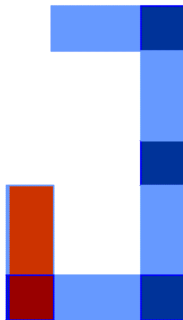
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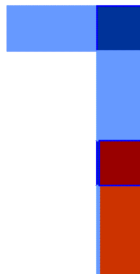
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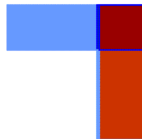
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Coreduction algorithm does not reduce an input to the end:

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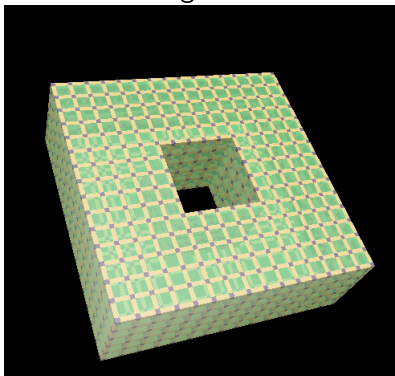
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Coreduction algorithm does not reduce an input to the end:



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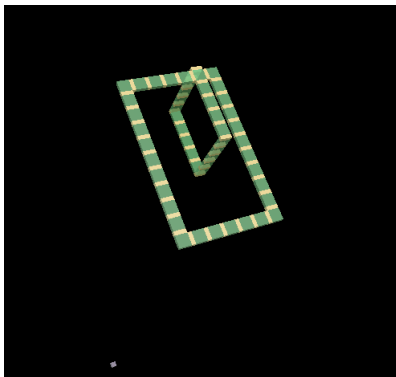
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- Coreduction algorithm is linear.

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- Coreduction algorithm is linear.
- We don't know if coreduction algorithm can reduce input set in each case, we don't have theoretical results.

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Summary

- Coreduction algorithm is linear.
- We don't know if coreduction algorithm can reduce input set in each case, we don't have theoretical results.
- Experiments based on cubical sets show that coreduction homology algorithm performs much better than other algorithms.

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*An S-complex  $X$  is vertexless if  $X_0 = \emptyset$ .*

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## Definition

*An S-complex  $X$  is vertexless if  $X_0 = \emptyset$ .*

## Theorem

*Coreduction algorithm returns a vertexless S-complex.*

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C. Delfinado and H. Edelsbrunner proposed almost linear time algorithm for simplicial complexes in  $\mathbb{R}^3$  embedded in  $\mathbb{S}^3$ .

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C. Delfinado and H. Edelsbrunner proposed almost linear time algorithm for simplicial complexes in  $\mathbb{R}^3$  embedded in  $\mathbb{S}^3$ .

- Complexity is  $O(n \log^* n)$ , where

$$k = \log^* n \Leftrightarrow k = \log(\log(\dots \log n)) - k \text{ times 'log'}$$

# Delfinado, Edelsbrunner algorithm

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- Complexity is  $O(n \log^* n)$ , where

$$k = \log^* n \Leftrightarrow k = \log(\log(\dots \log n)) - k \text{ times 'log'}$$

- Limited to subset of  $\mathbb{S}^3$ .



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C. Delfinado and H. Edelsbrunner proposed almost linear time algorithm for simplicial complexes in  $\mathbb{R}^3$  embedded in  $\mathbb{S}^3$ .

- Complexity is  $O(n \log^* n)$ , where

$$k = \log^* n \Leftrightarrow k = \log(\log(\dots \log n)) - k \text{ times 'log'}$$

- Limited to subset of  $\mathbb{S}^3$ .
- Computes only Betti numbers.

# Main result

A new algorithm for computing homology groups is given in the presentation.

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A new algorithm for computing homology groups is given in the presentation.

- Complexity is also  $O(n \log^* n)$ .

# Main result

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A new algorithm for computing homology groups is given in the presentation.

- Complexity is also  $O(n \log^* n)$ .
- Computes Betti numbers as well as generators.

# Main result

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A new algorithm for computing homology groups is given in the presentation.

- Complexity is also  $O(n \log^* n)$ .
- Computes Betti numbers as well as generators.
- No limit for the input dimension, but requires a special type of the input: 2 dimensional *pseudomanifold*

# Main result

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A new algorithm for computing homology groups is given in the presentation.

- Complexity is also  $O(n \log^* n)$ .
- Computes Betti numbers as well as generators.
- No limit for the input dimension, but requires a special type of the input: 2 dimensional *pseudomanifold*
- Our new algorithm together with coreduction algorithm as an initial step, computes homology groups for “surfaces” (pseudomanifolds), which can be embedded in multi dimensional space.

# Main result

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A new algorithm for computing homology groups is given in the presentation.

- Complexity is also  $O(n \log^* n)$ .
- Computes Betti numbers as well as generators.
- No limit for the input dimension, but requires a special type of the input: 2 dimensional *pseudomanifold*
- Our new algorithm together with coreduction algorithm as an initial step, computes homology groups for “surfaces” (pseudomanifolds), which can be embedded in multi dimensional space.

The algorithm works as an extension for coreduction homology algorithm.

# Pseudomanifold

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## Definition

*An S-complex  $X$  is a 2-pseudomanifold if*

- $X_q = \emptyset$  for  $q > 2$
- for each  $s \in X_1$  the cardinality of  $\text{cbd}_X(s)$  is exactly two
- for all  $a \in X_0$  and for all  $b_1 \neq b_2 \in X_1$  such that  $\kappa(b_i, a) \neq 0$  exists  $c \in X_2$  such that  $\kappa(c, b_i) \neq 0$ .

## Remark



*The last condition is to avoid:*



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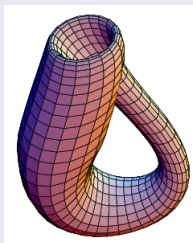
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## Example



*Klein Bottle.*

## Example

*Real projective plane.*

## Example

*Bing's House is not a 2-pseudomanifold.*

# 2-pseudomanifold example

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This examples are not in  $\mathbb{S}^3$ , Delfinado-Edelsbrunner algorithm can't be used.

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## Definition

*Two elements  $a, b$  of an  $S$ -complex  $X$  are adjacent if  $\kappa(a, b) \neq 0$  or  $\kappa(b, a) \neq 0$ .*

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## Definition

*Two elements  $a, b$  of an  $S$ -complex  $X$  are adjacent if  $\kappa(a, b) \neq 0$  or  $\kappa(b, a) \neq 0$ .*

Adjacent relation is symmetric on  $X$ . Its reflexive and transitive closure is an equivalence relation.

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## Definition

*Two elements  $a, b$  of an  $S$ -complex  $X$  are adjacent if  $\kappa(a, b) \neq 0$  or  $\kappa(b, a) \neq 0$ .*

Adjacent relation is symmetric on  $X$ . Its reflexive and transitive closure is an equivalence relation.

## Definition

*The equivalence classes of above relation will be referred to as connected components.*

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## Definition

*Two elements  $a, b$  of an S-complex  $X$  are adjacent if  $\kappa(a, b) \neq 0$  or  $\kappa(b, a) \neq 0$ .*

Adjacent relation is symmetric on  $X$ . Its reflexive and transitive closure is an equivalence relation.

## Definition

*The equivalence classes of above relation will be referred to as connected components.*

## Definition

*An S-complex is connected if it has exactly one connected component.*

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- $\mathcal{C}(X)$  the collection of connected components of  $X$ .

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### Summary

- $\mathcal{C}(X)$  the collection of connected components of  $X$ .
- $\mathcal{C}_p(X) := \{A \cap X_p \mid A \in \mathcal{C}(X)\}$



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## Lemma

*A connected component of an  $S$ -complex  $X$  is an  $S$ -complex.*

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## Lemma

*A connected component of an S-complex  $X$  is an S-complex.*

## Lemma

*Let  $X$  be an S-complex with connected components*

$$X^1, X^2, \dots, X^n$$

*Then*

$$H(X, \mathbb{Z}_2) = \bigoplus_{i=1}^n H(X^i, \mathbb{Z}_2)$$

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*If  $X$  is a connected 2-pseudomanifold then*

$$H_2(X, \mathbb{Z}_2) = [X_2]$$

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## Lemma

*If  $X$  is a connected 2-pseudomanifold then*

$$H_2(X, \mathbb{Z}_2) = [X_2]$$

## Corollary

*If  $X$  is a 2-pseudomanifold then*

$$H_2(X, \mathbb{Z}_2) = \bigoplus_{A \in \mathcal{C}_2(X)} [A].$$

# Main property

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## Theorem

*Let  $X$  be a vertexless 2-pseudomanifold and let  $a \in X_1$ .  
Consider the map*

$$\partial : H_2(X \setminus a, \mathbb{Z}_2) \ni [z] \rightarrow [\partial z] \rightarrow H_1(a, \mathbb{Z}_2).$$

*If  $\partial = 0$  then*

$$H_2(X, \mathbb{Z}_2) \cong H_2(X \setminus a, \mathbb{Z}_2) \quad (2)$$

$$H_1(X, \mathbb{Z}_2) \cong \mathbb{Z}_2 \oplus H_1(X \setminus a, \mathbb{Z}_2) \quad (3)$$

*If  $\text{im } \partial \cong \mathbb{Z}_2$  then*

$$H_2(X, \mathbb{Z}_2) \cong \ker \partial \quad (4)$$

$$H_1(X, \mathbb{Z}_2) \cong H_1(X \setminus a, \mathbb{Z}_2) \quad (5)$$

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Summary

The theorem leads to the following iterative algorithm for computing homology groups of 2-pseudomanifolds.

The Find-Union structure used in the algorithm is a standard data structure with add, find and union operations (see Cormen, Leiserson and Rivest “Introduction to algorithms”).

# Algorithm, [MJ, M.Mrozek - 2009]

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## Algorithm

```
function GetZ2Generators(S-complex  $X$ )
begin
    coreduction( $X$ );
     $S :=$  empty Find-Union structure;
    foreach  $b$  in  $X_2$  do  $S.Add(b)$ ;
    foreach  $a$  in  $X_1$  do begin
         $(b_1, b_2) := \text{cbd}(a)$ ;
        if  $S.Find(b_1) = S.Find(b_2)$  then
             $S.Add(a)$ ;
        else
             $S.Union(b_1, b_2)$ ;
        end;
    return  $S.Sets$ ;
end;
```

# Algorithm correctness

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**Theorem (MJ, M.Mrozek - 2009)**

*The algorithm called with a 2-pseudomanifold  $X$  returns a collection of sets  $\mathcal{S}$  such that*

$$H(X, \mathbb{Z}_2) = \bigoplus_{S \in \mathcal{S}} \sum_{s \in S} [s].$$



# Algorithm complexity

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**Theorem (MJ, M.Mrozek - 2009)**

*The algorithm runs in  $O(n \log^*(n))$  time, where  $n$  denotes the cardinality of the  $S$ -complex on input.*

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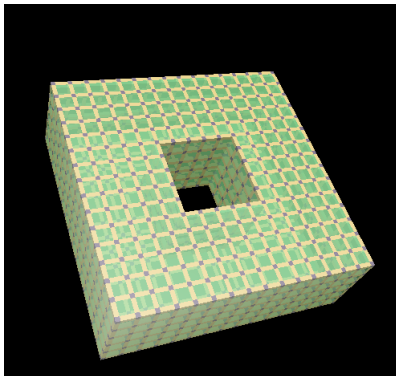
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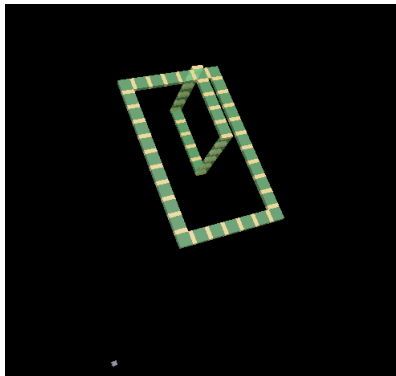
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Vertexless reduced torus:

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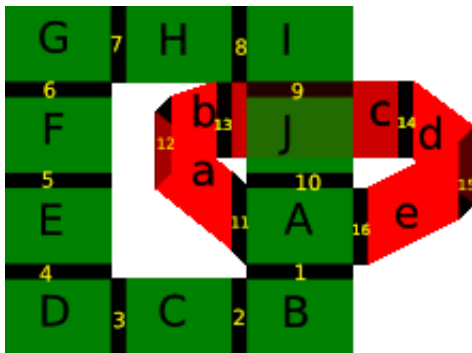
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$$X_0 = \emptyset$$

$$X_1 = \{1, 2, \dots, 16\}$$

$$X_2 = \{A, B, \dots, J, \\ a, b, \dots, e\}$$

# Initial step

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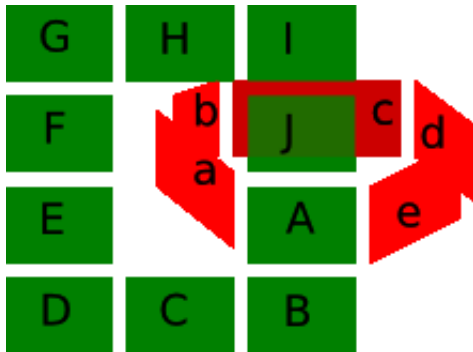
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$$S_1 = \emptyset$$

$$S_2 = \{\{A\}, \{B\}, \dots, \{J\}, \{a\}, \{b\}, \dots, \{e\}\}$$

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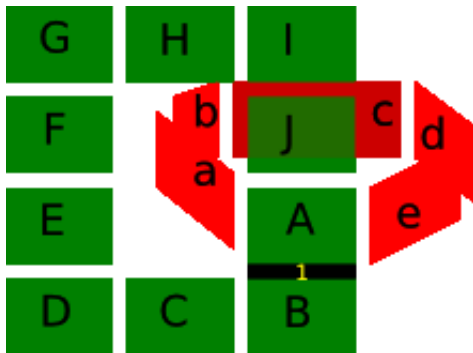
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$$S_1 = \emptyset$$

$$S_2 = \{\{A, B\}, \dots, \{J\} \\ \{a\}, \{b\}, \dots, \{e\}\}$$

# Considered element 10

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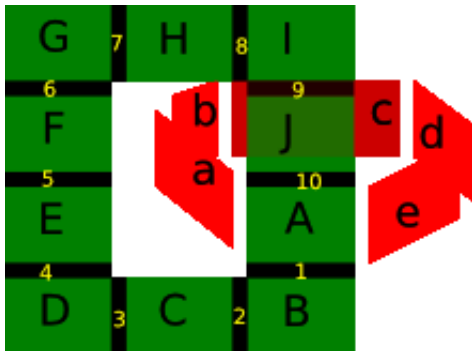
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$$S_1 = \{\{10\}\}$$

$$S_2 = \{\{A, B, \dots, J\} \\ \{a\}, \{b\}, \dots, \{e\}\}$$

# Considered element 16

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### Pseudomanifold

#### Main result

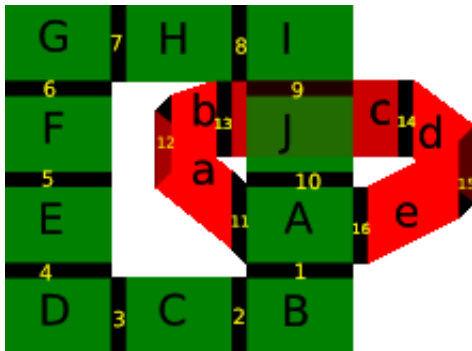
#### Definitions

#### S-complex properties

#### Algorithm

#### Limitation

### Summary



$$S_1 = \{\{10\}, \{16\}\}$$

$$S_2 = \{\{A, B, \dots, J, a, b, \dots, e\}\}$$



# Pseudomanifold limitation

It is obvious from coreduction algorithm we cannot get pseudomanifold for each input data:

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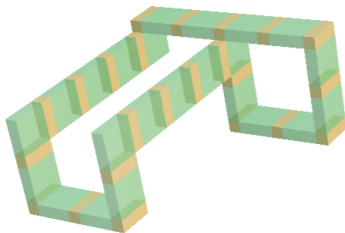
Algorithm

**Limitation**

Summary

It is obvious from coreduction algorithm we cannot get pseudomanifold for each input data:

Reduced Bing's house:



# Extended coreduction

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In coreduction algorithm we use empty set as a boundary for vertices.

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In coreduction algorithm we use empty set as a boundary for vertices. It gives a possibility to start coreductions, because we could have first pair for coreduction.

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In coreduction algorithm we use empty set as a boundary for vertices. It gives a possibility to start coreductions, because we could have first pair for coreduction. When coreduction algorithm ends, the input  $S$ -complex doesn't have any vertices.

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In coreduction algorithm we use empty set as a boundary for vertices. It gives a possibility to start coreductions, because we could have first pair for coreduction. When coreduction algorithm ends, the input  $S$ -complex doesn't have any vertices. We could remove an edge and start the algorithm again.

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Summary

In coreduction algorithm we use empty set as a boundary for vertices. It gives a possibility to start coreductions, because we could have first pair for coreduction. When coreduction algorithm ends, the input  $S$ -complex doesn't have any vertices. We could remove an edge and start the algorithm again. Work on this is in progress. The main problem is in time complexity for getting generators.

# References

## Homology of 2-manifolds

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