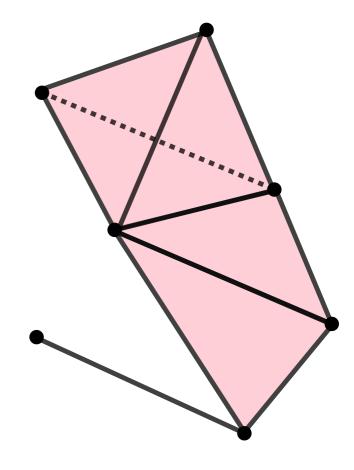
Def: Let $P = \{p_1, \dots, p_n\}$ be a (finite) set. An abstract simplicial complex K with vertex set P is a set of subsets of P satisfying the two conditions:

- (i) the elements of P belong to K,
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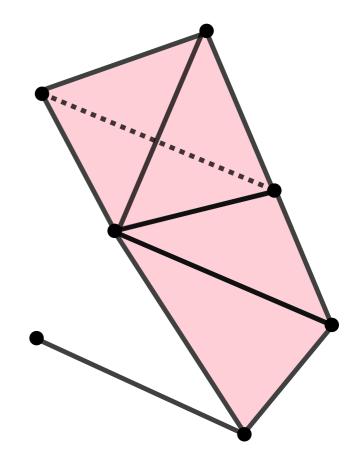
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IMPORTANT

Simplicial complexes can be seen at the same time as geometric/topological spaces (good for topological/geometrical inference) and as combinatorial objects (abstract simplicial complexes, good for computations).

Def: A realization of an abstract simplicial complex K is a geometric simplicial complex K' who is isomorphic to K, i.e., there exists a bijection

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such that $\sigma \in K \iff f(\sigma) \in K'$.

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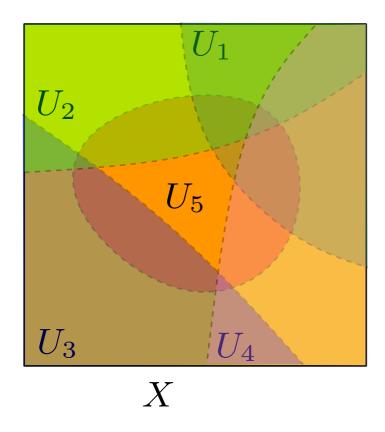
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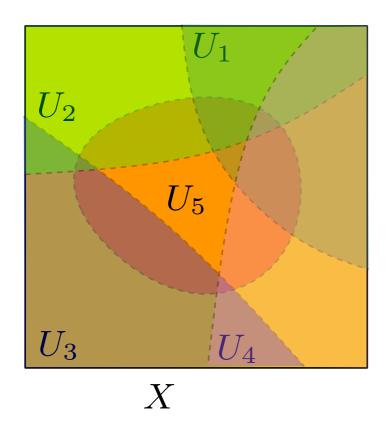
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Abstract simplicial complexes and their realizations are homeomorphic.

Def: An open cover of a topological space X is a collection $\mathcal{U} = (U_i)_{i \in I}$ of open subsets $U_i \subseteq X$, $i \in I$ where I is a set, such that $X \subseteq \bigcup_{i \in I} U_i$.



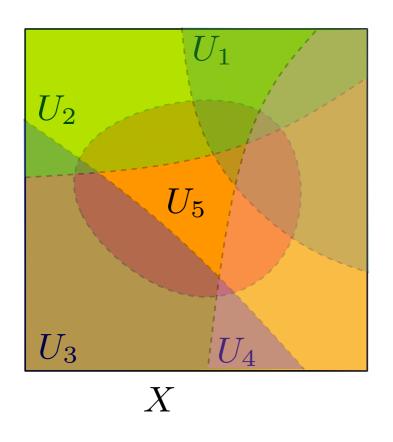
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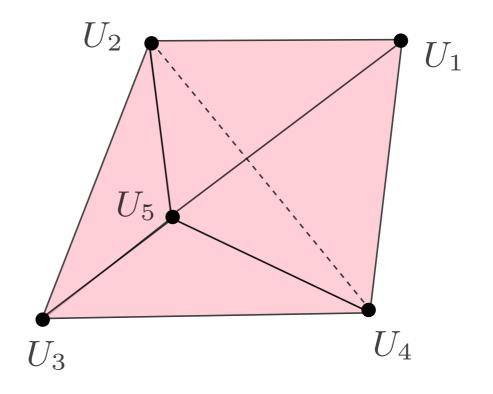


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Def: Given a cover of a topological space X, $\mathcal{U} = (U_i)_{i \in I}$, its nerve is the abstract simplicial complex $C(\mathcal{U})$ whose vertex set is \mathcal{U} and s.t.

$$\sigma = [U_{i_0}, U_{i_1}, \dots, U_{i_k}] \in C(\mathcal{U})$$
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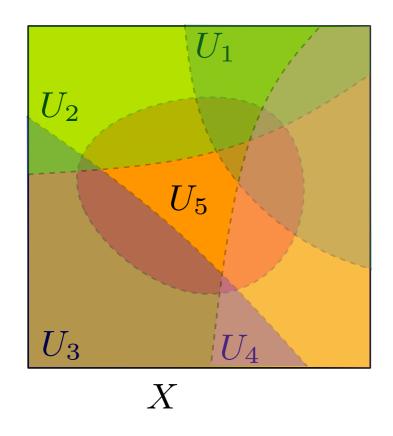


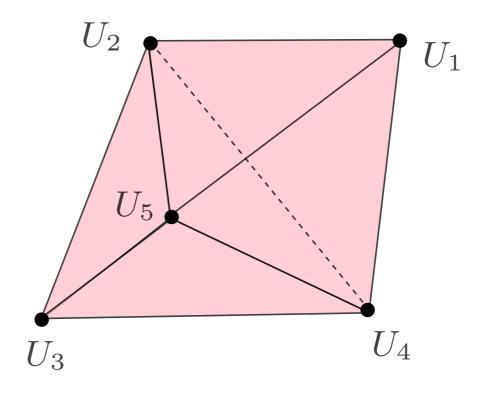
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[On the imbedding of systems of compacta in simplicial complexes, Borsuk, Fund. Math., 1948]





The Nerve Theorem: Let $\mathcal{U} = (U_i)_{i \in I}$ be a finite open cover of a subset X of \mathbb{R}^d such that any intersection of the U_i 's is either empty or contractible. Then X and $C(\mathcal{U})$ are homotopy equivalent.

In particular, every convex set is contractible.

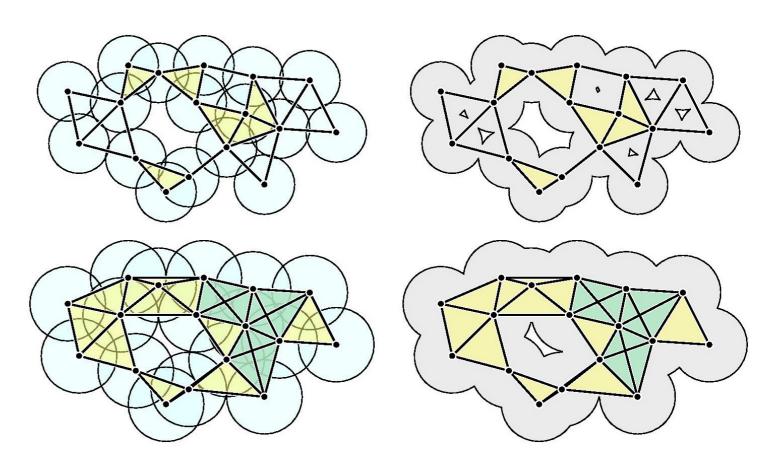
Def: Given a point cloud $P = \{P_1, \dots, P_n\} \subset \mathbb{R}^d$, its Čech complex of radius r > 0 is the abstract simplicial complex C(P, r) s.t. vert(C(P, r)) = P and

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Q: Does the Nerve Theorem apply to Čech complexes?



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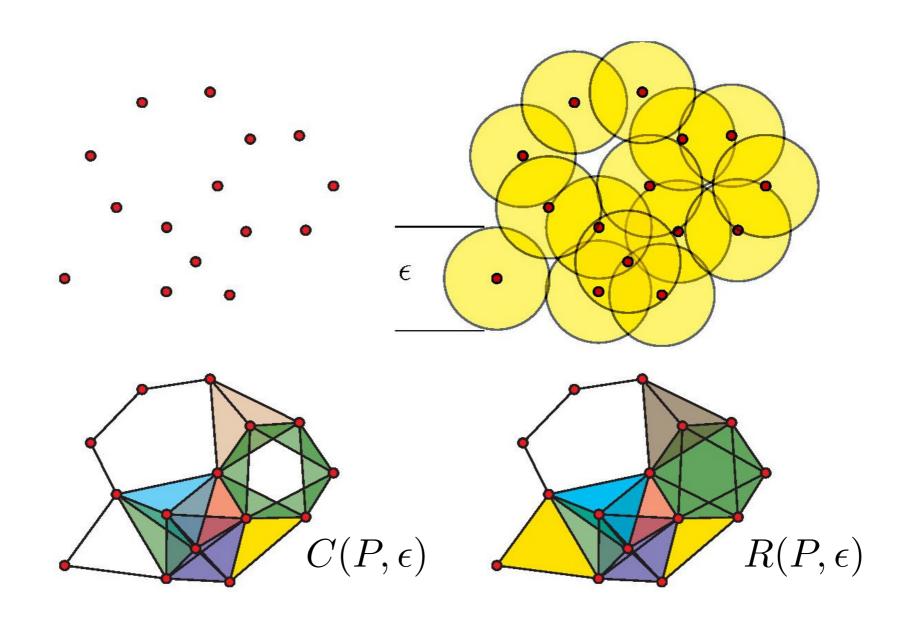
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Prop: $R(P, r/2) \subseteq C(P, r) \subseteq R(P, r)$.

Q: Prove it.

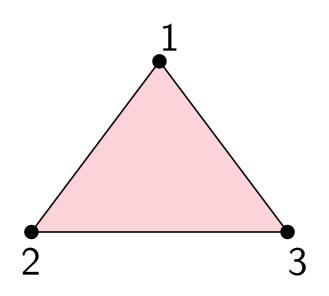
Storing simplicial complexes

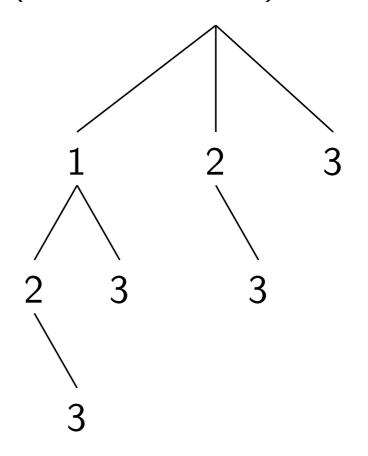
[The Simplex Tree: An Efficient Data Structure for General Simplicial Complexes, Boissonnat, Maria, Algorithmica, 2014]

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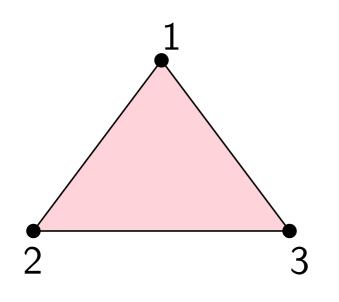
Idea: store sorted simplices in a prefix tree (also called *trie*).



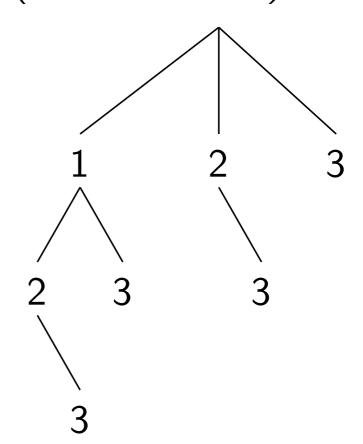


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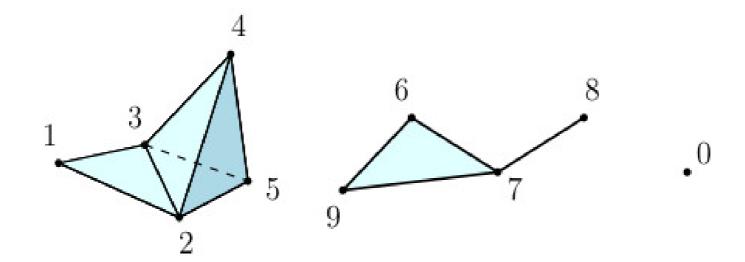
This is called the *simplex tree*.



It allows to store all simplices explicitly without storing all adjacency relations, while maintaining low complexity for basic operations.

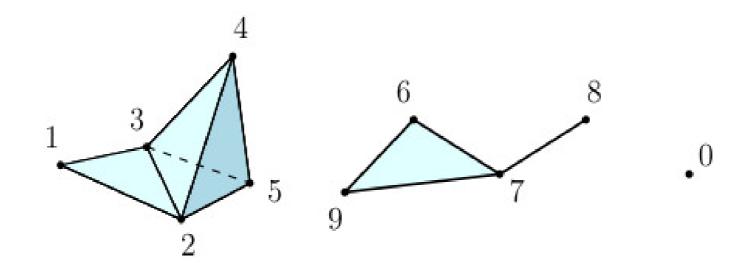
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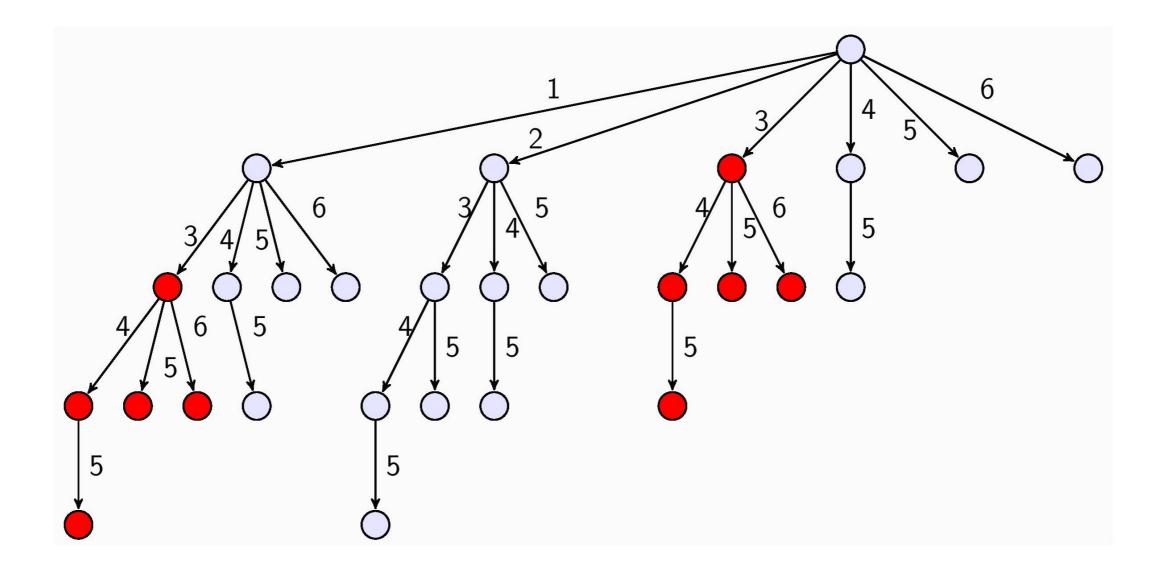


Number of nodes in simplex tree = number of simplices

Depth of simplex tree = 1 + dimension of complex

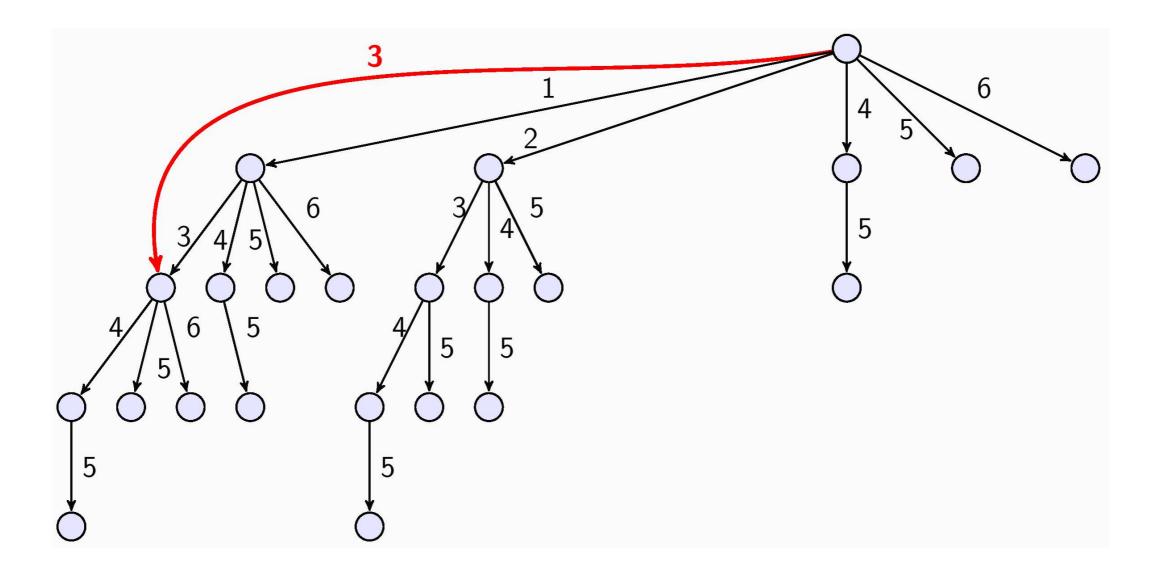
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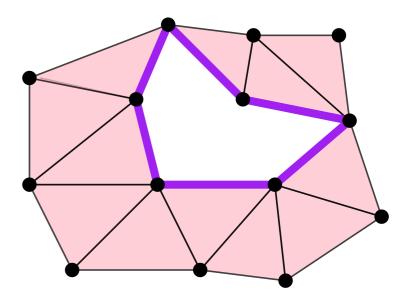
Pb 2: Looking for homotopy equivalences/homeomorphisms/isotopies is extremely difficult. Are there mathematical quantities that are invariant to homotopy equivalences **and** easy to compute?

A: The *holes*, encoded in the *homology groups* H_k , $k \in \mathbb{N}$

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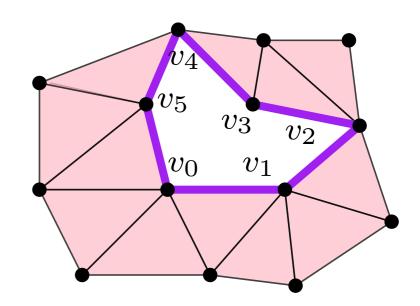
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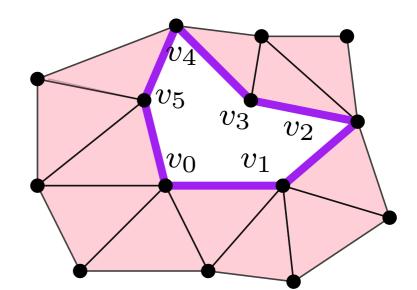


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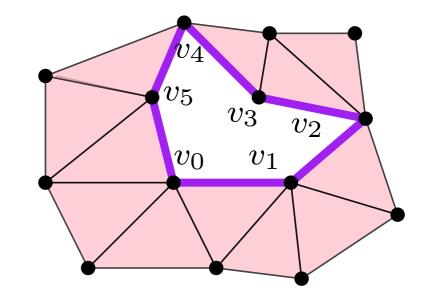


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But what about higher dimensional holes (like the inside of a tetrahedron)?



A: A hole in dimension d is a simplicial complex in which each (d-1)-simplex appears an even number of times.

Def: A d-chain is a formal sum of d-simplices with coefficients in $\mathbb{Z}/2\mathbb{Z}$.

$$C = [v_0, v_1] + [v_1, v_2] + [v_2, v_3] + [v_3, v_4] + [v_4, v_5] + [v_5, v_0].$$

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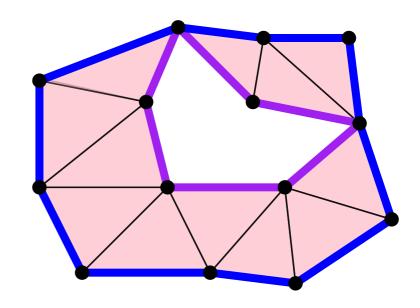
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Pb: Cycles are not holes!!



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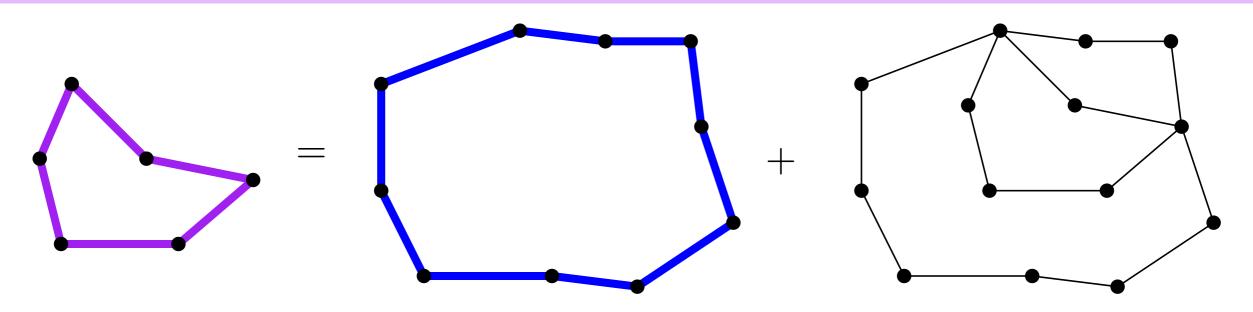
Q: Prove it.

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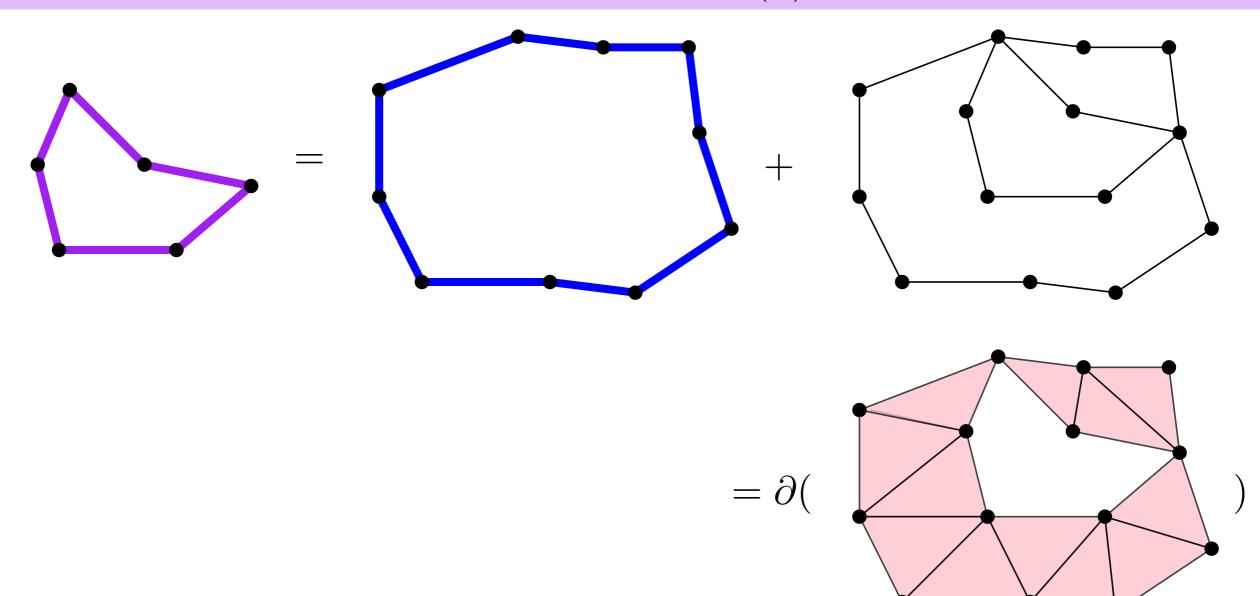
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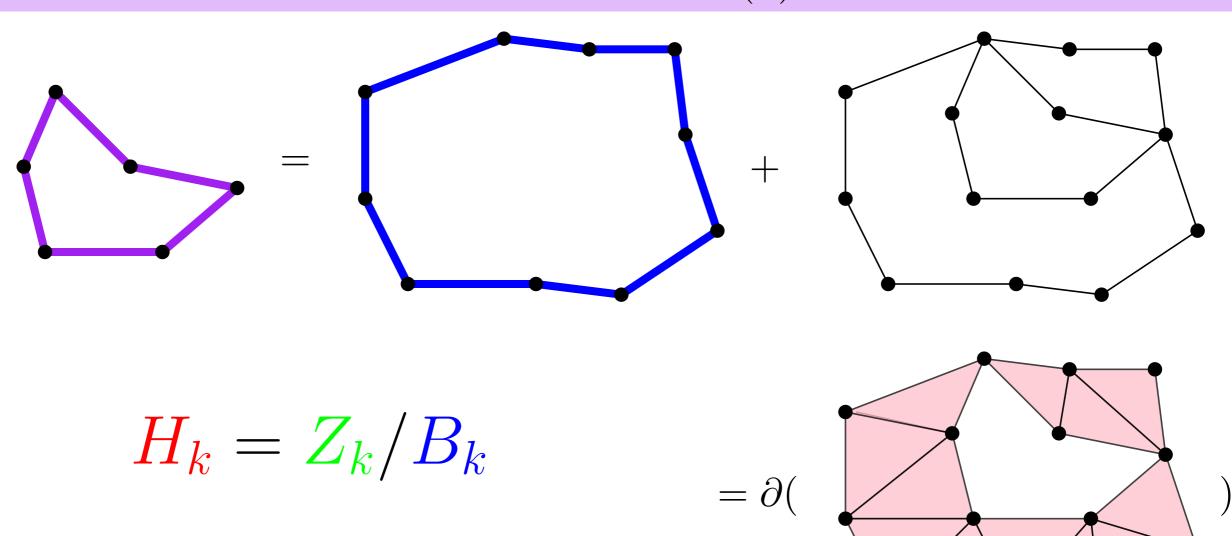
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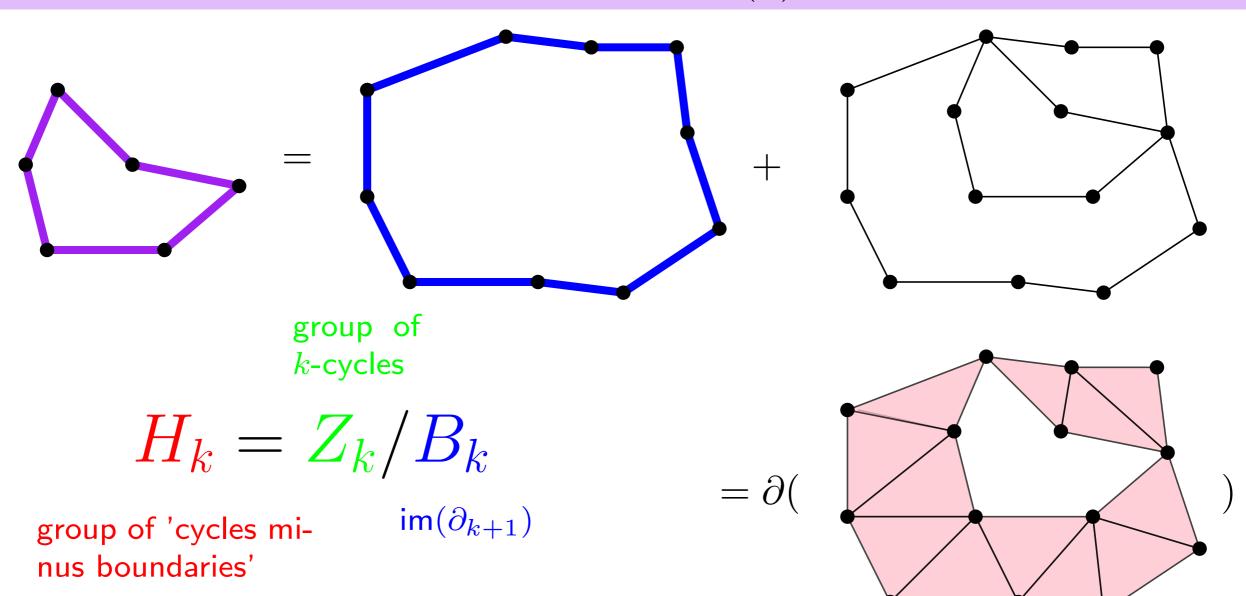
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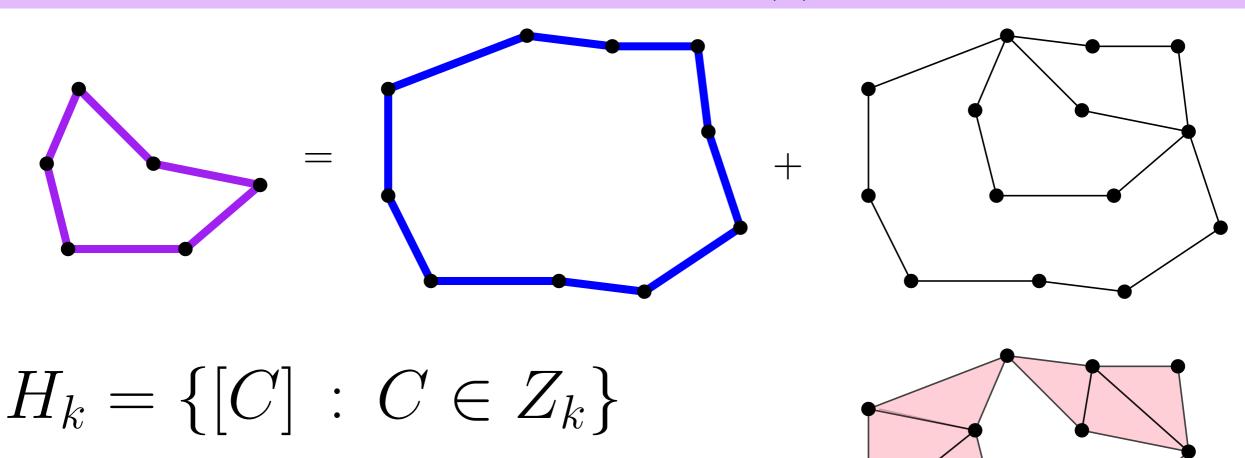


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$$H_k = \{ [C] : C \in Z_k \}$$
 where

$$[C] = \{C' : C \sim C'\}$$

 H_k is a group (vector space) in which each element is an equivalence class of cycles associated to the same hole.

Def: The dimension of H_k is called the *Betti number* β_k .

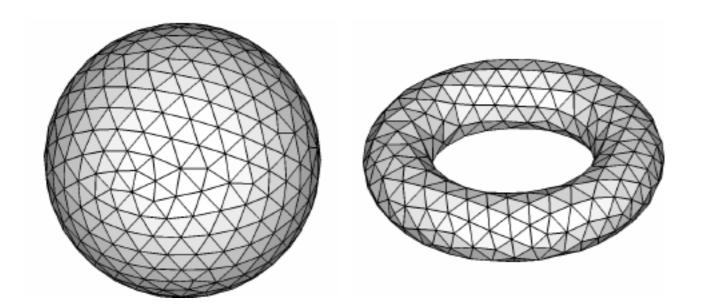
Minimum number of (classes of) cyles needed to create a basis, i.e., to be able to write *any* cycle as a linear combination of cycles in the basis.

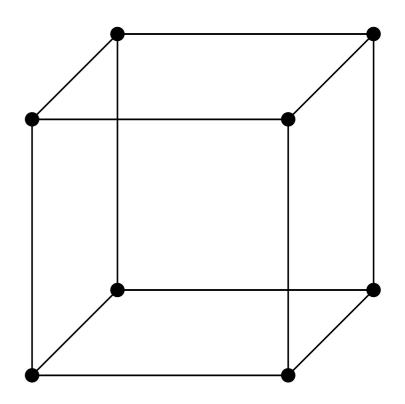
 β_0 counts the connected components, β_1 counts the loops, β_2 counts the cavities, and so on...

 H_k is a group (vector space) in which each element is an equivalence class of cycles associated to the same hole.

Def: The dimension of H_k is called the *Betti number* β_k .

Q: What are the Betti numbers of:

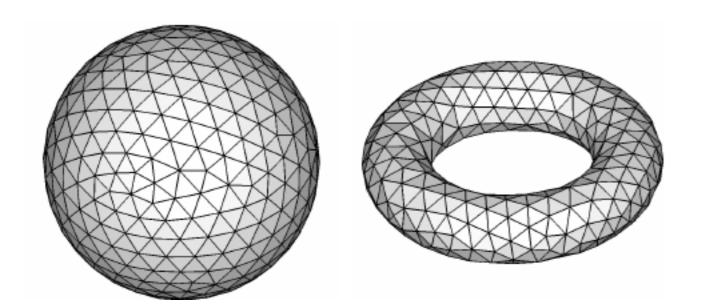


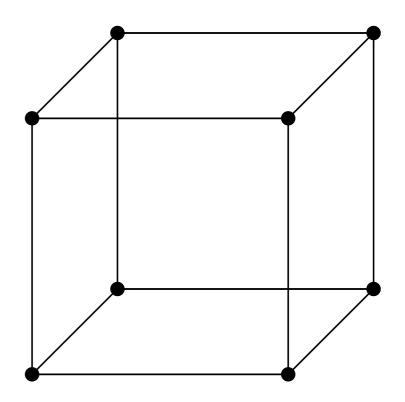


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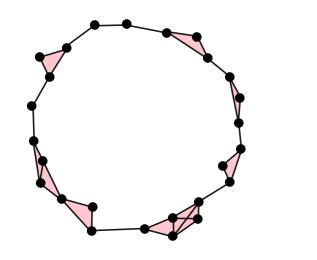


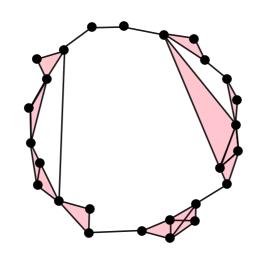
The whole point of homology groups and Betti numbers is that they satisfy:

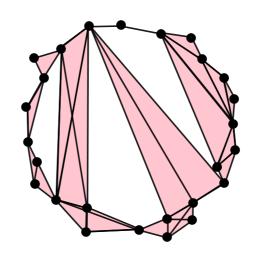
$$H_k(X) \not\sim H_k(Y) \Longrightarrow X \not\sim Y$$

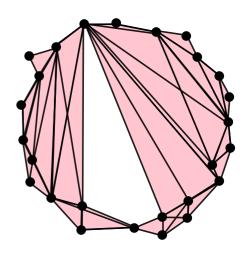
Algorithms to compute the homology groups of a simplicial complex work by decomposing the simplicial complex, with a so-called *filtration*.

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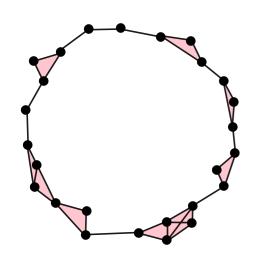


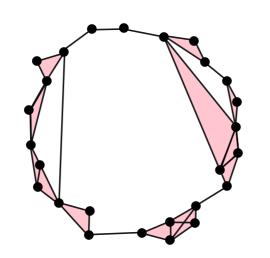


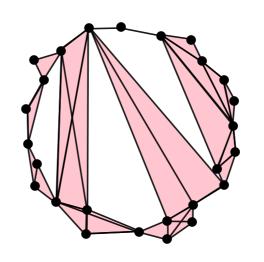


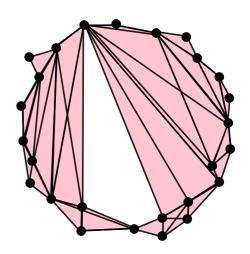
Def: A filtered simplicial complex S is a family $\{S_a\}_{a\in\mathbb{R}}$ of subcomplexes of some fixed simplicial complex S s.t. $S_a\subseteq S_b$ for any $a\leq b$.

Algorithms to compute the homology groups of a simplicial complex work by decomposing the simplicial complex, with a so-called *filtration*.









Def: A filtered simplicial complex S is a family $\{S_a\}_{a\in\mathbb{R}}$ of subcomplexes of some fixed simplicial complex S s.t. $S_a\subseteq S_b$ for any $a\leq b$.

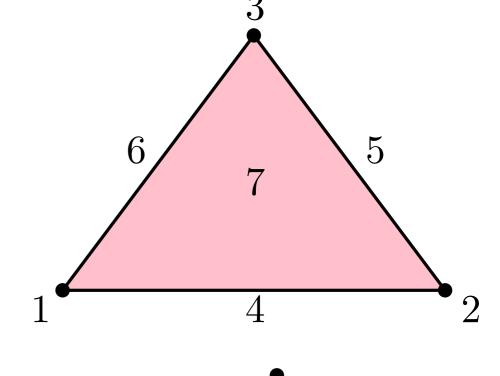
Def: Let f be a real valued function defined on the vertices of K. For $\sigma = [v_0, \ldots, v_k] \in K$, let $f(\sigma) = \max_{i=0,\ldots,k} f(v_i)$, and order the simplices of K in increasing order w.r.t. the function f values (and break ties with dimension in case some simplices have the same function value).

Q: Show that this is a filtration.

Input: simplicial filtration

Homology can be computed by using the fact that each simplex is either:

positive, i.e., it creates a new homology class negative, i.e., it destroys an homology class

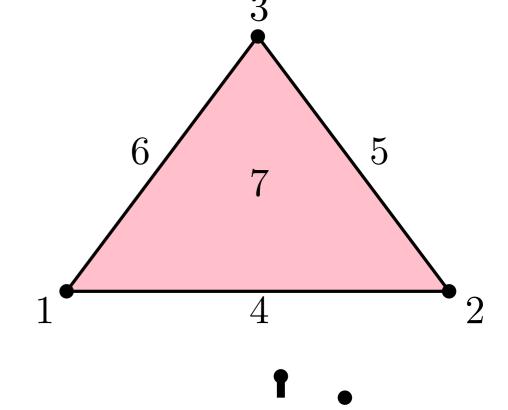


1

Input: simplicial filtration

Homology can be computed by using the fact that each simplex is either:

positive, i.e., it creates a new homology class negative, i.e., it destroys an homology class



1

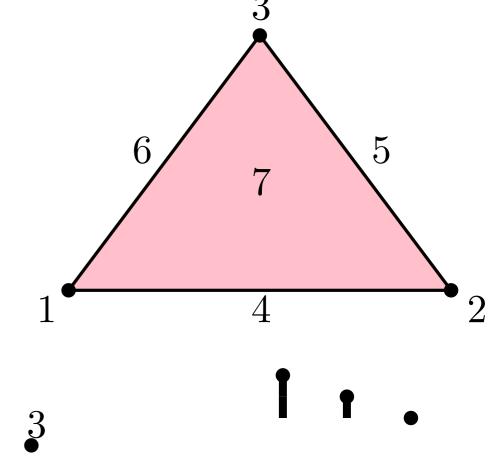
1

2

Input: simplicial filtration

Homology can be computed by using the fact that each simplex is either:

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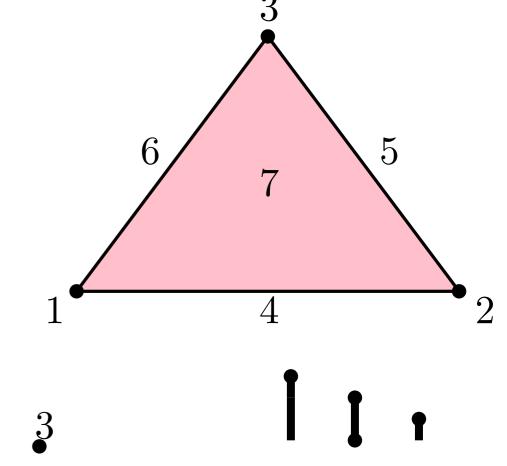


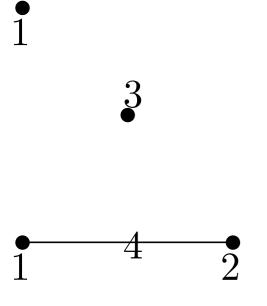
 $\stackrel{ullet}{1}$ $\stackrel{ullet}{1}$ $\stackrel{ullet}{2}$ $\stackrel{ullet}{1}$ $\stackrel{ullet}{2}$

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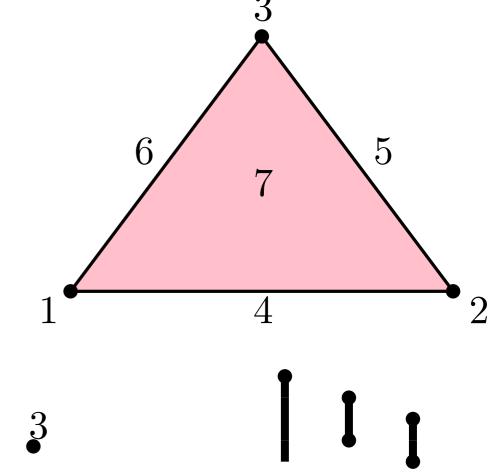


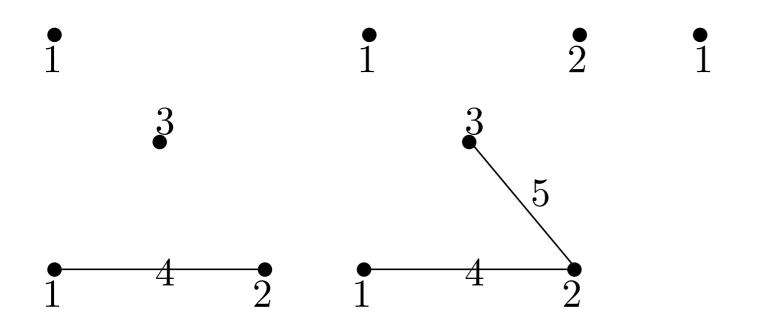


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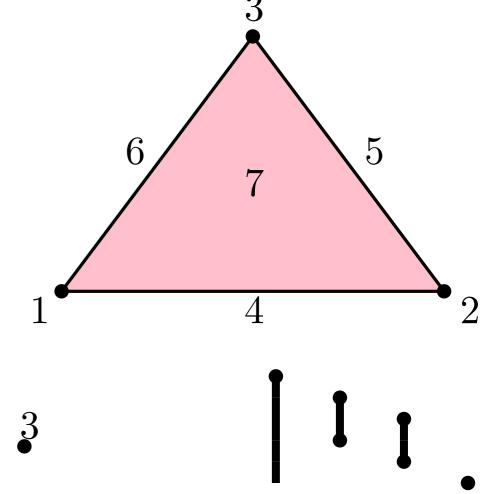


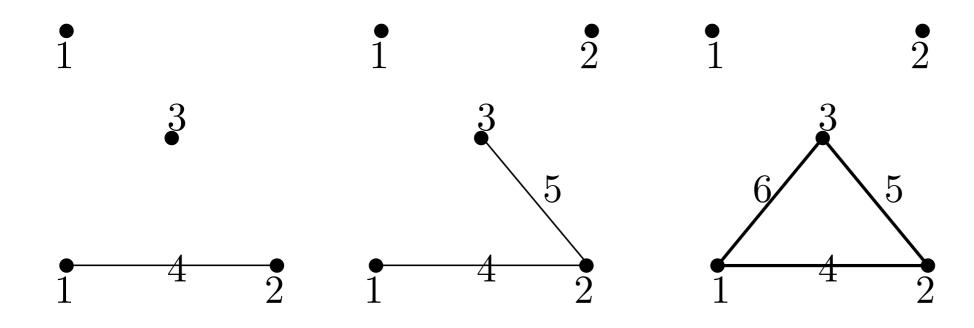


Input: simplicial filtration

Homology can be computed by using the fact that each simplex is either:

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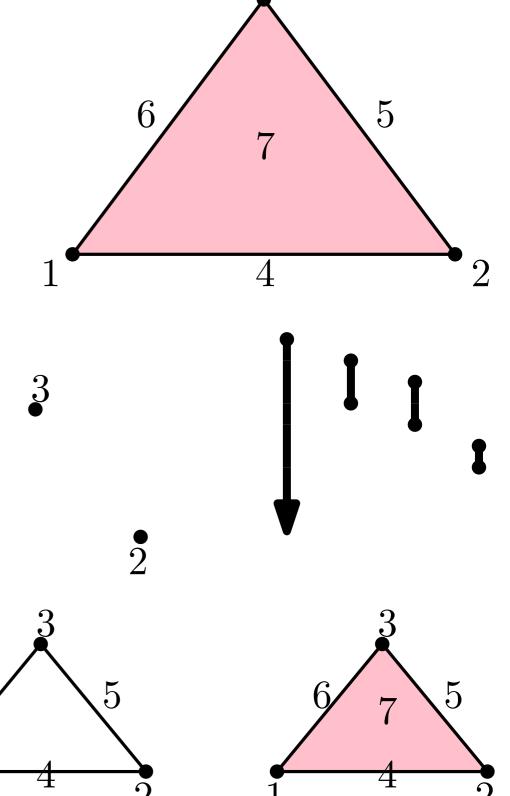


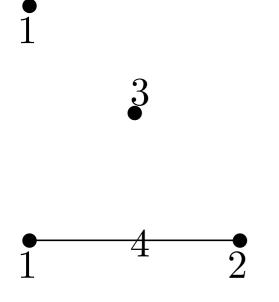


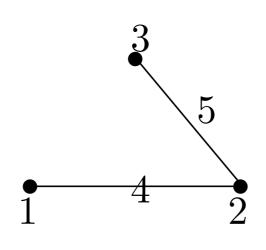
Input: simplicial filtration

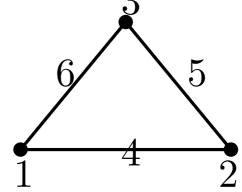
Homology can be computed by using the fact that each simplex is either:

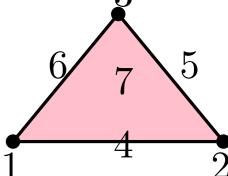
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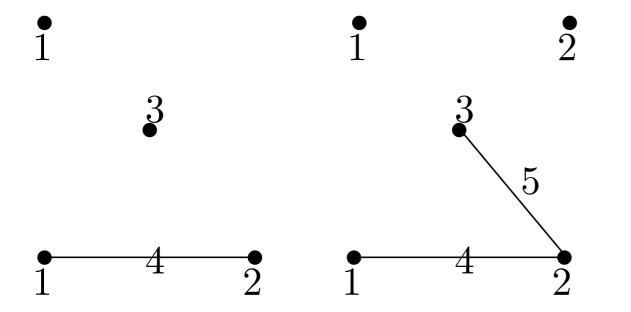


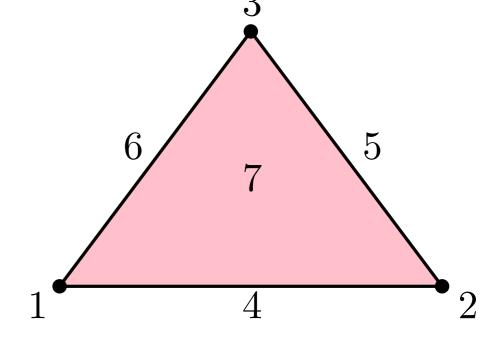
Input: simplicial filtration

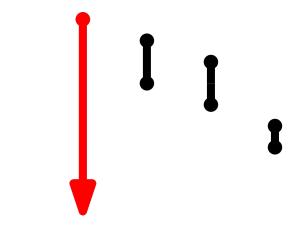
Homology can be computed by using the fact that each simplex is either:

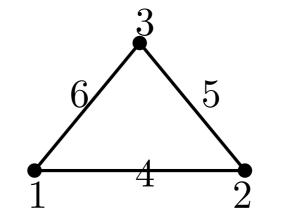
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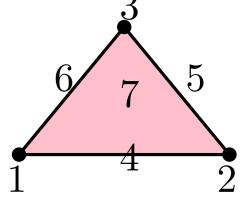
The Betti number is equal to the number of bars that are still alive when the full complex is reached in the filtration







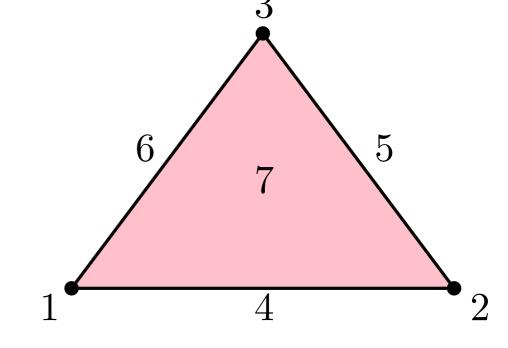




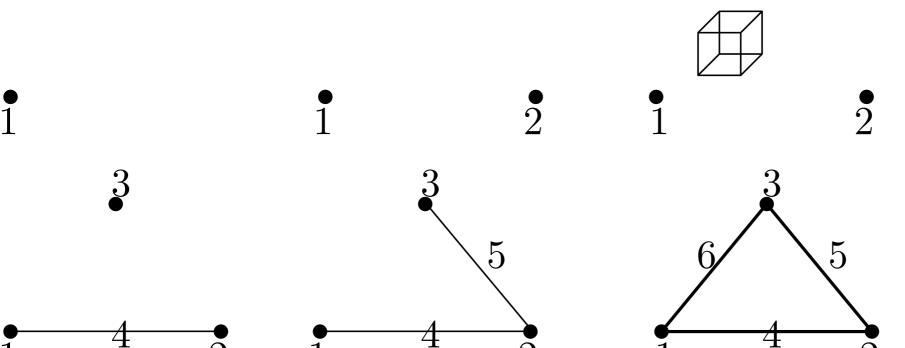
Input: simplicial filtration

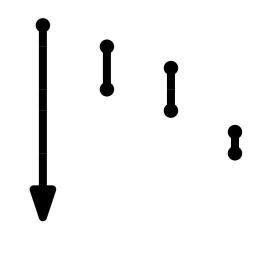
Homology can be computed by using the fact that each simplex is either:

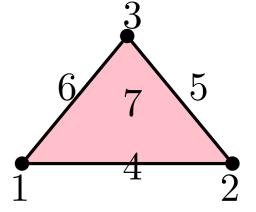
positive, i.e., it creates a new homology class negative, i.e., it destroys an homology class



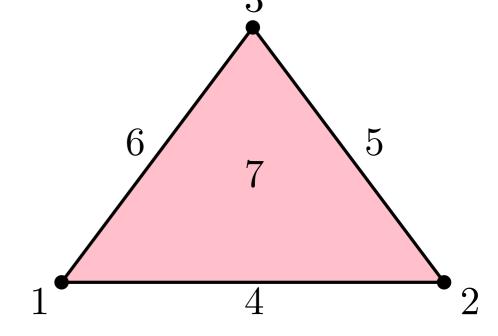
Q: Do the same for the homology of the cube.



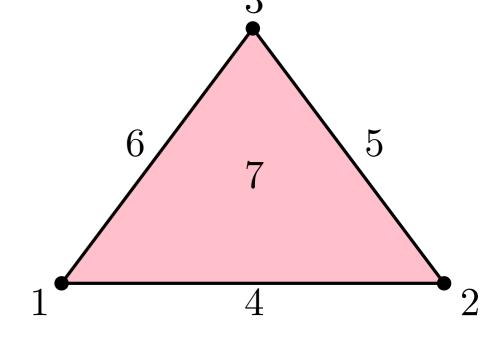




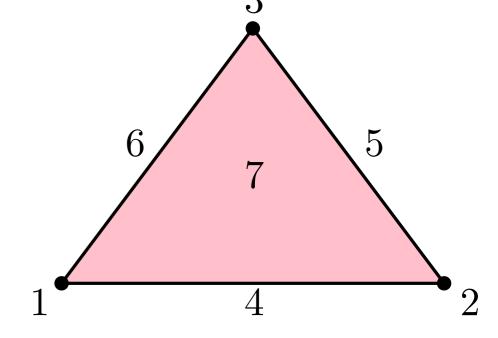
	1	2	3	4	5	6	7
1							
2							
2 3 4 5 6							
4							
5							
6							
7							



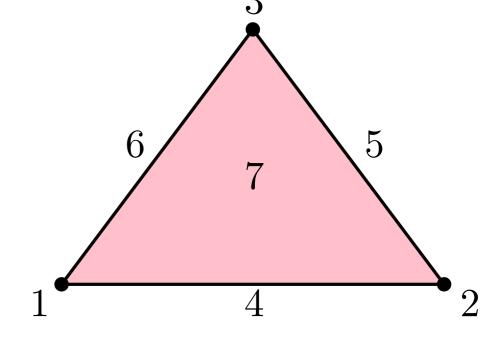
	1	2	3	4	5	6	7
1				•			
2 3				•			
4							
4 5 6							
6							
7							



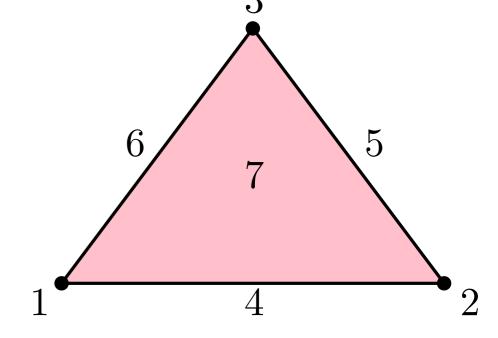
	1	2	3	4	5	6	7
1				•			
2 3 4 5 6				•	•		
3					•		
4							
5							
6							
7							



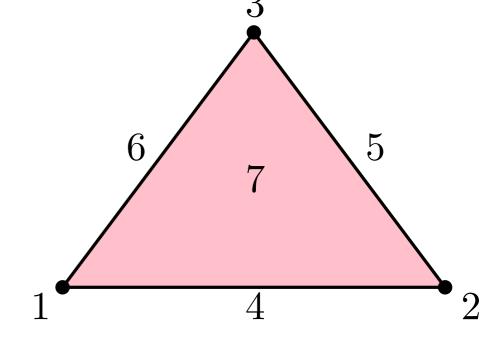
	1	2	3	4	5	6	7
1				•		•	
2				•	•		
3					•	•	
4							
5							
1 2 3 4 5 6							
7							



	1	2	3	4	5	6	7
1				•		•	
2				•	•		
2 3 4 5 6					•	•	
4							•
5							•
6							•
7							



	1	2	3	4	5	6	7
1				•		•	
				•	•		
3					•	•	
4							•
5							•
2 3 4 5 6							•
7							



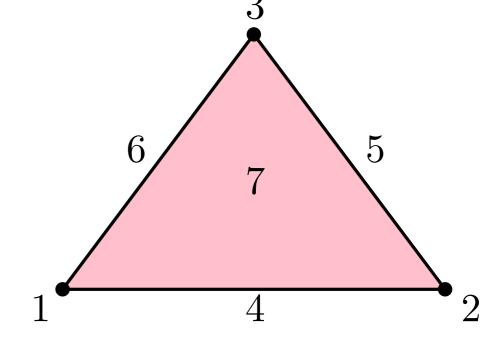
for
$$j=1$$
 to m do:

while
$$\exists k < j \text{ s.t. } low(k) == low(j) \text{ do:}$$

 $col(j) = col(j) + col(k)$

Input: simplicial filtration given as *boundary matrix*

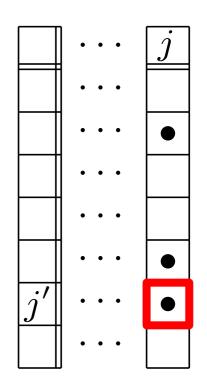
	1	2	3	4	5	6	7
1				•		•	
2 3 4 5 6				•	•		
3					•	•	
4							•
5							•
6							•
7							



for j=1 to m do:

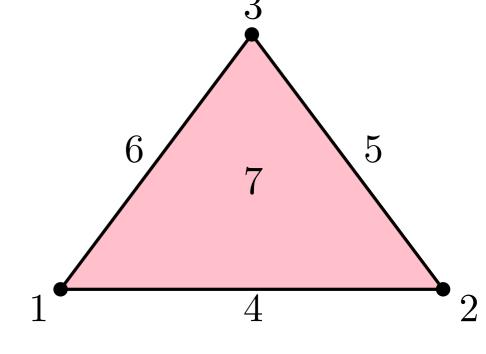
$$col(j) = col(j) + col(k)$$

$$\mathsf{low}(j) = j'$$



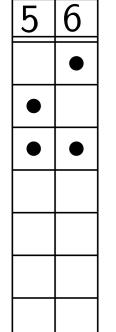
Input: simplicial filtration given as *boundary matrix*

	1	2	3	4	5	6	7
1				•		•	
2				•	•		
3					•	•	
4							•
5							•
6							•
7							



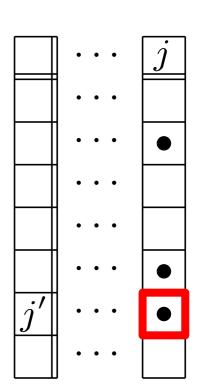
for j=1 to m do:

$$col(j) = col(j) + col(k)$$



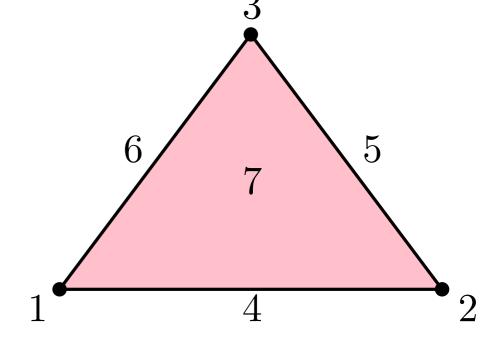
$$6 = 6 + 5$$

$$\mathsf{low}(j) = j'$$



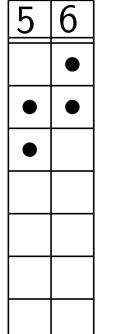
Input: simplicial filtration given as *boundary matrix*

	1	2	3	4	5	6	7
1				•		•	
2				•	•	•	
3					•		
4							•
5							•
4 5 6							•
7							



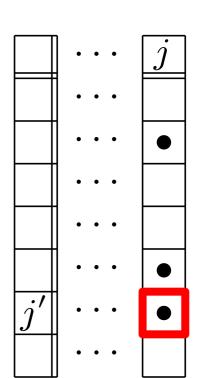
for j=1 to m do:

$$col(j) = col(j) + col(k)$$



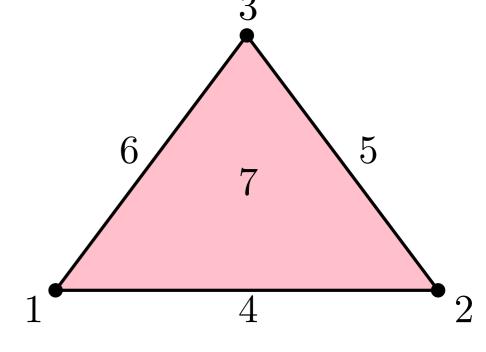
$$6 = 6 + 5$$

$$\mathsf{low}(j) = j'$$



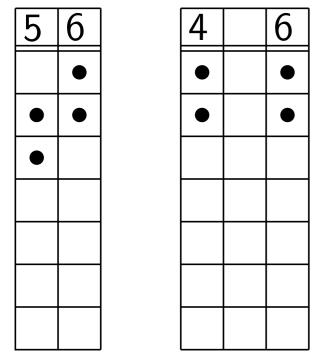
Input: simplicial filtration given as *boundary matrix*

	1	2	3	4	5	6	7
1				•		•	
2				•	•	•	
3					•		
4							•
4 5 6							•
6							•
7							

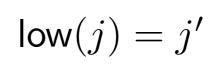


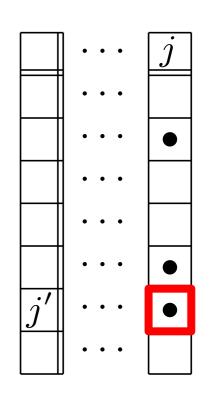
for j=1 to m do:

$$col(j) = col(j) + col(k)$$



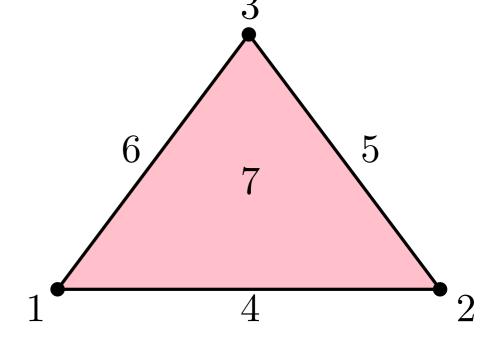
6	=	6+	5
6	=	6+	4





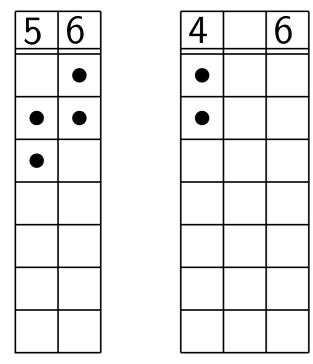
Input: simplicial filtration given as *boundary matrix*

	1	2	3	4	5	6	7
1				•			
2				•	•		
3					•		
4							•
456							•
6							•
7							

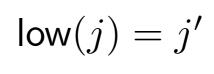


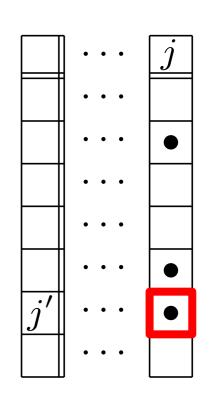
for j=1 to m do:

$$col(j) = col(j) + col(k)$$



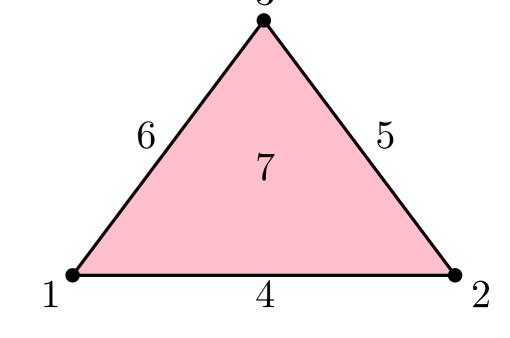
6	=	6+	-5
6	_	64	_1





Input: simplicial filtration

Output: boundary matrix

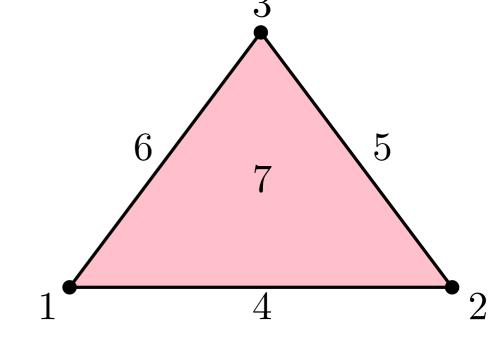


	1	2	3	4	5	6	7
1				*		*	
2				*	*		
3					*	*	
4							*
5							*
6							*
7							

Input: simplicial filtration

Output: boundary matrix

reduced to column-echelon form

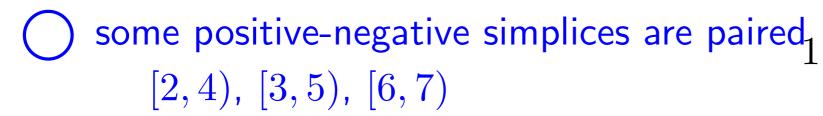


	1	2	3	4	5	6	7
$\boxed{1}$				*		*	
3				*	*		
					*	*	
$\boxed{4}$							*
5							*
6							*
7							

	1	2	3	4	5	6	7
1				*			
3				1	*		
					1		
4							*
5							*
6							1
7							

Input: simplicial filtration

Output: boundary matrix reduced to column-echelon form





	1	2	3	4	5	6	7
1				*		*	
$\begin{bmatrix} 2 \\ 3 \end{bmatrix}$				*	*		
					*	*	
$\boxed{4}$							*
5							*
6							*
7							

	1	2	3	4	5	6	7
1				*			
2				1	*		
3					1		
4							*
5							*
6							1
7							

5

Input: simplicial filtration

Output: boundary matrix

reduced to column-echelon form

Q: Complexity?

Input: simplicial filtration

Output: boundary matrix

reduced to column-echelon form

Q: Complexity?

PLU factorization:

- Gaussian elimination
- fast matrix multiplication (divide-and-conquer)
- random projections?

Input: simplicial filtration

Output: boundary matrix reduced to column-echelon form

Q: Complexity?

PLU factorization:

Gaussian elimination

```
    PLEX / JavaPLEX (http://appliedtopology.github.io/javaplex/)
    Dionysus (http://www.mrzv.org/software/dionysus/)
    Perseus (http://www.sas.upenn.edu/~vnanda/perseus/)
    Gudhi (http://gudhi.gforge.inria.fr/)
    PHAT (https://bitbucket.org/phat-code/phat)
    DIPHA (https://github.com/DIPHA/dipha/)
    CTL (https://github.com/appliedtopology/ctl)
```

Q: Triangulate and compute homology of dunce cap:

