Applied Computational Group Theory?

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Are  $\pi_1 X$  and  $\pi_2 X$  practical tools for computational topology?

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Part I: The Fundamental Group (with P. Dlotko, M. Mrozek)

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Part I: The Fundamental Group (with P. Dlotko, M. Mrozek)

Every protein has a representation as an amino acid chain.



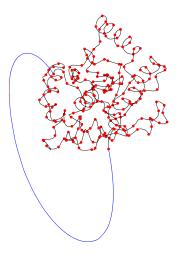
#### Anfinsen's Dogma

This representation determines the 3-D structure of the protein.



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#### Protein Data Base: image of *H. Sapiens 1xd3* data



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Protein ends joined to form an embedding  $K: S^1 \longrightarrow \mathbb{R}^3$ .

## Pure cubical complex representation of *H. Sapiens 1xd3*



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#### GAP system for computatiponal algebra

$$G := \pi_1(\mathbb{R}^3 \setminus K) \cong \langle x, y \mid y^{-1}x^{-1}yxyx^{-1}y^{-3}x^{-1}yxyx^{-1}y^{-1}x \rangle$$

```
gap> K:=ReadPDBfileAsPureCubicalComplex("1XD3.pdb");;
gap> G:=KnotGroup(K);;
#I there are 2 generators and 1 relator of length 14
gap> RelatorsOfFpGroup(G);
[ f2^-1*f1^-1*f2*f1*f2*f1^-1*f2^-3*f1^-1*f2*f1*f2*
f1^-1*f2^-1*f1 ]
```

## What can we do with a group presentation?

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#### What can we do with a group presentation?

EXAMPLE For  $N \lhd G$ ,  $G/N \cong C_5$  and Q = N/[[N, N], N] we could compute  $H_3(BQ, \mathbb{Z}) = (\mathbb{Z}_3)^6 \oplus \mathbb{Z}_{192}$ 

```
gap> N:=LowIndexSubgroupsFpGroup(G,5)[4];;
```

```
gap> Q:=NilpotentQuotient(N,2);;
```

```
gap> GroupHomology(Q,3);
```

```
[ 3, 3, 3, 3, 3, 3, 192 ]
```

$$\mathit{Inv}(\mathit{K}) = \{ H_1(\mathit{N},\mathbb{Z}) \ : \ \mathit{N} \leq \mathit{G} := \pi_1(\mathbb{R}^3 \setminus \mathit{K}), \ |\mathit{G} : \mathit{N}| \leq 5 \}$$

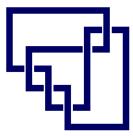
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distinguishes between all prime knots with  $\leq$  10 crossings.

$$Inv(K) = \{ H_1(N,\mathbb{Z}) : N \leq G := \pi_1(\mathbb{R}^3 \setminus K), |G:N| \leq 5 \}$$

distinguishes between all prime knots with  $\leq$  10 crossings.

This invariant shows that the H. Sapiens 1xd3 knot is



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A single thickening of the 1XD3 knot K changes its isotopy type to an embedding  $K' : S^1 \vee S^1 \to \mathbb{R}^3$ 

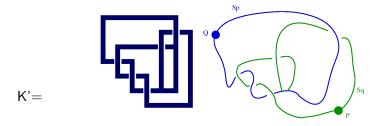
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A single thickening of the 1XD3 knot K changes its isotopy type to an embedding  $K': S^1 \vee S^1 \to \mathbb{R}^3$  and  $\pi_1(\mathbb{R}^3 \setminus K')$  suggests:

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A single thickening of the 1XD3 knot K changes its isotopy type to an embedding  $K' : S^1 \vee S^1 \to \mathbb{R}^3$  and  $\pi_1(\mathbb{R}^3 \setminus K')$  suggests:



A few extra thickenings contribute no further isotopy changes. So perhaps the 1XD3 knot is actually a trefoil.

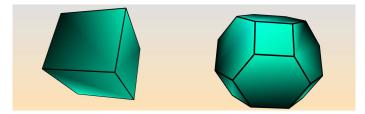


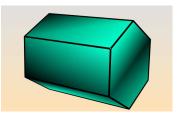
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### A representation of proteins (and other Euclidean data)

Choose a lattice  $L \subseteq \mathbb{R}^n$  and determine

$$D_L = \{x \in \mathbb{R}^n : ||x|| \le ||x - v|| \ \forall v \in L\} .$$

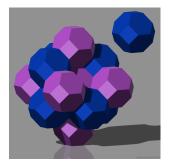




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Any finite set  $\Lambda \subset L$  determines an *L*-complex

$$X = \bigcup_{\lambda \in \Lambda} D_L + \lambda$$



which we represent as a binary array

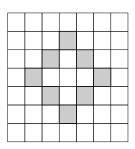
 $(a_{\lambda})_{\lambda\in\Lambda}$ 

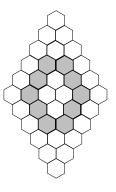
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 $a_{\lambda} = 1$  if  $\lambda \in \Lambda$ ,  $a_{\lambda} = 0$  otherwise.

#### One advantage to permutahedral complexes

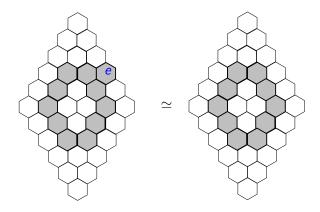
They are always topological manifolds, and so their complements behaves nicely.





#### Second advantage to permutahedral complexes

Permutahedron has at most  $2^{n+1} - 2$  neighbours (compared to  $3^n - 1$  for the cube) so for  $n \le 4$  we cheaply compute retracts



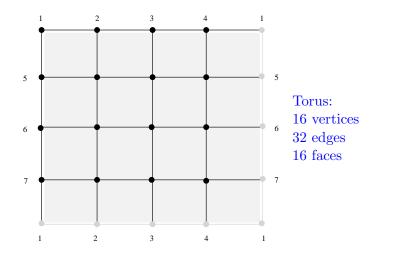
because  $e \in S$  with  $|S| < 2^{2^{n+1}-2}$ .

A zig-zag homotopy retract

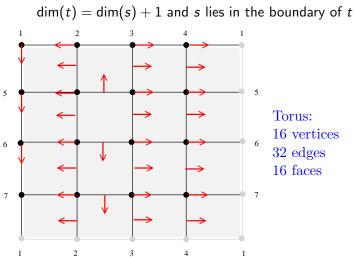
$$X \stackrel{\simeq}{\hookrightarrow} X_1 \stackrel{\simeq}{\leftarrow} X_2 \stackrel{\simeq}{\hookrightarrow} X_3 \cdots \stackrel{\simeq}{\leftarrow} Y$$

```
gap> K:=ReadPDBfileAsPureCubicalComplex("1XD3.pdb");;
gap> X:=ComplementOfPureCubicalComplex(K);;
gap> Size(X);
14692851
gap> Y:=ZigZagContractedPureCubicalComplex(X);;
gap> Size(Y);
74649
```

## Computing fundamental groups of finite regular CW-spaces



A discrete vector field on a regular CW-space X is a collection of arrows  $s \rightarrow t$  where

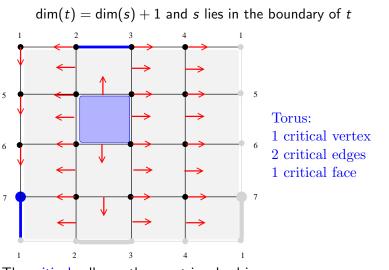


s, t are cells and any cell is involved in at most one arrow

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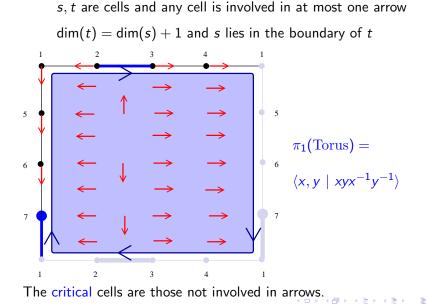


The critical cells are those not involved in arro

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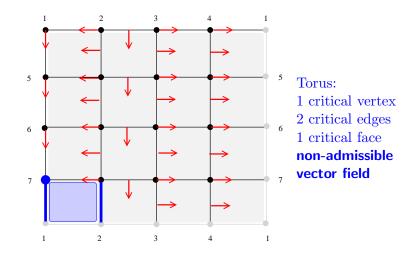
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Algorithm produces a presentation for the fundamental group of a regular CW-space with admissible discrete vector field.



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Computing low-index groups of a finitely presented group G

Index n subgroup  $H \leq G$  corresponds to a homomorphism

$$G \rightarrow S_n$$

into the group of permutations of  $X = \{gH \mid g \in G\}$ .

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Only finitely many such homomorphisms.

Index n subgroups  $H \leq G$  are finitely presented (Reidemeister-Schreier).

#### Multiplication in a nilpotent group G

Use power-commutator presentations

$$\langle x, y, z \mid x^2 = 1, y^2 = z, z^2 = 1, x^{-1}yxy^{-1} = z \rangle$$

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and GAP or Magma's fast rewrite rules for such presentations.

# Computing homology $H_n(BG,\mathbb{Z})$ of a nilpotent group G

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Implement theoretical descriptions of BG for abelian G.

Computing homology  $H_n(BG,\mathbb{Z})$  of a nilpotent group G

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For G of class 2  $[G,G] \to G \to G/[G,G]$ 

construct BG from spaces B([G, G]) and B(G/[G, G]) by homological perturbation techniques involving contracting discrete vector fields on universal covers.

Computing homology  $H_n(BG,\mathbb{Z})$  of a nilpotent group G

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For G of nilpotency class c use recursion on

$$\gamma_c G \to G \to G/\gamma_c G$$
 .

An application

 $H_4(B\mathbb{M}_{24},\mathbb{Z})=0$ 

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gap> GroupHomology(MathieuGroup(24),4);
[]

An application

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[ ]

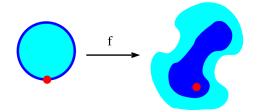
 $H^3(B\mathbb{M}_{24},\mathbb{U}(1))=\mathbb{Z}_{12}$ 

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Part II: The Second Homotopy Group (joint work with Le Van Luyen)

For spaces  $Y \subset X$  and  $D^2 = \{x \in \mathbb{R}^2 : ||x|| \le 1\}$  define

 $\pi_2(X,Y) = \{f: D^2 \rightarrow X : f(S^1) \subset Y\}/homotopy$ 

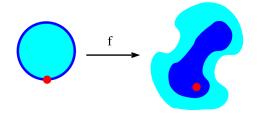


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There is a "restriction" homomorphism

$$\partial \colon \pi_2(X,Y) \to \pi_1(Y)$$

and  $g \in \pi_1(Y)$  acts canonically on  $f \in \pi_2(X, Y)$ .



Theorem (JHC Whitehead): There is an exact sequence of groups

$$\pi_2(Y) \to \pi_2(X) \to \pi_2(X,Y) \stackrel{\partial}{\longrightarrow} \pi_1(Y) \to \pi_1(X)$$

in which  $\partial$  is a *crossed module*:

A crossed module is a group homomorphism  $\partial \colon M \to G$  with action  $(g, m) \mapsto^g m$  statisfying

$$\triangleright \ \partial({}^{g}m) = g \ \partial(m) g^{-1}$$

$$\blacktriangleright \ \partial^m m' = m m' m^{-2}$$

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We define

$$\pi_1(\partial) = G/\mathrm{image}\,\partial \qquad \pi_2(\partial) = \ker\,\partial \;.$$

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Taking  $Y = X^1$  we get Whitehead's functor

Ho(regular CW – spaces)  $\longrightarrow \Sigma^{-1}$ (crossed modules) which is faithful on homotopy types X with  $\pi_n X = 0$  for  $n \neq 1, 2$ .  $\Sigma^{-1}$  is localization with respect to "quasi-isomorphisms"

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Let

$$B(M \stackrel{\partial}{\longrightarrow} G)$$

denote a CW-space with  $\pi_n X = 0$  for  $n \neq 1, 2$  that maps to  $\partial$ .

#### Two algebraic examples of crossed modules

 $\partial: M \to Aut(M), m \mapsto \{x \mapsto mxm^{-1}\}$ for any group M.  $\pi_1(\partial) = Out(M), \pi_2(\partial) = Z(M).$ 

 $\partial \colon M \hookrightarrow G$ 

for any normal subgroup  $M \leq G$ .  $\pi_1(\partial) = G/M$ ,  $\pi_2(\partial) = 0$ .

Computing  $H_n(B(M \xrightarrow{\partial} G), \mathbb{Z})$  in GAP

$$H_5(B(D_{32} \rightarrow Aut(D_{32})), \mathbb{Z}) \cong (\mathbb{Z}_2)^5 \oplus \mathbb{Z}_8$$

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```
gap> M:=DihedralGroup(64);;
gap> C:=AutomorphismGroupAsCatOneGroup(M);;
gap> Size(C); #Size(M) * Size(Aut(M))
32768
gap> Homology(C,5);
[ 2, 2, 2, 2, 2, 8 ]
gap>
```

A morphism of crossed modules is a commutative diagram



with  $\phi_1$ ,  $\phi_2$  group homomorphisms satisfying

$$\phi_2(^g m) = {}^{(\phi_1 g)} \phi_2(m)$$

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It is a quasi-isomorphism if it induces isomorphisms

$$\pi_n(\partial) \cong \pi_n(\partial') , \qquad n=1,2.$$

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,  $n = 1, 2.$ 

Two crossed modules  $\partial$ ,  $\partial''$  are quasi-isomorphic if there exists a sequence of quasi-isomorphisms:

$$\partial \to \partial_1 \leftarrow \partial_2 \to \cdots \leftarrow \partial_k \to \partial''$$

There are 49487365422 different groups (i.e. homotopy 1-types) of order 1024.

Question: Define the order of  $\partial: M \to G$  to be |M||G|. How many quasi-isomorphism types of crossed module of order 16 are there?

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E & Le: By finding explicit quasi-isomorphisms, there are at most 51 quasi-isomorphism types. The invariants

 $\pi_1(\partial), \ \pi_2(\partial), \ H_2(\partial, \mathbb{Z}), \ H_3(\partial, \mathbb{Z})$ 

establish at least 49 quasi-isomorphism types of crossed modules of order 16.

Computing the homology of  $M \stackrel{\partial}{\longrightarrow} G$ 

- 1. The cellular chain complex  $C_*(B(\partial))$  has an algebraic description using the language of simplicial sets.
- 2. By the Homological Perturbation Lemma and discrete vector fields we need only compute a much smaller homotopic chain complex  $C_* \simeq C_*(B(\partial))$ .
- 3. Coreduction can be applied to obtain an even smaller chain complex  $D_* \simeq C_*$ .

# A curiosity about coreduction

The crossed module

$$\partial \colon \mathbb{Z}_2 \to 0$$

yields a homotopy 2-type  $B = B(\partial)$  with

$$\pi_2(B) = \mathbb{Z}_2, \ \pi_k(B) = 0 \text{ for } k \neq 2.$$

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gap> B:=EilenbergMacLaneComplex(CyclicGroup(2),2,11);; gap> C:=ChainComplex(B);; gap> List([0..11],CK!.dimension); [ 1, 1, 2, 4, 8, 16, 32, 64, 128, 256, 512, 1024 ]

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```
gap> D:=CoreducedChainComplex(C);;
gap> List([0..10],D!.dimension);
[ 1, 0, 1, 1, 2, 3, 5, 8, 13, 21, 34 ]
```

# $\mathcal{N}$ : (crossed modules) $\longrightarrow$ (simplicial groups)

Given  $\partial: M \to G$  we consider the category

$$\begin{array}{ll} A = M \ltimes G \\ s \colon A \to A, & (m,g) \mapsto (1,g) \\ t \colon A \to A, & (m,g) \mapsto (1,\partial(m)g) \\ \circ \colon A \times_G A \to A, & ((m,g), (m',g') \mapsto (m, (\partial m)^{-1}g') \end{array}$$

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s, t,  $\circ$  are group homomorphisms and A is a category internal to the category of groups.

The nerve  $\mathcal{N}(A)$  is thus a simplicial group.

# $\begin{array}{c} B: \mbox{ (crossed modules)} & \xrightarrow{\mathcal{N}} \mbox{ (simplicial groups)} \\ & & & & \\ & & & & \\ & & & \\ &$

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$$\begin{array}{c} B: (\text{crossed modules}) \xrightarrow{\mathcal{N}} (\text{simplicial groups}) \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ & & & \\ F: (\text{sets}) \xrightarrow{\Delta} (\text{simplicial sets}) \xrightarrow{\Delta} (\text{simplicial sets}) \end{array}$$

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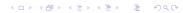
B: (crossed modules)  $\xrightarrow{\mathcal{N}}$  (simplicial groups)  $\mathcal{N}$ (bisimplicial sets)  $\xrightarrow{\Delta}$  (simplicial sets)  $F: (sets) \longrightarrow (free abelian groups)$  $C_*(B(\partial: M \to G))$  is the total complex of the bicomplex:  $\longrightarrow F\mathcal{N}_2\mathcal{N}_2(A) \longrightarrow F\mathcal{N}_2\mathcal{N}_1(A) \longrightarrow F\mathcal{N}_2\mathcal{N}_0(A)$  $\longrightarrow F\mathcal{N}_1\mathcal{N}_2(A) \longrightarrow F\mathcal{N}_1\mathcal{N}_1(A) \longrightarrow F\mathcal{N}_1\mathcal{N}_0(A)$  $\longrightarrow F\mathcal{N}_0\mathcal{N}_2(A) \longrightarrow F\mathcal{N}_0\mathcal{N}_1(A) \longrightarrow F\mathcal{N}_0\mathcal{N}_0(A)$ 



We could replace each column by

$${\it R}_{*}^{\mathcal{N}_{j}({\it A})}\otimes_{\mathbb{Z}\mathcal{N}_{j}({\it A})}\mathbb{Z}$$

where  $R_*^{\mathcal{N}_j(A)}$  is an arbitrary free  $\mathbb{Z}\mathcal{N}_j(A)$ -resolution of  $\mathbb{Z}$ .



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$${\sf R}^{\mathcal{N}_j({\sf A})}_*\otimes_{\mathbb{Z}\mathcal{N}_j({\sf A})}\mathbb{Z}$$

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$$\mathsf{R}^{\mathcal{N}_{j}(\mathcal{A})}_{*}\otimes_{\mathbb{Z}\mathcal{N}_{j}(\mathcal{A})}\mathbb{Z}^{*}$$

where  $R_*^{\mathcal{N}_j(A)}$  is an arbitrary free  $\mathbb{Z}\mathcal{N}_j(A)$ -resolution of  $\mathbb{Z}$ . But the horizontally induced maps won't square to zero if the resolutions aren't functorial.

Homological Perturbation Lemma solves this problem by providing a filtered complex

 $R^{\mathcal{N}_*(A)}_*\otimes_{\mathbb{Z}\mathcal{N}_*(A)}\mathbb{Z}$ 

#### A homotopy equivalence data

$$(L,d) \stackrel{\stackrel{\rho}{\rightarrow}}{\rightarrow} (M,d), h$$
 (\*)

consists of chain complexes L, M, quasi-isomorphisms i, p and a homotopy ip - 1 = dh + hd. A perturbation on (\*) is a homomorphism  $\epsilon \colon M \to M$  of degree -1 such that  $(d + \epsilon)^2 = 0$ .

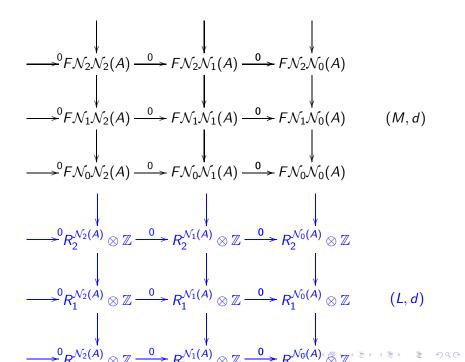
PERTURBATION LEMMA: If  $A = (1 - \epsilon h)^{-1} \epsilon$  exists then

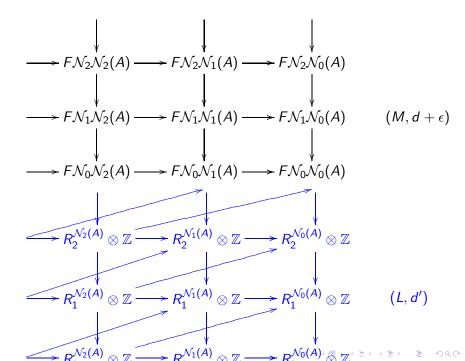
$$(L, d') \xrightarrow{p'}{i'} (M, d + \epsilon), h'$$
 (\*\*)

is a homotopy equivalence data where

$$i'=i+hAi, \ p'=p+pAh, \ h'=h+hAh, \ d'=d+pAi$$
 .

M. Crainic, "On the perturbation lemma, and deformations", 2004





# Persistent homology of crossed modules

$$B = B(\partial \colon M \to G)$$
  
$$\pi_i = \pi_i B$$

$$\cdots \hookrightarrow [[\pi_2, \pi_1], \pi_1] \hookrightarrow [\pi_2, \pi_1] \hookrightarrow \pi_2$$

$$\rightarrow \cdots \pi_1 / [[[\pi_1, \pi_1], \pi_1] \rightarrow \pi_1 / [\pi_1, \pi_1]$$

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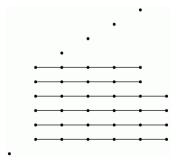
$$\to \dots \pi_1 / [[[\pi_1, \pi_1], \pi_1] \to \pi_1 / [\pi_1, \pi_1]$$
  
induce a sequence of homotopy 2-types

$$\rightarrow B_{-2} \rightarrow B_{-1} \rightarrow B \rightarrow B_1 \rightarrow B_2 \rightarrow \cdots$$

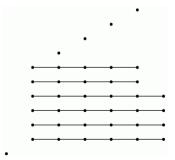
whose degree k homology is a homotopy invariant bar code for B.

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# $H_3(B_*,\mathbb{Z}_2)$ barcode for $B = B(C_{32} \rightarrow Aut(C_{32}))$



# $H_3(B_*,\mathbb{Z}_2)$ barcode for $B = B(C_{32} \rightarrow Aut(C_{32}))$



THANK YOU

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