

PERSISTENT MATCHMAKING

ULRICH BAUER

TUM

COMPUTATIONAL MATHEMATICS  
SEMINAR

JAGIELLONIAN UNIVERSITY

MAR 25, 2021

# RIPSER: COMPUTING VIETORIS-RIPS PERSISTENCE

new version released (v 1.2)

- roughly 2x faster / less memory
- handles  $H^1$  of a data set (COVID genomes) with 93k points (136 trillion simplices)

# PERSISTENT HOMOLOGY

• . . . →



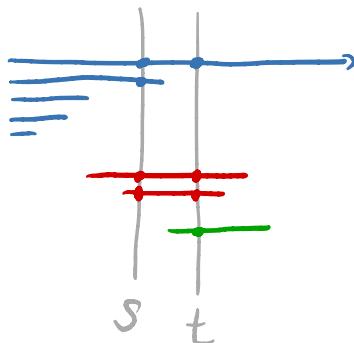
Filtration  $K_\cdot : K_1 \hookrightarrow K_2 \hookrightarrow K_3 \hookrightarrow \dots \hookrightarrow K_s \hookrightarrow \dots$

} Homology  $H_\ast$

Persistence  $H_\ast(K_\cdot) : H_\ast(K_1) \rightarrow \dots \rightarrow H_\ast(K_s) \rightarrow \dots$   
module

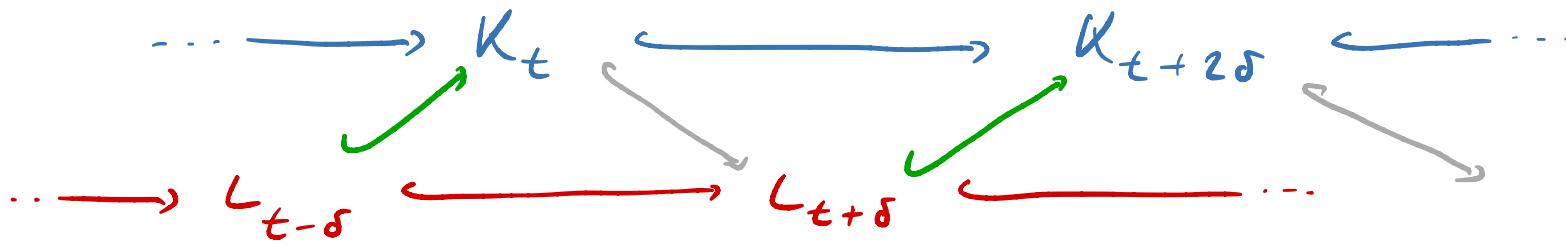
} Barcode  $B$

Matching  
diagram

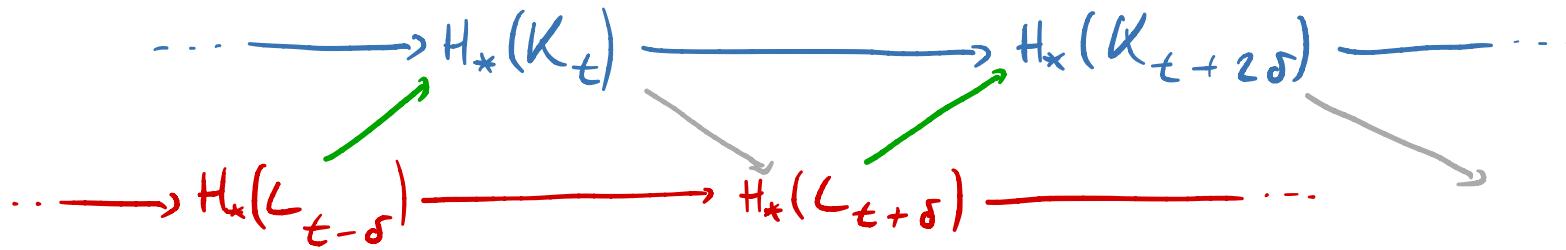


# INDUCED MATCHINGS

Two filtrations:



Interleaving of persistent homology:



$H_*(L_\cdot) \xrightarrow{\varphi} H_*(K_{\cdot+\delta})$  induces

matching of barcodes

$$B(H_*(L_\cdot)) \hookleftarrow B(\text{im } \varphi) \hookrightarrow B(H_*(K_\cdot))$$



—

# ALGEBRAIC STABILITY OF BARCODES

Theorem [Chazal et al. '09] [B, Lesnick '14]

A  $\delta$ -interleaving of persistence modules  
~~admits~~ induces a  $\delta$ -matching of their barcodes.

Applications: (extending classical stability)

- weaker tameness assumptions
- filtrations of two different domains
- Gromov-Hausdorff stability (Vietoris-Rips barcode)

# PERSISTENCE COMPUTATION : THE BASICS

$(X, d)$  : metric space

$$\text{Rips}_t(X) = \{\emptyset \neq \sigma \subseteq X \mid \text{diam } \sigma \leq t\}$$

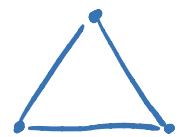
→ Rips filtration

→ refine (lexicographically) to simplexwise

$D$  : p-boundary matrix (wrt. ordered simplices)

Matrix reduction : compute  $R = D \cdot V$

- $R$  reduced : unique pivots
- $V$  upper triangular, full rank



$$\begin{pmatrix} 1 & & \\ 1 & 1 & \\ 1 & & \end{pmatrix} = \begin{pmatrix} 1 & 1 & 1 \\ 1 & 1 & 1 \\ 1 & 1 & 1 \end{pmatrix} \cdot \begin{pmatrix} 1 & & \\ 1 & 1 & \\ 1 & & \end{pmatrix} \quad (\text{in } \mathbb{Z}/(2))$$

R            D            V

Columns of  $R$  and some of  $V$  form a compatible basis for persistent homology:

$R_i$  generates homology (at index  $i$  = pivot  $R_i$ )

$V_j$  kills homology (at index  $j$ )

If  $R_i = 0$  :

- $V_i$  generates homology (at index i)
- if  $i = \text{pivot } R_j$  : we don't need  $V_i$
- else :  $V_i$  is essential cycle

Clearing (Chen, Kerber 2011) :

- avoid unneeded computations
- requires reducing boundary matrices  
in decreasing dimensions
- unavailable in top dimension

## Persistent cohomology

- vector space dual to homology
- same barcode
- computed by reducing coboundary matrices  
(transpose boundary matrix & reverse order)  
in increasing dimension
- unavailable in dimension 0  
(where Kruskal's MST algorithm is used)

Filtration  $K_1 \hookrightarrow K_2 \hookrightarrow \dots \hookrightarrow K_m = A$

Cochains  $C^*(K_1) \leftarrow C^*(K_2) \leftarrow \dots \leftarrow C^*(K_m)$

Relative filt.  $C^*(A, K_1) \leftarrow C^*(A, K_2) \leftarrow \dots \leftarrow C^*(A, K_m)$

Reducing the coboundary matrix actually computes persistent relative cohomology!

[dSMV11] : absolute / relative barcodes determine each other.

Consider short exact sequence

$$C_*(K_*) \hookrightarrow C_*(A) \rightarrow C_*(A, K_*)$$

of filtered chain complexes.

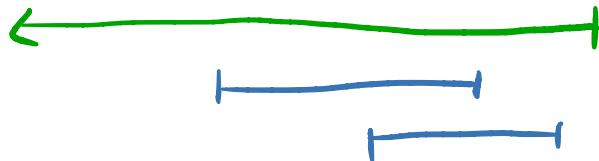
→ long exact sequence

$$\begin{matrix} & H_d(K_*) \rightarrow H_d(A) \rightarrow H_d(A, K_*) \\ \curvearrowleft & H_{d-1}(K_*) \xrightarrow{\eta} H_{d-1}(A) \rightarrow \dots \end{matrix}$$

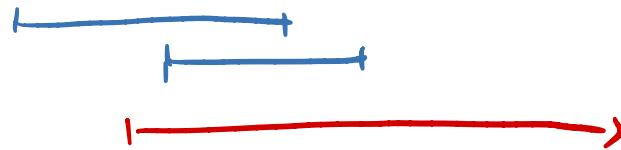
- splits at  $H_{d-1}(K_*) = \underbrace{H_{d-1}(K_*)}_{\text{im } \eta}^\infty \oplus \underbrace{H_{d-1}(K_*)}_{\text{im } \partial}^+$   
and at  $H_d(A, K_*) = \underbrace{H_d(A, K_*)}_{H_d(A, K_*)^*}^* \oplus \underbrace{H_d(A, K_*)}_{H_d(A, K_*)^{-\infty}}$

Example

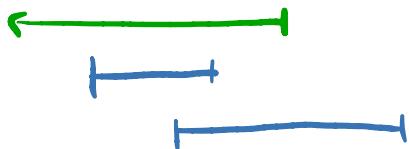
$$H_d(A, K.)$$



$$H_{d-1}(K.)$$



$$H_{d-1}(A, K.)$$



⋮

# COMPUTING IMAGE BARCODES

Now consider a pair of filtrations:

$$K_1 \hookrightarrow K_2 \hookrightarrow \dots \hookrightarrow K_m = A$$

$$\uparrow \quad \uparrow$$

$$f : L_0 \hookrightarrow K_0$$

$$L_1 \hookrightarrow L_2 \hookrightarrow \dots \hookrightarrow L_m = A$$

Compute  $\text{im } H_*(f)$ ?

- Long exact sequence is functorial
- But image may not be an exact sequence

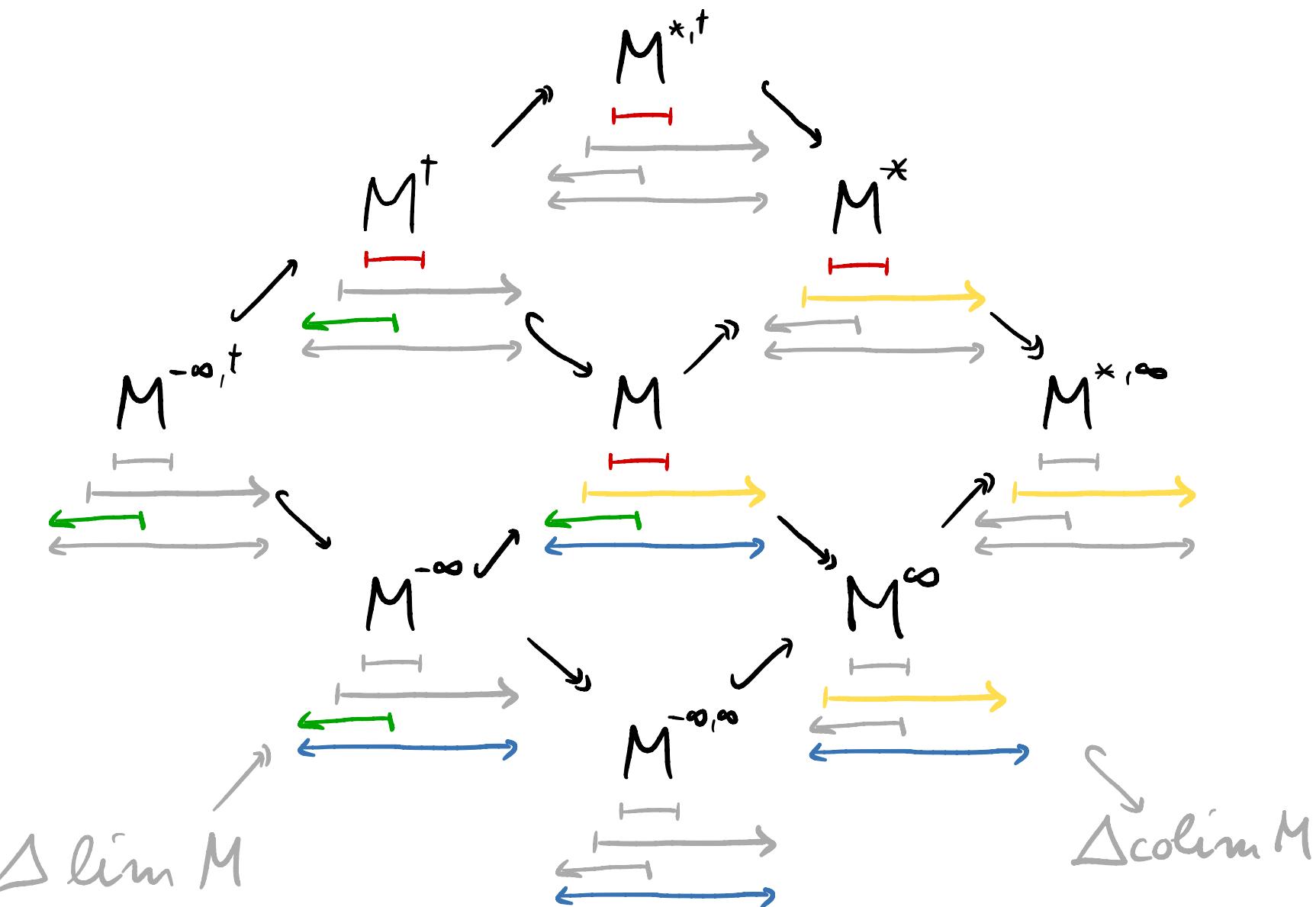
Previous work :

- [Edels., Harer, Kozu '09] :
  - $L_t = K_t \cap L$  (one function)
  - no clearing
- [Skraba, Vej.-Joh '13]
  - for presentations of persistence modules

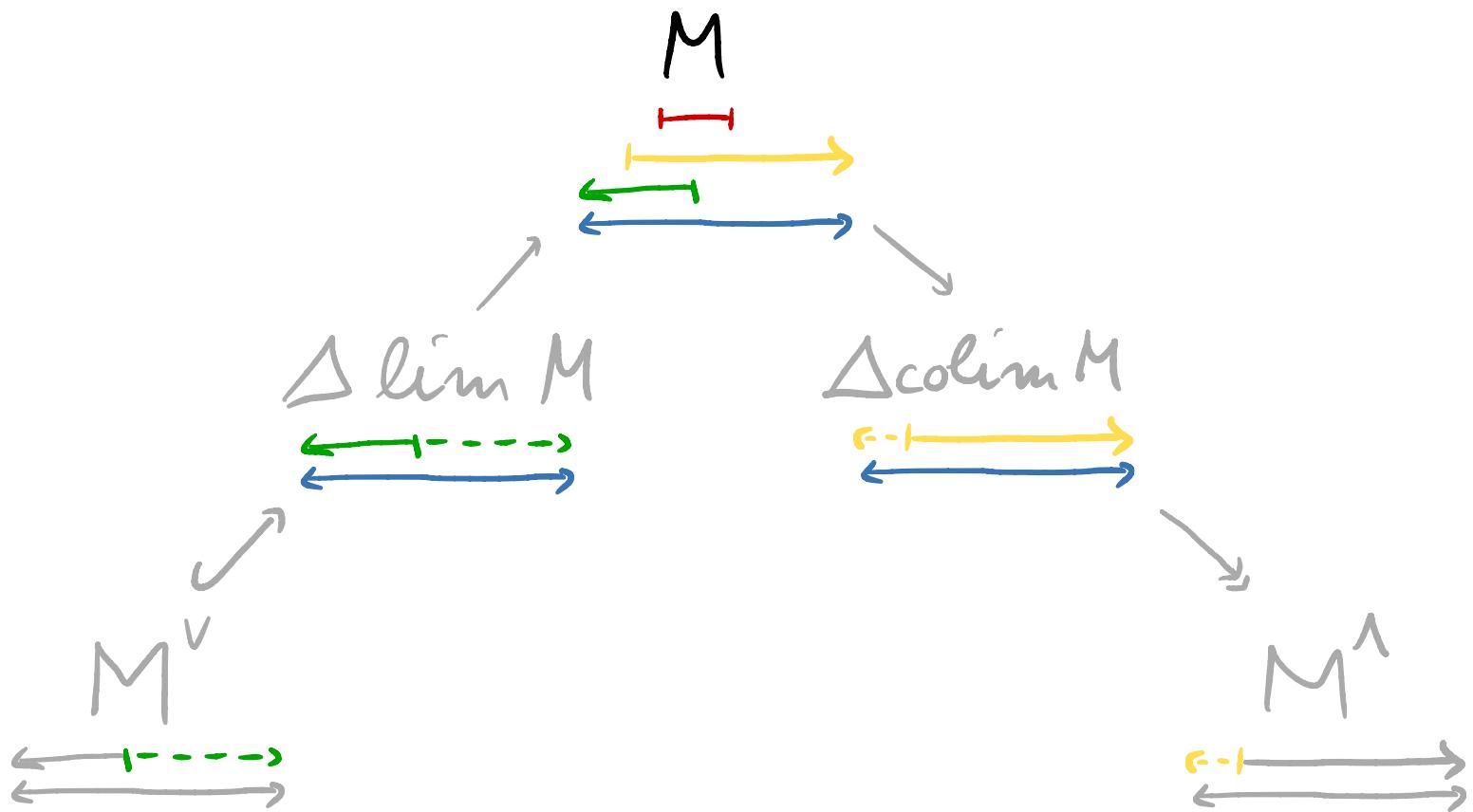
This talk:

- [B, Schmahl '20]
  - $K$ ,  $L$  arbitrary filtrations of  $A$
  - Clearing & cohomology

# LIFESPAN FUNCTORS

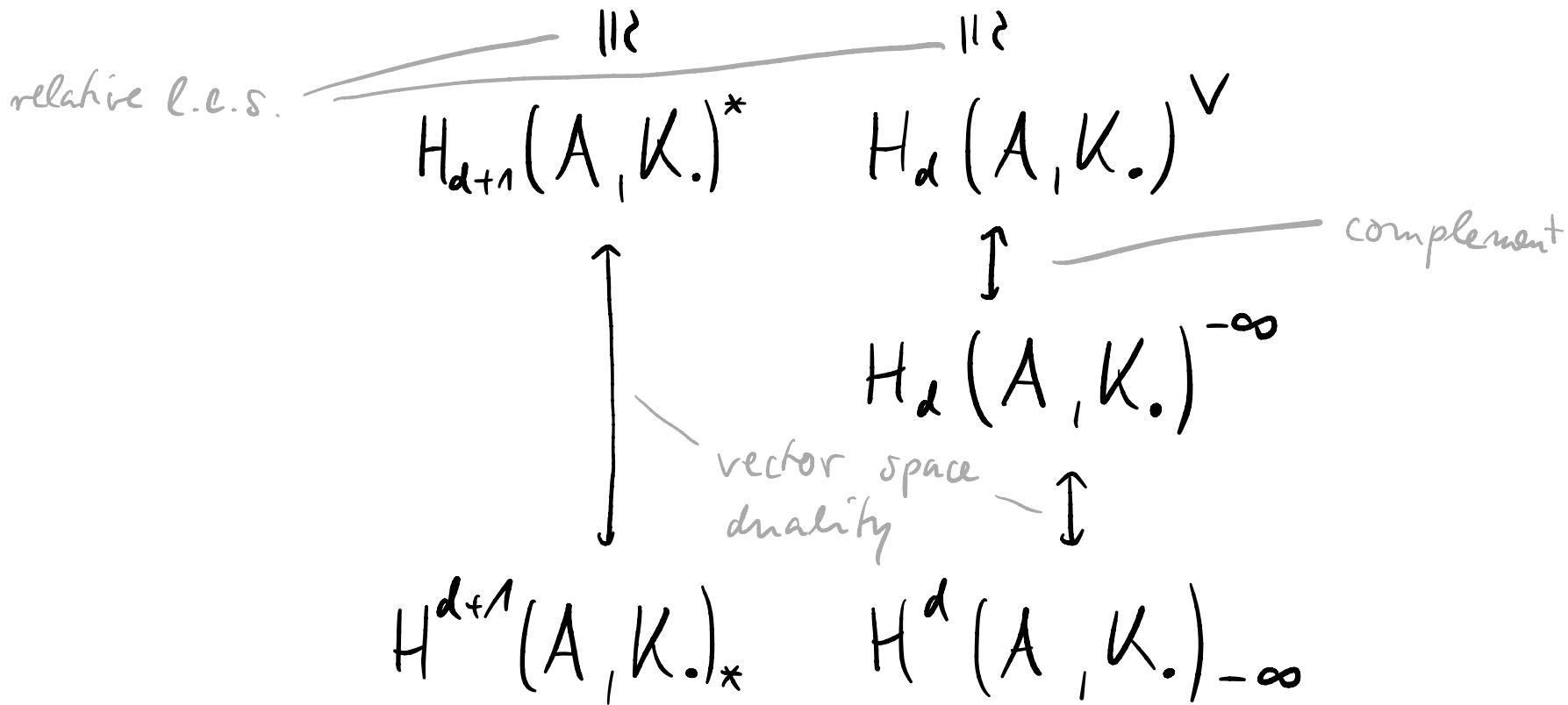


# COMPLEMENTS



$$H_d(K_{\cdot}) \doteq H_d(K_{\cdot})^+ \oplus H_d(K_{\cdot})^-$$

interval decomposition  
not natural



$$f : L_+ \hookrightarrow K_+ \quad \phi : (A, L_+) \hookrightarrow (A, K_+)$$

$$\text{im } H_d(f) \cong (\text{im } H_d(f))^+ \oplus (\text{im } H_d(f))^\infty$$

requires  $\text{colim } H_d(f)$  mono — ||2 ||2

$$\text{im}(H_d(f)^+) \quad H_d(L_+)^{\infty}$$

relative l.e.s. ————— ||2 ||2

$$\text{im}(H_{d+1}(\phi)^*) \quad H_d(A, L)^{\vee}$$

requires  $\lim H_{d+1}(\phi)$  epi ————— ||2 ↓

$$(\text{im } H_{d+1}(\phi))^* \quad H_d(A, L_+)^{-\infty}$$

# COMPUTING IMAGE PERSISTENCE

$$\begin{array}{ccc} & \xrightarrow{L} & \\ & \downarrow D^L & \\ L & & K \\ & \downarrow D^{im} & \end{array}$$

Reduce to  $R^L = D^L V^L$        $R^{im} = D^{im} V^{im}$

Theorem [B, Schmahl]

$$B(\text{im } H_k(L_+ \hookrightarrow K_+)) =$$

$$\{(i, j) \mid i = \text{pivot } R^{im}{}_j\}$$

$$\cup \{(i, \infty) \mid R^L{}_{ii} = 0, i \in \text{pivots } R^L\}$$

# CLEARING FOR IMAGE PERSISTENCE

Homology:

$$R^L = D^L V^L$$

{

$$H_*(f)^\infty$$

$$R^{im} = D^{im} V^{im}$$

{

$$H_*(f)^+$$

$$R^K = D^K V^K$$

{

$$H_*(\phi)^\infty$$

Cohomology:

$$S^L = W^L D^L$$

{

$$H^*(\phi)_\infty$$

$$S^{im} = W^{im} D^{im}$$

{

$$H^*(f)_+$$

$$S^K = W^K D^K$$

{

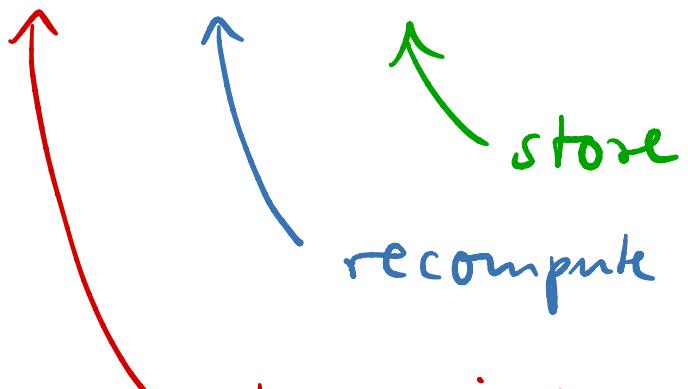
$$H^*(f)_\infty$$

& clearing  $R^{im}$

& clearing  $S^{im}$

# IMPLICIT REDUCTION

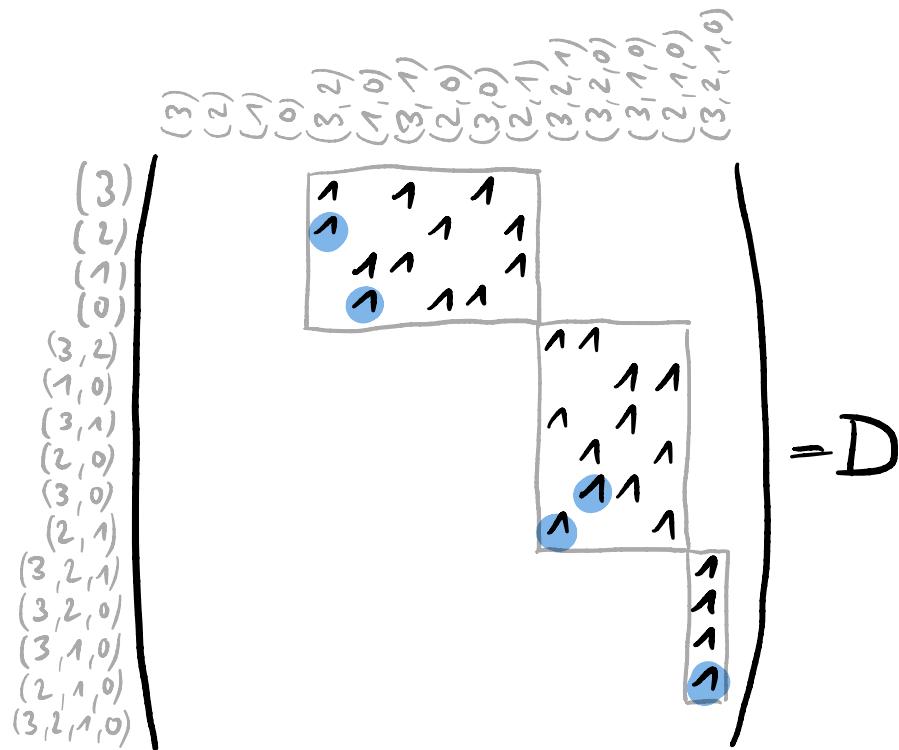
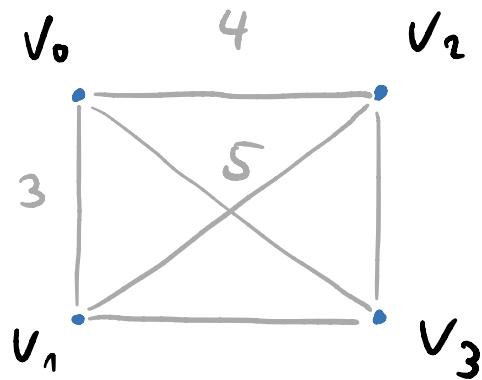
$$R = D \cdot V$$



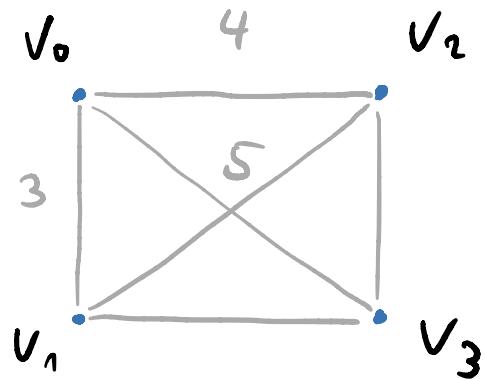
store pivots (unless apparent)  
recompute other entries

# APPARENT PAIRS

(how simplices find the perfect partner)



# APPARENT ZERO PAIRS



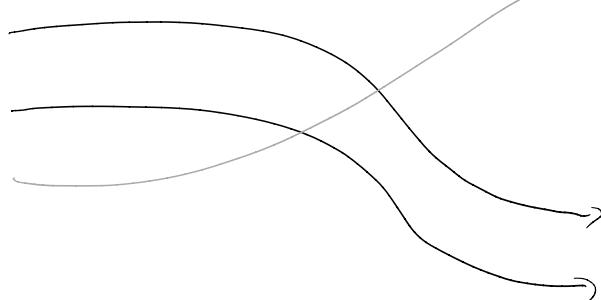
$(3, 2, 0) : \text{diam} = 5$

facets (lex order):

$(2, 0)$

$(3, 0)$

$(3, 2)$

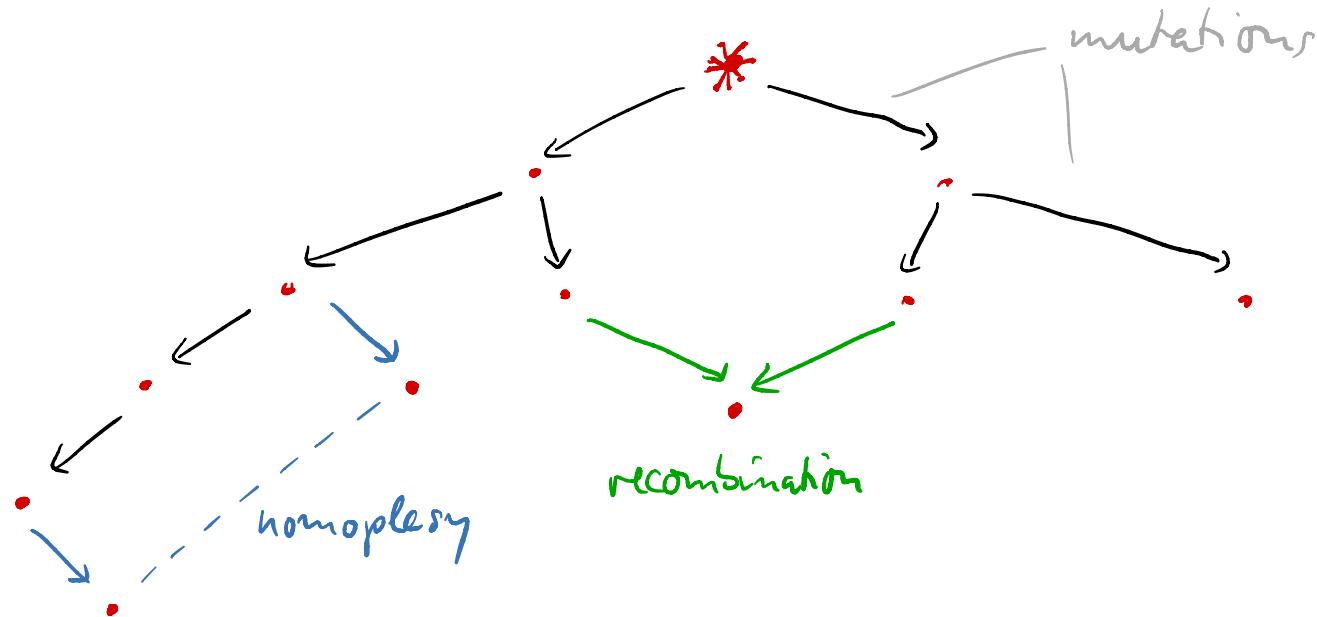


1
1
1

$(3, 2) :$	3
$(1, 0) :$	3
$(3, 1) :$	4
$(2, 0) :$	4
$(3, 0) :$	5
$(2, 1) :$	5

# TOPOLOGY OF VIRAL EVOLUTION

[Chen et al. 2013]



Example (Mutation E484K) :

- prevalence still low (15.1.21: <1%)
- but appears in several persistent cycles

# ACKNOWLEDGEMENTS

Induced matchings

joint with M. Lesnick (U Albany)

doi: 10.20382/jocg.v6i2a9

doi: 10.1007/978-3-030-43408-3\_3

Image persistence

joint with M. Schmahl (U Heidelberg)

arxiv: 2012.12881

Ripses

ripses.org

Cor'd

joint with: M. Bleher, L. Hahn, A. Ott (U Heidelberg)

J. Patiño-Galindo, R. Rabadian (Columbia U)

M. Carrère (INRIA Sophia-Antipolis)

Funding: DFG (SFB-TR 109)

# AN EXCURSION INTO THE WILDERNESS

Given a finite indexing poset  $P$ .

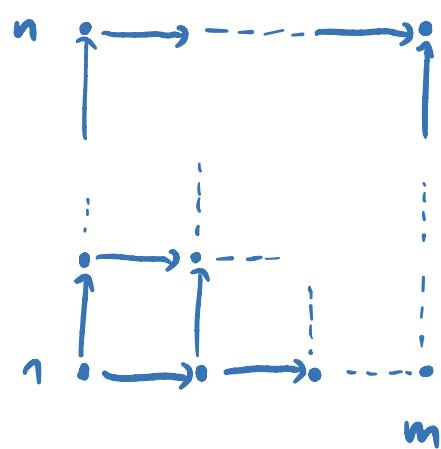
Indecomposable diagrams of  $K$ -vector spaces  
with shape  $P$ ? ( $K$  algebra. closed)

3 cases (representation types) :

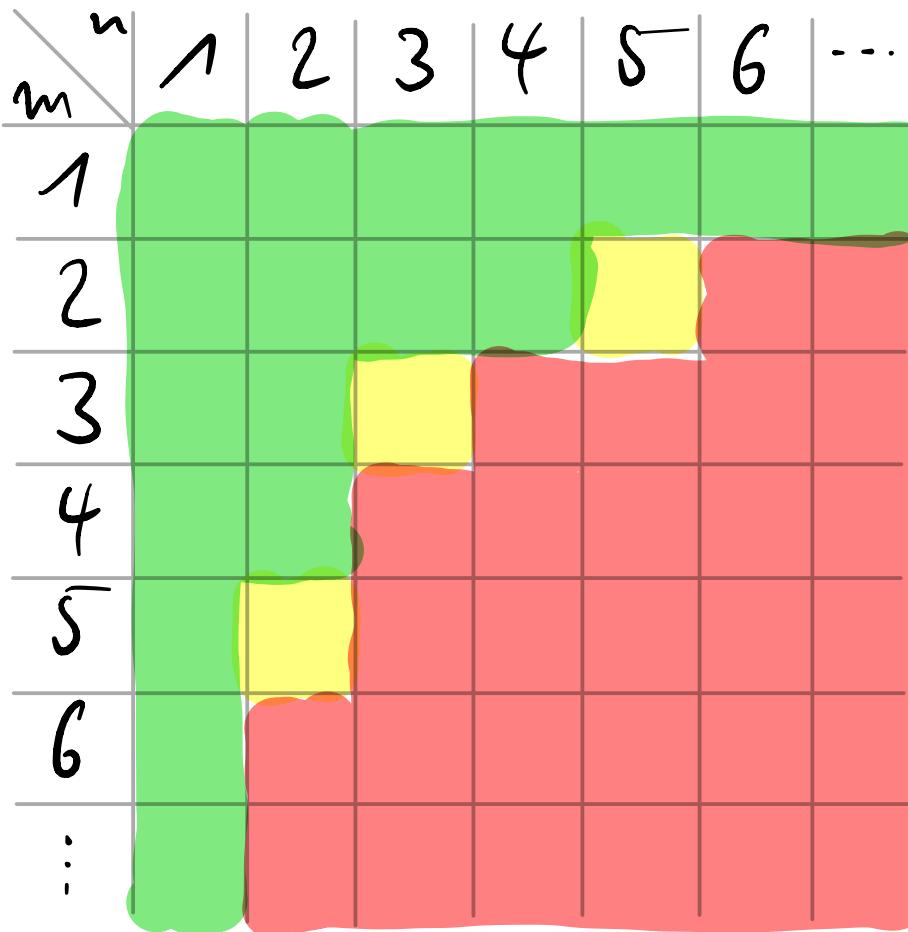
- (a) A finite list finite type
- (b) A finite list (of 1-param. families) tame
- (c) It's complicated. wild

(as complicated as modules over any finite-dim. algebra; undecidable theory)

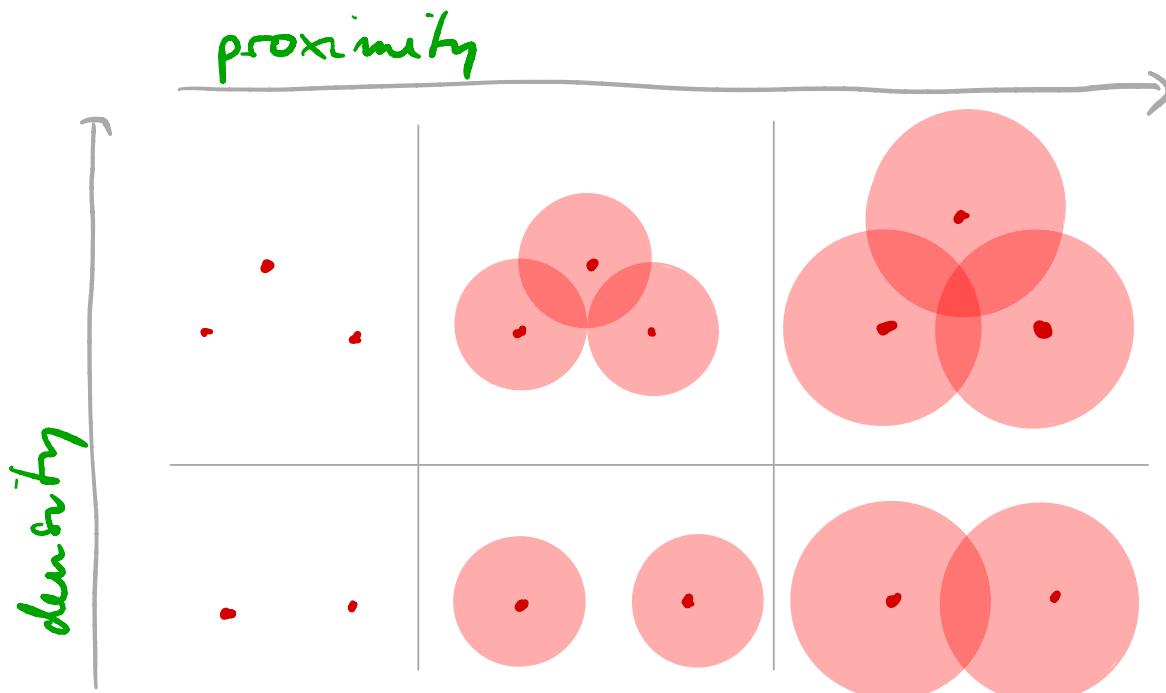
# REPRESENTATION TYPES OF COMMUTATIVE GRIDS



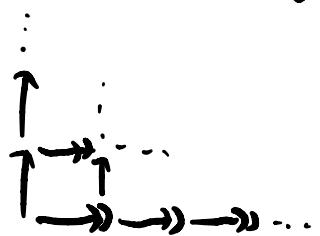
finite }  
 tame } :  $(m-1)(n-1)$  {  
 wild }  $<$  =  $>$  4



# GRID DIAGRAMS FROM CLUSTERING



apply  $H_0 \rightsquigarrow$  diagram of the form



(horizontal : surjective)

$$\begin{array}{ccc}
 K^3 & \xrightarrow{(111)} & K \longrightarrow K \\
 \left(\begin{matrix} 1 & 0 \\ 0 & 1 \\ 0 & 0 \end{matrix}\right) \uparrow & (11) \uparrow & \uparrow \\
 K^2 & \longrightarrow & K^2 \xrightarrow{(11)} K
 \end{array}$$

$$\cong \boxed{\begin{array}{ccc}
 K & \longrightarrow & K \longrightarrow K \\
 \uparrow & & \uparrow \\
 K & \longrightarrow & K \longrightarrow K
 \end{array}} \oplus \boxed{\begin{array}{ccc}
 K & \longrightarrow & 0 \longrightarrow 0 \\
 \uparrow & \uparrow & \uparrow \\
 K & \longrightarrow & K \longrightarrow 0
 \end{array}} \oplus \boxed{\begin{array}{ccc}
 K & \longrightarrow & 0 \longrightarrow 0 \\
 \uparrow & \uparrow & \uparrow \\
 0 & \longrightarrow & 0 \longrightarrow 0
 \end{array}}$$

Lemma  $\text{Rep}^{\rightarrow}(m, 2)$  is finite type.

# EPIC GRIDS & WILD THINGS

Theorem [B, Botnan, Oppermann, Steen '19]

*Corollary*  $\text{Rep}^{\rightarrow}(m, n)$  is

finite type

Tame

wild

} for  $(m-1)(m-2)$  { < = > } 4 .