

Super-Polynomial Growth of the Generalized Persistence Diagram

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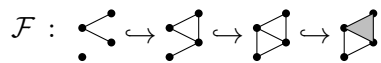
Dept. of Mathematical Sciences, KAIST

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Introduction

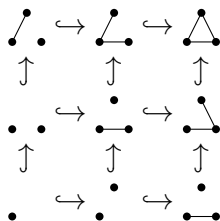
(1-parameter) Persistent Homology v.



$$H_0 : \mathbb{F}^2 \rightarrow \mathbb{F} \rightarrow \mathbb{F} \rightarrow \mathbb{F}$$

$$H_1 : 0 \rightarrow \mathbb{F} \rightarrow \mathbb{F}^2 \rightarrow \mathbb{F}$$

Multi-parameter Persistent Homology



Persistence module
over a **poset**

Persistent Homology

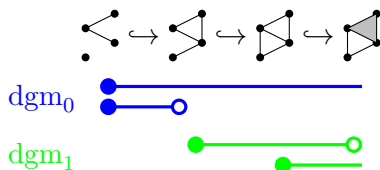
v.

Multi-parameter Persistent Homology

Persistence Diagram (equivalently, **Barcode**)

[Edelsbrunner, Letscher, Zomorodian '02],

[Carlsson, Zomorodian '05]



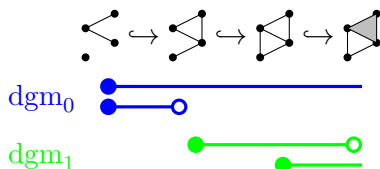
dgm_m denotes the PD obtained via $H_m(-; \mathbb{F})$.

?

Persistent Homology

Persistence Diagram (equivalently, Barcode)

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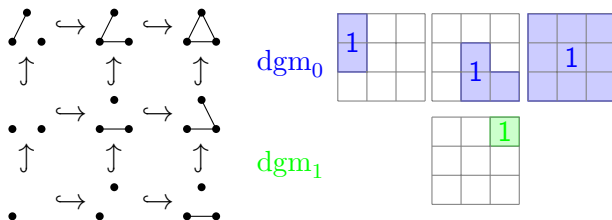
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v.

Multi-parameter Persistent Homology

Generalized Persistence Diagram (GPD)

[Patel '16], [Kim, Mémoli '19],
[Asashiba, Escobar, Nakashima, Yoshiwaki '19]



Here, dgm_m denotes the GPD obtained via $H_m(-; \mathbb{F})$.

Persistent Homology

Persistence Diagram

“viewed as the function”

$$\mathrm{dgm}_M([a, b]) = \text{mult. of } \mathbb{I}_{[a, b]} \text{ in } M$$

[Edelsbrunner, Letscher, Zomorodian '02],

[Carlsson, Zomorodian '05]

Rank Invariant

$$\mathrm{rk}_M([a, b]) = \mathrm{rank}(M_a \rightarrow M_b)$$

[Carlsson, Zomorodian '09]

$$\Rightarrow \mathrm{rk}_M([a, b]) = \sum_{\substack{J \in \mathrm{Int}(\mathbb{Z}) \\ J \supseteq [a, b]}} \mathrm{dgm}_M(J)$$

v.

Multi-parameter Persistent Homology

Generalized Rank Invariant (GRI)

$$\mathrm{rk}_M : \mathrm{Int}(P) \rightarrow \mathbb{Z}_{\geq 0}$$

$$\mathrm{rk}_M(I) = \mathrm{rank}(\varprojlim M|_I \rightarrow \varinjlim M|_I)$$

[Kinser '08], [Kim, Mémoli '19], [AENY '19],

[Chamber, Letscher '19]

The **GPD** is defined as the **Möbius inverse** of the GRI on intervals.

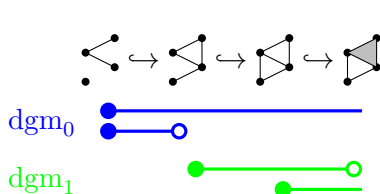
$$\mathrm{rk}_M(I) = \sum_{\substack{J \in \mathrm{Int}(P) \\ J \supseteq I}} \mathrm{dgm}_M(J)$$

[Patel '16], [Kim, Mémoli '19], [AENY '19]

Intuitively, the GRI counts how many features persist throughout an interval, and the GPD tries to count how many features persist **exactly** on the interval (not beyond).

Persistent Homology

Persistence Diagram



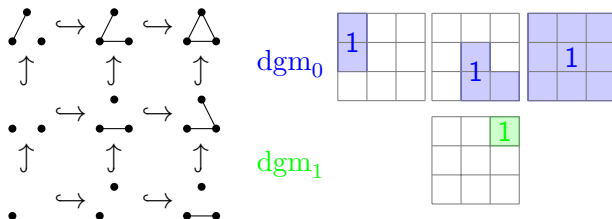
Remark

For a simplicial filtration $\mathcal{F}$ containing N simplices, then, for each $m \in \mathbb{Z}_{\geq 0}$, $dgm_m(\mathcal{F})$ contains **at most** N bars.

v.

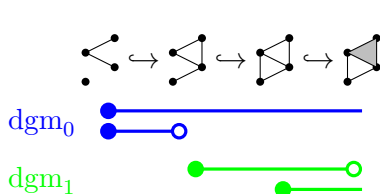
Multi-parameter Persistent Homology

Generalized Persistence Diagram (GPD)



Persistent Homology

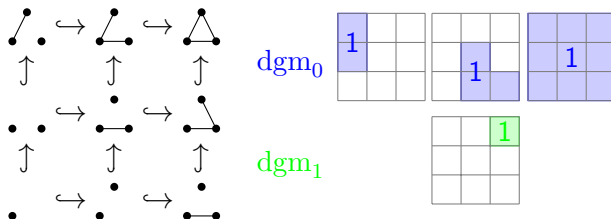
Persistence Diagram



v.

Multi-parameter Persistent Homology

Generalized Persistence Diagram (GPD)



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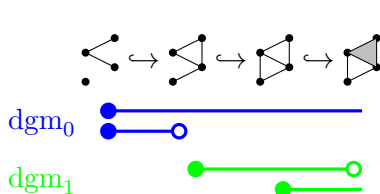
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Question

What about the multi-parameter case?
Is there an analogy with the 1-parameter setting?

Persistent Homology

Persistence Diagram



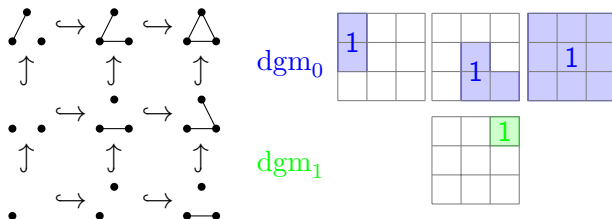
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v.

Multi-parameter Persistent Homology

Generalized Persistence Diagram (GPD)



Question

What about the multi-parameter case?
Is there an analogy with the 1-parameter setting?

And our answer is **no**.

Main Theorem 1

Theorem (Kim, Kim, L., SoCG 2025)

*For filtrations $\mathcal{F}$ over $\mathbb{R}^d$ ($d \geq 2$), there does **not** exist a polynomial in the number of simplices in $\mathcal{F}$, that bounds $\|\text{dgm}_m(\mathcal{F})\|$ for arbitrary $\mathcal{F}$.*

$\|\text{dgm}_m(\mathcal{F})\|$ denotes the **size** (the cardinality of the support) of the m^{th} GPD.

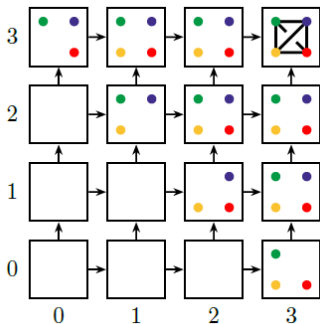
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Proof. (i) $d = 2$ case



$m = 0, d = 2, \mathcal{F}_3 (n = 3)$

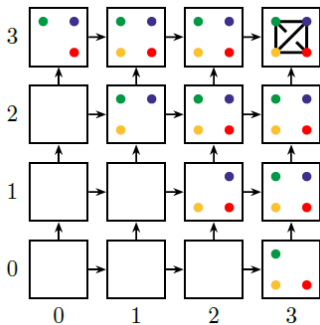
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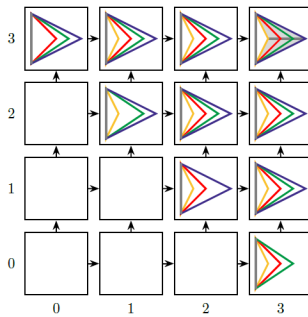
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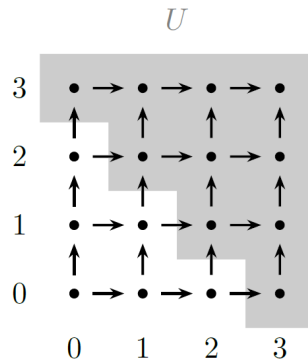
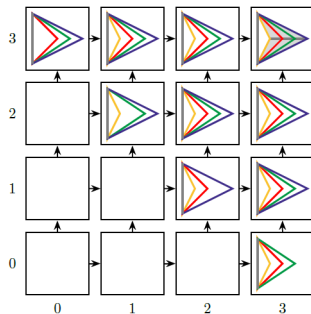
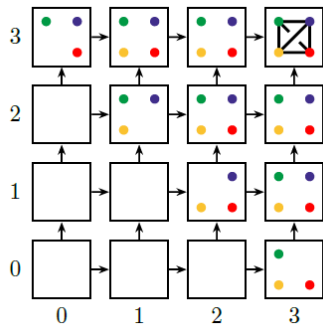


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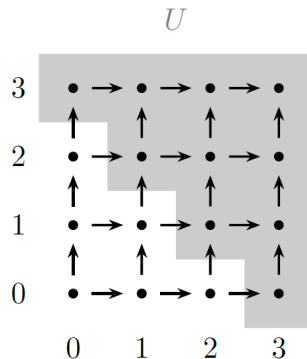
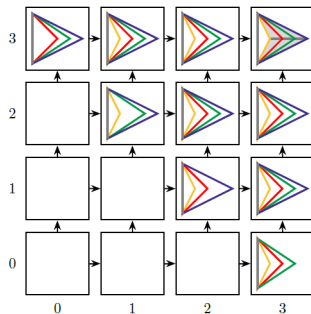
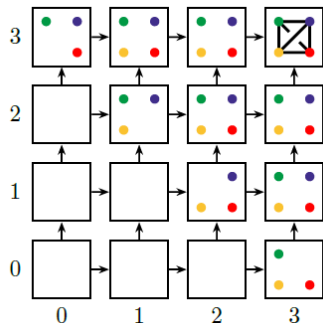
$m = 1, d = 2, \mathcal{F}_3 (n = 3)$

Main Theorem 1



Step 1. If J is not a proper subset of U , then $\text{rk}_M(J) = 0$.

Main Theorem 1

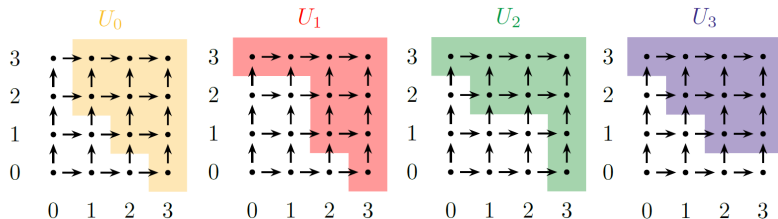
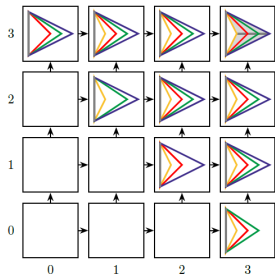
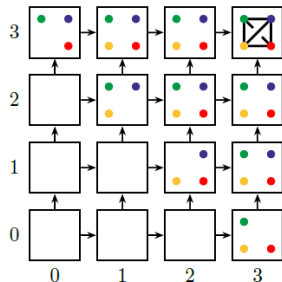


Step 1. If J is not a proper subset of U , then $\text{rk}_M(J) = 0$.

- None of the features persist over the entire U

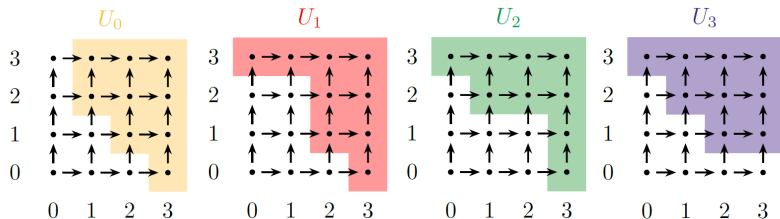
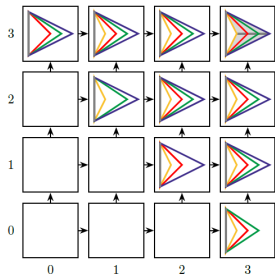
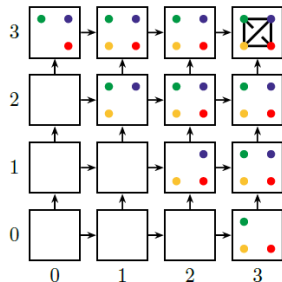
$\Rightarrow \text{dgm}_M(J) = 0$; don't need to consider the interval J

Main Theorem 1



Step 2. For $i \in [n]$, $\text{rk}_M(U_i) = 1 \Rightarrow \text{dgm}_M(U_i) = 1$

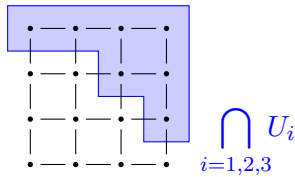
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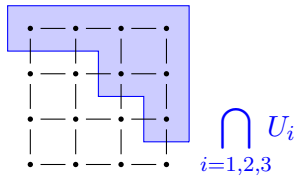
- Only the yellow feature persists throughout U_0 .
- Similarly, for all other intervals U_i
- $1 = \text{rk}_M(U_i) = \sum_{J \supseteq U_i} \text{dgm}_M(J) = \text{dgm}_M(U_i)$
from Step 1

Main Theorem 1



Step 3. For $\emptyset \neq S \subseteq [n]$, $\text{dgm}_M(\bigcap_{i \in S} U_i) = (-1)^{|S|+1} \neq 0$.

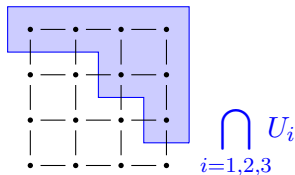
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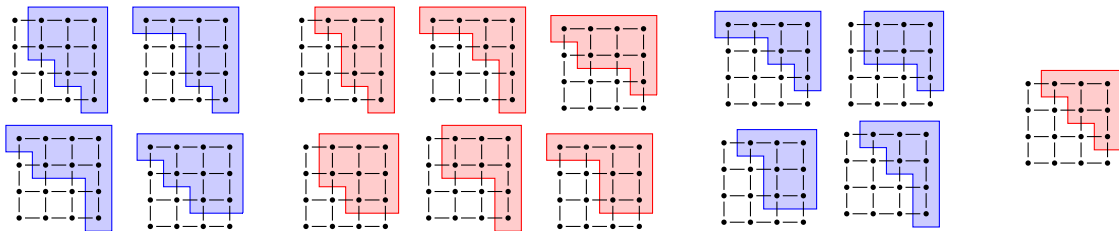
- Step 2 is actually the case of $|S| = 1$.
- The result comes from Step 1, 2, and a lemma [AENY '19].

Main Theorem 1



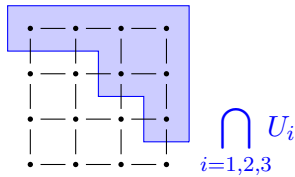
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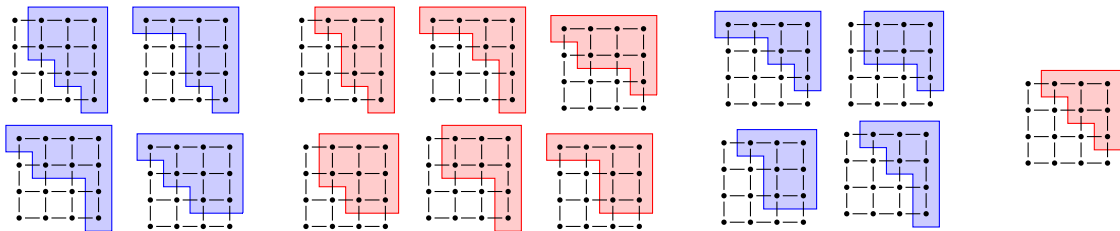
These are the intervals with nonzero GPD values identified through the previous steps.

Main Theorem 1



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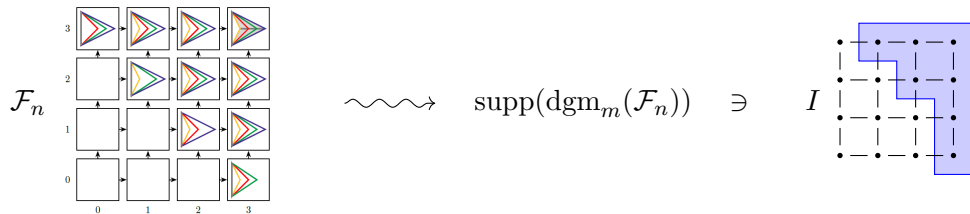


These are the intervals with nonzero GPD values identified through the previous steps.

We have $||\text{dgm}_m(\mathcal{F}_n)|| \geq 2^{n+1} - 1 \notin O(n^{(m+2)k}) = O(N^k)$.

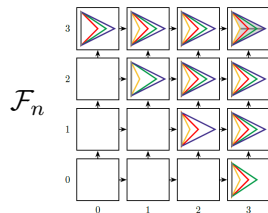
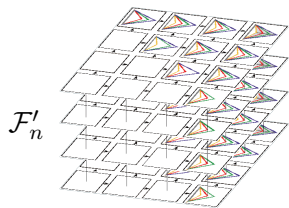
Main Theorem 1

(ii) What if $d > 2$?



Main Theorem 1

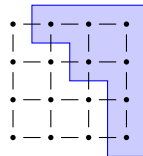
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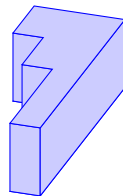
$\text{supp}(\text{dgm}_m(\mathcal{F}_n))$



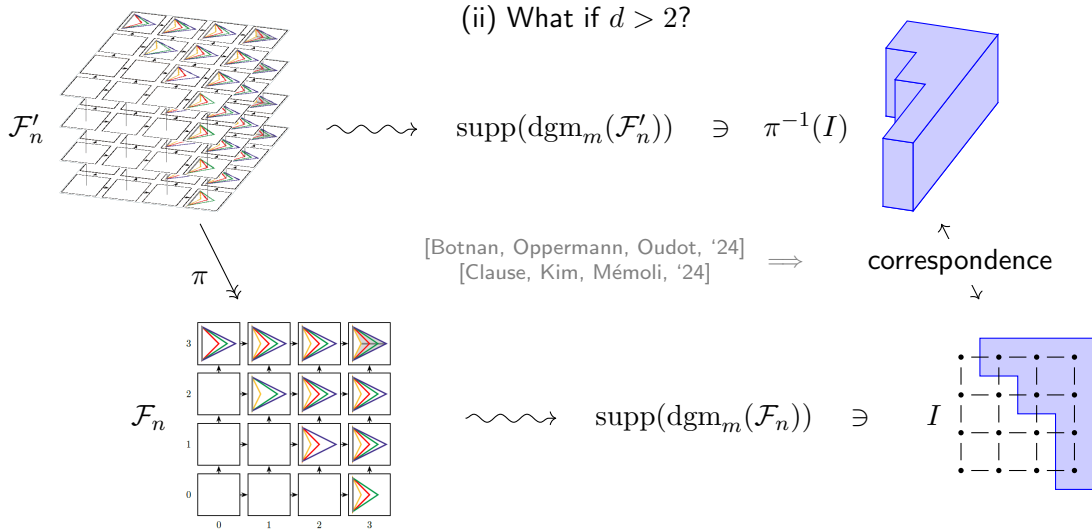
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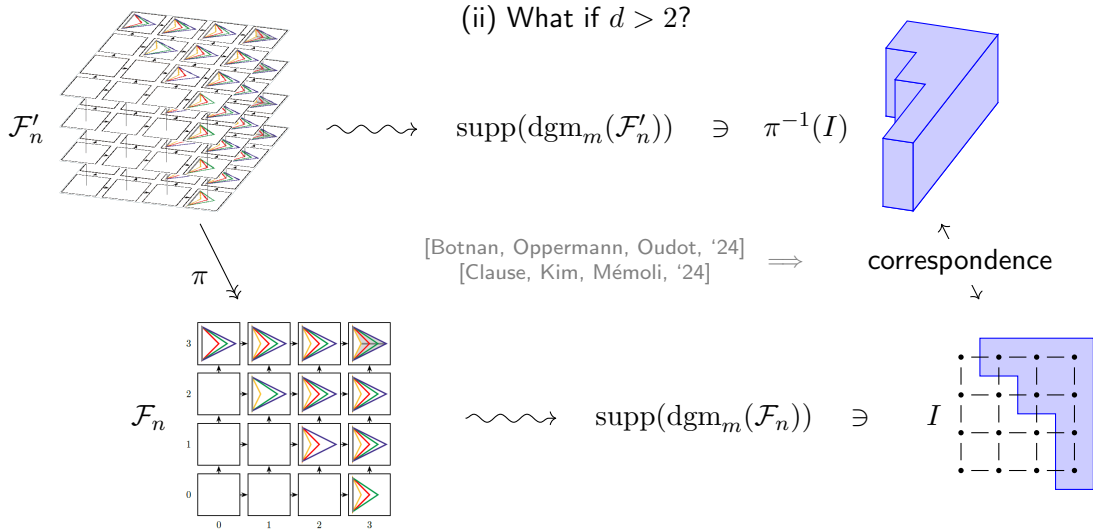
$\pi^{-1}(I)$



Main Theorem 1



Main Theorem 1



We have $\|\text{dgm}_m(\mathcal{F}'_n)\| = \|\text{dgm}_m(\mathcal{F}_n)\| \geq 2^{n+1} - 1 \notin O(n^{(m+2)k}) = O(N^k)$.

Main Theorem 2

Question

Can GPDs with super-polynomial size also arise from common 2-parameter filtrations?

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Answer

Yes, considering the 1st GPD of a sublevel-Rips, a sublevel-Čech, a degree-Rips, or a degree-Čech filtration.

These 4 types of filtration are well-known constructions of 2-parameter filtrations.
We will take a look at the case of sublevel-Rips filtrations.

Main Theorem 2: Sublevel-Rips case

Theorem (Kim, Kim, L., SoCG 2025)

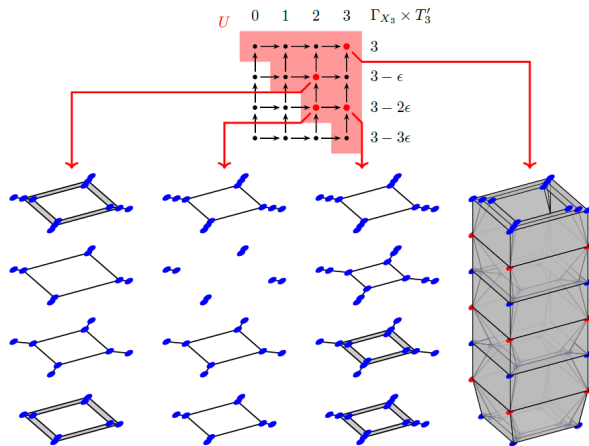
*For sublevel-Rips filtrations $\mathcal{F}$, there does **not** exist a polynomial in the number of simplices in $\mathcal{F}$, that bounds $\|\mathrm{dgm}_1(\mathcal{F})\|$ for any $\mathcal{F}$.*

Definition [Carlsson, Zomorodian '09]

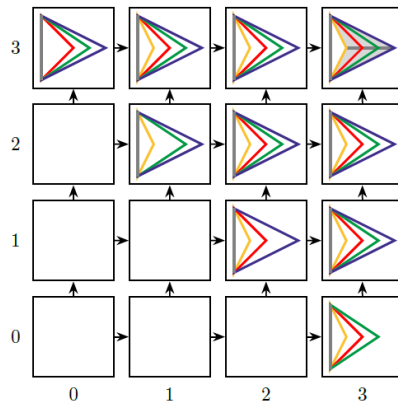
Given a metric space (X, d) and a function $\gamma : X \rightarrow \mathbb{R}$, the **sublevel-Rips** filtration is a filtration $\mathcal{F}$ on $\mathbb{R}^2$, which for each $(r, s) \in \mathbb{R}^2$, $\mathcal{F}_{(r,s)}$ is the Vietoris-Rips complex formed by a sublevel set $\{x \in X : \gamma(x) \leq r\}$, at scale s .

Main Theorem 2: Sublevel-Rips case

The (restricted) filtration when $n = 3$.



cf. $\mathcal{F}_3$ in Main theorem 1



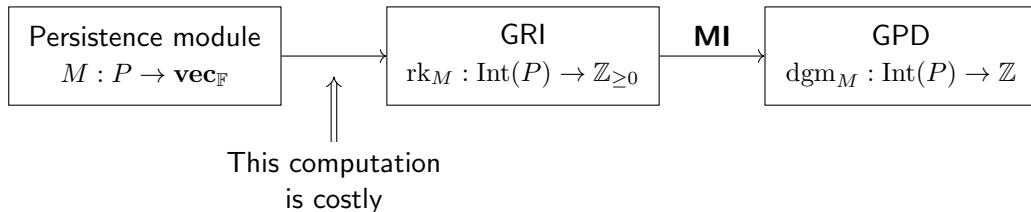
We adopt a similar strategy with Main theorem 1, by drawing on the results of [Gülen, McCleary '23].

Corollary (Kim, Kim, L., SoCG 2025)

For an arbitrary $[n]^d$ -module M ($d \geq 2$), computing $\mathrm{dgm}_M(I)$ via the Möbius inversion of the GRI of M requires at least $\Theta(2^n)$ in time in general (even the GRI is given).

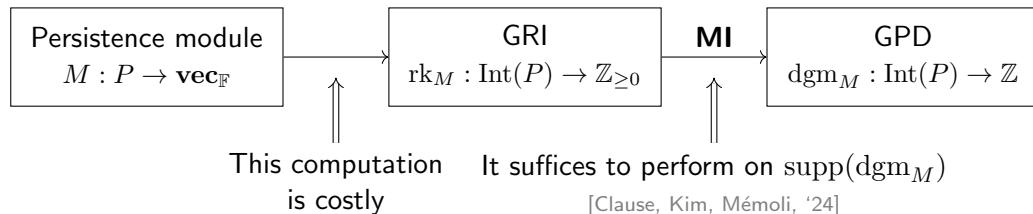
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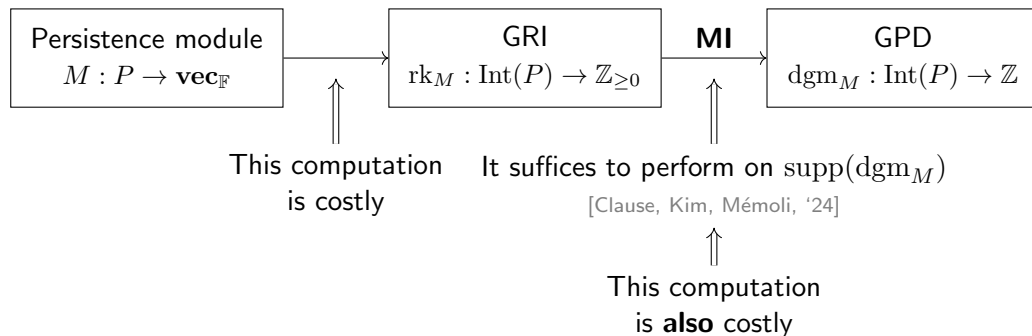
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Future directions

- Developing an output-sensitive algorithm [Morozov, Patel '23]
- Efficiently approximating the GPD [Carrière, Kim, Kim '24]

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Thank you for your attention!