

# WAVE TOP-K RANDOM-D FAMILY SEARCH

Interactive exploration of a space of structured patterns

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# Plan de l'exposé

1. Context and stakes
2. Method presentation
3. Experiences and evaluation

# Subgraph mining and chemoinformatics

## Data :

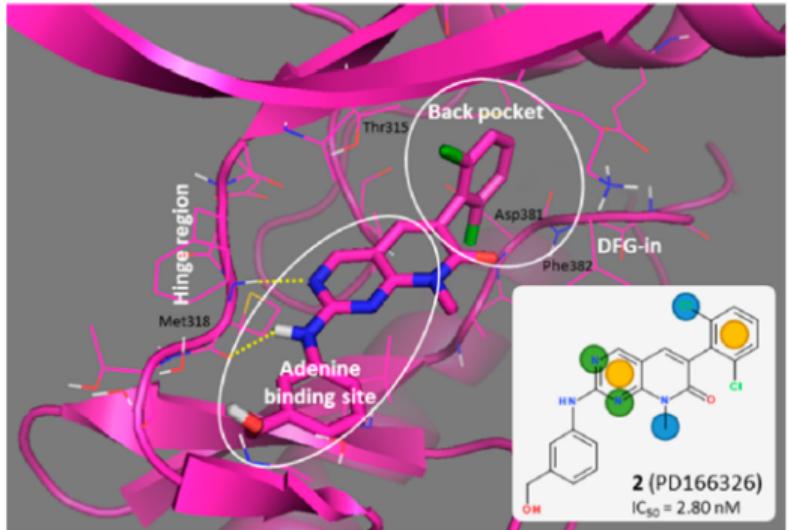
- ▶ 1 receptor : BCR-ABL
- ▶ 1485 molecules

## Structured pattern :

- ▶ 112 363 pharmacophores
- ▶ 15 533 equivalence classes

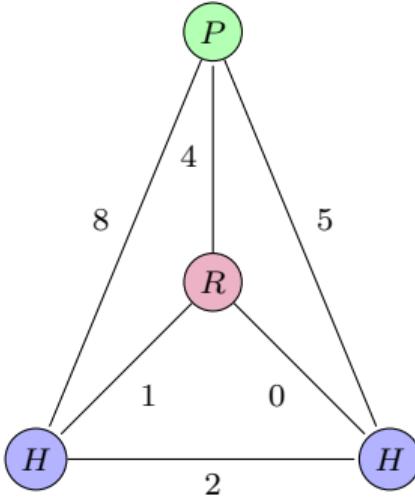
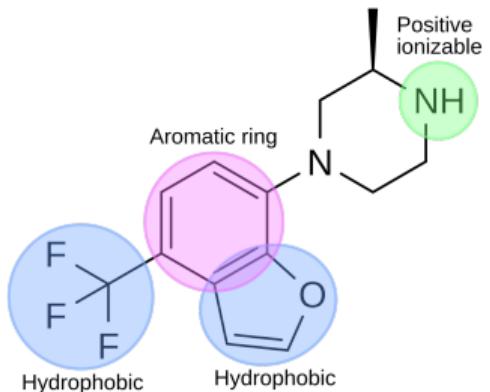
## Method goal :

Guide an experts' exploration of the pharmacophores set.



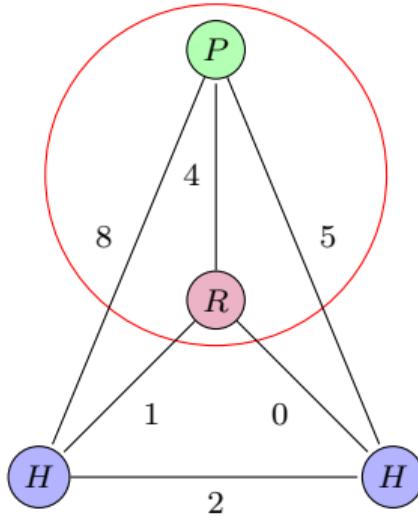
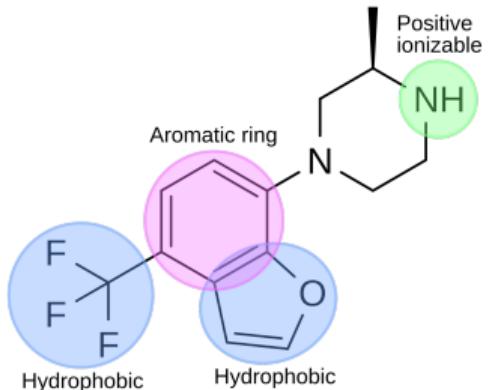
J.-P. METIVIER, B. CUISSART, R. BUREAU, and A. LEPAILLEUR.

# From molecules to pharmacophores (1)



- ▶ Pharmacophoric features = pharmacophore vertices.
- ▶ Distances between features = labels of the pharmacophore edges.

# From molecules to pharmacophores (2)



- ▶ The subgraph of a pharmacophore is a pharmacophore.
- ▶ We can build a partial ordered graph from the pharmacophore set.

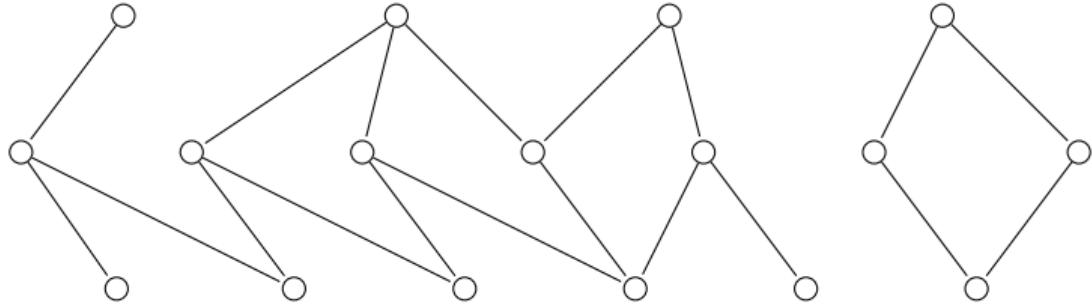
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# Proposed interactions

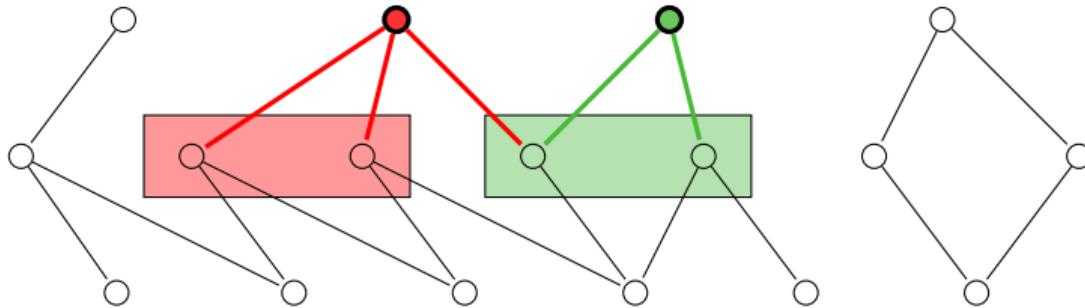
Interaction	Consequences	Color
Accepted	Prioritized area & Weights augmentation	Green
Interested	Weights augmentation	Blue
Uncertain	—	Purple
Not-interested	Weights diminution	Orange
Rejected	Exclusion area & Weights diminution	Red

# Example on a partially ordered graph



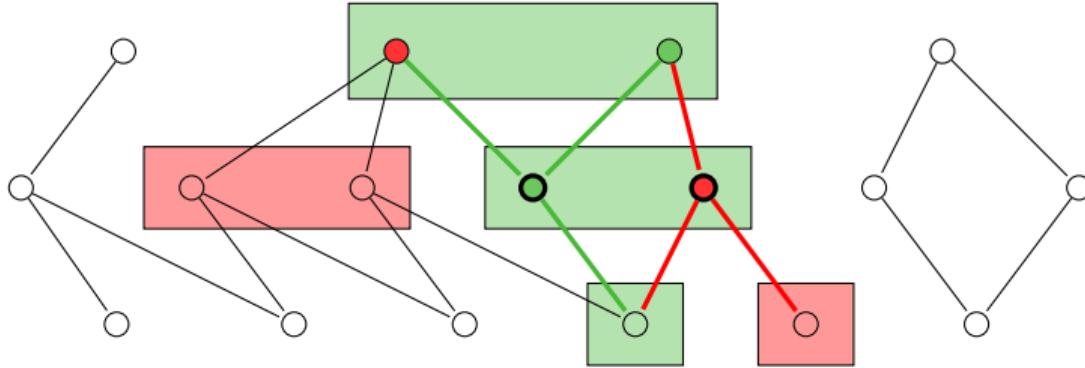
- ▶ Partially ordered graph as an input of the method.
- ▶ Each vertex is a pharmacophore.

## Definition of the search areas (1)



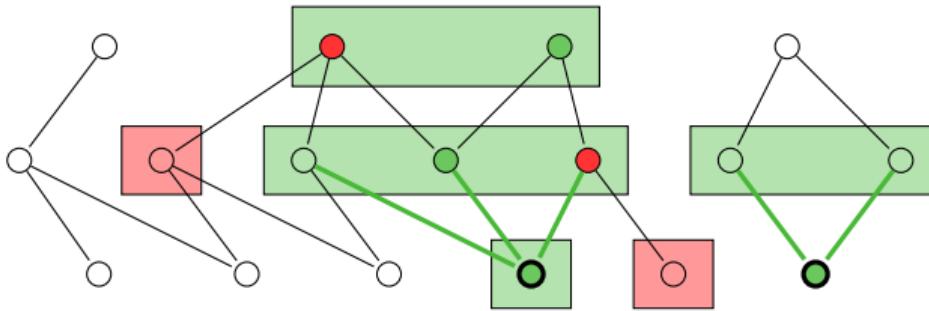
- ▶ **Interactions** : rejected (red) and accepted (green).
  - ▶ Exclusion area in red, prioritized area in green.
  - ▶ In case of **conflict**, the prioritized areas win.
  - ▶ Areas are only defined on the **directs descendants**.

## Definition of the search areas (2)



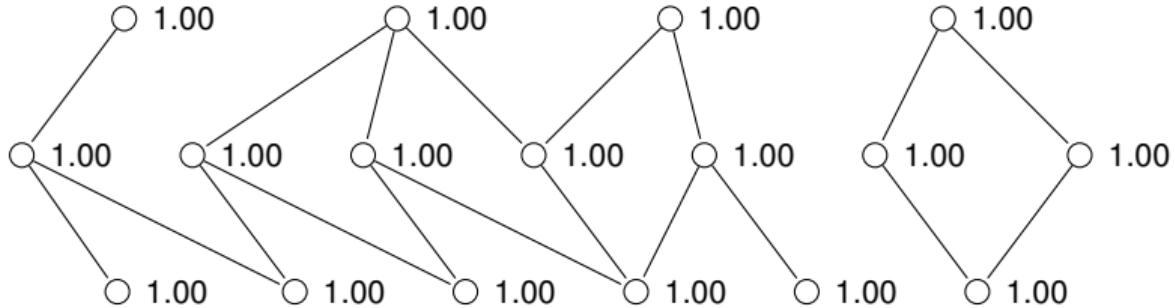
- ▶ Excluded and prioritized areas are defined on direct ancestors and descendants.
- ▶ There is not enough pharmacophores available in prioritized areas, we must offer a pharmacophore from an unbiased area.

## Definition of the search areas (3)



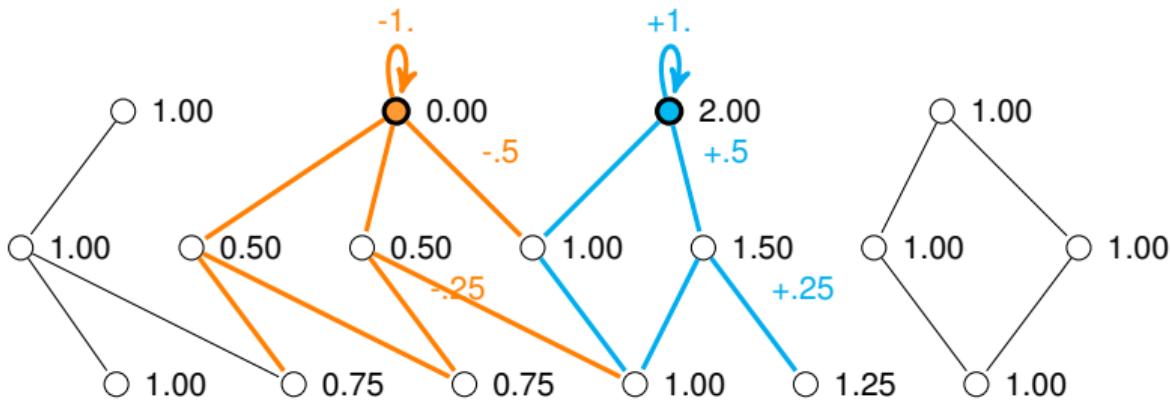
- ▶ Prioritized areas are defined on the direct ancestors.
- ▶ The search has reached the last layer. It goes up to the more generic patterns.
- ▶ **Question** : how to we choose the pharmacophores to propose ?

# Weights diffusion (1)



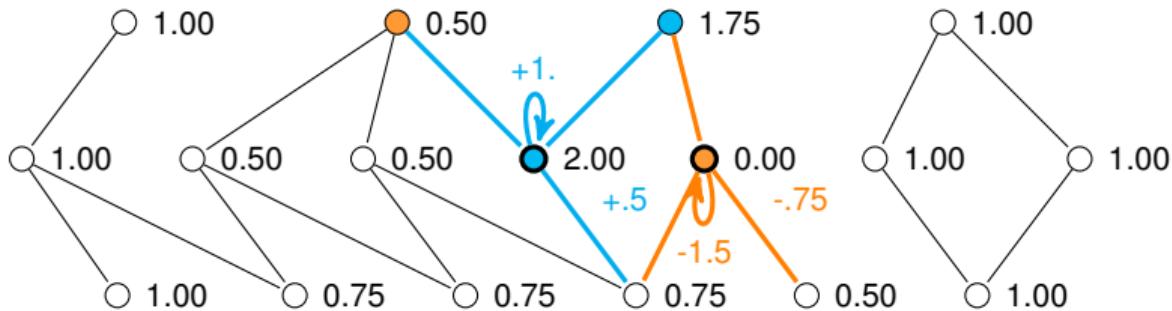
- ▶ The POG carries no interaction, every pharmacophores have the same weight.

## Weight diffusion (2)



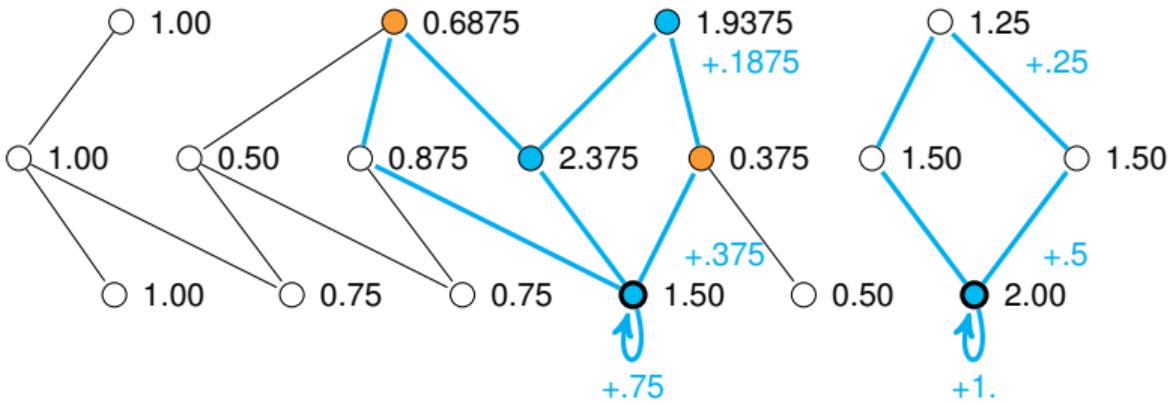
- ▶ **Interactions** : rejected or not-interested (orange) and accepted or interested (blue).
- ▶ Diffusing to the descendants :  $-1$  for the negative interest and  $+1$  for the positive interest.
- ▶ The raw interest diffused decreased with the increase of the distance from the origin of the interaction.

# Weight diffusion (3)



- ▶ Diffusing interest to ancestors and descendants.
- ▶ Diffused interest is the **weight** of the pharmacophore carrying the interaction.

# Weight diffusion (4)



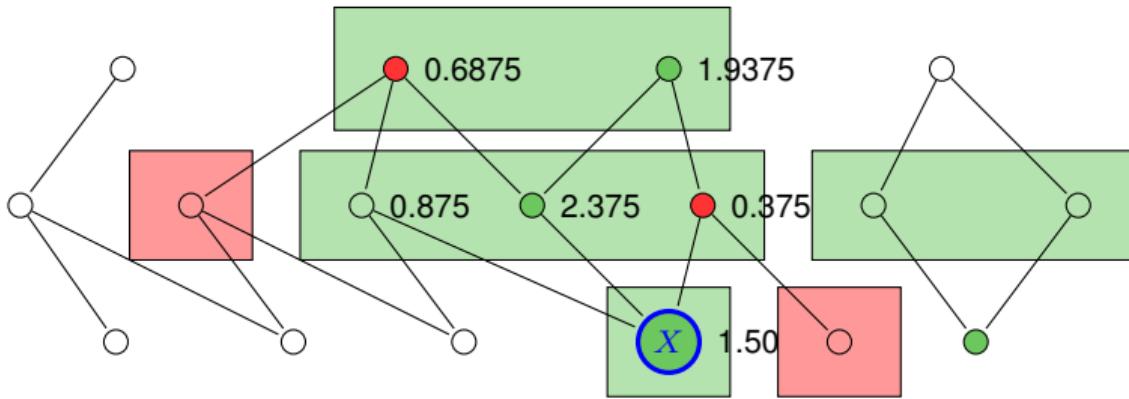
- ▶ Diffusing interest to the ancestors.
- ▶ The raw interest diffused decreased with the increase of the distance from the origin of the interaction.

# Computing potential interest (1)

$$f_p(v, \mathbb{G}) = \sum_{i=0}^k (Weight(L_i^+(v, \mathbb{G})) + Weight(L_i^?(v, \mathbb{G})) + (Weight(L_i^*(v, \mathbb{G})) - |Weight(L_i^+(v, \mathbb{G})) - Weight(L_i^-(v, \mathbb{G}))|)) * \frac{1}{2^i} \quad (1)$$

- ▶  $Weight(L_i^+(v, \mathbb{G}))$  is the **exploitation** heuristic.
- ▶  $Weight(L_i^?(v, \mathbb{G}))$  is the **exploration** heuristic.
- ▶  $(Weight(L_i^*(v, \mathbb{G})) - |Weight(L_i^+(v, \mathbb{G})) - Weight(L_i^-(v, \mathbb{G}))|)$  is the **ambiguity** heuristic.
- ▶  $1/2^i$  is the distance modifier.

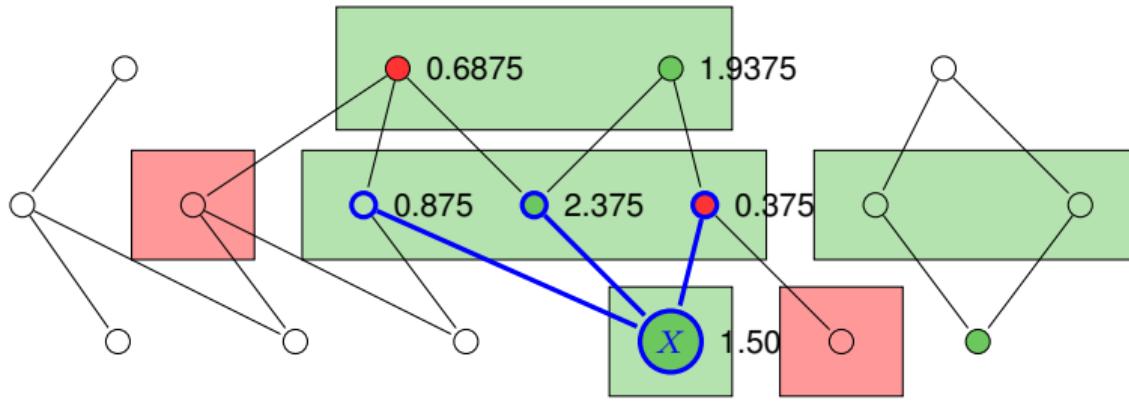
# Computing potential interest (2)



$$f_p(X) = 1.50$$

(2)

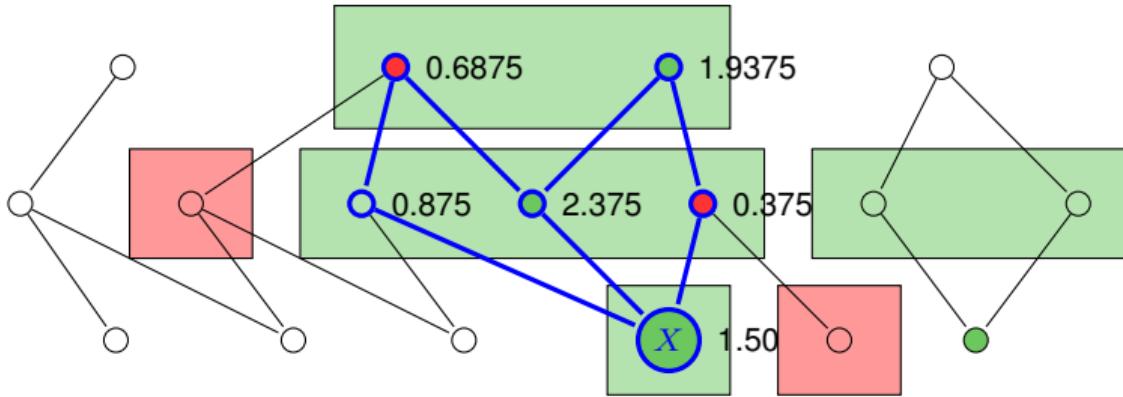
# Computing potential interest (3)



$$f_p(X) = 1.50 \tag{3}$$

$$+ (2.375 + 0.875 + (2.375 + 0.375 - |2.375 - 0.375|)) * \frac{1}{2} \tag{4}$$

## Computing potential interest (4)



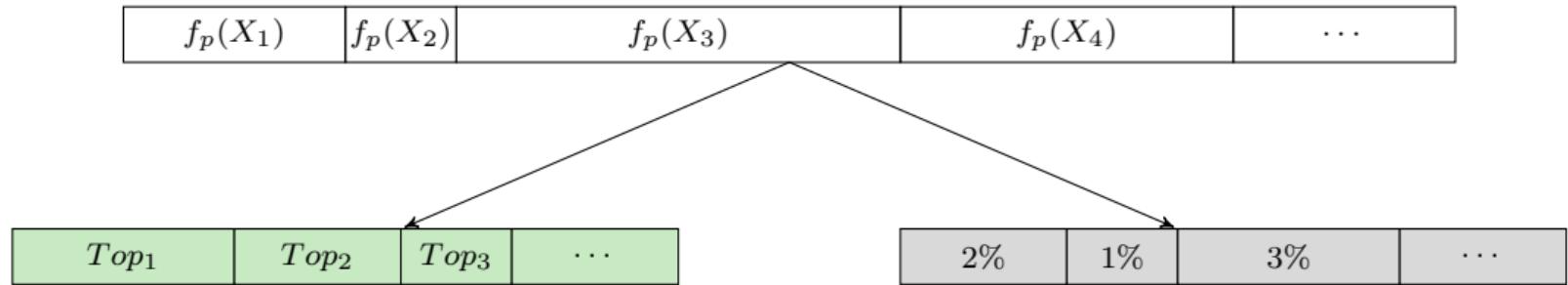
$$f_p(X) = 1.50 \quad (5)$$

$$+ (2.375 + 0.875 + (2.375 + 0.375 - |2.375 - 0.375|)) * \frac{1}{2} \quad (6)$$

$$+ (1.9375 + (1.9375 + 0.6875 - |1.9375 - 0.6875|)) * \frac{1}{4} \quad (7)$$

# Potential interest and sampling

Unexplored layer of the partially ordered graph containing the pharmacophores.



Prioritized area (Drawing ' $Top - k$ ' patterns)

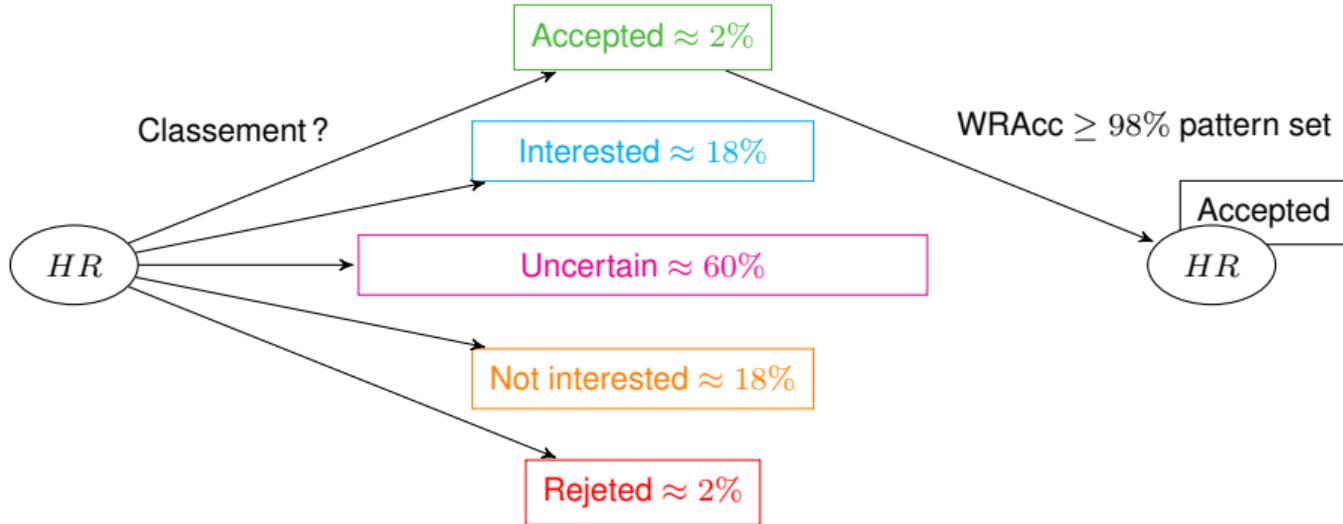
Unbiased area (Drawing ' $Random - d$ ' patterns)

- ▶ Prioritized area : we select patterns with the **highest potential interest**.
- ▶ Unbiased area : potential interest is converted into a **selection probability**.

# Plan de l'exposé

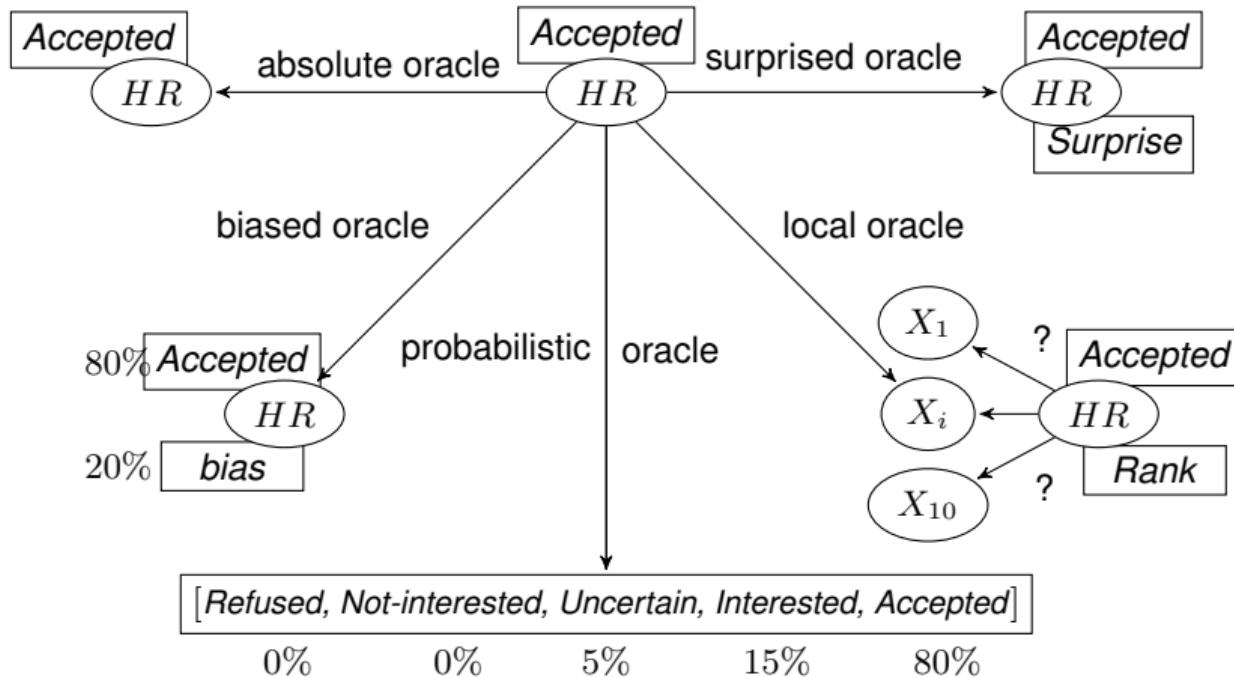
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# Oracles definition (1)



- ▶ The *HR* WRAcc is compared to the WRAcc of the studied pharmacophore set.

## Oracles definition (2)



# Recall graphs

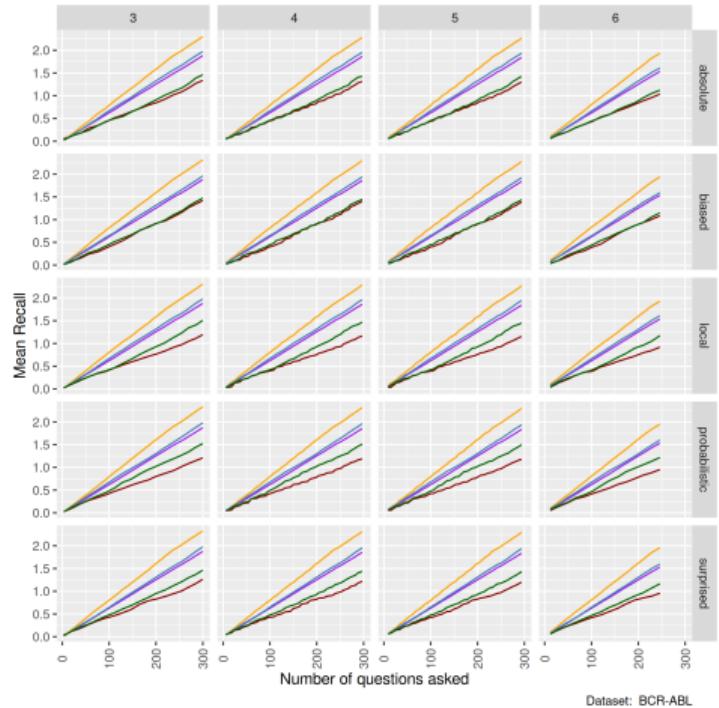


FIGURE – Random Sampling

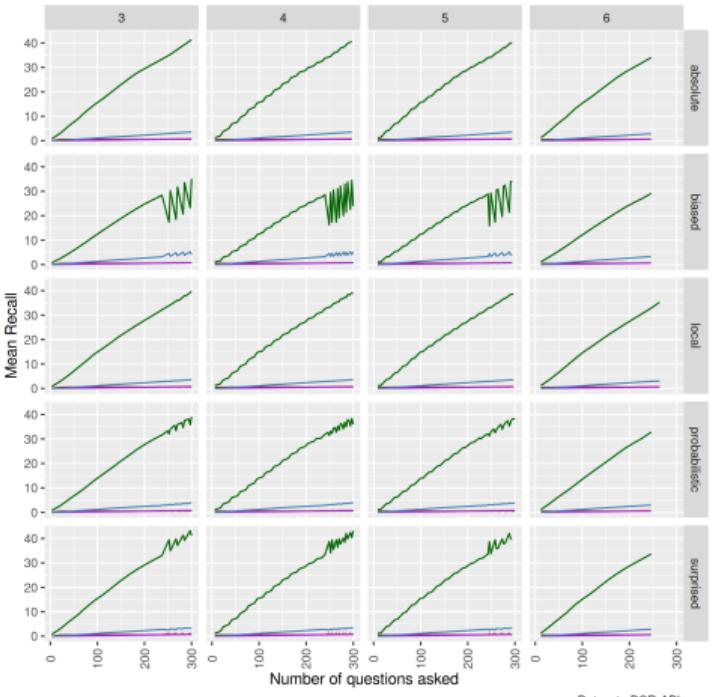
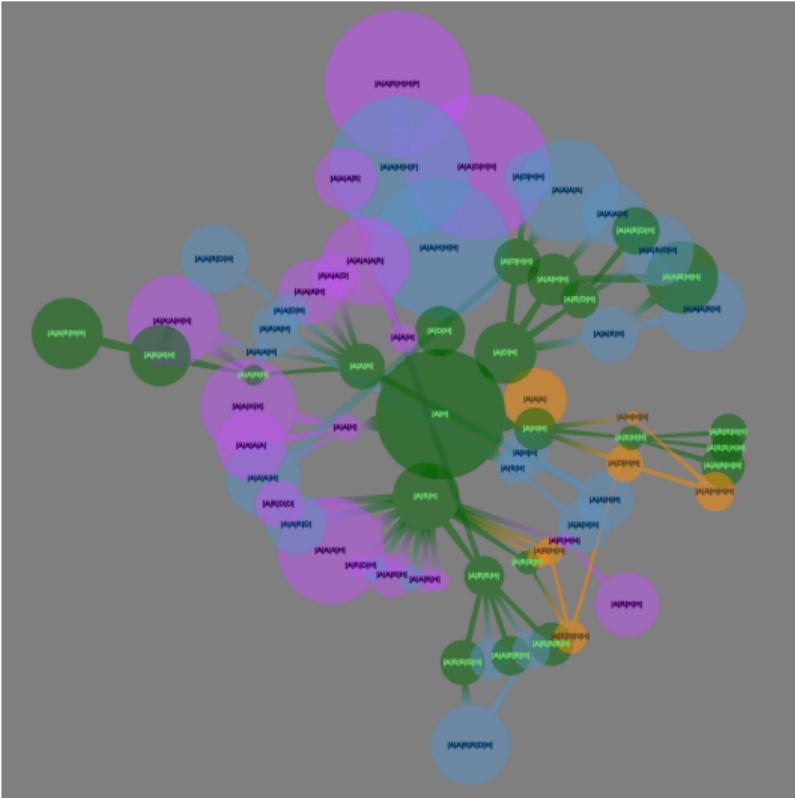


FIGURE – WTRFS

# Interests' partially ordered graph



# Conclusion and future works

## Conclusion

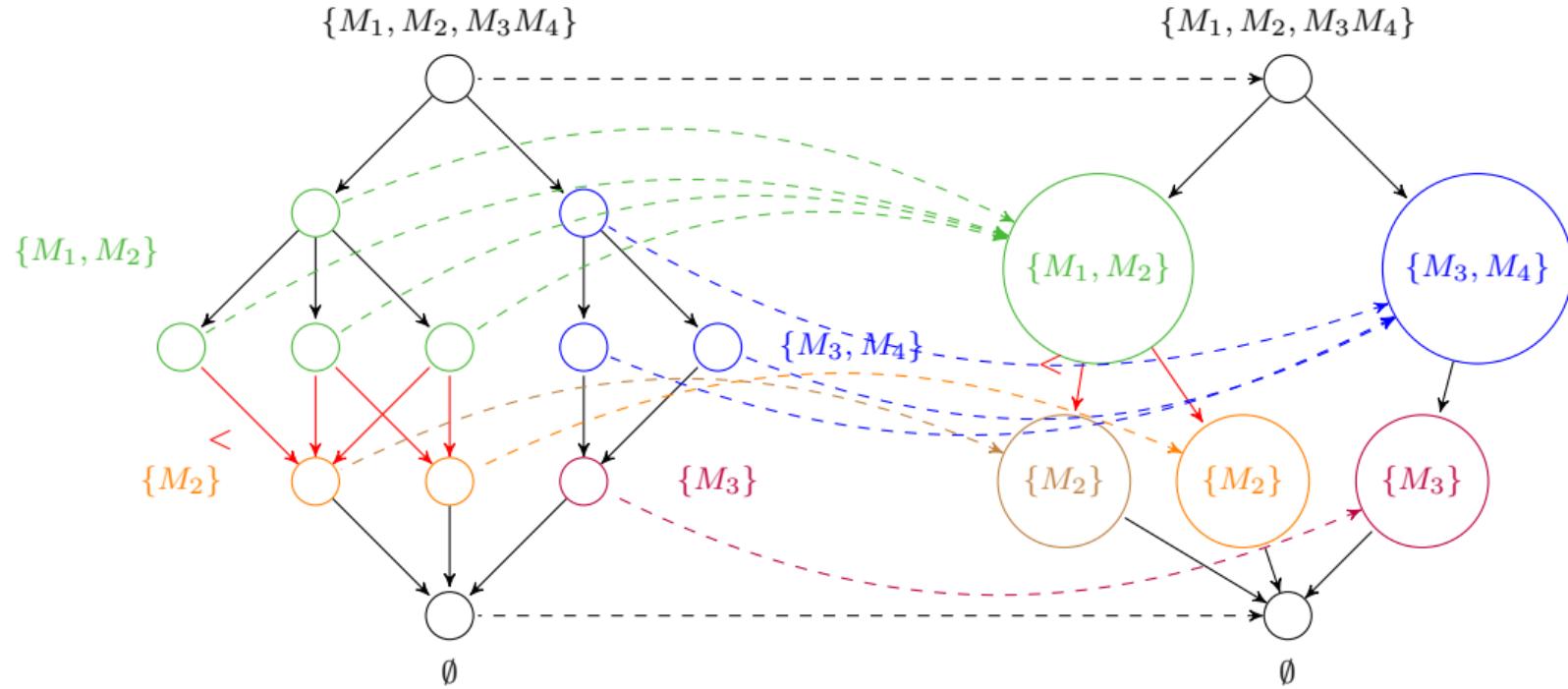
- ▶ Creation of the Interests' partial order graph.
- ▶ Interactive exploration without descriptor.
- ▶ New ideas for the evaluation process.

## Future works

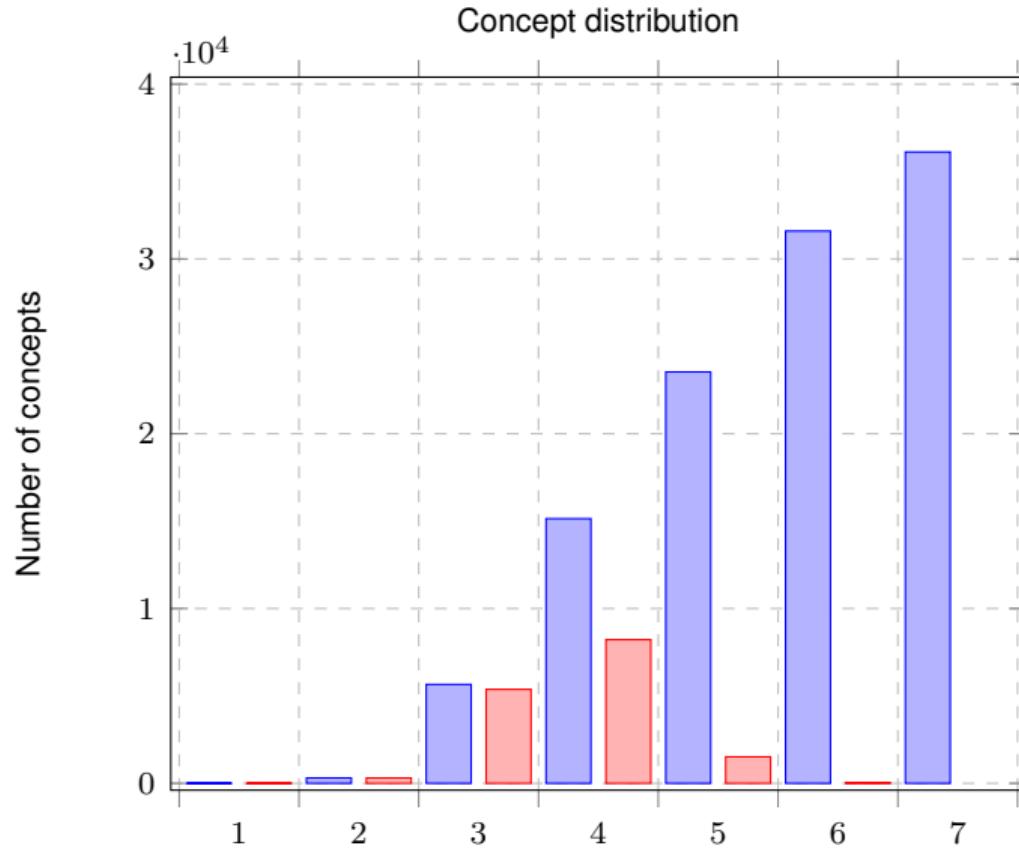
- ▶ Working on the visual representation (GUI and more).
- ▶ Adapting the algorithm to a mining task.
- ▶ Creation of new heuristics.

# Annexes

# Structured Equivalence Classes



# BCR-ABL : patterns distribution by layers



# Case study : selecting an outstanding pattern

