

# M-MODULES AND ROBINSON DISSIMILARITIES

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# Dissimilarity

## Definition

$(X, d)$  **dissimilarity space**:

- ▶ for all  $x \in X$ ,  $d(x, x) = 0$ ,
- ▶ for all  $x, y \in X$  distinct,  $d(x, y) = d(y, x) > 0$ .

$d(x, y)$ : the **dissimilarity** or **distance** between  $x$  and  $y$ ; how much  $x$  and  $y$  are dissimilar.

No triangular inequality ( $d(x, y) + d(y, z) \geq d(x, z)$ ).

Equivalent to nonnegative symmetric square matrices (rows and columns indexed by  $X$ ) with zeros exactly on the diagonal.

# Robinson matrix

## Definition

$M$  **Robinson matrix**:  $M$  is nonnegative symmetric square with zeros exactly on the diagonal, and each row (or column) is bitonic:

$$m_{i,1} \geq m_{i,2} \geq \dots \geq m_{i,i} = 0$$

$$0 = m_{i,i} \leq m_{i,i+1} \leq \dots \leq m_{i,n}$$

$$\begin{pmatrix} 0 & 1 & 4 & 5 & 5 & 6 \\ 1 & 0 & 2 & 4 & 5 & 5 \\ 4 & 2 & 0 & 3 & 3 & 4 \\ 5 & 4 & 3 & 0 & 1 & 2 \\ 5 & 5 & 3 & 1 & 0 & 1 \\ 6 & 5 & 4 & 2 & 1 & 0 \end{pmatrix}$$

# Robinson space

A dissimilarity space  $(X, d)$  is **Robinson** if there is a permutation on  $X$  such that the associated matrix, with rows and columns ordered along that permutation, is Robinson.

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>
<i>a</i>	0	5	1	4	2	5
<i>b</i>	5	0	6	2	4	1
<i>c</i>	1	6	0	5	4	5
<i>d</i>	4	2	5	0	3	1
<i>e</i>	2	4	4	3	0	3
<i>f</i>	5	1	5	1	3	0

	<i>c</i>	<i>a</i>	<i>e</i>	<i>d</i>	<i>f</i>	<i>b</i>
<i>c</i>	0	1	4	5	5	6
<i>a</i>	1	0	2	4	5	5
<i>e</i>	4	2	0	3	3	4
<i>d</i>	5	4	3	0	1	2
<i>f</i>	5	5	3	1	0	1
<i>b</i>	6	5	4	2	1	0

# Decision problem

## Question

Given a dissimilarity space  $(X, d)$  with  $X$  finite, decide whether  $(X, d)$  is Robinson.

## Definition

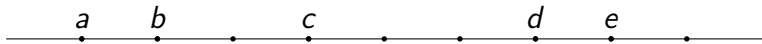
**Compatible order:** order for the rows and columns which makes the matrix Robinson.

## Question

Given a Robinson space, find a (all) compatible order(s).

# Example: line distances

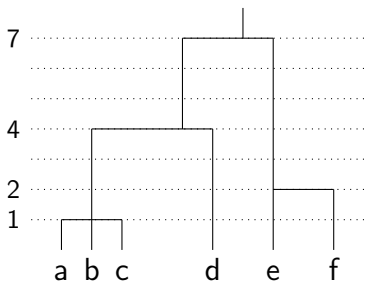
**Line distances:** take  $X \subset \mathbb{R}$ , and  $d(x, y) = |x - y|$ . The (restriction of the) usual order  $<$  on  $\mathbb{R}$  is a compatible order.



	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>
<i>a</i>	0	1	3	6	7
<i>b</i>	1	0	2	5	6
<i>c</i>	3	2	0	3	4
<i>d</i>	6	5	3	0	1
<i>e</i>	7	6	4	1	0

# Example: ultrametrics

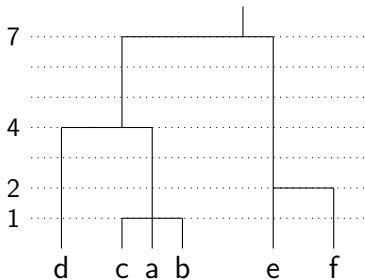
**Ultrametrics:** take  $X$  the leaves of a tree, where all leaves are at the same depth. Let  $d(x, y)$  be the height of the least common ancestor of  $x$  and  $y$ . A left-to-right ordering of the leaves is a compatible order.



	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>
<i>a</i>	0	1	1	4	7	7
<i>b</i>	1	0	1	4	7	7
<i>c</i>	1	1	0	4	7	7
<i>d</i>	4	4	4	0	7	7
<i>e</i>	7	7	7	7	0	2
<i>f</i>	7	7	7	7	2	0

# Compatible orders in ultrametrics

- ▶ Any reordering of the children of each node of the tree gives a different compatible order.
- ▶ Actually, any compatible order is obtainable like this.
- ▶ Thus the tree encodes the set of compatible orders.



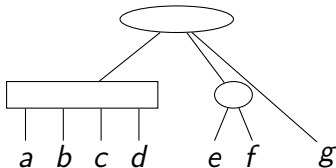
	<i>d</i>	<i>c</i>	<i>a</i>	<i>b</i>	<i>e</i>	<i>f</i>
<i>d</i>	0	4	4	4	7	7
<i>c</i>	4	0	1	1	7	7
<i>b</i>	4	1	0	1	7	7
<i>a</i>	4	1	1	0	7	7
<i>e</i>	7	7	7	7	0	2
<i>f</i>	7	7	7	7	2	0



# PQ-trees

## Definition

**PQ-tree on  $X$** : tree with leaves  $X$ , and each internal node is either a **P-node** (Permutation-node, circle), or a **Q-node** (rectangle).



# Represented orders

**Equivalence on PQ-trees.** PQ-trees are **equivalent**:

- ▶ by reordering *arbitrarily* the order of children of P-nodes (any permutation),

$$P(\alpha_1, \dots, \alpha_k) \leftrightarrow P(\alpha_{\sigma(1)}, \dots, \alpha_{\sigma(k)})$$

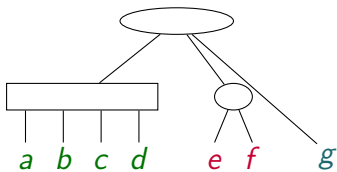
- ▶ by reversing the order of children of Q-nodes.

$$Q(\alpha_1, \alpha_2, \dots, \alpha_k) \leftrightarrow Q(\alpha_k, \dots, \alpha_2, \alpha_1)$$

## Definition

An order on  $X$  is **represented** by a PQ-tree  $\mathcal{T}$  if it is the left-to-right order of leaves of a PQ-tree equivalent to  $\mathcal{T}$ .

# Represented orders (example)

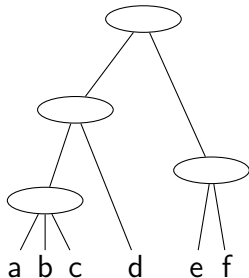
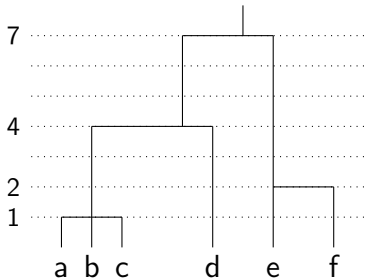


24 represented orders:

- ▶ *abcdefg*,
- ▶ *dcbaefg*,
- ▶ *efgdcba*,
- ▶ *gabcdfe*,
- ▶ *fedcbag*,
- ▶ *abcdgfe*,...

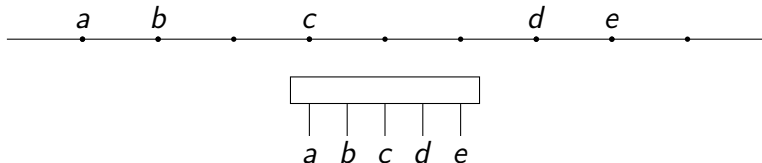
# Compatible orders of an ultrametric

For an ultrametric, making each node a P-node gives a representation of the set of compatible orders.



# Compatible order of a line distance

For a line distance, having a single Q-node, with children in the same order as on the line gives a representation of the set of compatible orders.



# The consecutive-ones property

Let  $M$  be a  $\{0, 1\}$ -matrix.

## Question

Can we permute the columns of  $M$ , such that in each row, the 1s are consecutive?

$$\begin{pmatrix} 0 & 1 & 1 & 1 & 0 & 0 \\ 1 & 1 & 1 & 0 & 0 & 0 \\ 0 & 0 & 1 & 1 & 1 & 1 \\ 1 & 1 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 1 & 1 & 1 \end{pmatrix}$$

# The consecutive-ones property

## Theorem (Booth and Lueker)

*The set of permutations for which the 1s are consecutive is either empty, or representable by a PQ-tree.*

Algorithm:

1. start with  $\mathcal{T} := P(1, \dots, n)$ ,
2. for each row, add to  $\mathcal{T}$  the constraint that the 1s of that row must be consecutive (or fail).

# Applying Booth and Lueker to Robinson

## Definition

The **ball** with center  $x \in X$  and radius  $r \in \mathbb{N}$  is

$$B(x, r) := \{y \in X : d(x, y) \leq r\}.$$

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>
<i>a</i>	0	1	3	4	6	8
<i>b</i>	1	0	1	4	4	7
<i>c</i>	3	1	0	1	2	6
<i>d</i>	4	4	2	0	1	4
<i>e</i>	6	4	2	1	0	2
<i>f</i>	8	7	6	4	2	0

<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>
0	0	1	0	0	0
0	1	1	1	0	0
0	1	1	1	1	0
1	1	1	1	1	0
1	1	1	1	1	1

Balls with **center c**



# Applying Booth and Lueker to Robinson

## Theorem (Mirkin and Rodin)

*A dissimilarity space is Robinson if and only if the incidence matrix of balls has the consecutive-ones property.*

Furthermore, compatible orders are exactly those orders where the 1s are consecutive.

## Theorem

*The set of compatible orders of a Robinson space is representable by a PQ-tree.*

Also, gives a polynomial-time algorithm.

# Deciding whether a dissimilarity is Robinson.

Let  $n = |X|$ .

- ▶ Mirkin and Rodin, 1984:  $O(n^4)$ ,
- ▶ Chepoi and Fichet, 1997, divide-and-conquer:  $O(n^3)$ ,
- ▶ Atkins, Boman, Hendrickson, 1998, spectral method:  $O(nT(n) + n^2 \log n)$ ,
- ▶ Seston, 2008, threshold graphs:  $O(n^2 \log n)$ ,
- ▶ Fortin and Pr ea, 2014, PQ-trees:  $O(n^2)$ ,
- ▶ Laurent and Seminaroti, 2017, LexBFS:  $O(n^2 + nm \log n)$ .

$O(n^2)$  is optimal (size of the input).

# Goals

- ▶ To find a simpler  $O(n^2)$  algorithm, efficient in practice, that do not use Booth and Lueker algorithm,
- ▶ To study the correspondance between PQ-trees and mmodule trees (and ultrametrics).

This talk:

1. Introduction to Robinson spaces and PQ-trees (done),
2. Mmodules and their relations to PQ-trees,
3. Flat Robinson spaces.

# Mmodules

## Definition

**Mmodule**  $M \subseteq X$ : for each  $x, y \in M$  and each  $z \notin M$ ,  $d(x, z) = d(y, z)$ .

An mmodule is a set of elements indistinguishable from elements outside the set. Example:  $\{d, e\}$ .

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>
<i>a</i>	0	1	3	4	4
<i>b</i>		0	2	4	4
<i>c</i>			0	2	2
<i>d</i>				0	1
<i>e</i>					0

# Mmodules

- ▶  $X$  is an mmodule, so is any one-element set and  $\emptyset$  (trivial mmodules).
- ▶ Reminiscent of modules in graph theory. Mmodule = metric-module or matrix-module.
- ▶ Maximal modules in graphs can be computed with a partition refinement technique.
- ▶ Known as *clans* in symmetric 2-structure (Erhenfeucht and Rozenberg).

# Some properties of mmodules

## Lemma

Let  $M_1, M_2$  be mmodules, then

- (i)  $M_1 \cap M_2$  is an mmodule,
- (ii) if  $M_1 \cap M_2 \neq \emptyset$ , then  $M_1 \cup M_2$  is an mmodule,
- (iii) if  $M_1 \cap M_2 = \emptyset$ , then  $d(x, y)$  is constant for  $x \in M_1$ ,  $y \in M_2$ .

## Lemma

Let  $M_1, M_2$  be distinct maximal mmodules (maximal by inclusion distinct from  $X$ ), then

- (i) if  $M_1 \cap M_2 \neq \emptyset$ ,  $M_1 \cup M_2 = X$ ,
- (ii) if  $M$  is an mmodule contained in  $M_1 \cup M_2$ , then  $M \subseteq M_1$  or  $M \subseteq M_2$ .

# Partitions and copartition

## Lemma

*The maximal modules  $\mathcal{M}_{\max}$  are either a partition of  $X$ , or their complements are a partition of  $X$  (that is, they are a copartition of  $X$ ).*

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>
<i>a</i>	0	1	2	2	4	4
<i>b</i>		0	2	2	4	4
<i>c</i>			0	1	3	3
<i>d</i>				0	3	3
<i>e</i>					0	2
<i>f</i>						0

$\mathcal{M}_{\max}$ : ab, cd, ef

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>
<i>a</i>	0	2	4	4	4	4
<i>b</i>		0	4	4	4	4
<i>c</i>			0	2	4	4
<i>d</i>				0	4	4
<i>e</i>					0	3
<i>f</i>						0

$\mathcal{M}_{\max}$ : abcd, cdef, abef

# The mmodule tree

## Lemma

There is a unique tree, the *mmodule tree*, with leaves  $X$ , and inner nodes labelled  $\cup$  and  $\cap$ , such that

- (i) if a node  $\alpha$  is a  $\cup$ -node, its arity is at least 3, and for any child  $\beta$  of  $\alpha$ ,  $X(\beta)$  is an mmodule (partition case);
- (ii) if a node  $\alpha$  is a  $\cap$ -node, its arity is at least 2, and for any children  $\beta_1, \dots, \beta_k$  of  $\alpha$ ,  $X(\beta_1), \dots, X(\beta_k)$  is an mmodule (copartition case);
- (iii) any proper mmodule appears exactly once as in (i) or (ii).

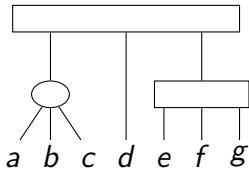
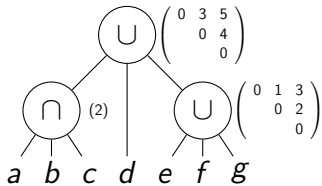
$X(\beta)$ : set of leaves with ancestor  $\beta$ .

This holds for any dissimilarity space (not just Robinson).

The order of childrens does not matter.



# Example of mmodule tree



	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>	<i>g</i>
<i>a</i>	0	2	2	3	5	5	5
<i>b</i>		0	2	3	5	5	5
<i>c</i>			0	3	5	5	5
<i>d</i>				0	4	4	4
<i>e</i>					0	1	3
<i>f</i>						0	2
<i>g</i>							0

# Mmodule tree and PQ-tree

Erhenfeucht, Gabow, MacConnell, Sullivan 1994:  
 $O(|X|^2)$ -time algorithm to build the mmodule tree.

## Question

For Robinson spaces, are the mmodule tree and PQ-tree identical? Or at least can we build the PQ-tree from the mmodule tree?

At least, **the order of children of Q-nodes matters**, while the order of children of  $\cap$ -nodes does not.

## Question

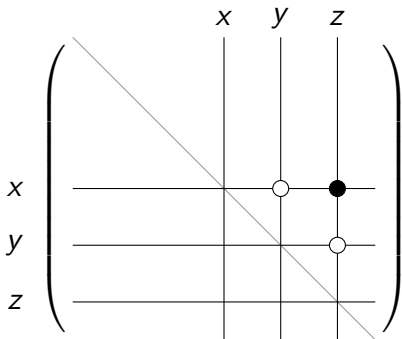
When restricted to a Robinson dissimilarity whose PQ-tree is a single Q-node, can we find the compatible order efficiently?

# An alternative definition for Robinson

## Lemma

$(X, d)$  is a Robinson space if and only if there is an order  $<$  such that for any  $x < y < z$ ,

$$\max\{d(x, y), d(y, z)\} \leq d(x, z).$$

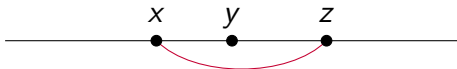


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$$\max\{d(x, y), d(y, z)\} \leq d(x, z).$$



# Block

## Definition

**Block** of a set of permutations/orders on  $X$ : subset  $B \subseteq X$  such that the elements of  $B$  are consecutive (an **interval**) in any of these permutations.

## Lemma

*Given a PQ-tree on  $X$  with block  $B$ ,*

- (i) either there is a node  $\alpha$  with  $B = X(\alpha)$ ,*
- (ii) or there is a Q-node  $\alpha = Q(\beta_1, \dots, \beta_k)$  such that  $B = X(\beta_i) \cup X(\beta_{i+1}) \cup \dots \cup X(\beta_j)$ .*

# PQ-nodes are mmodules

## Lemma

*Let  $\alpha$  be a node of the PQ-tree for  $(X, d)$ , then  $X(\alpha)$  is an mmodule.*

**Proof.** Let  $x, y \in X(\alpha)$ ,  $z \notin X(\alpha)$ , and  $<$  compatible order with  $x < y < z$ . Then  $d(y, z) \leq d(x, z)$ .

Reversing the order of  $X(\alpha)$  in  $<$  gives a compatible order  $<'$  with  $y <' x <' z$ . Then  $d(x, z) \leq d(y, z)$ .

Thus  $d(x, z) = d(y, z)$ .

# Characterization of PQ-nodes

## Theorem

*$M \subseteq X$  is a block and an mmodule iff there is a node  $\alpha$  in the PQ-tree such that  $M = X(\alpha)$ .*

**Proof:** show that for  $Q(\beta_1, \dots, \beta_k)$ , for  $i < j$  with  $(i, j) \neq (1, k)$ ,  $X(\beta_i) \cup \dots \cup X(\beta_j)$  is not an mmodule.

# Flat Robinson spaces

## Definition

**Flat Robinson space:** a Robinson space having only two compatible orders (reverse from each other).

Example: line distances are flat.

Flat Robinson space have PQ-tree reduced to a single internal node of type Q.

## Corollary

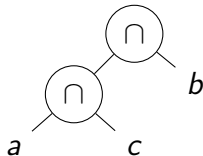
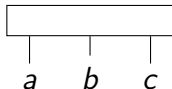
*If all the modules are trivial ( $X$  and one-element sets), then  $(X, d)$  is flat and its PQ-tree has a single node of type Q.*



# Flat Robinson spaces have single node?

Is the converse true? Are the mmodules of flat Robinson spaces always trivial? **No!**

D	a	b	c
a	0	1	2
b		0	1
c			0



## Consequence

PQ-tree and mmodule tree are not similar!

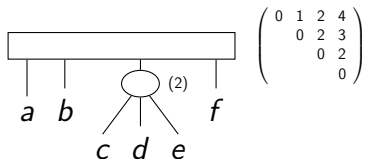
# Conical node

## Definition

**conical node:** a Q-node  $Q(\beta_1, \dots, \beta_k)$  with a (unique) child  $\beta_i$  and  $\delta$  such that  $d(\beta_i, \beta_j) = \delta$  for each  $j \neq i$ .

**apex child:** the child  $\beta_i$  in that case.

**split child:**  $\beta_i$  when it is a P-node with associated value  $\delta$ .



	a	b	c	d	e	f
a	0	1	2	2	2	4
b		0	2	2	2	3
c			0	2	2	2
d				0	2	2
e					0	2
f						0

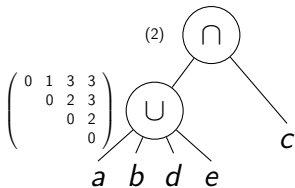
# Conical nodes and flat Robinson spaces

## Lemma

If  $(X, d)$  is flat Robinson space,

- (i) either all its mmodule are trivial,
- (ii) or there is  $p \in X$  and  $\delta$  with  $d(p, x) = \delta$  for all  $x \in X \setminus \{p\}$ .  $X \setminus \{p\}$  is the only non-trivial mmodule. Also  $p$  is not in a diametral pair.

In case (ii), the Q-root is conical, with leaf  $p$  apex.



	$a$	$b$	$c$	$d$	$e$
$a$	0	1	2	3	3
$b$		0	2	2	3
$c$			0	2	2
$d$				0	2
$e$					0

# Special $\cap$ -node and large child

## Definition

**Special  $\cap$ -node:** a  $\cap$ -node whose associated value is less than the diameter of one of its child.

**Large child:** the child (unique for Robinson space) of a  $\cap$ -node whose diameter is more than the  $\cap$ -node associated value

Conical  $Q$ -node and special  $\cap$ -node are the only bad cases, for which the correspondance

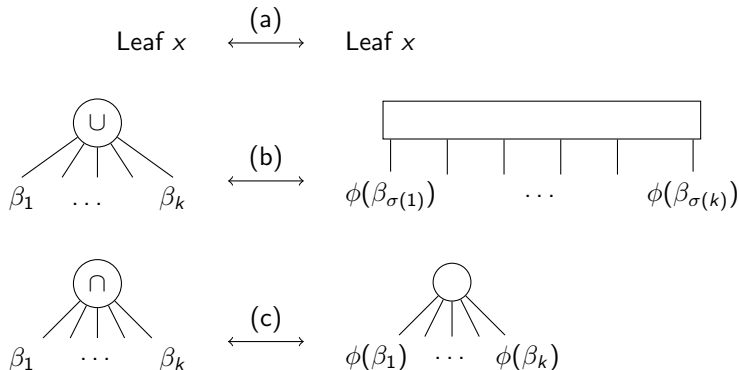
$\cap$ -node  $\leftrightarrow$  P-node

U-node  $\leftrightarrow$  Q-node

does not work.

# From mmodule trees to PQ-trees and back

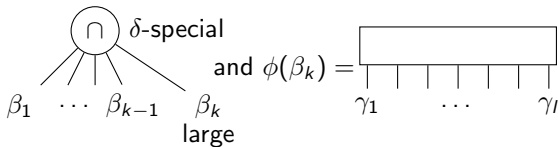
Easy cases (no special node, no conical node):



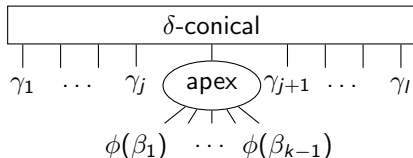
In case (b), it requires  $\sigma$ .

# From mmodule trees to PQ-trees and back

Bad case: special  $\cap$ -node to conical Q-node.

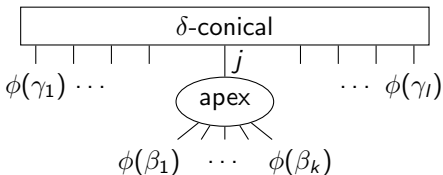


translates into

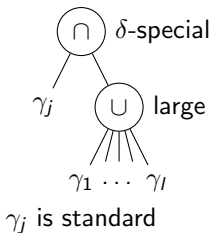


# From mmodule trees to PQ-trees and back

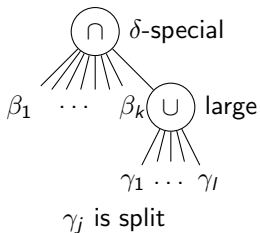
Bad case: conical Q-node to special  $\cap$ -node.



translates into



or



This gives:

## Theorem

*Given a Robinson space, one can build the module tree from the PQ-tree, and the PQ-tree from the module tree in time  $O(|X|)$  (without counting the time spent to order children of Q-nodes).*



# Solving the flat Robinson case

It remains to show how to order the children of Q-node. More generally: find the compatible order (up to reversal) for a flat Robinson space.

1. choose arbitrarily a pivot  $p$ ,
2. sort vertices by their *proximity* to  $p$ ,
3. choose for each vertex its side, left or right of  $p$ .

Step 2 is based on the partition refinement algorithm (used to build the mmodule tree).

# The partition refinement algorithm

(used by Erhenfeucht et al. to build the mmodule tree)

**Input:** a partition  $\mathcal{P}$  of  $X$

**Output:** a partition  $\mathcal{P}'$ , refining  $\mathcal{P}$  (for all  $S' \in \mathcal{P}'$ , there is  $S \in \mathcal{P}$  with  $S' \subseteq S$ ), where each  $S' \in \mathcal{P}'$  is an mmodule.

**Idea:** Keep for each part  $S$  a set of candidate *distinguishers*  $Z_S$ .

**Invariant:** For any  $x, y \in S$ , if there is  $z \in X \setminus S$  with  $d(x, z) \neq d(y, z)$ , then  $z \in Z_S$ .

**Procedure:** Iteratively pick  $z$  in some  $Z_S$ , and refine  $S$  depending on the distances from  $z$ .

# The partition refinement algorithm

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>	<i>g</i>
<i>a</i>	0	2	2	4	4	4	5
<i>b</i>		0	2	4	4	4	5
<i>c</i>			0	3	3	4	4
<i>d</i>				0	1	2	2
<i>e</i>					0	2	2
<i>f</i>						0	2
<i>g</i>							0

$$\{(S, Z_S) : S \in \mathcal{P}\} = \{(abc, defg), (defg, abc)\}$$

# The partition refinement algorithm

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>	<i>g</i>
<i>a</i>	0	2	2	4	4	4	5
<i>b</i>		0	2	4	4	4	5
<i>c</i>			0	3	3	4	4
<i>d</i>				0	1	2	2
<i>e</i>					0	2	2
<i>f</i>						0	2
<i>g</i>							0

$\{(abc, defg), (defg, abc)\}$

$\rightarrow \{(ab, cefg), (c, abefg)(defg, abc)\}$

# The partition refinement algorithm

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>	<i>g</i>
<i>a</i>	0	2	2	4	4	4	5
<i>b</i>		0	2	4	4	4	5
<i>c</i>			0	3	3	4	4
<i>d</i>				0	1	2	2
<i>e</i>					0	2	2
<i>f</i>						0	2
<i>g</i>							0

$\{(ab, cefg), (c, abefg)(defg, abc)\}$

$\rightarrow \{(ab, \emptyset), (c, abefg), (defg, abc)\}$

# The partition refinement algorithm

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>	<i>g</i>
<i>a</i>	0	2	2	4	4	4	5
<i>b</i>		0	2	4	4	4	5
<i>c</i>			0	3	3	4	4
<i>d</i>				0	1	2	2
<i>e</i>					0	2	2
<i>f</i>						0	2
<i>g</i>							0

$\{(ab, \emptyset), (c, \emptyset)(defg, abc)\}$

$\rightarrow \{(ab, \emptyset), (c, \emptyset), (def, gbc), (g, defbc)\}$

# The partition refinement algorithm

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>	<i>g</i>
<i>a</i>	0	2	2	4	4	4	5
<i>b</i>		0	2	4	4	4	5
<i>c</i>			0	3	3	4	4
<i>d</i>				0	1	2	2
<i>e</i>					0	2	2
<i>f</i>						0	2
<i>g</i>							0

$\{(ab, \emptyset), (c, \emptyset), (def, gbc), (g, defbc)\}$

$\longrightarrow \{(ab, \emptyset), (c, \emptyset), (def, c), (g, defbc)\}$

# The partition refinement algorithm

	a	b	c	d	e	f	g
a	0	2	2	4	4	4	5
b		0	2	4	4	4	5
c			0	3	3	4	4
d				0	1	2	2
e					0	2	2
f						0	2
g							0

$\{(ab, \emptyset), (c, \emptyset), (def, c), (g, defbc)\}$

$\longrightarrow \{(ab, \emptyset), (c, \emptyset), (de, f), (f, de)(g, defbc)\}$



# The partition refinement algorithm

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>	<i>g</i>
<i>a</i>	0	2	2	4	4	4	5
<i>b</i>		0	2	4	4	4	5
<i>c</i>			0	3	3	4	4
<i>d</i>				0	1	2	2
<i>e</i>					0	2	2
<i>f</i>						0	2
<i>g</i>							0

$\{(ab, \emptyset), (c, \emptyset), (de, f), (f, de)(g, defbc)\}$

$\{(ab, \emptyset), (c, \emptyset), (de, \emptyset), (f, \emptyset)(g, \emptyset)\}$

# The partition refinement algorithm

	<i>a</i>	<i>b</i>	<i>c</i>	<i>d</i>	<i>e</i>	<i>f</i>	<i>g</i>
<i>a</i>	0	2	2	4	4	4	5
<i>b</i>		0	2	4	4	4	5
<i>c</i>			0	3	3	4	4
<i>d</i>				0	1	2	2
<i>e</i>					0	2	2
<i>f</i>						0	2
<i>g</i>							0

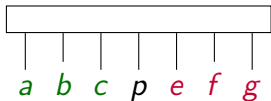
$$\mathcal{P}' = \{ab, c, de, f, g\}$$

# $p$ -proximity order

## Definition

$p$ -proximity order for a compatible order  $<$ : a total order  $\prec$  for some  $p \in X$ , such that:

1.  $p$  is the minimum,
2. for all  $q \in X$ ,  $\{q\} \cup \{x \in X : x \prec q\}$  is an interval of  $<$ .



- ▶  $p \prec c \prec b \prec e \prec a \prec f \prec g$ ,
- ▶  $p \prec e \prec c \prec f \prec b \prec a \prec g$ ,
- ▶  $p \prec e \prec f \prec g \prec c \prec b \prec a$ ,
- ▶  $p \prec c \prec e \prec b \prec f \prec g \prec a, \dots$

# $p$ -proximity order

$p \prec x \prec y$  is equivalent to saying that  $y$  is not between  $p$  and  $x$  in  $\prec$ .

## Lemma

Let  $x, y \in X \setminus \{p\}$ ,  $y$  is not between  $p$  and  $x$  if

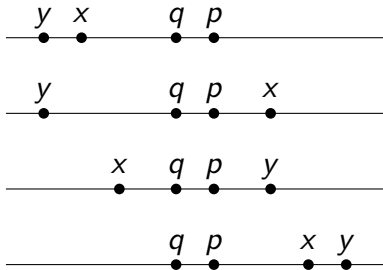
- (i) either  $d(p, x) < d(p, y)$ ;
- (ii) or  $d(p, x) = d(p, y)$  and there is  $q \in X$  with  $q \prec x$ ,  $q \prec y$  and  $d(q, x) < d(q, y)$ ;
- (iii) or  $d(p, x) = d(p, y)$  and there is  $q \in X$  with  $x \prec q$ ,  $y \prec q$  and  $d(y, q) < d(x, q)$ .

Case (ii):  $q$  is an *in-pivot*, case (iii):  $q$  is an *out-pivot*.

# Proof

Case (ii). Let  $x, y \in X \setminus \{p\}$  with  $d(p, x) = d(p, y)$ . Let  $q \in X$  with  $q \prec x$ ,  $q \prec y$  and  $d(q, x) < d(q, y)$ . Assume  $q < p$ .

Possible cases:



In any case,  $y$  is not between  $p$  and  $x$ .

# Computing a $p$ -proximity order

Modify the partition refinement algorithm:

$$(S, Z) \implies (S, \text{In}_S, \text{Out}_S) \text{ with } \text{In}_S \prec S \prec \text{Out}_S$$

When refining  $S$  with in-pivot  $q \in \text{In}_S$ : partition  $S$  into

$$S_1 \cup S_2 \cup \dots \cup S_k \text{ with } d(q, S_1) < d(q, S_2) < \dots < d(q, S_k)$$

Set  $S_1 \prec S_2 \prec \dots \prec S_k$ .

Set  $\text{In}_{S_i} = S_1 \cup \dots \cup S_{i-1} \cup \text{In}_S \setminus \{q\}$ .

Set  $\text{Out}_{S_i} = S_{i+1} \cup \dots \cup S_k \cup \text{Out}_S$ .

Slightly more complicated rules when  $q \in \text{Out}_S$ .

# Computing a $p$ -proximity order

	$a$	$b$	$c$	$p$	$e$	$f$
$a$	0	2	4	5	6	6
$b$		0	2	3	4	6
$c$			0	3	4	6
$p$				0	3	5
$e$					0	1
$f$						0

$$(p, \emptyset, \emptyset) \prec (abcef, p, \emptyset)$$

$$(p, \emptyset, \emptyset) \prec (bce, \emptyset, af) \prec (af, bce, \emptyset)$$

# Computing a $p$ -proximity order

	$a$	$b$	$c$	$p$	$e$	$f$
$a$	0	2	4	5	6	6
$b$		0	2	3	4	6
$c$			0	3	4	6
$p$				0	3	5
$e$					0	1
$f$						0

$$(p, \emptyset, \emptyset) \prec (bce, \emptyset, af) \prec (af, bce, \emptyset)$$

$$(p, \emptyset, \emptyset) \prec (c, \emptyset, bef) \prec (b, c, ef) \prec (e, cb, f) \prec (af, bce, \emptyset)$$



# Computing a $p$ -proximity order

	$a$	$b$	$c$	$p$	$e$	$f$
$a$	0	2	4	5	6	6
$b$		0	2	3	4	6
$c$			0	3	4	6
$p$				0	3	5
$e$					0	1
$f$						0

$$(p, \emptyset, \emptyset) \prec (c, \emptyset, \emptyset) \prec (b, \emptyset, \emptyset) \prec (e, \emptyset, \emptyset) \prec (af, bce, \emptyset)$$

$$(p, \emptyset, \emptyset) \prec (c, \emptyset, \emptyset) \prec (b, \emptyset, \emptyset) \prec (e, \emptyset, \emptyset) \prec (a, ce, f) \prec (f, ace, \emptyset)$$

# Computing a $p$ -proximity order

	$a$	$b$	$c$	$p$	$e$	$f$
$a$	0	2	4	5	6	6
$b$		0	2	3	4	6
$c$			0	3	4	6
$p$				0	3	5
$e$					0	1
$f$						0

$$(p, \emptyset, \emptyset) \prec (c, \emptyset, \emptyset) \prec (b, \emptyset, \emptyset) \prec (e, \emptyset, \emptyset) \prec (a, \emptyset, \emptyset) \prec (f, \emptyset, \emptyset)$$

$$p \prec c \prec b \prec e \prec a \prec f$$

# How to use the $p$ -proximity order

- ▶ Choose  $p$  that is not apex.
- ▶ Refine  $\{\{p\}, X \setminus \{p\}\}$ . Because no module (except possibly  $X \setminus \{\text{apex}\}$ ), the partition contains one-element sets only.
- ▶ Obtain a  $p$ -proximity order  $\prec$ .

$$p \prec e_1 \prec e_2 \prec \dots \prec e_k$$

## Next goal

Partition  $X \setminus \{p\}$  into two sides  $L \cup R$ : elements at the left (resp. right) of  $p$  in  $\prec$ .

$$\prec + L \cup R \implies \prec$$

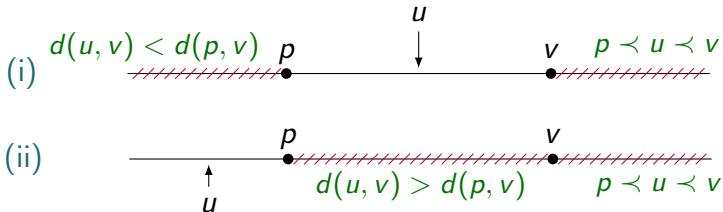
# Choosing side

## Lemma

Let  $\prec$  be a  $p$ -proximity order, and  $u \prec v$ .

- (i)  $d(u, v) < d(p, v)$  implies  $\text{side}(u) = \text{side}(v)$ ,
- (ii)  $d(u, v) > d(p, v)$  implies  $\text{side}(u) \neq \text{side}(v)$ .

## Proof.



# The constraint graph $G$

Define the constraint graph  $G = (V, E)$ :

$$V = X \setminus \{p\}$$

$$E = \{(u, v) : u \prec v \wedge d(u, v) \neq d(p, v)\}$$

If  $G$  has a single connected component: pick a side for an arbitrary  $x \in X \setminus \{p\}$ , propagate to deduce  $L$  and  $R$ .

# The component graph $H$

## Definition

**Tangled components**  $C, C'$ :  $C$  and  $C'$  are not comparable under  $\prec$ .

Define the component graph  $H = (K, F)$ :

$$K = \{C : C \text{ connected component of } G\}$$

$$F = \{(C, C') : C, C' \text{ are tangled}\}$$

# Analysis of tangled components

Suppose  $C, C'$  are tangled. We may assume  $x, z \in C, y \in C'$  with  $xz \in E$  and  $x \prec y \prec z$ .

- ▶  $d(p, y) = d(x, y)$  (as  $xy \notin E$ ),
- ▶  $d(p, z) = d(y, z)$  (as  $yz \notin E$ ).

## Lemma

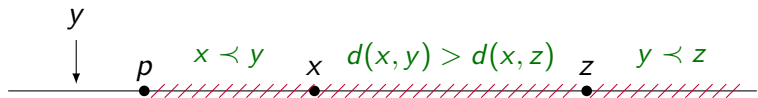
- (i) if  $d(x, z) < d(p, z)$ , then  $\text{side}(x) = \text{side}(z) \neq \text{side}(y)$ ,
- (ii) if  $d(x, z) > d(p, z)$ , then  $\text{side}(x) \neq \text{side}(z) = \text{side}(y)$

Thus if  $H$  has a single connected component: pick a side for an arbitrary  $x \in X \setminus \{p\}$ , propagate to deduce  $L$  and  $R$ .

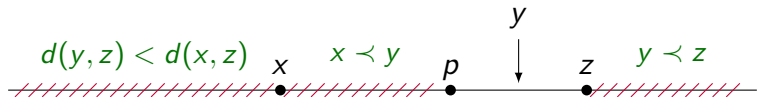
# Tangled lemma proof

We have  $d(x, y) = d(p, y)$  and  $d(y, z) = d(p, z)$  and  $x \prec y \prec z$ .

(i) If  $d(x, z) < d(p, z)$ , then  $\text{side}(x) = \text{side}(z)$ , and  $d(x, z) < d(p, z) = d(y, z)$ .



(ii) If  $d(x, z) > d(p, z)$ , then  $\text{side}(x) \neq \text{side}(z)$ , and  $d(y, z) = d(p, z) < d(x, z)$ .





# Last missing piece

Now we just need:

## Lemma

*H is connected.*

**Proof.** Let  $m$  be the maximum of  $\prec$  and  $M$  be  $p$  plus the vertices not determined by  $\text{side}(m)$ .

Let  $x, y \in M, z \in X \setminus M$ .

$x \prec z$  and  $y \prec z$  (as no entanglement here).

Then  $xz, yz \notin E$  thus  $d(x, z) = d(p, z) = d(y, z)$ .

$M$  is an mmodule.

As  $(X, d)$  is flat (and  $m$  is not apex),  $M = \{p\}$ .

# Algorithmically

```
1: let  $m \in X$ , maximum for  $\prec$ 
2: let  $L = []$ ,  $R = [m]$ ,  $Undecided = \text{reverse}(X \setminus \{p, m\})$ 
3: for  $q \in X \setminus \{p\}$  in decreasing order for  $\prec$  do
4:   let  $Skipped = []$ 
5:   for  $x \in Undecided$  from first to last do
6:     if  $d(x, q) = d(p, q)$  then
7:        $Skipped \leftarrow x \cdot Skipped$ 
8:     else
9:       if  $d(x, q) < d(p, q) \Leftrightarrow q \in L$  then
10:         $L \leftarrow x \cdot L$ ,  $R \leftarrow Skipped \uparrow R$ 
11:       else
12:         $R \leftarrow x \cdot R$ ,  $L \leftarrow Skipped \uparrow L$ 
13:        $Skipped \leftarrow []$ 
14:    $Undecided \leftarrow \text{reverse}(Skipped)$ 
15: return  $\text{reverse}(L) \uparrow [p] \uparrow R$ 
```

# proximity order + side bipartition

Flexible framework:

- ▶ build mmodule tree, translate to PQ-tree using flat Robinson ordering,
- ▶ build ordering of  $p$ -copoints (maximal mmodules not containing  $p$ ), recurse on copoints and build PQ-tree,
- ▶ contract  $p$ -copoints to get a flat Robinson quotient space, merge compatible orders of copoints with order of quotient space.

Available implementation for the last method.

# Open problems

- ▶  $o(n^2)$  with additional information (promise to be Robinson, minimum spanning tree for  $d, \dots$ ),
- ▶ extension to circular Robinson,
- ▶ extension to other topologies than the line,
- ▶ 3D-matrices.