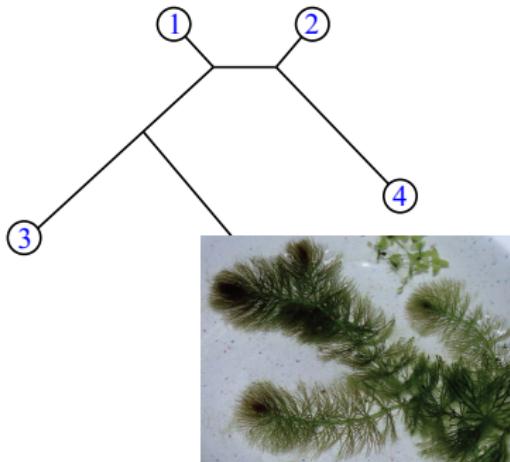
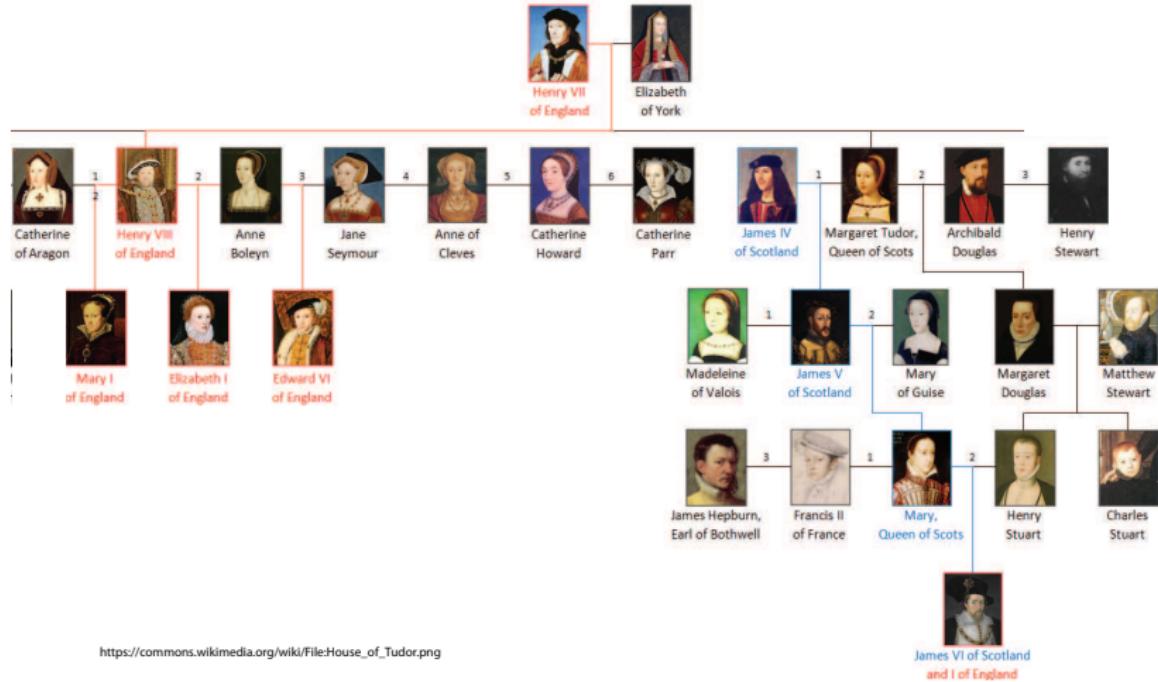


# Clades and tubes: facets of graph associahedra and phylogenetic polytopes.

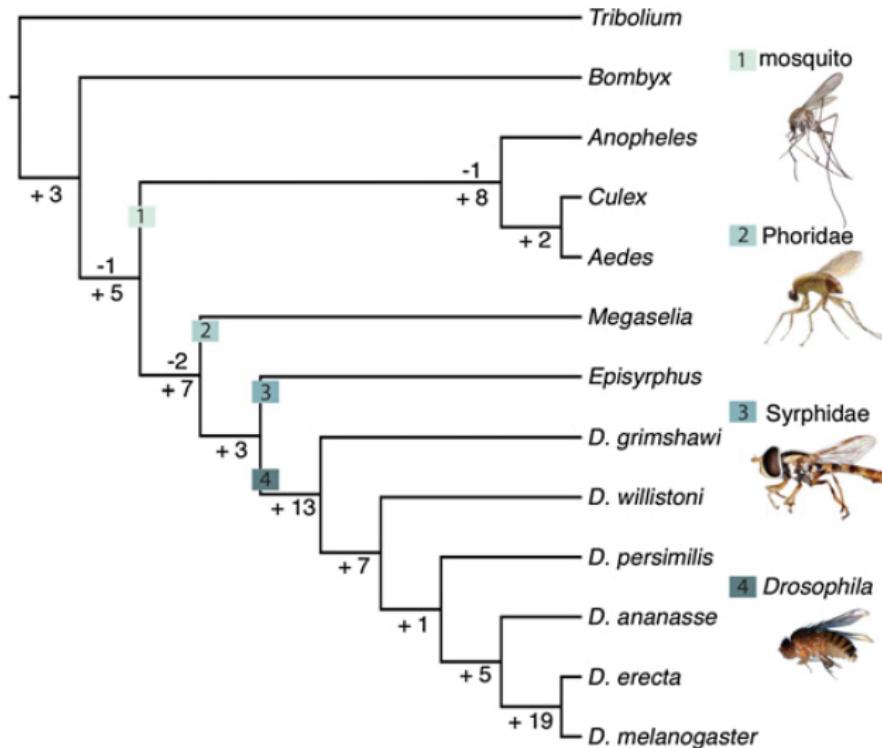
S. Forcey, L. Keefe, W. Sands. U. Akron.  
S. Devadoss. U. San Diego



# Trees

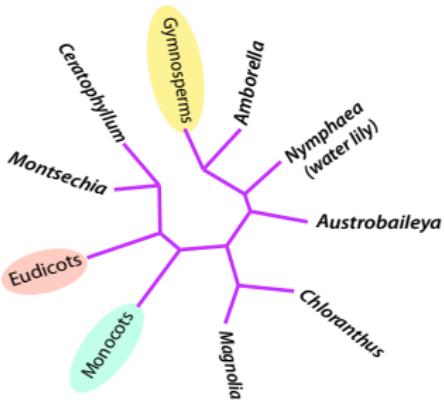
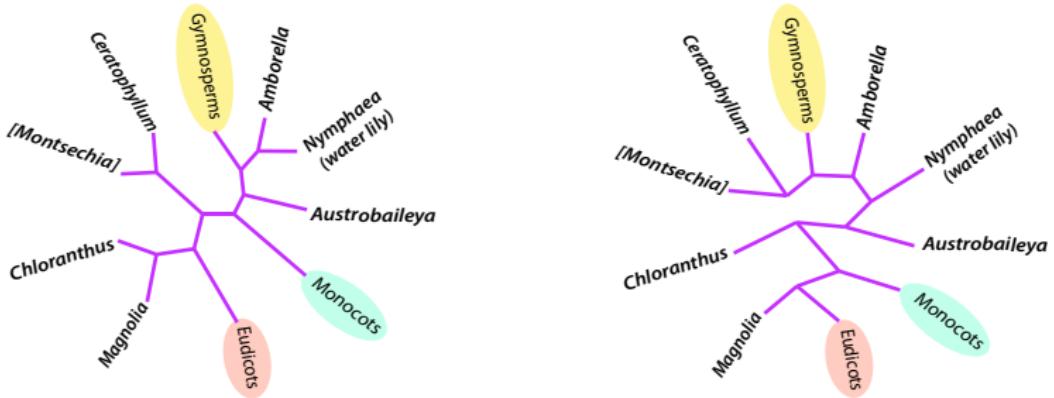


# Trees



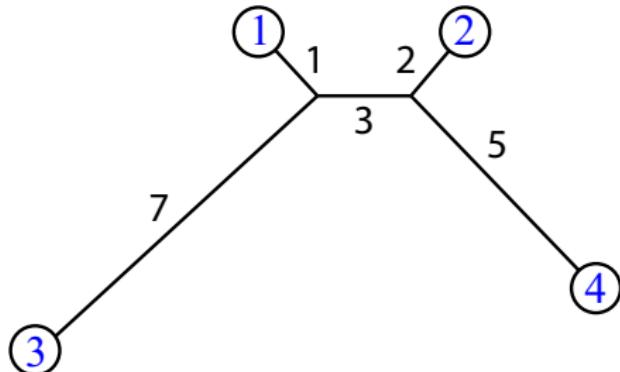
Episodic radiations in the fly tree of life, Wiegmann et.al. PNAS 2011

# Trees

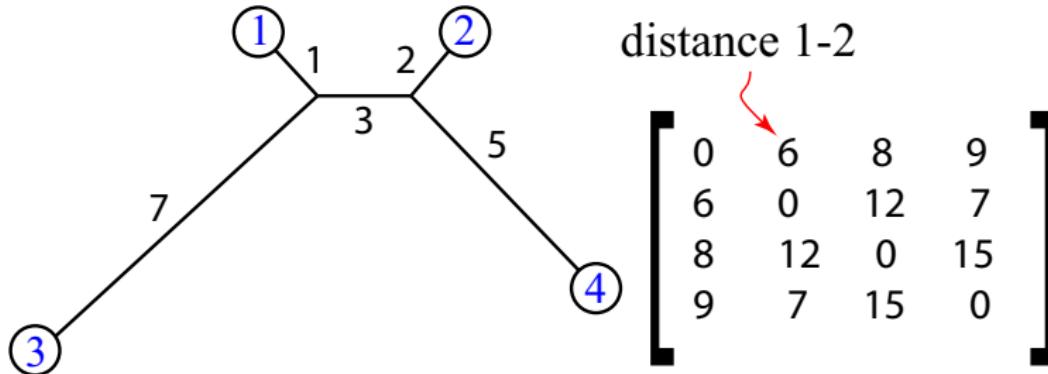


Premalatha Kalagara, CC BY-SA 4.0, <https://commons.wikimedia.org/w/index.php?curid=37953761> Christian Fischer, CC, <https://commons.wikimedia.org/wiki/File:CeratophyllumSubmersum.jpg>  
Josep Renalias, CC, [https://commons.wikimedia.org/wiki/File:Magn%C3%ADa\\_verbenia.JPG](https://commons.wikimedia.org/wiki/File:Magn%C3%ADa_verbenia.JPG) https://commons.wikimedia.org/wiki/File:Amborella\_trichopoda\_(3065968016)\_fragment.jpg

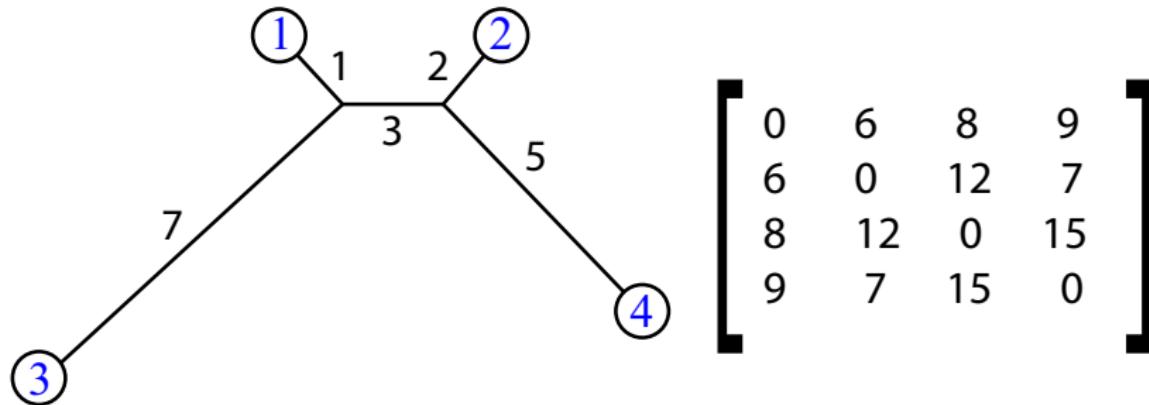
## The Balanced minimal evolution method: ex. tree metric.



# The Balanced minimal evolution method: ex. tree metric.



## The Balanced minimal evolution method: ex. tree metric.



$$\mathbf{d} = \langle 6, 8, 9, 12, 7, 15 \rangle$$

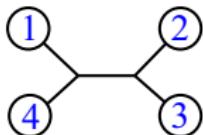
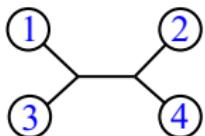
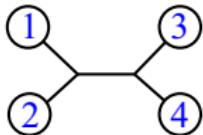
Now: if we are given **d**, (experiment, measurement), can we recover the original tree?

# The Balanced minimal evolution method: ex. tree metric.

$$x(t)_{ij} = 2^{(n-2-l_{ij})}$$

$t$

$x(t)$



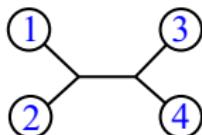
# The Balanced minimal evolution method: ex. tree metric.

$$x(t)_{ij} = 2^{(n-2-l_{ij})}$$

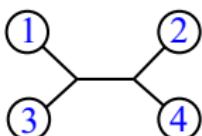
$t$

$x(t)$

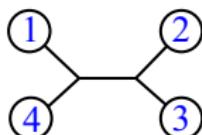
$d \cdot x(t)$



$\langle 2, 1, 1, 1, 1, 2 \rangle$



$\langle 1, 2, 1, 1, 2, 1 \rangle$



$\langle 1, 1, 2, 2, 1, 1 \rangle$

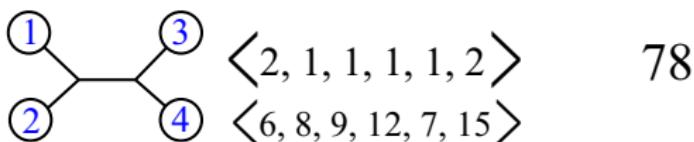
## The Balanced minimal evolution method: ex. tree metric.

$$x(t)_{ij} = 2^{(n-2-l_{ij})}$$

*t*

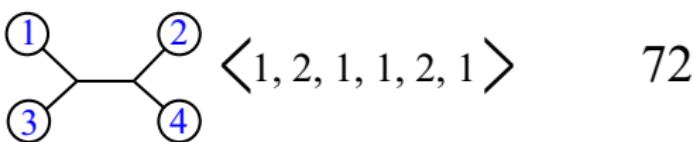
*x(t)*

*d*·*x(t)*



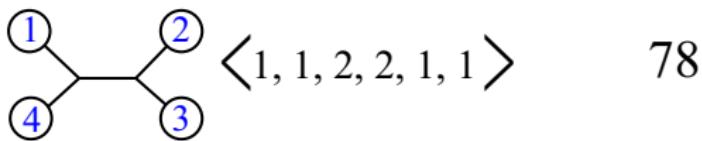
$\langle 2, 1, 1, 1, 1, 2 \rangle$   
 $\langle 6, 8, 9, 12, 7, 15 \rangle$

78



$\langle 1, 2, 1, 1, 2, 1 \rangle$

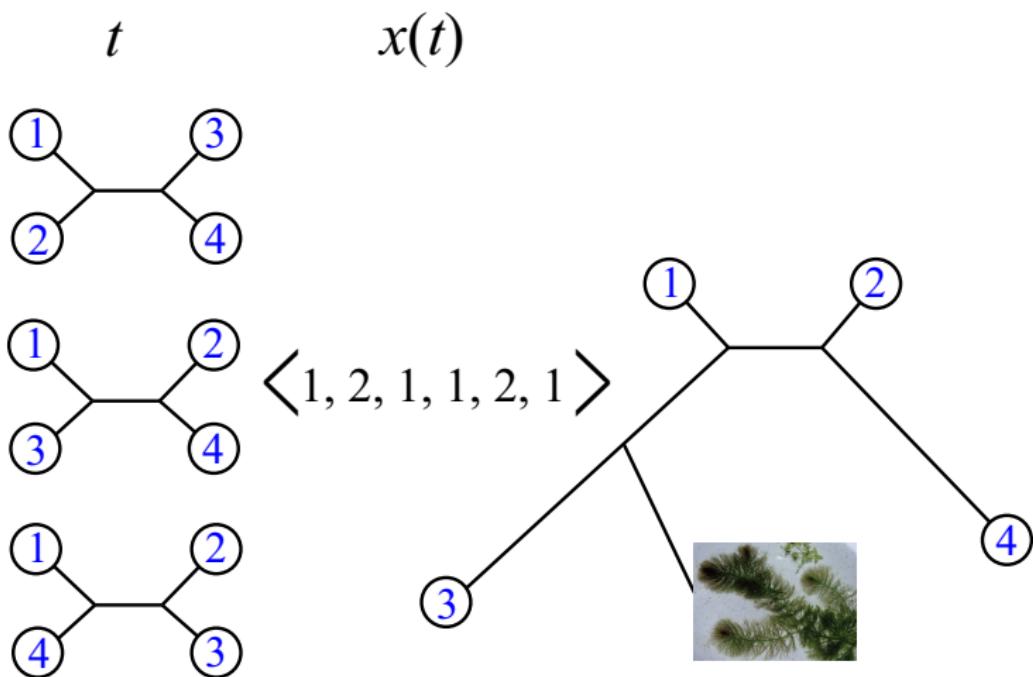
72



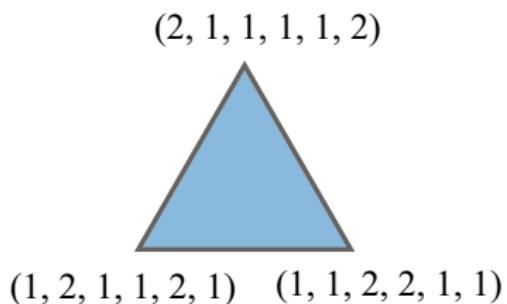
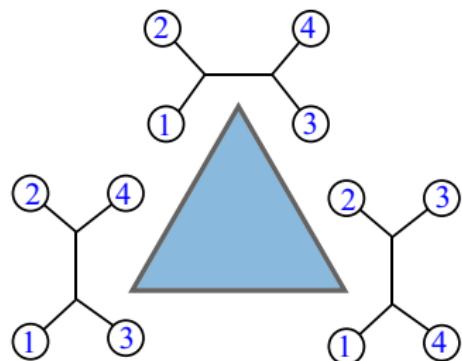
$\langle 1, 1, 2, 2, 1, 1 \rangle$

78

# The Balanced minimal evolution method: ex. tree metric.



# The Balanced minimal evolution polytope $\mathcal{P}_4$ .



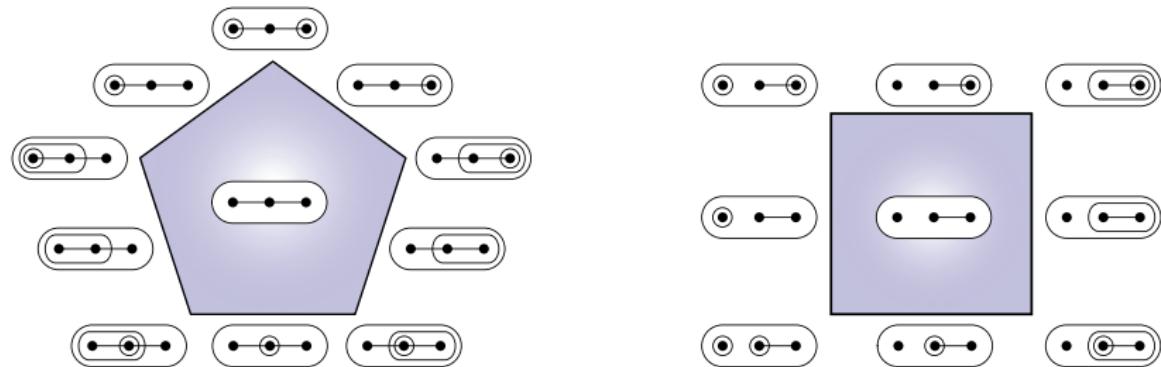
# Statistics.

- Dimensions (start  $n = 3$ ):  $0, 2, 5, 9, 14, \dots, \binom{n}{2} - n$
- Number of Vertices in  $n^{th}$  polytope:  $1, 3, 15, 105, \dots, (2n - 5)!!$
- Number of Facets:  $0, 3, 52, 90262\dots$  OPEN
- $f$ -vectors:  $1, 3, 3, 1, 15, 105, 250, 210, 52, 1, 105, 5460\dots$

## Definitions.

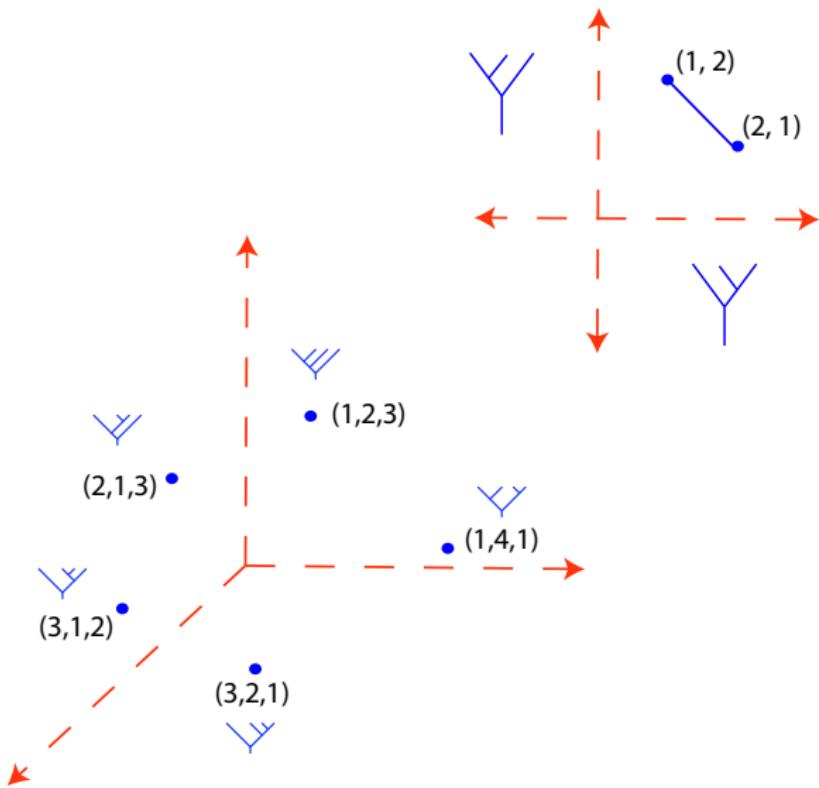
- A *clade* is a sub-tree of a phylogenetic tree which is a connected component after deleting a single interior edge. (It contains all the leaves of a single ancestor, for rooted trees).
- A *cherry* is a clade with only two leaves.
- A pair of *intersecting cherries*  $\{a, b\}$  and  $\{b, c\}$  have intersection in one leaf  $b$ , and thus cannot exist both on the same tree.
- A *caterpillar* is a tree with only two cherries.
- A *split* of the set of  $n$  leaves for our phylogenetic trees is a partition of the leaves into two parts, one part called  $S$  with  $m$  leaves and another with the remaining  $n - m$  leaves. A tree *displays* a split if each part makes up the leaves of a *clade*.
- A *tube* is a connected subgraph. A clade is a specialized tube. A *tubing* is a set of nested or disconnected tubes. Any set of clades on a rooted tree form a tubing.

# Definitions.

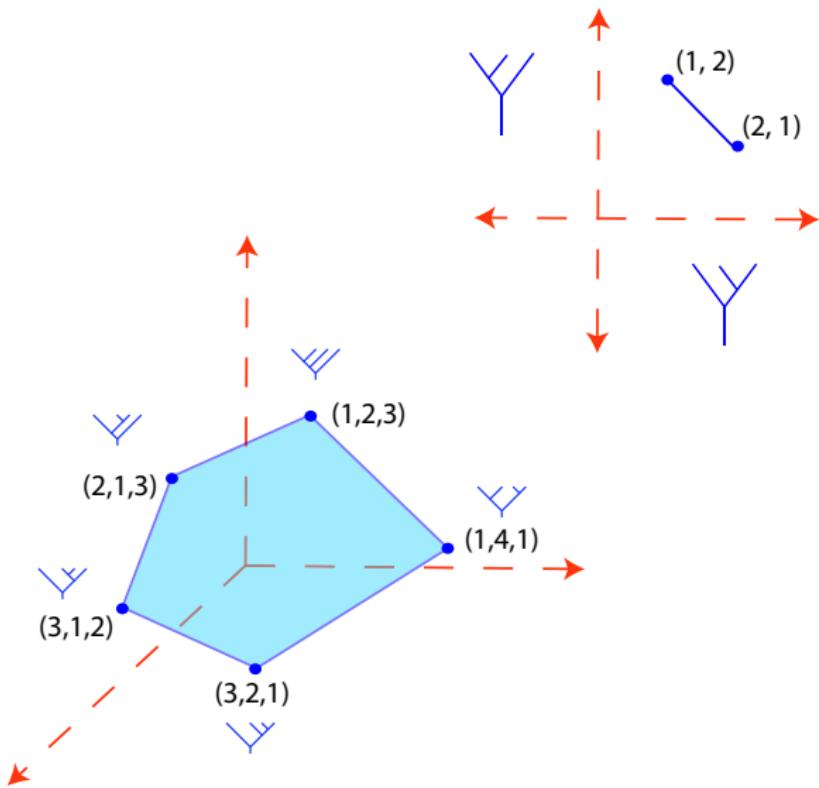


Graph associahedra of a path and a disconnected graph. The pentagon is the classical associahedron, Stasheff's polytope.

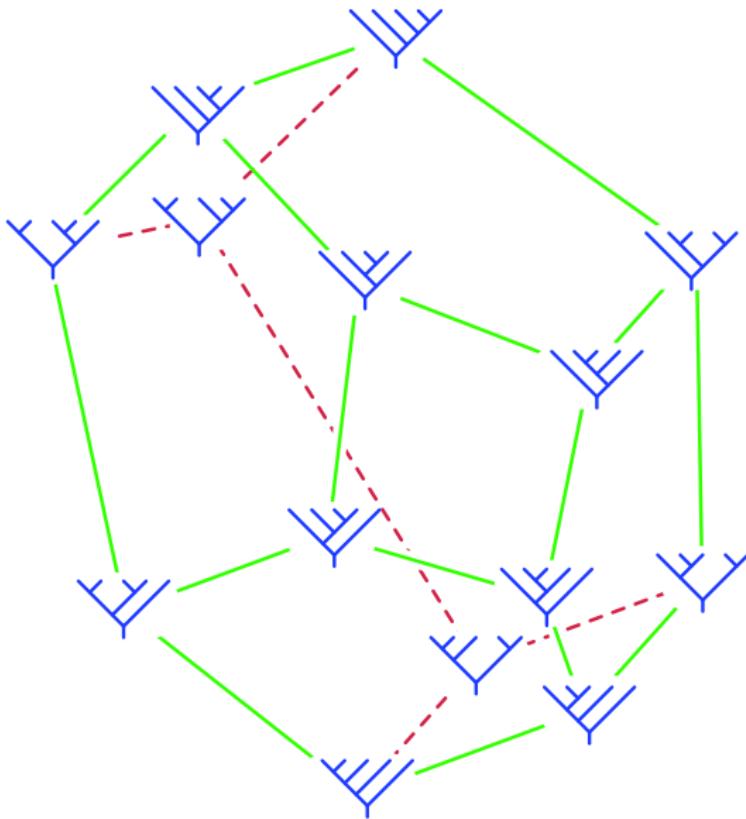
# A convex hull of binary trees.



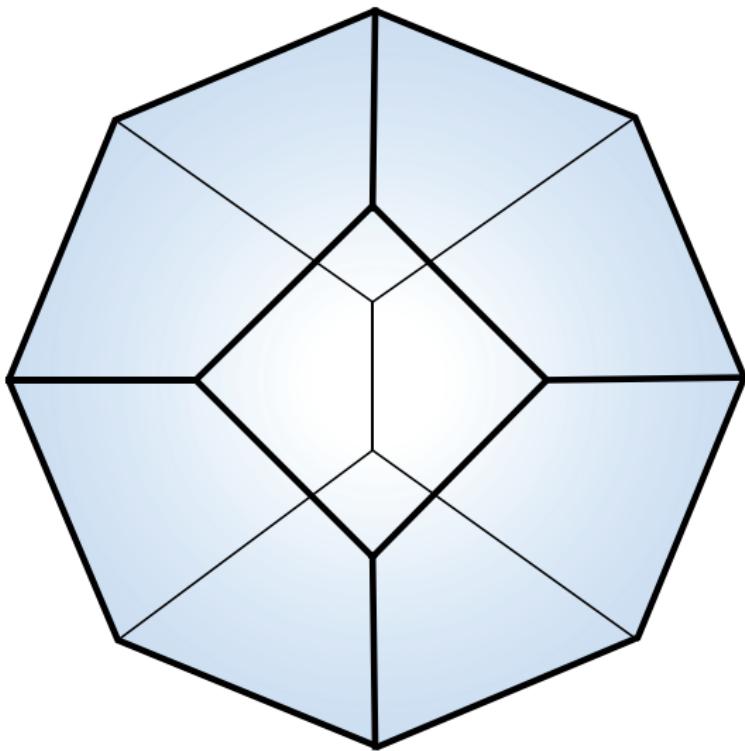
# A convex hull of binary trees.



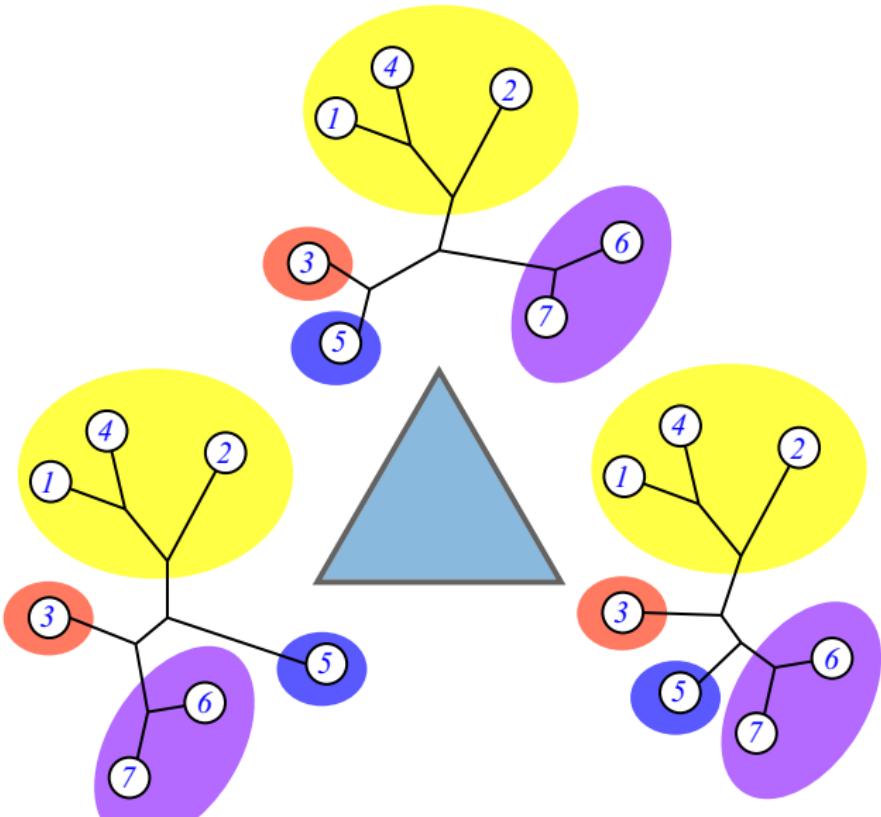
# A convex hull of binary trees.



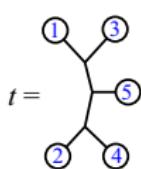
# The associahedron.



# Clade face



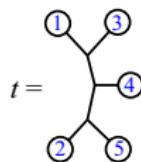
# The Balanced minimal evolution polytope $\mathcal{P}_5$ .



$$\vec{c}(t) = (1, 4, 1, 2, 1, 4, 2, 1, 2, 2)$$

$\{(4,1,1,2,1,1,2,4,2,2)$   
 $(4,2,1,1,2,1,1,2,2,4)$   
 $(4,1,2,1,1,2,1,2,4,2)$   
 $(2,1,4,1,2,2,2,1,4,1)$   
 $(2,2,2,2,1,4,1,1,4,1)$   
 $(1,4,1,2,1,4,2,1,2,2)$   
 $(1,2,1,4,2,4,1,2,2,1)$   
 $(2,1,1,4,2,2,2,4,1,1)$

$$\vec{f} = \langle 15, 105, 250, 210, 52, 1 \rangle$$

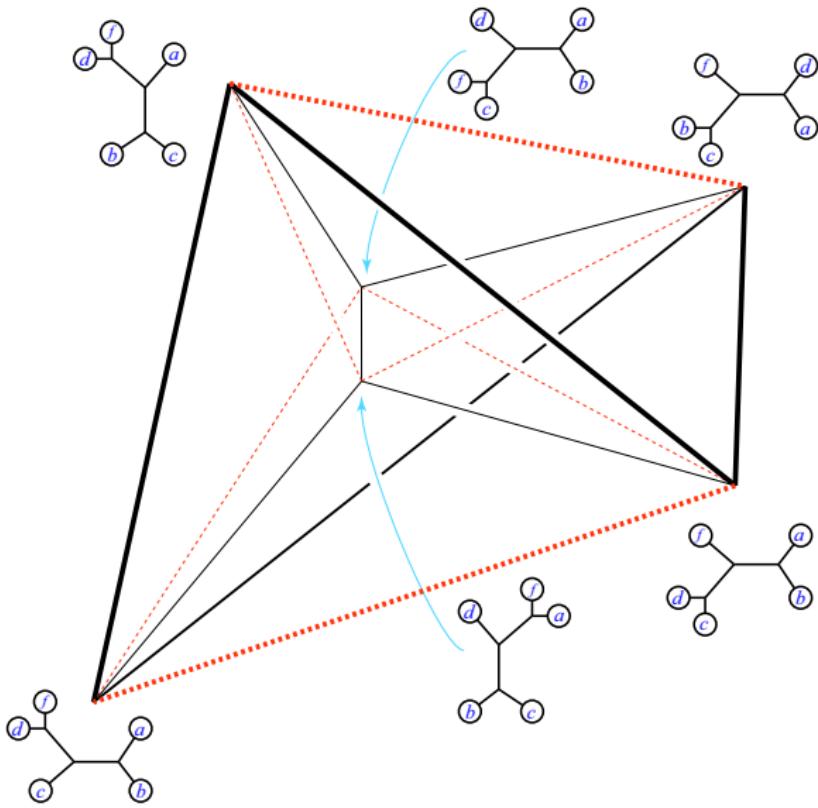


$$\vec{c}(t) = (1, 4, 2, 1, 1, 2, 4, 2, 1, 2)$$

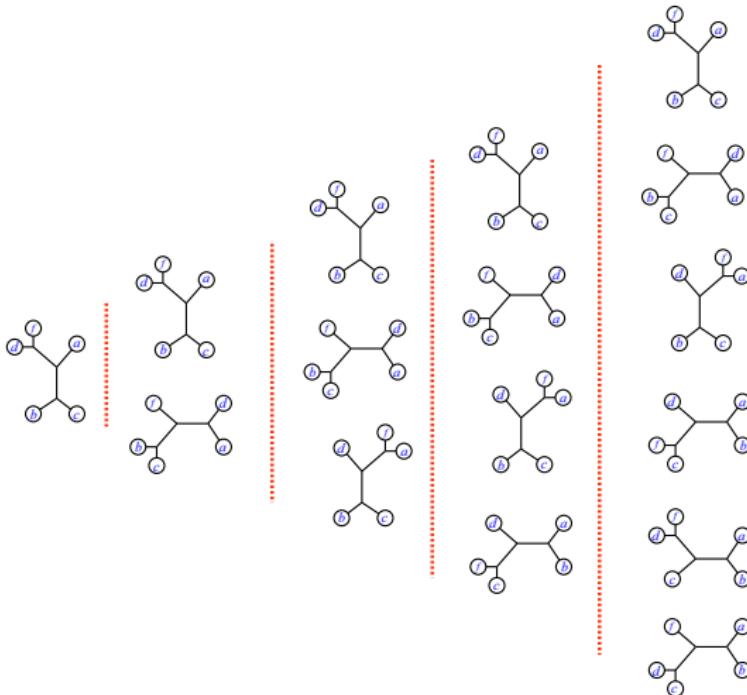
$(1,1,2,4,4,2,1,2,1,2)$   
 $(1,1,4,2,4,1,2,1,2,2)$   
 $(2,2,2,2,4,1,1,1,1,4)$   
 $(2,4,1,1,2,2,2,1,1,4)$   
 $(1,4,2,1,1,2,4,2,1,2)$   
 $(1,2,4,1,2,1,4,2,2,1)$   
 $(2,2,2,2,1,1,4,4,1,1)\}$

**Figure:** Two sample vertex trees of  $\mathcal{P}_5$  with their respective coordinates shown beneath, followed by all 15 vertex points calculated for  $n=5$ , and the  $f$ -vector for  $\mathcal{P}_5$  as found by polymake.

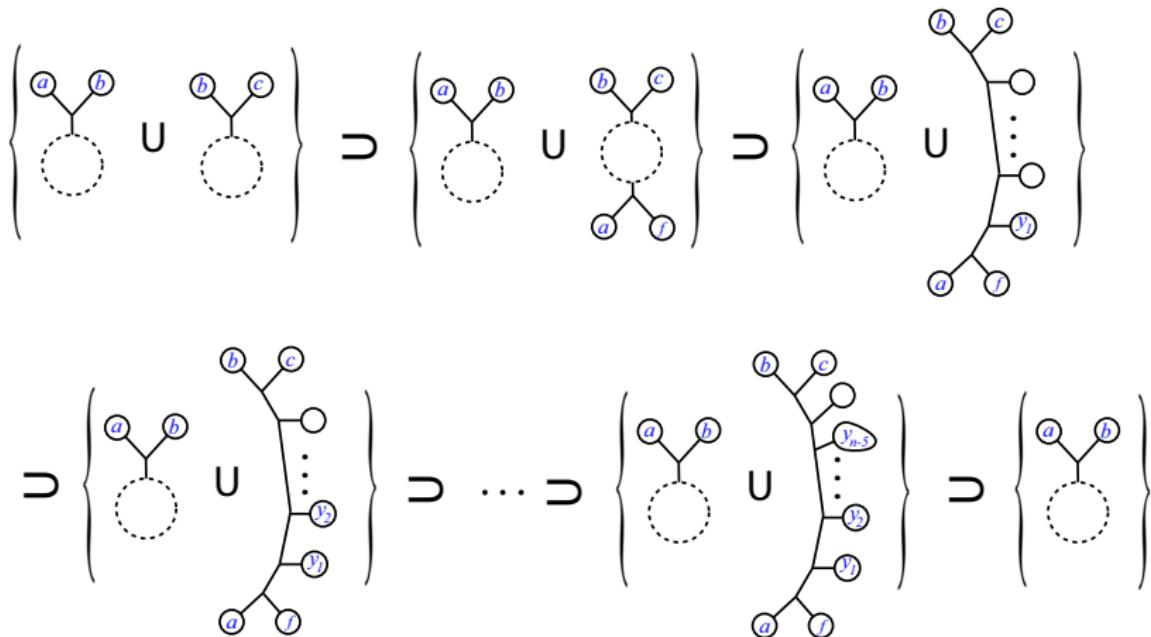
Intersecting cherries facet:  $x_{ab} + x_{bc} - x_{ac} \leq 8$ .



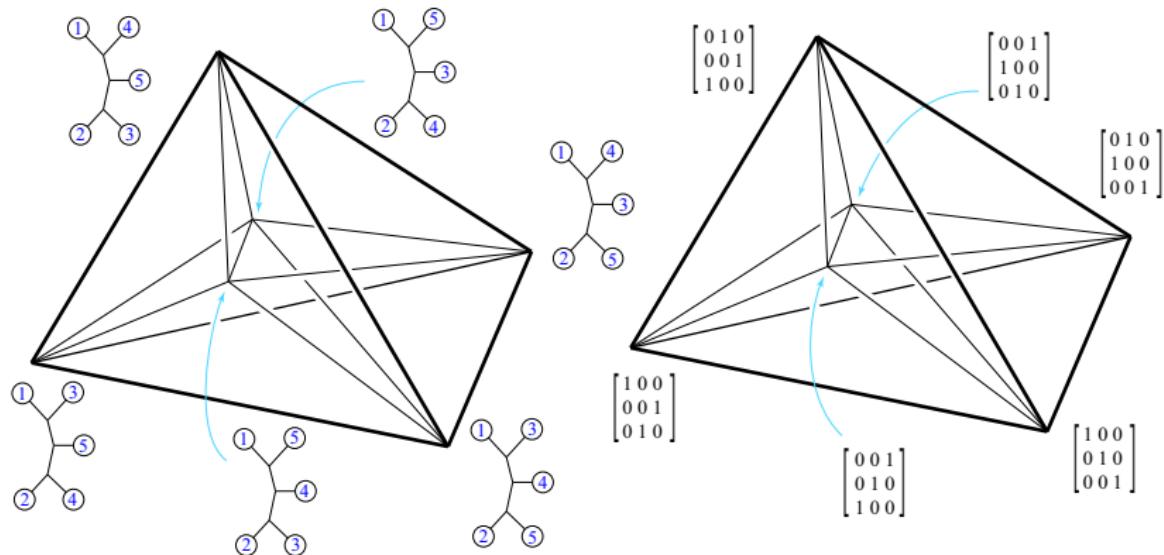
Intersecting cherry flag:  $x_{ab} + x_{bc} - x_{ac} \leq 8$ .



Intersecting cherries facet flag:  $x_{ab} + x_{bc} - x_{ac} \leq 2^{n-3}$ .

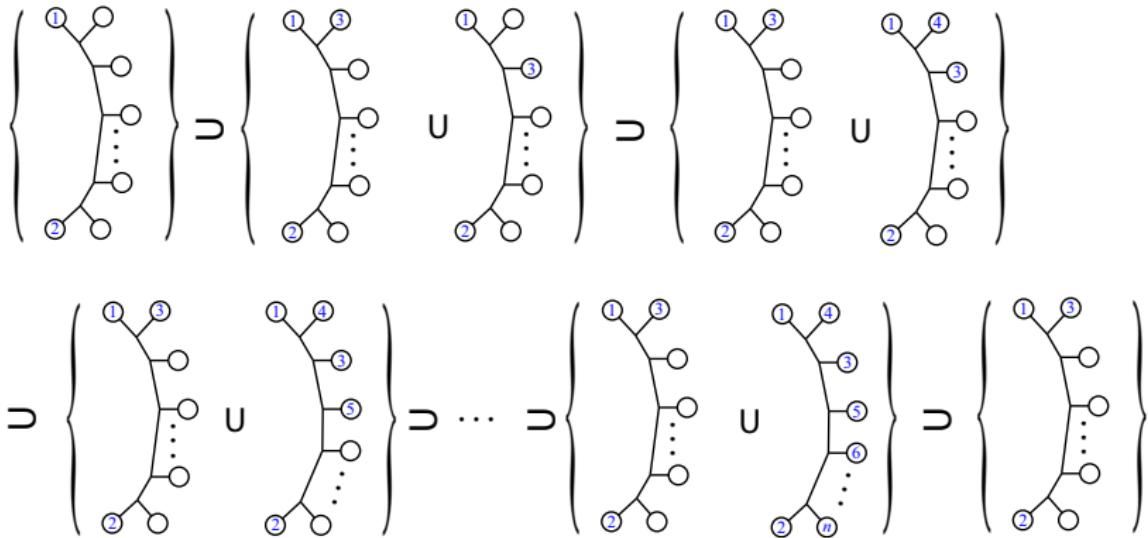


Caterpillar facet:  $x_{ab} \geq 1$ .

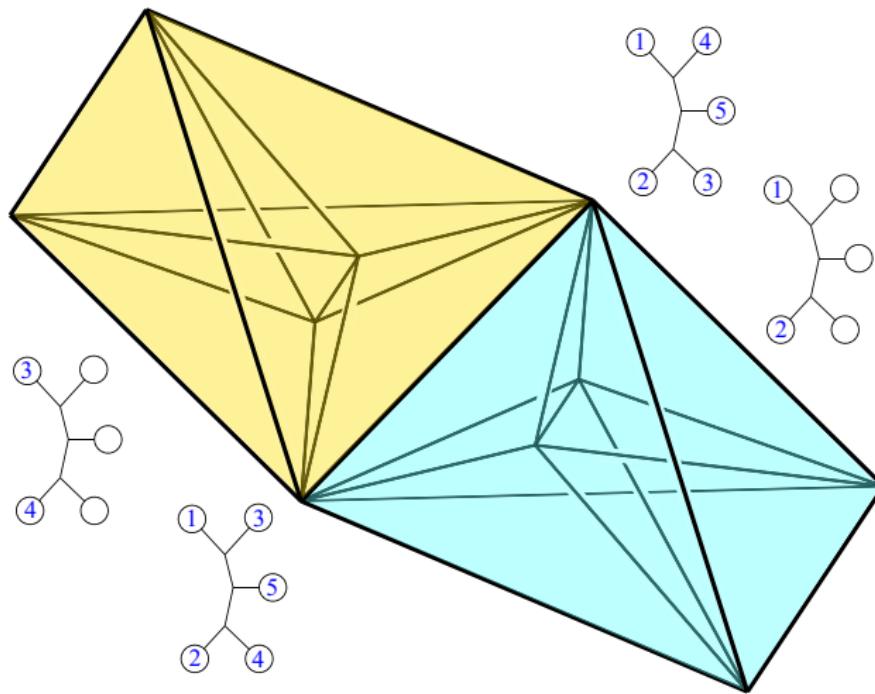


**Figure:** On the left is a facet of  $\mathcal{P}_5$  with each vertex labeled by the caterpillar tree. On the right is the Birkhoff polytope  $B(3)$  with vertices labeled by the corresponding permutation matrices.

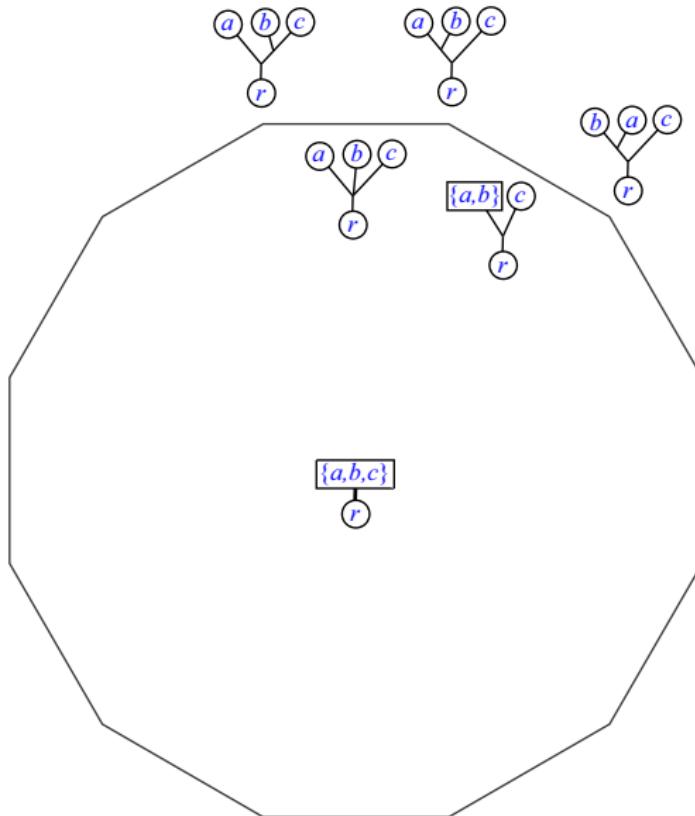
Caterpillar flag:  $x_{ab} \geq 1$ .



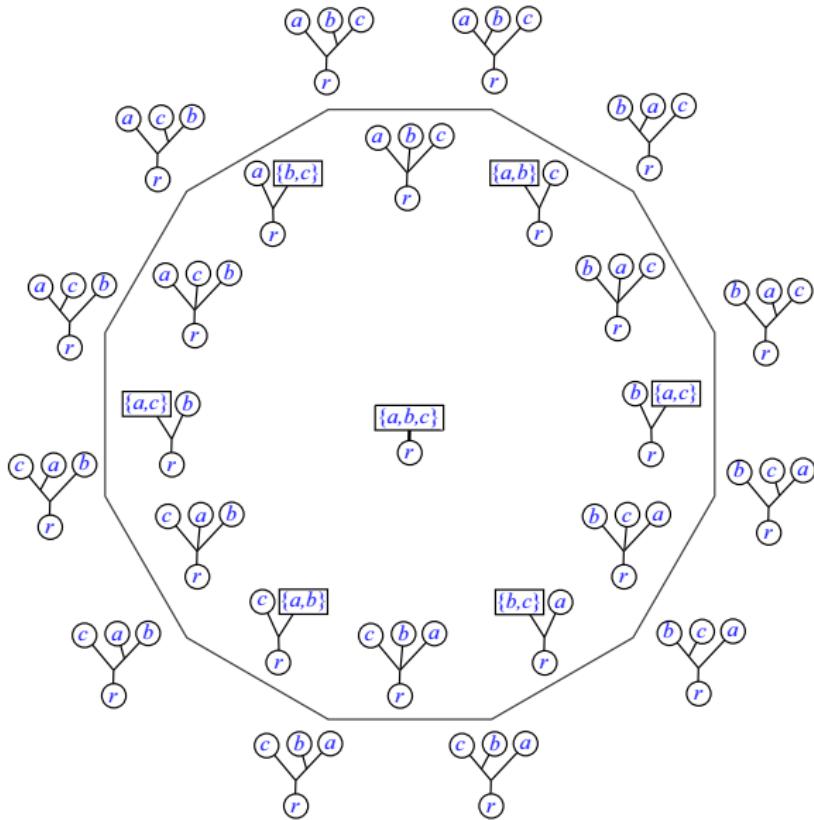
# Intersection.



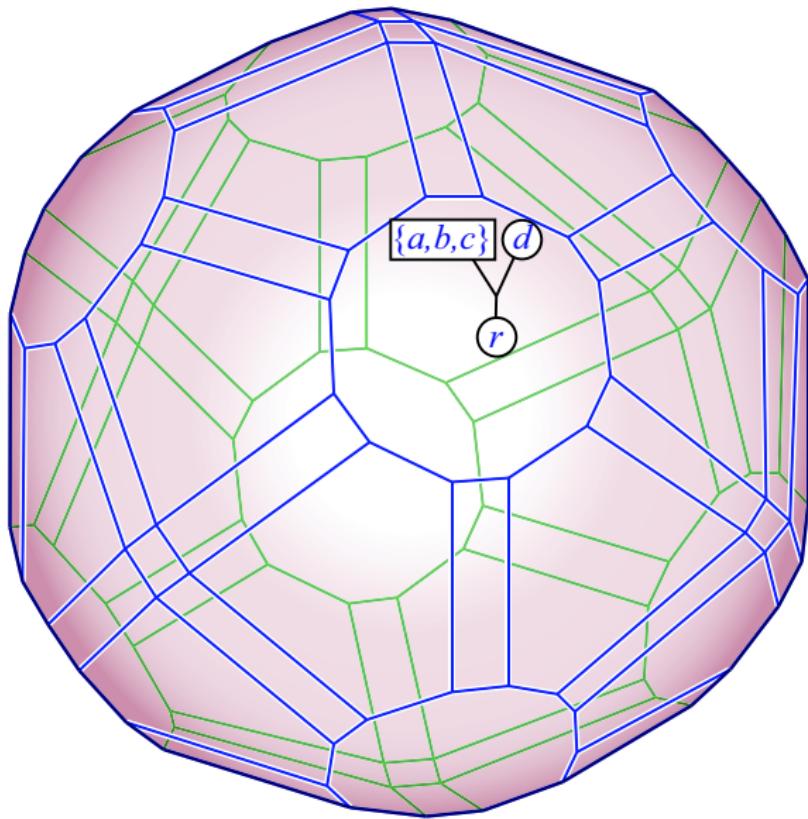
# Permutoassociahedron $\mathcal{KP}_2$



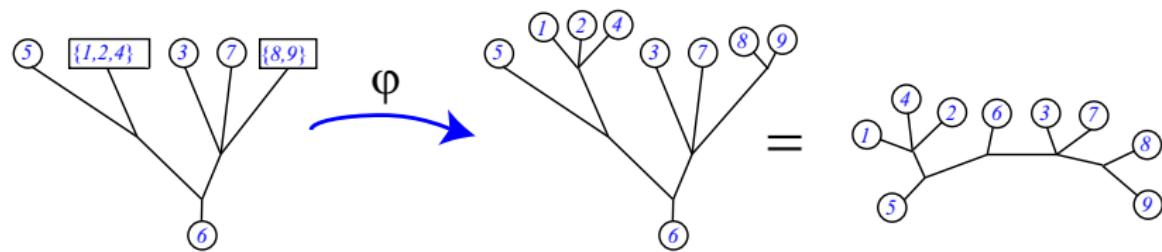
# Permutoassociahedron $\mathcal{KP}_2$



# Permutoassociahedron $\mathcal{KP}_3$



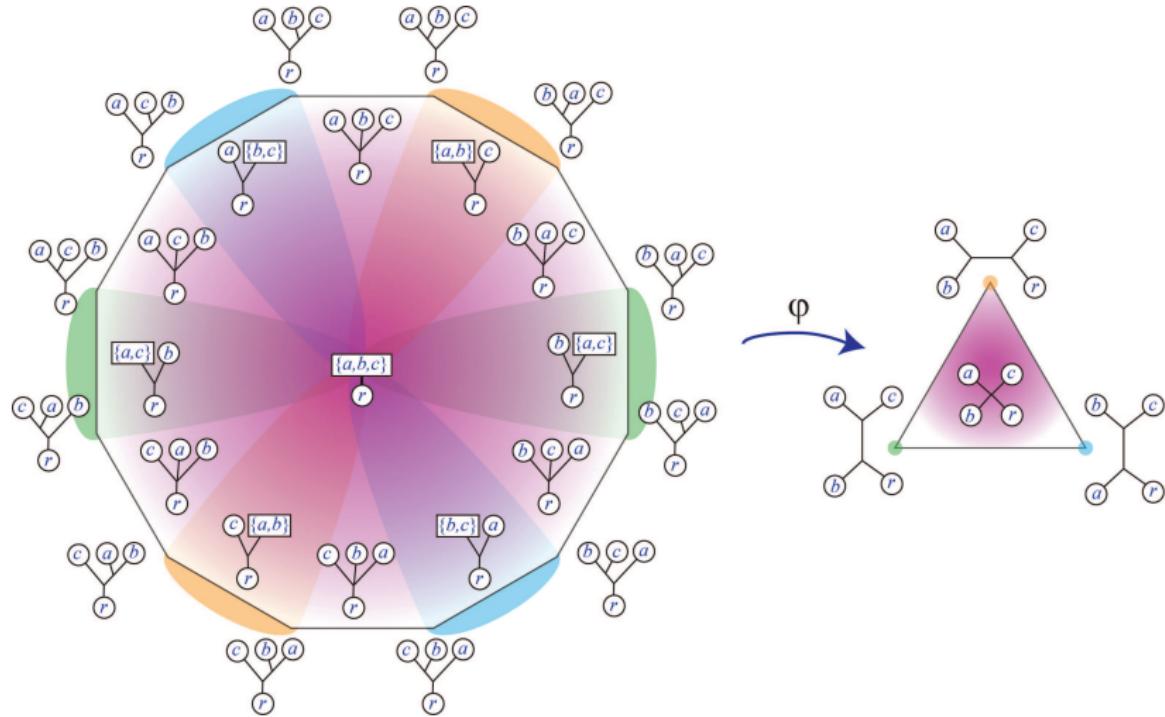
# Projection to BME(n)



## Theorem

If  $x \leq y$  as faces in the face lattice of  $\mathcal{KP}_n$ , then  $\varphi(x) \leq \varphi(y)$  as faces in the face lattice of  $\mathcal{P}_n$ , the BME polytope.

# Projection to BME(2)



Now we show how the target of the map  $\varphi$  is actually the BME polytope.

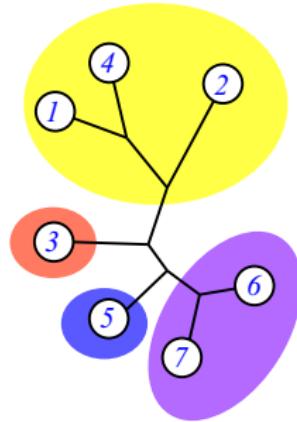
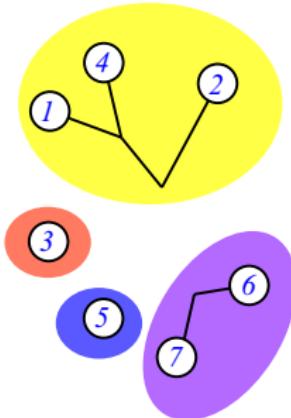
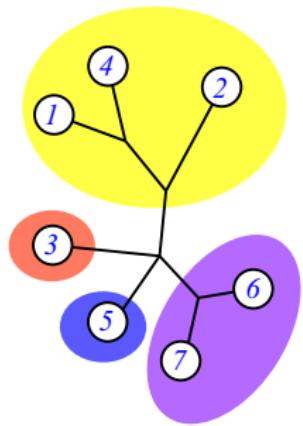
### Theorem

*For each non-binary phylogenetic tree  $t$  with  $n$  leaves there is a corresponding face  $F(t)$  of the BME polytope  $\mathcal{P}_n$ . The vertices of  $F(t)$  are the binary phylogenetic trees which are refinements of  $t$ .*

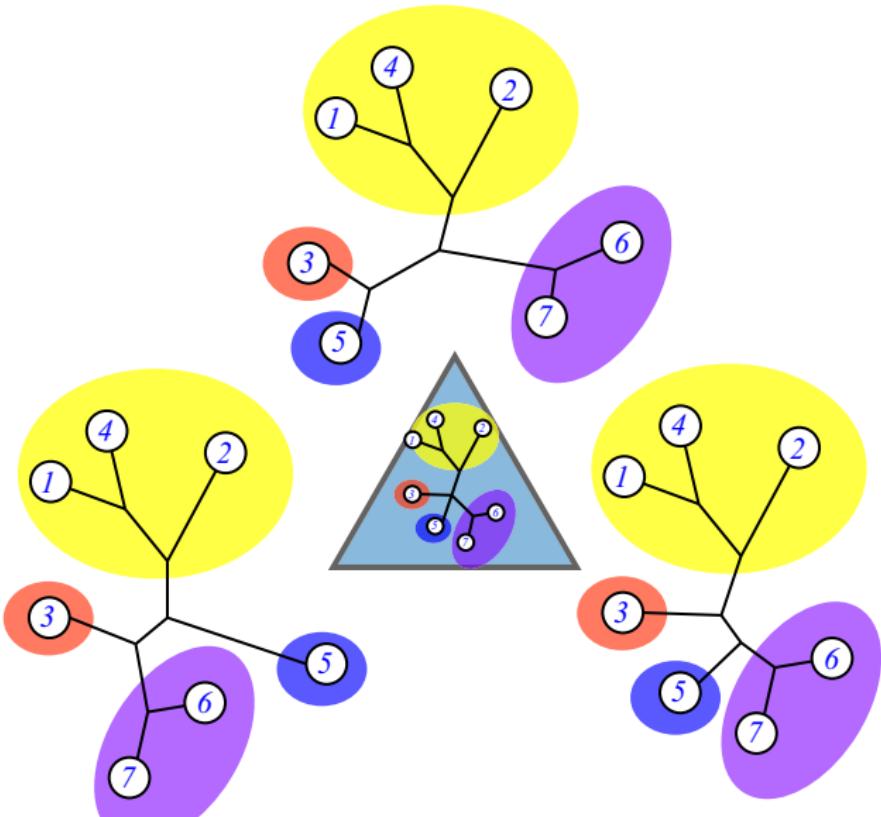
## Theorem

For  $t$  an  $n$ -leaved phylogenetic tree with exactly one node  $\nu$  of degree  $m > 3$ , the tree face  $F(t)$  is precisely the clade-face  $F_{C_1, \dots, C_p}$ , defined in [H,H,Y], corresponding to the collection of clades  $C_1, \dots, C_p$  which result from deletion of  $\nu$ . Thus  $F(t)$  is combinatorially equivalent to the smaller dimensional BME polytope  $\mathcal{P}_m$ .

# Clade face



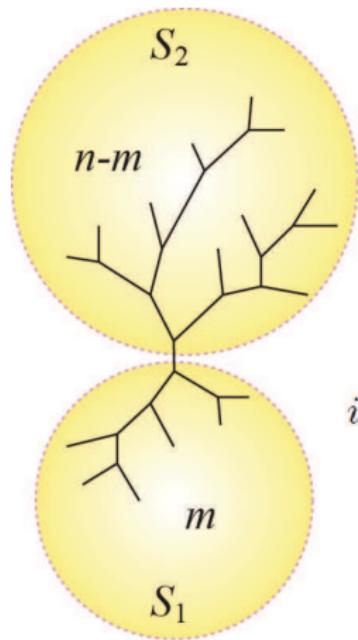
# Clade face



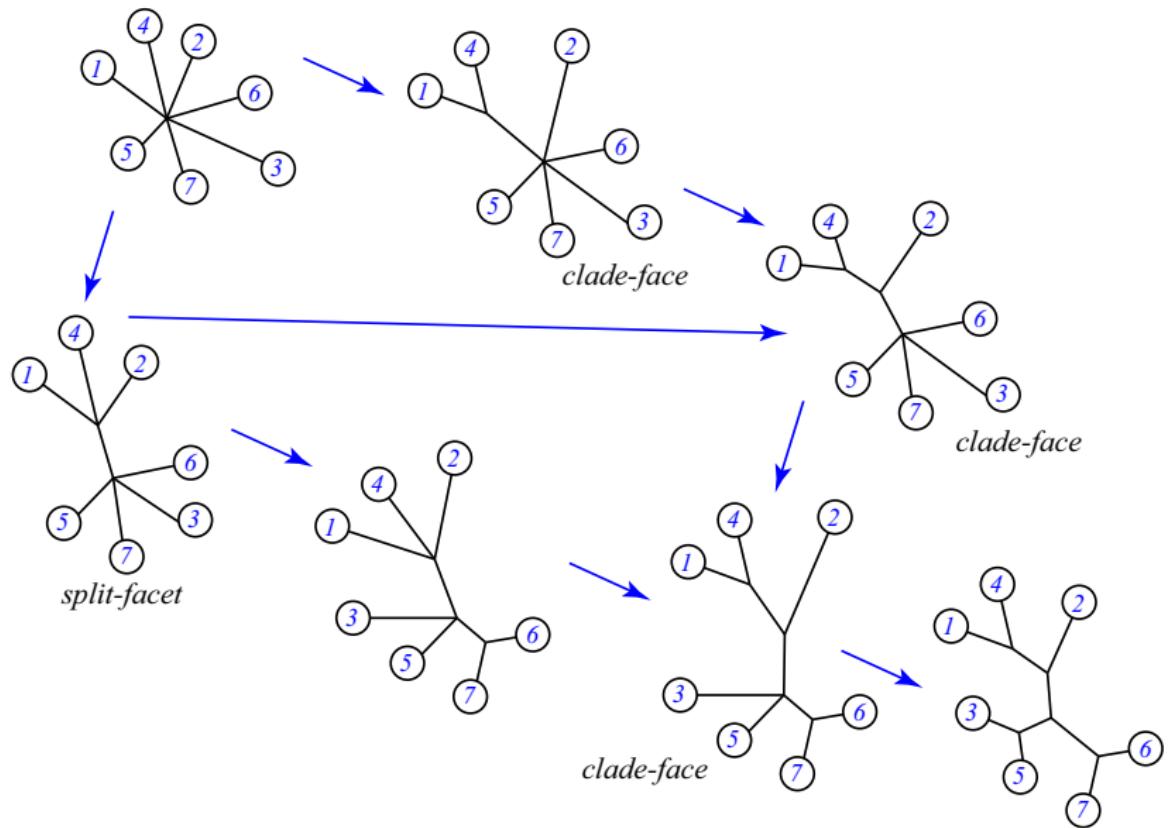
## Theorem

*Let  $t$  be a phylogenetic tree with  $n > 5$  leaves which has exactly two nodes  $\nu$  and  $\mu$ , with degrees both larger than 3. Then the trees which refine  $t$  are the vertices of a facet of the BME polytope  $\mathcal{P}_n$ .*

## Split faces; split facets.



$$\sum_{i < j, \text{ leaves } i, j \in S_1} x_{ij} \leq (m - 1)2^{n-3}$$



**Figure:** Examples of chains in the lattice of tree-faces of the BME polytope  $\mathcal{P}_9$ .

## Definitions

A *split network* is a collection of splits of a set of leaves.

A *split network diagram* represents each split with a set of parallel edges.

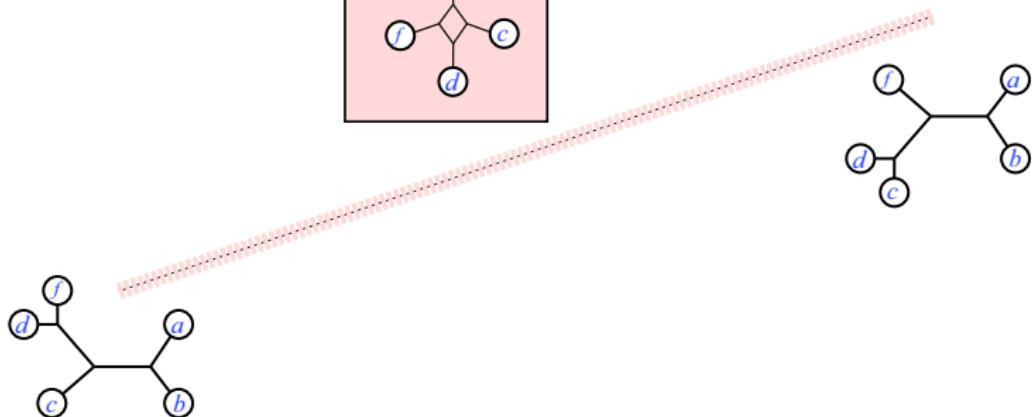
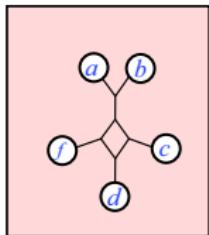
A *circular split network*, also known as a *planar split network*, is a network whose diagram can be drawn on the plane without crossing edges.

A network of *compatible* splits is one whose diagram is a tree.

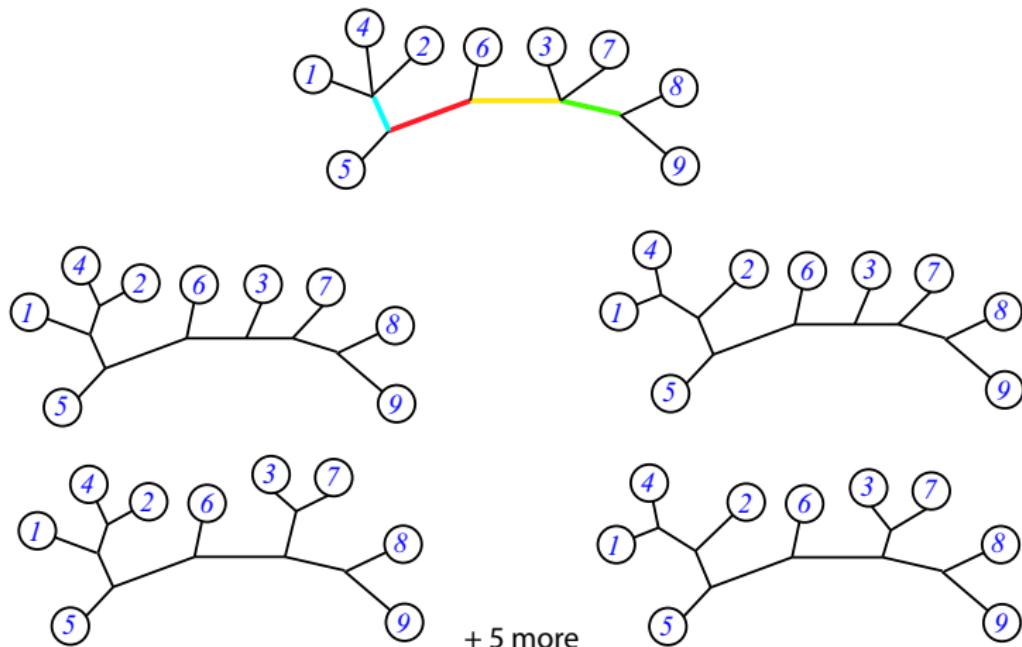
A *binary* split network is one whose diagram has vertices of degree three (or one, for the leaves) only.

# Q1: Split faces; split facets.

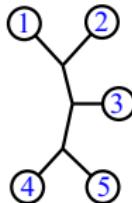
Question 1. Which split networks correspond to faces  
(and especially facets)  
of the Balanced Minimal Evolution polytope?



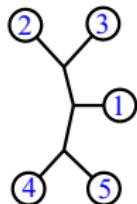
# A1. any set of compatible splits.



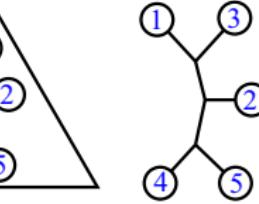
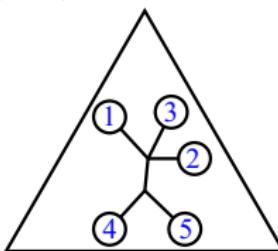
# A1. any set of compatible splits.



$$x(t) = (4, 2, 1, 1, 2, 1, 1, 2, 2, 4)$$

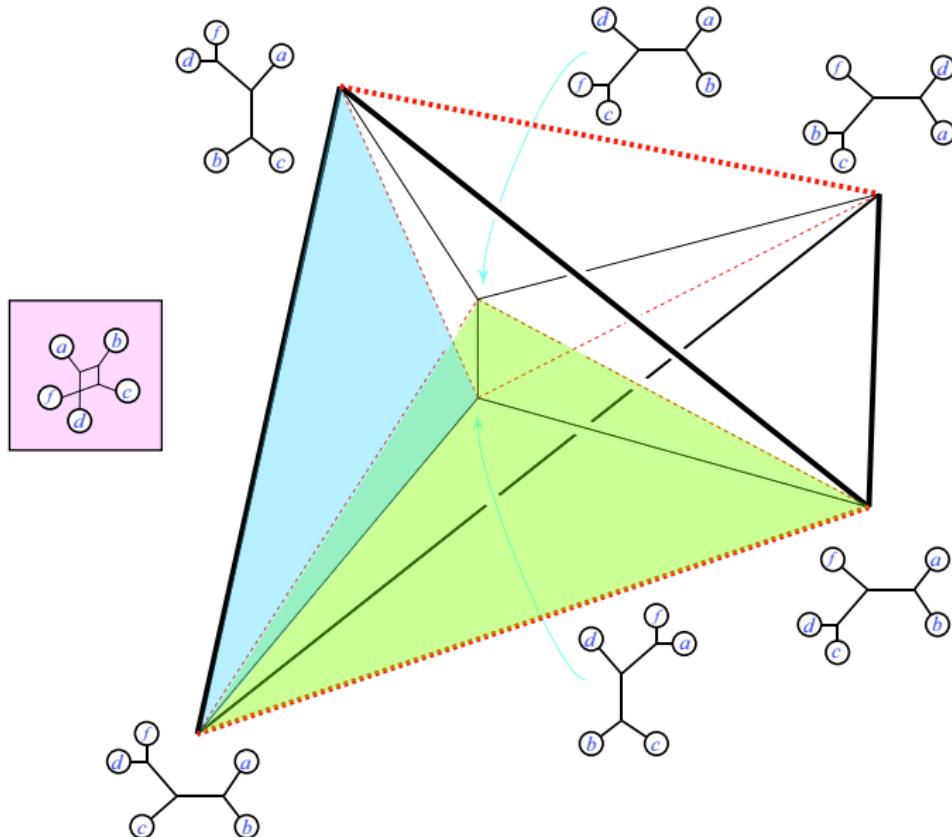


$$x(t) = (2, 2, 2, 2, 4, 1, 1, 1, 1, 4)$$

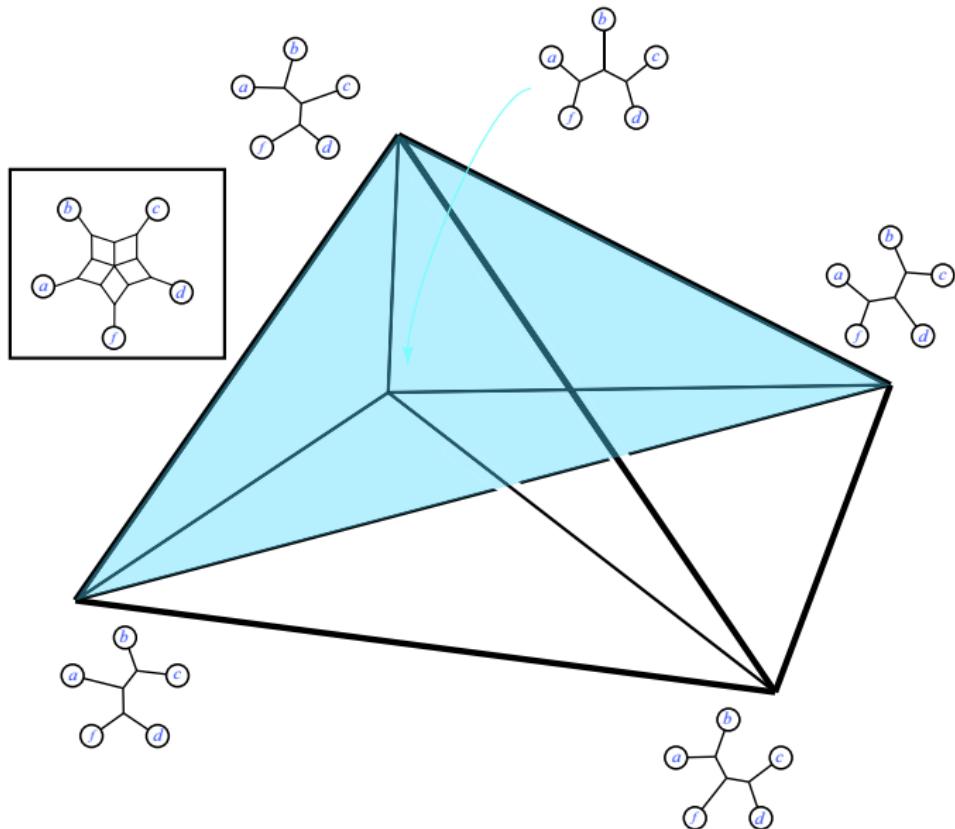


$$x(t) = (2, 4, 1, 1, 2, 2, 2, 1, 1, 4)$$

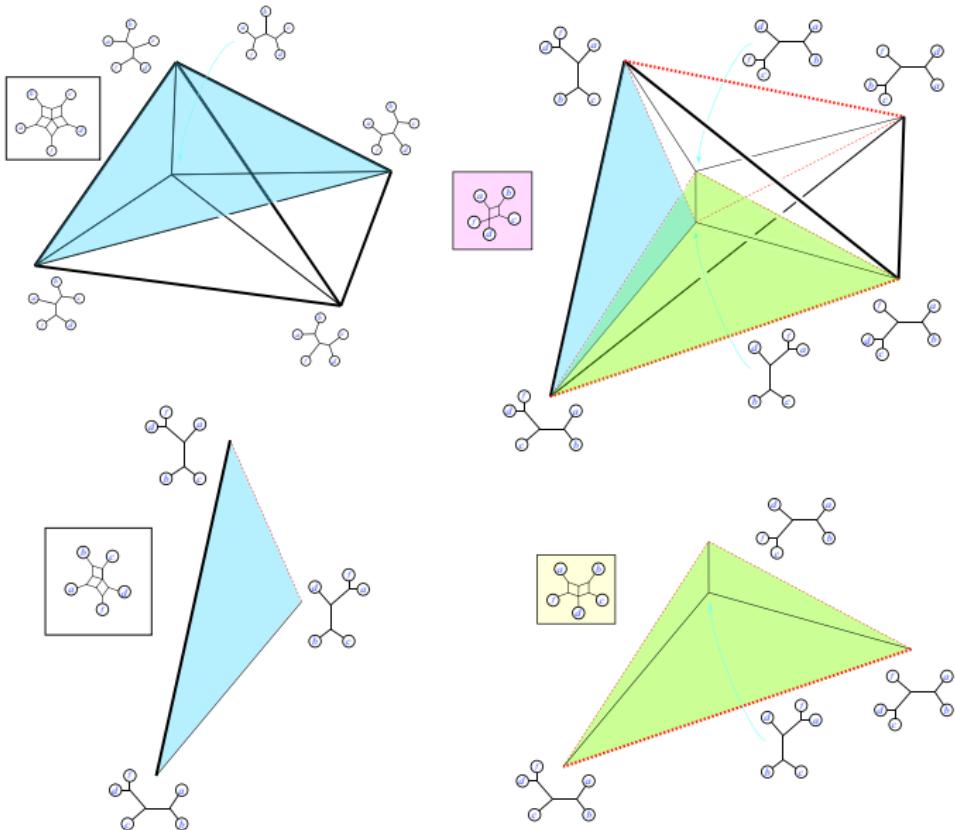
## A1. Intersecting cherry splits



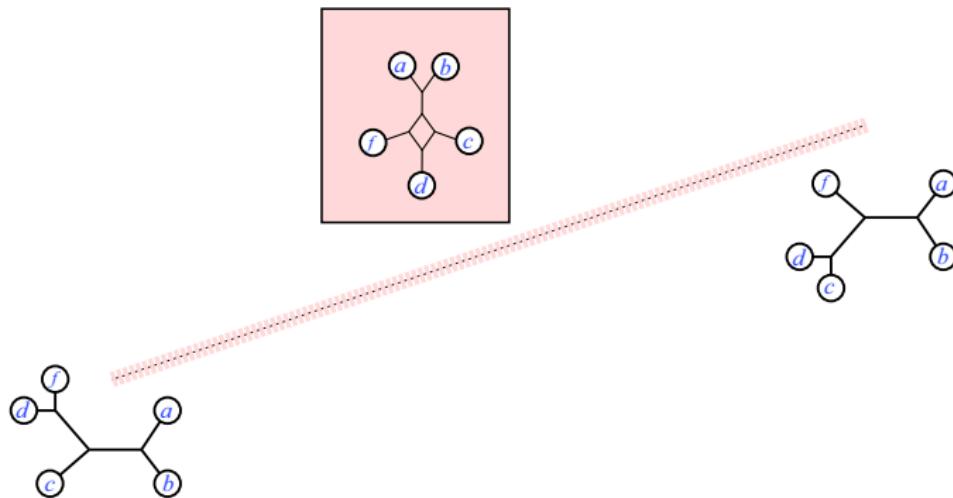
# A1: Cyclic splits for $n = 5$



# A1: Four split networks.

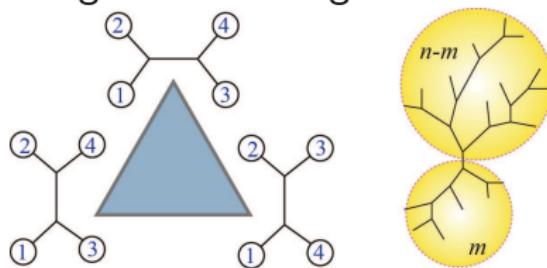


# A1: Nearest Neighbor Interchange.

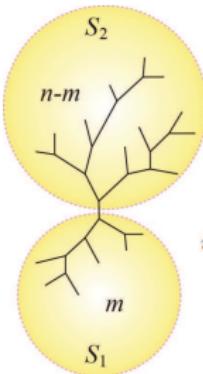


## Q2: Split faces; split facets.

Question 2. If we use branch and bound to optimize on the region bounded by split faces of the BME polytope, are we guaranteed to get a valid tree?



# Splitohedron.



$$\sum_{i < j, \text{ leaves } i, j \in S_1} x_{ij} \leq (m-1)2^{n-3}$$

Theorem: the Splitohedron is a bounded polytope that is a relaxation of the BME polytope.

Proof: The split-faces include the cherries where the inequality is  $x_{ij} \leq 2^{n-3}$ , and the caterpillar facets have the inequality  $x_{ij} \geq 1$ , thus the resulting intersection of halfspaces is a bounded polytope since it is inside the hypercube  $[1, 2^{n-3}]^{\binom{n}{2}}$ .

# Features of the BME polytope $\mathcal{P}_n$

number of species	dim. of $\mathcal{P}_n$	vertices of $\mathcal{P}_n$	facets of $\mathcal{P}_n$	facet inequalities (classification)	number of facets	number of vertices in facet
3	0	1	0	-	-	-
4	2	3	3	$x_{ab} \geq 1$	3	2
				$x_{ab} + x_{bc} - x_{ac} \leq 2$	3	2
5	5	15	52	$x_{ab} \geq 1$ (caterpillar)	10	6
				$x_{ab} + x_{bc} - x_{ac} \leq 4$ (intersecting-cherry)	30	6
				$x_{ab} + x_{bc} + x_{cd} + x_{df} + x_{fa} \leq 13$ (cyclic ordering)	12	5
				$x_{ab} \geq 1$ (caterpillar)	15	24
6	9	105	90262	$x_{ab} + x_{bc} - x_{ac} \leq 8$ (intersecting-cherry)	60	30
				$x_{ab} + x_{bc} + x_{ac} \leq 16$ (3, 3)-split	10	9
				$x_{ab} \geq 1$ (caterpillar)	$\binom{n}{2}$	$(n-2)!$
$n$	$\binom{n}{2} - n$	$(2n-5)!!$	?	$x_{ab} + x_{bc} - x_{ac} \leq 2^{n-3}$ (intersecting-cherry)	$\binom{n}{2}(n-2)$	$2(2n-7)!!$
				$x_{ab} + x_{bc} + x_{ac} \leq 2^{n-2}$ ( $m, 3$ )-split, $m \geq 3$	$\binom{n}{3}$	$3(2n-9)!!$
				$\sum_S x_{ij} \leq (m-1)2^{n-3}$ ( $m, n-m$ )-split $S$ , $m > 2, n > 5$	$2^{n-1} - \binom{n}{2}$ $-n-1$	$(2(n-m)-3)!!$ $\times (2m-3)!!$

# Splitohedron.

```
polytope > print $p->VERTICES;
```

```
1 1 2 1 4 2 4 1 2 2 1  
1 1 2 4 1 2 1 4 2 2 1  
1 1 4 2 1 1 2 4 2 1 2  
1 1 1 2 4 4 2 1 2 1 2  
1 1 1 4 2 4 1 2 1 2 2  
1 1 4 1 2 1 4 2 1 2 2  
1 2 1 4 1 2 2 2 1 4 1  
1 8/3 4/3 8/3 4/3 4/3 4/3 8/3 8/3 8/3 4/3  
1 2 1 1 4 2 2 2 4 1 1  
1 4/3 4/3 8/3 8/3 8/3 4/3 4/3 8/3 4/3 8/3  
1 4/3 8/3 4/3 8/3 8/3 4/3 4/3 8/3 4/3 8/3  
1 4 1 2 1 1 2 1 2 4 2 2  
1 8/3 4/3 4/3 8/3 8/3 4/3 4/3 8/3 4/3 8/3  
1 8/3 4/3 8/3 4/3 8/3 4/3 4/3 8/3 4/3 8/3  
1 2 2 2 2 4 1 1 1 1 4  
1 8/3 8/3 4/3 4/3 4/3 8/3 4/3 4/3 8/3 8/3  
1 8/3 8/3 4/3 4/3 4/3 8/3 4/3 8/3 4/3 8/3  
1 2 4 1 1 2 2 2 1 1 4  
1 4/3 4/3 8/3 8/3 8/3 4/3 8/3 8/3 4/3 4/3  
1 4/3 8/3 4/3 8/3 4/3 8/3 8/3 4/3 8/3 4/3
```

```
1 2 2 2 2 1 1 4 4 1 1  
1 2 2 2 2 1 4 1 1 4 1  
1 4/3 8/3 8/3 4/3 8/3 4/3 8/3 4/3 8/3 4/3  
1 4/3 8/3 8/3 4/3 4/3 8/3 8/3 4/3 8/3 4/3  
1 4 1 1 2 1 1 2 4 2 2  
1 8/3 4/3 4/3 8/3 8/3 4/3 4/3 8/3 4/3 8/3  
1 8/3 4/3 8/3 4/3 8/3 4/3 4/3 8/3 4/3 8/3  
1 2 2 2 2 4 1 1 1 1 4  
1 8/3 8/3 4/3 4/3 4/3 8/3 4/3 4/3 8/3 8/3  
1 8/3 8/3 4/3 4/3 4/3 8/3 4/3 8/3 4/3 8/3  
1 2 4 1 1 2 2 2 1 1 4  
1 4/3 4/3 8/3 8/3 8/3 4/3 8/3 8/3 4/3 4/3  
1 4/3 8/3 4/3 8/3 4/3 8/3 8/3 4/3 8/3 4/3
```

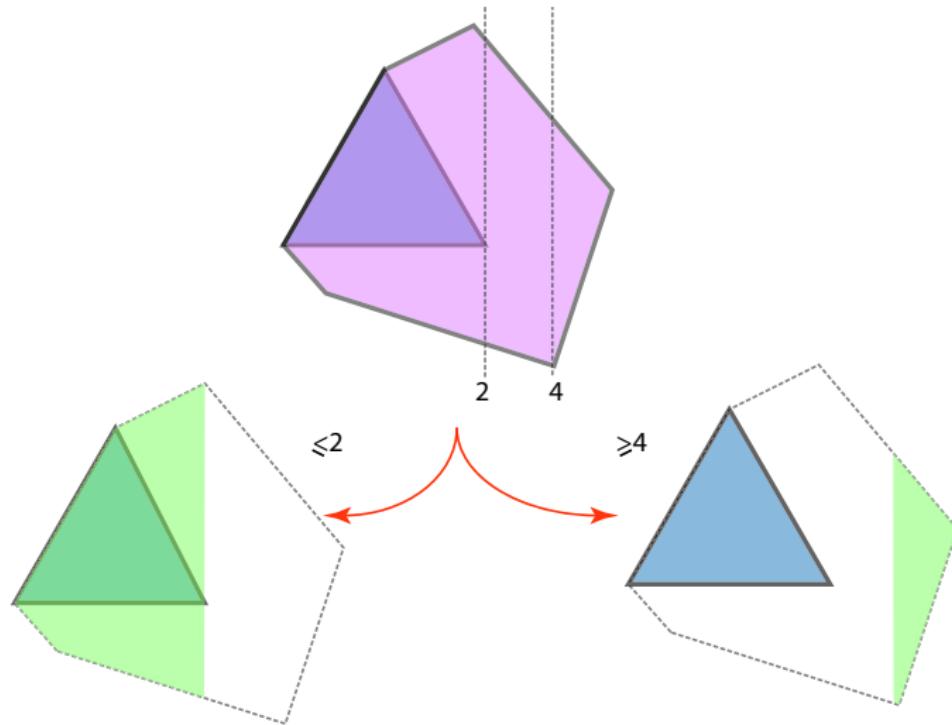
# Splitohedron.

```
polytope > print $p->VERTICES;
```

```
1 1 2 1 4 2 4 1 2 2 1  
1 1 2 4 1 2 1 4 2 2 1  
1 1 4 2 1 1 2 4 2 1 2  
1 1 1 2 4 4 2 1 2 1 2  
1 1 1 4 2 4 1 2 1 2 2  
1 1 4 1 2 1 4 2 1 2 2  
1 2 1 4 1 2 2 2 1 4 1  
1 8/3 4/3 8/3 4/3 4/3 4/3 8/3 8/3 8/3 4/3  
1 2 1 1 4 2 2 2 4 1 1  
1 4/3 4/3 8/3 8/3 8/3 4/3 4/3 8/3 4/3  
1 4/3 8/3 4/3 8/3 8/3 4/3 4/3 4/3 8/3  
1 4 1 2 1 1 2 1 2 4 2  
1 4 2 1 1 2 1 1 2 2 4  
1 8/3 4/3 4/3 8/3 4/3 8/3 4/3 8/3 4/3
```

```
1 2 2 2 2 1 1 4 4 1 1  
1 2 2 2 2 1 4 1 1 4 1  
1 4/3 8/3 8/3 4/3 4/3 8/3 4/3 8/3 4/3 8/3  
1 4/3 8/3 8/3 4/3 4/3 8/3 8/3 4/3 8/3 4/3  
1 4 1 1 2 1 1 2 4 2 2  
1 8/3 4/3 4/3 8/3 8/3 4/3 4/3 8/3 4/3 8/3  
1 8/3 4/3 8/3 4/3 8/3 4/3 4/3 8/3 4/3 8/3  
1 2 2 2 2 4 1 1 1 4  
1 8/3 8/3 4/3 4/3 4/3 8/3 4/3 8/3 4/3 8/3  
1 8/3 8/3 4/3 4/3 4/3 8/3 4/3 8/3 4/3 8/3  
1 2 4 1 1 2 2 2 1 1 4  
1 4/3 4/3 8/3 8/3 8/3 4/3 8/3 8/3 4/3 4/3  
1 4/3 8/3 4/3 8/3 4/3 8/3 8/3 4/3 8/3 4/3
```

BnB.



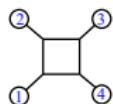
## A2: So far so good!

- We tested up to  $n = 10$ , with and without noise.
- Results are completely accurate...
- We need to find a way to break it! MatLab code available: [http://www.math.uakron.edu/~sf34/class\\_home/research.htm](http://www.math.uakron.edu/~sf34/class_home/research.htm)

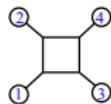
Next.

For any circular split system  $S$ ,  $\mathbf{x}(S)$  is a vector whose  $ij$ -component is the number of circular orderings consistent with that system for which  $i$  and  $j$  are adjacent.

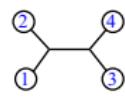
We define a *binary split system* to be a circular split system whose graph has only vertices of degree 3. Examples together with their vectors  $\mathbf{x}(S)$  are illustrated here:



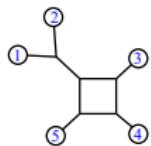
$$(1, 0, 1, 1, 0, 1)$$



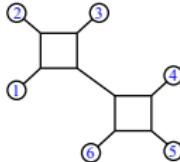
$$(1, 1, 0, 0, 1, 1)$$



$$(2, 1, 1, 1, 1, 2)$$



$$(2, 1, 0, 1, 1, 0, 1, 1, 0, 1)$$



$$(2, 0, 1, 0, 1, 2, 0, 0, 0, 1, 1, 0, 0, 2, 2)$$

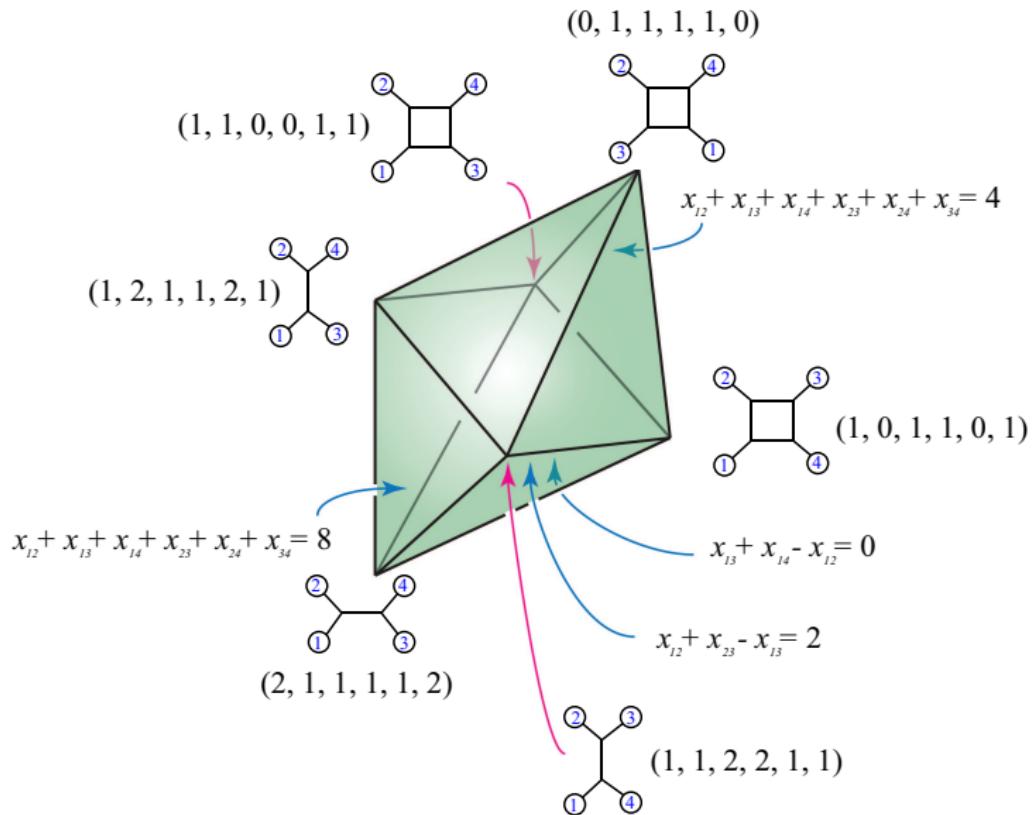
Next.

We propose an extension of the BME polytope which is the convex hull of all vectors  $\eta(S)$  for binary split systems  $S$  on a set of size  $n$ . An example for  $n = 4$  is seen on the next slide.

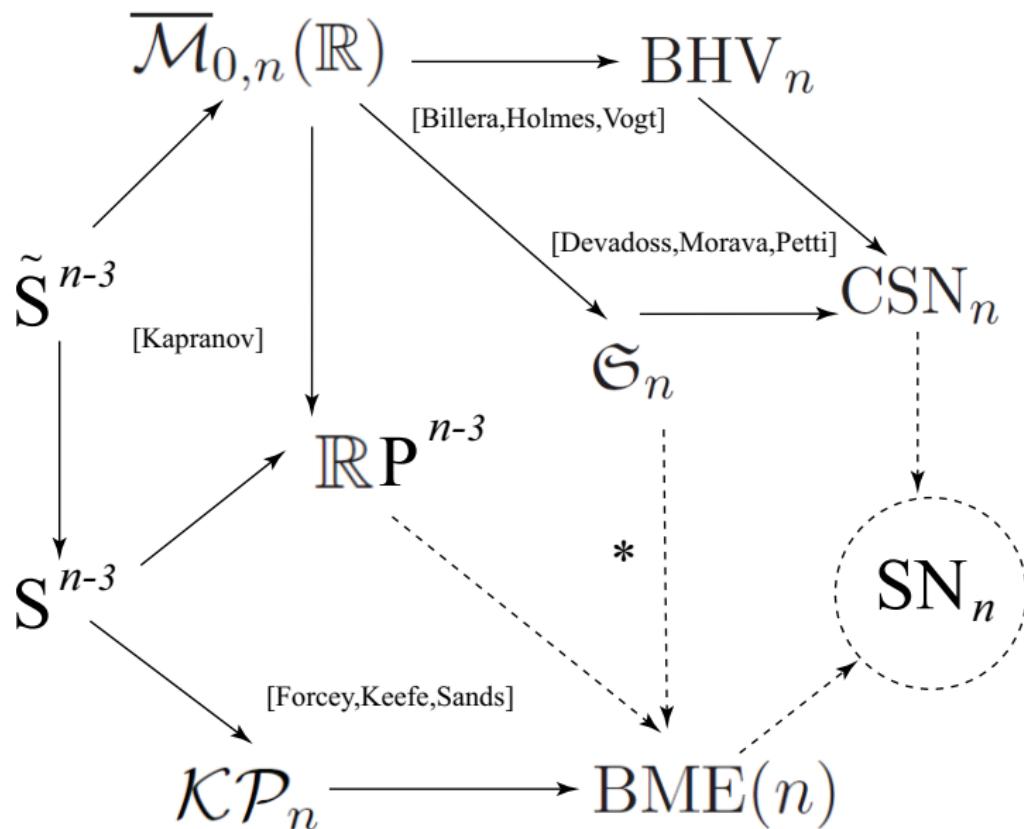
This new polytope has vertices corresponding to all the binary split systems.

These binary split systems come in two varieties: the binary phylogenetic trees and the split systems for which any split is incompatible with at most one other split.

Next.



Next.



Thanks!

Questions and comments?

Advertisement:

<http://www.math.uakron.edu/~sf34/hedra.htm>