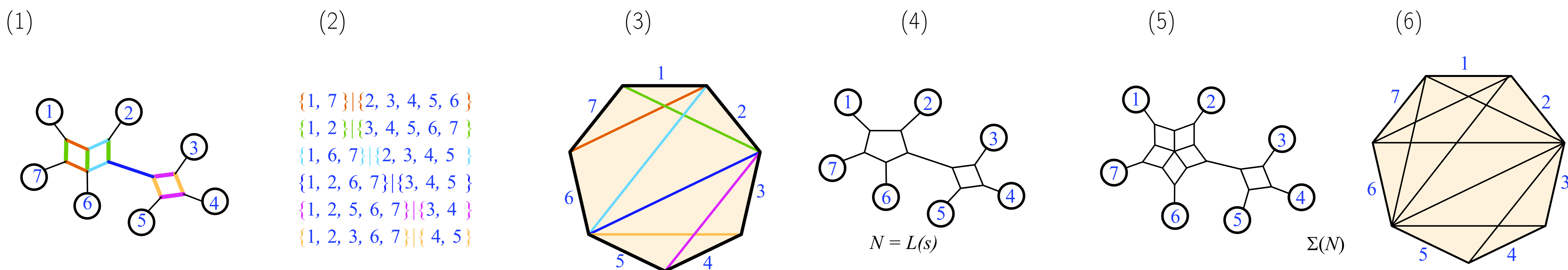


# Split Network Polytopes and Network Spaces

Satyan Devadoss *University of San Diego*, Cassandra Durell *University of Akron*, Stefan Forcey† *University of Akron*

## Definitions with examples: Split Networks and Level-1 Phylogenetic Networks.



Above, we see (1) a *circular split network*  $s$ , which represents the collection of *non-trivial splits*  $A|B$  in the *split system* (2). The same system is represented by (3) the subdivided polygon. Next there is pictured (4) a *level-1 phylogenetic network*  $N$  as defined in [3], which is found by forgetting the internal structure of  $s$  via the map  $L$ . From any level-1 network we can recover a split network  $\Sigma(N)$  which has all the splits corresponding to *minimal cuts* in  $N$  as shown in two representations: (5) and (6).

## Definition: New Polytopes.

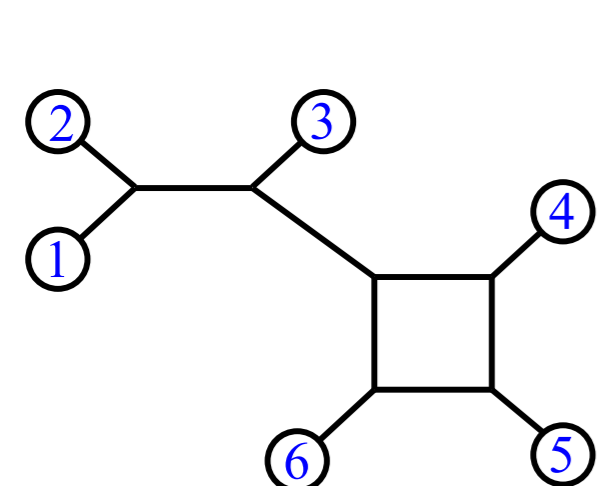
The vector  $\mathbf{x}(s)$  is defined to have lexicographically ordered components  $x_{ij}(s)$  for each unordered pair of distinct leaves  $i, j \in [n]$  as follows:

$$x_{ij}(s) = \begin{cases} 2^{k-b_{ij}} & \text{if there exists a representation of } s \text{ with } i, j \text{ adjacent,} \\ 0 & \text{otherwise.} \end{cases}$$

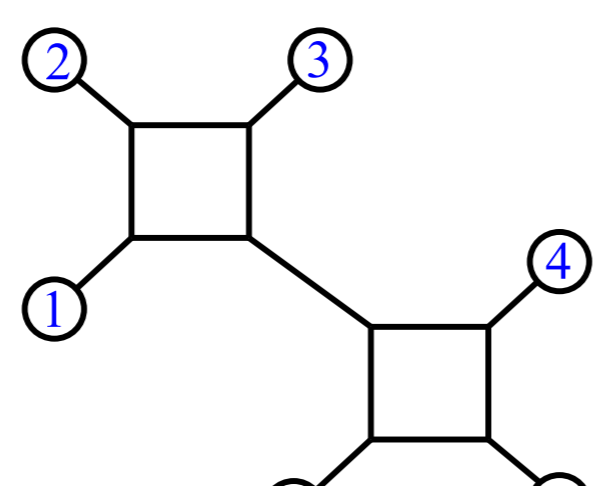
where  $k$  is the number of bridges in  $s$  and  $b_{ij}$  is the number of bridges crossed on any path from  $i$  to  $j$ .

The convex hull of all the vectors  $\mathbf{x}(s)$  for  $s$  any binary level-1 network with  $n$  leaves and  $k$  nontrivial bridges is the *level-1 network polytope*  $\text{BME}(n, k)$ . It has dimension  $\binom{n}{2} - n$ . When  $k = 0$  it is the Symmetric Travelling Salesman polytope  $\text{STSP}(n)$ .

## Examples: vertex vectors $\mathbf{x}(s)$ .



(4, 2, 1, 0, 1, 2, 1, 0, 1, 2, 0, 2, 4, 0, 4)



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

## Theorem: Counting vertices.

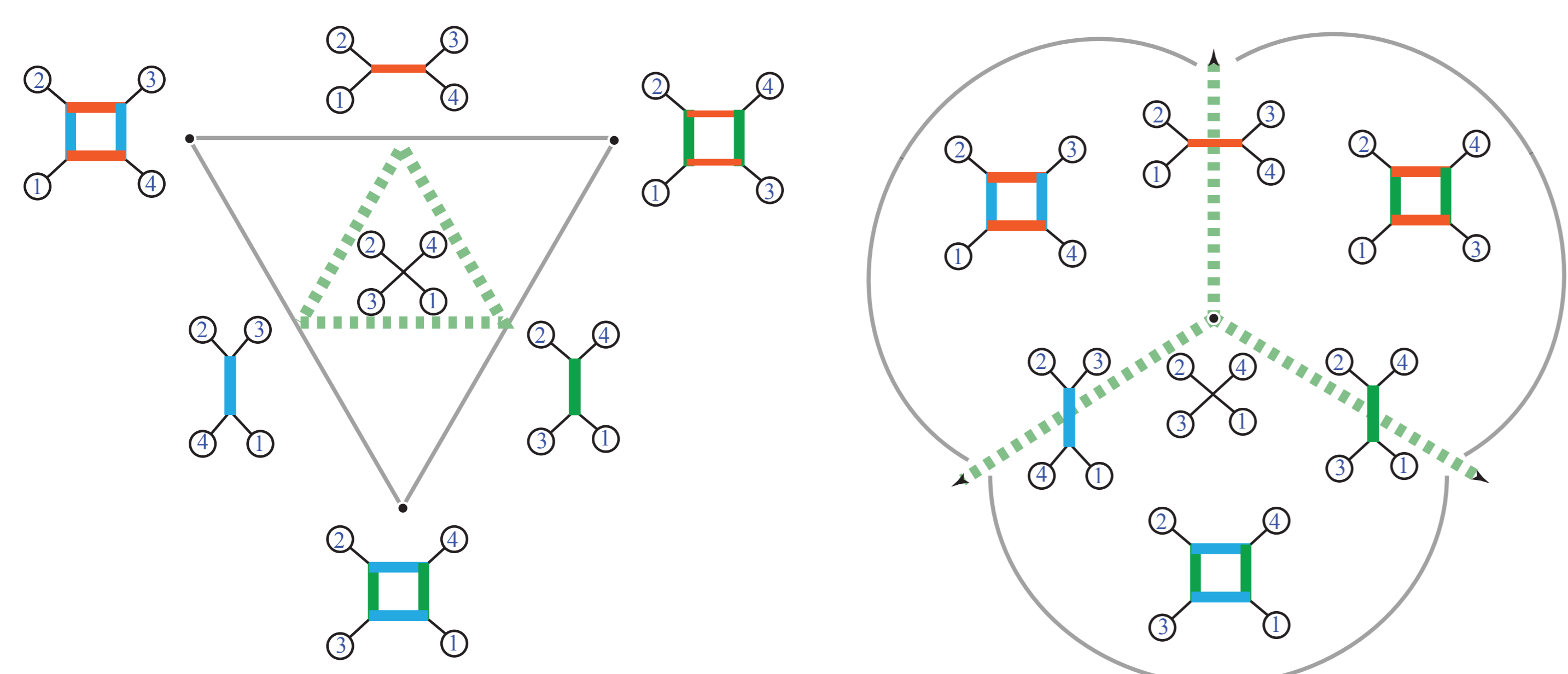
The number of vertices of  $\text{BME}(n, k)$ , and thus the number of binary level-1 networks with  $n$  leaves and  $k$  non-trivial bridges, with  $0 \leq k \leq n - 3$  is:

$$v(n, k) = T(n, k) \frac{(n-1)!}{2^{k+1}}$$

where  $T(n, k)$  gives the components of the face vector of the associahedron  $K(n)$ , which allows the simpler count:

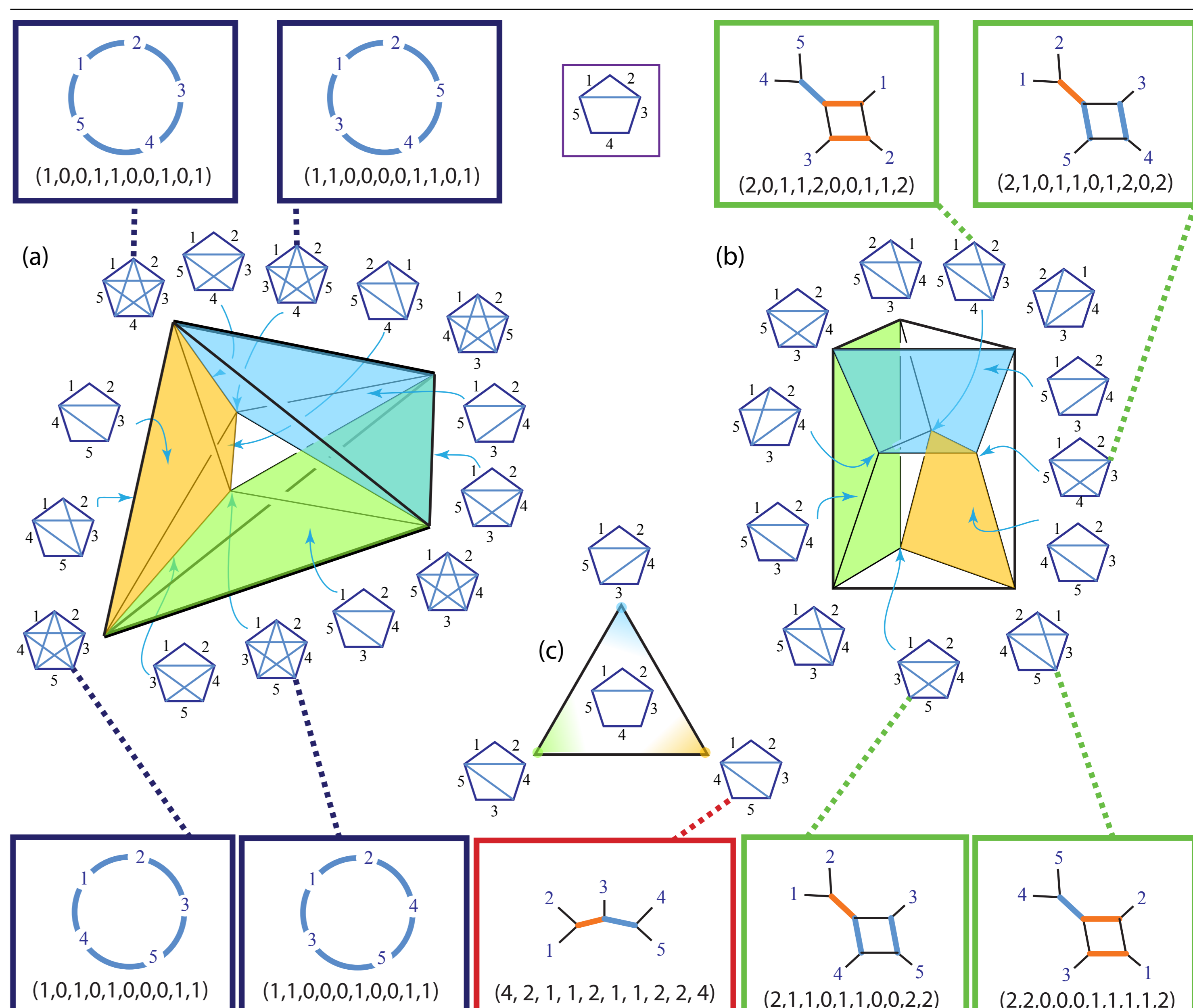
$$v(n, k) = \binom{n-3}{k} \frac{(n+k-1)!}{(2k+2)!!}$$

## Example : Duality...



BME(4,0) on the left is dual to CSN(4) on the right. BME(4,1) is dual to BHV(4); these are the restrictions to trees shown by the dashed lines.

## Facets: new and old polytopes.



A split facet (a) in  $\text{STSP}(5) = \text{BME}(5,0)$ , a facet (b) in  $\text{BME}(5,1)$ , and a face (c) in  $\text{BME}(5,2) = \text{BME}(5)$ . All three correspond to the same split, pictured at the top.

## Definition: New Spaces.

A *weighted split network* has non-negative real values assigned to each split. The geometric space  $\text{CSN}_n$  of metric circular split networks with  $n$  labeled leaves is defined in [2]. A vector of weights, or edge lengths in the network:  $(l_1, \dots, l_{n(n-3)/2})$  specifies a point in the orthant  $[0, \infty)^{n(n-3)/2}$ . Orthants are glued along their common boundaries by identifying equivalent networks (with edge lengths of 0). When we restrict to trees we recover the Billera-Holmes-Vogtman space of phylogenetic trees  $\text{BHV}(n)$ , [1].

## ...Theorem: Duality.

There exists a poset injection:  $f : \mathcal{L}(\text{BHV}_n) \rightarrow \mathcal{L}(\text{BME}(n, n-3)^\Delta)$ . In particular the  $(2n-5)!!$  top-dimensional cells of  $\text{BHV}_n$  map to the  $(2n-5)!!$  vertices of  $\text{BME}(n) = \text{BME}(n, n-3)$ .

The polytope  $\text{STSP}(n)$  has a complex of subfaces which is the dual image of a projection  $f$  from  $\text{CSN}_n$ . The  $(n-1)!/2$  orthants of  $\text{CSN}_n$  map to the vertices of  $\text{STSP}(n)$ . See Example to the left.

- [1] L. Billera, S. Holmes, and K. Vogtmann, *Geometry of the space of phylogenetic trees*, Adv. in Appl. Math. **27** (2001), no. 4, 733–767. MR 1867931
- [2] S. Devadoss and S. Petti, *A space of phylogenetic networks*, SIAM Journal on Applied Algebra and Geometry **1** (2017), 683–705.
- [3] P. Gambette, K. T. Huber, and G. E. Scholz, *Uprooted phylogenetic networks*, Bulletin of Mathematical Biology **79** (2017), no. 9, 2022–2048.