

Matching & Vertex CoverRecall: Theorem: Hall

Let G be a bipartite graph with bipartition (A, B) .
 Then G contains a matching covering A iff

$$|N(S)| \geq |S| \quad \forall S \subseteq A$$

↳ set of neighbours of vertices in S .

A set of vertices $X \subseteq V(G)$ is a vertex cover if every edge of G has an end in X .



not a vertex cover



is a vertex cover

$\tau(G)$ is the minimum size of a vertex cover in G .
 $\nu(G)$ is the maximum size of a matching in G .

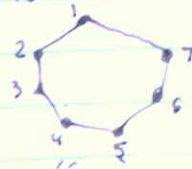
Ex. Complete graphs:

 K_n $n=5$ 

$$\tau(K_n) = n-1$$

$$\nu(K_n) = \lfloor \frac{n}{2} \rfloor$$

* In general, $\nu(G) \leq \frac{|V(G)|}{2}$.

Ex. Cycles: C_{2k+1} odd number vertices $k=3$ 

$$\nu(C_{2k+1}) = k$$

$$\tau(C_{2k+1}) = k+1$$

$$\tau(C_{2k+1}) \leq k+1$$

- If v_1, \dots, v_{2k+1} are vertices of C_{2k+1} in order then
 $X = \{v_1, v_3, \dots, v_{2k+1}\}$ is a vertex cover

$$\tau(C_{2k+1}) \geq k+1$$

- because every vertex "covers" only 2 edges, to cover all $2k+1$ edges, we need at least $\lceil \frac{2k+1}{2} \rceil = k+1$ vertices.

Lemma:For any graph G ,

$$\nu(G) \leq \tau(G) \leq 2\nu(G)$$

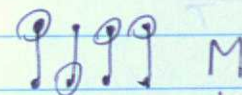
Proof:

$$\nu(G) \leq \tau(G):$$

Let M be a matching in G s.t. $|M| = \nu(G)$

Let X be a vertex cover in G s.t. $|X| = \tau(G)$

$$|X| \geq |M|$$



Every edge $e \in M$ has an end in X and different edges in M must have different vertices of X corresponding to them.

$$\tau(G) \leq 2\nu(G):$$

Let M be a matching in G with $|M| = \nu(G)$

We need to find a vertex cover of G with $\leq 2\nu(G) = 2|M|$ vertices.



Let X be the set of all end of edges in M .
 Then X is a vertex cover.

If not, and e is an edge not covered by X . Then $M \cup \{e\}$ is a matching in G . This contradicts the maximality of M .

Theorem: König

In a bipartite graph G . $\nu(G) = \tau(G)$.

Proof:

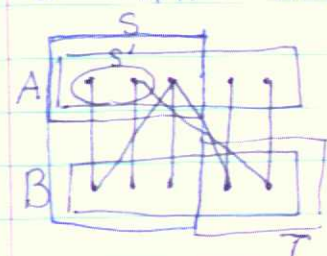
By Lemma, $\nu(G) \leq \tau(G)$

Now, we need to show $\nu(G) \geq \tau(G)$.

Let X be a vertex cover in G . $|X| = \tau(G)$

Our goal is to show that there exists a matching in G with $|M| \geq |X|$.

Let (A, B) be a bipartition of G , and let $S = X \cap A$ and $T = X \cap B$



We find a matching M_1 covering S such that the ends of edges of M_1 in B are in T .

If such a matching does not exist.

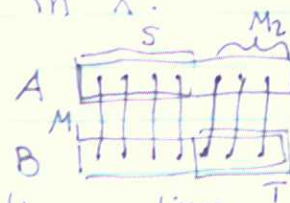
Then by Hall's theorem there exists $S' \subseteq S$ s.t. $|S'| > |N(S')|$ where here $N(S')$ denotes neighbours of S' in $B \setminus T$.

Let $X' = (X \setminus S') \cup N(S')$. $|X'| < |X|$. X' is a vertex cover of G .

If an edge does not have an end in X' then it has an end in S' and so its second end is either in $N(S')$ or T , still in X' .

So M_1 exists.

Symmetrically, there exists a matching M_2 , covering T matching vertices of $A \setminus S$



$$|M_1| = |S|, |M_2| = |T|$$

$M = M_1 \cup M_2$ is a matching. $|M| = |S| + |T| = |X|$.

Ex. Given a 0-1 matrix M .

a	1	0	0	1	0	0
b	1	0	0	0	0	1
c	0	0	0	1	0	1
d	0	1	1	0	1	1
e	1	0	0	1	0	1
	1	2	3	4	5	6

-rank of a matrix is the size of a maximum collection of linearly independent rows (or columns)

$\tau(M)$ (cover number of M) is the size of minimum collection of lines (rows and/or columns) such that deleting them results in all zero matrix.

• $\tau(M) \leq 4$ (in this example)

$$\text{rank}(M) \leq \tau(M)$$

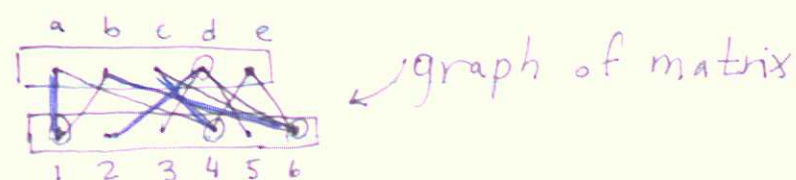
term rank: $\nu(M)$ is the size of the maximum collection of 1's such that no two of them share a row or column.

• $\nu(M) \geq 4$ (in this example)

$$\nu(M) \leq \tau(M)$$

$\nu(M)$ is the lower bound on the maximum possible rank of a matrix obtained from M by replacing 1's by some real numbers. And $\tau(M)$ is an upper bound.

Theorem: König - Egervary
 $\tau(M) = \nu(M)$ for any 0-1 matrix M



Proof:

Given M construct a bipartite graph G with bipartition (R, C) where R is the set of rows of M , and C is the set of columns of M , and $r \in R$ is adjacent to $c \in C$ iff the entry of M on the intersection of r & c is 1.

$\tau(G) = \tau(M)$: deleted rows and columns are vertex covers

$\nu(G) = \nu(M)$: "independent" set of ones correspond to matchings

Phylogenetic Trees

Given a set of species s_1, s_2, \dots, s_m with characteristics c_1, c_2, \dots, c_n .

We want to understand the evolution of these species from a common ancestor.

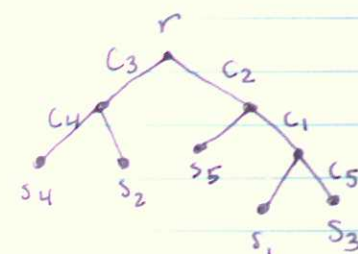
Ex.

	c_1	c_2	c_3	c_4	c_5
s_1	1	1	0	0	0
s_2	0	0	1	0	0
s_3	1	1	0	0	1
s_4	0	0	1	1	0
s_5	0	1	0	0	0

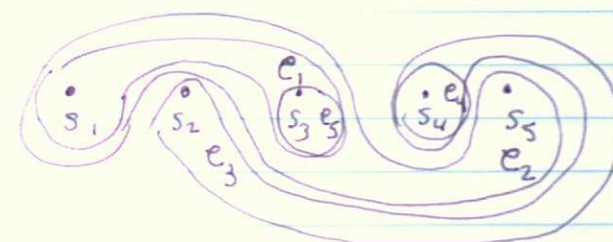
A perfect phylogenetic tree T for these species. T has a root r and m leaves corresponding to s_1, \dots, s_m .

For each characteristic c_i , there is exactly one edge of T labelled by c_i .

Species s_i has a characteristic c_j iff some edge or a path from r to s_i is labelled c_j .



When does a perfect phylogenetic tree exist?
 For each characteristic c_i let \mathcal{C}_i be the set of all species having characteristic c_i .



For any two sets \mathcal{C}_i and \mathcal{C}_j , we will have $\mathcal{C}_i \subseteq \mathcal{C}_j$ or $\mathcal{C}_j \subseteq \mathcal{C}_i$ or $\mathcal{C}_i \cap \mathcal{C}_j = \emptyset$.

This condition is necessary for the existence of the tree. It is also sufficient.

Suppose we want the tree fitting the data in the best way possible.

c_i and c_j are conflicting if $\mathcal{C}_i \cap \mathcal{C}_j \neq \emptyset$ but $\mathcal{C}_i \not\subseteq \mathcal{C}_j$, $\mathcal{C}_j \not\subseteq \mathcal{C}_i$.

To eliminate conflicting pairs, we find minimum vertex cover in the graph of conflicting pairs, and ignore the corresponding characteristics.