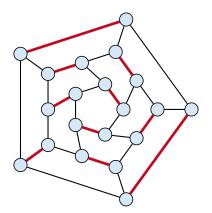
Part V

Matchings

Matching

- ▶ Input: undirected graph G = (V, E).
- ▶ $M \subseteq E$ is a matching if each node appears in at most one edge in M.
- Maximum Matching: find a matching of maximum cardinality



16 Bipartite Matching via Flows

Which flow algorithm to use?

- Generic augmenting path: $\mathcal{O}(m \operatorname{val}(f^*)) = \mathcal{O}(mn)$.
- Capacity scaling: $\mathcal{O}(m^2 \log C) = \mathcal{O}(m^2)$.
- Shortest augmenting path: $O(mn^2)$.

For unit capacity simple graphs shortest augmenting path can be implemented in time $\mathcal{O}(m\sqrt{n})$.

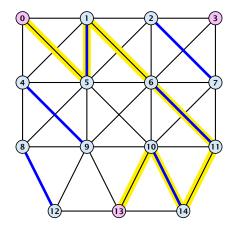
Definitions.

- Given a matching M in a graph G, a vertex that is not incident to any edge of M is called a free vertex w.r..t. M.
- For a matching M a path P in G is called an alternating path if edges in M alternate with edges not in M.
- An alternating path is called an augmenting path for matching M if it ends at distinct free vertices.

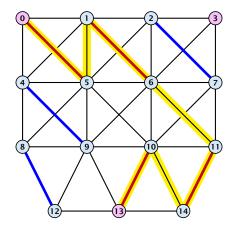
Theorem 89

A matching M is a maximum matching if and only if there is no augmenting path w.r.t. M.

Augmenting Paths in Action



Augmenting Paths in Action



Proof.

- \Rightarrow If M is maximum there is no augmenting path P, because we could switch matching and non-matching edges along P. This gives matching $M' = M \oplus P$ with larger cardinality.
- \Leftarrow Suppose there is a matching M' with larger cardinality. Consider the graph H with edge-set $M' \oplus M$ (i.e., only edges that are in either M or M' but not in both).

Each vertex can be incident to at most two edges (one from M and one from M'). Hence, the connected components are alternating cycles or alternating path.

As |M'| > |M| there is one connected component that is a path P for which both endpoints are incident to edges from M'. P is an alternating path.

Algorithmic idea:

As long as you find an augmenting path augment your matching using this path. When you arrive at a matching for which no augmenting path exists you have a maximum matching.

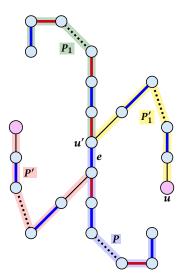
Theorem 90

Let G be a graph, M a matching in G, and let u be a free vertex w.r.t. M. Further let P denote an augmenting path w.r.t. M and let $M' = M \oplus P$ denote the matching resulting from augmenting M with P. If there was no augmenting path starting at u in M then there is no augmenting path starting at u in M'.

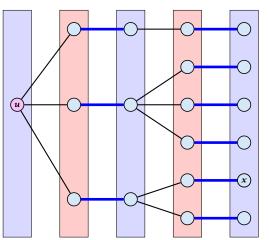
The above theorem allows for an easier implementation of an augmenting path algorithm. Once we checked for augmenting paths starting from u we don't have to check for such paths in future rounds.

Proof

- Assume there is an augmenting path P' w.r.t. M' starting at u.
- If P' and P are node-disjoint, P' is also augmenting path w.r.t. M (∮).
- Let u' be the first node on P' that is in P, and let e be the matching edge from M' incident to u'.
- u' splits P into two parts one of which does not contain e. Call this part P_1 . Denote the sub-path of P' from u to u' with P'_1 .
- $P_1 \circ P_1'$ is augmenting path in M (3).



Construct an alternating tree.

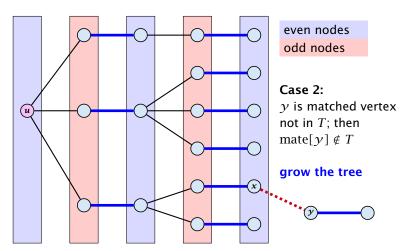


even nodes odd nodes

Case 1: *y* is free vertex not contained in *T*

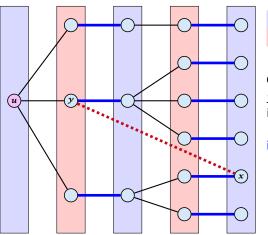
you found alternating path

Construct an alternating tree.





Construct an alternating tree.

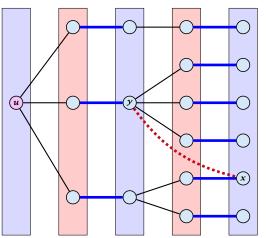


even nodes odd nodes

Case 3: *y* is already contained in *T* as an odd vertex

ignore successor y

Construct an alternating tree.



even nodes odd nodes

Case 4:

y is already contained in T as an even vertex

can't ignore ${m y}$

does not happen in bipartite graphs



```
Algorithm 49 BiMatch(G, match)
 1: for x \in V do mate[x] \leftarrow 0;
 2: r \leftarrow 0; free \leftarrow n;
 3: while free \ge 1 and r < n do
 4:
    r \leftarrow r + 1
 5: if mate[r] = 0 then
           for i = 1 to n do parent[i'] \leftarrow 0
 6:
          Q \leftarrow \emptyset; Q. append(r); aug \leftarrow false;
 7:
 8:
           while aug = false and Q \neq \emptyset do
9:
              x \leftarrow Q. dequeue();
10:
               for y \in A_x do
11:
                  if mate[\gamma] = 0 then
12:
                      augm(mate, parent, y);
13:
                      aua ← true:
                      free \leftarrow free - 1:
14:
15:
                  else
16:
                      if parent[y] = 0 then
                         parent[y] \leftarrow x;
17:
18:
                         Q. enqueue(mate[y]);
```

contain a step by step explanation. graph $G = (S \cup S', E)$ $S = \{1, ..., n\}$

The lecture slides

 $S' = \{1', \dots, n'\}$

Weighted Bipartite Matching/Assignment

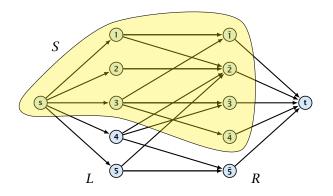
- ▶ Input: undirected, bipartite graph $G = L \cup R, E$.
- ▶ an edge $e = (\ell, r)$ has weight $w_e \ge 0$
- find a matching of maximum weight, where the weight of a matching is the sum of the weights of its edges

Simplifying Assumptions (wlog [why?]):

- ightharpoonup assume that |L| = |R| = n
- ▶ assume that there is an edge between every pair of nodes $(\ell, r) \in V \times V$
- can assume goal is to construct maximum weight perfect matching

Theorem 91 (Halls Theorem)

A bipartite graph $G = (L \cup R, E)$ has a perfect matching if and only if for all sets $S \subseteq L$, $|\Gamma(S)| \ge |S|$, where $\Gamma(S)$ denotes the set of nodes in R that have a neighbour in S.



Halls Theorem

Proof:

- Of course, the condition is necessary as otherwise not all nodes in S could be matched to different neighbours.
- \Rightarrow For the other direction we need to argue that the minimum cut in the graph G' is at least |L|.
 - Let S denote a minimum cut and let $L_S \stackrel{\text{\tiny def}}{=} L \cap S$ and $R_S \stackrel{\text{\tiny def}}{=} R \cap S$ denote the portion of S inside L and R, respectively.
 - ▶ Clearly, all neighbours of nodes in L_S have to be in S, as otherwise we would cut an edge of infinite capacity.
 - ► This gives $R_S \ge |\Gamma(L_S)|$.
 - ▶ The size of the cut is $|L| |L_S| + |R_S|$.
 - ▶ Using the fact that $|\Gamma(L_S)| \ge L_S$ gives that this is at least |L|.

Algorithm Outline

Idea:

We introduce a node weighting \vec{x} . Let for a node $v \in V$, $x_v \in \mathbb{R}$ denote the weight of node v.

Suppose that the node weights dominate the edge-weights in the following sense:

$$x_u + x_v \ge w_e$$
 for every edge $e = (u, v)$.

- Let $H(\vec{x})$ denote the subgraph of G that only contains edges that are tight w.r.t. the node weighting \vec{x} , i.e. edges e = (u, v) for which $w_e = x_u + x_v$.
- Try to compute a perfect matching in the subgraph $H(\vec{x})$. If you are successful you found an optimal matching.

Algorithm Outline

Reason:

▶ The weight of your matching M^* is

$$\sum_{(u,v)\in M^*} w_{(u,v)} = \sum_{(u,v)\in M^*} (x_u + x_v) = \sum_v x_v \ .$$

Any other perfect matching M (in G, not necessarily in $H(\vec{x})$) has

$$\sum_{(u,v)\in M} w_{(u,v)} \leq \sum_{(u,v)\in M} (x_u + x_v) = \sum_v x_v \ .$$

Algorithm Outline

What if you don't find a perfect matching?

Then, Halls theorem guarantees you that there is a set $S \subseteq L$, with $|\Gamma(S)| < |S|$, where Γ denotes the neighbourhood w.r.t. the subgraph $H(\vec{x})$.

Idea: reweight such that:

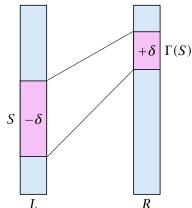
- the total weight assigned to nodes decreases
- the weight function still dominates the edge-weights

If we can do this we have an algorithm that terminates with an optimal solution (we analyze the running time later).

Changing Node Weights

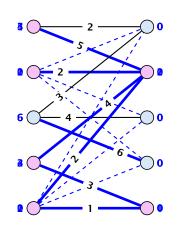
Increase node-weights in $\Gamma(S)$ by $+\delta$, and decrease the node-weights in S by $-\delta$.

- Total node-weight decreases.
- ▶ Only edges from S to $R \Gamma(S)$ decrease in their weight.
- Since, none of these edges is tight (otw. the edge would be contained in $H(\vec{x})$, and hence would go between S and $\Gamma(S)$) we can do this decrement for small enough $\delta>0$ until a new edge gets tight.



Edges not drawn have weight 0.

$$\delta = 1 \ \delta = 1$$



Analysis

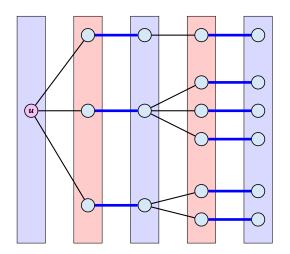
How many iterations do we need?

- One reweighting step increases the number of edges out of S by at least one.
- Assume that we have a maximum matching that saturates the set $\Gamma(S)$, in the sense that every node in $\Gamma(S)$ is matched to a node in S (we will show that we can always find S and a matching such that this holds).
- ▶ This matching is still contained in the new graph, because all its edges either go between $\Gamma(S)$ and S or between L-S and $R-\Gamma(S)$.
- Hence, reweighting does not decrease the size of a maximum matching in the tight sub-graph.

Analysis

- We will show that after at most n reweighting steps the size of the maximum matching can be increased by finding an augmenting path.
- This gives a polynomial running time.

Construct an alternating tree.



Analysis

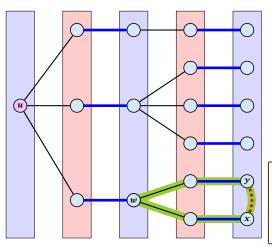
How do we find *S*?

- Start on the left and compute an alternating tree, starting at any free node u.
- ▶ If this construction stops, there is no perfect matching in the tight subgraph (because for a perfect matching we need to find an augmenting path starting at *u*).
- The set of even vertices is on the left and the set of odd vertices is on the right and contains all neighbours of even nodes.
- ▶ All odd vertices are matched to even vertices. Furthermore, the even vertices additionally contain the free vertex u. Hence, $|V_{\rm odd}| = |\Gamma(V_{\rm even})| < |V_{\rm even}|$, and all odd vertices are saturated in the current matching.

Analysis

- The current matching does not have any edges from $V_{\rm odd}$ to $L \setminus V_{\rm even}$ (edges that may possibly be deleted by changing weights).
- After changing weights, there is at least one more edge connecting $V_{\rm even}$ to a node outside of $V_{\rm odd}$. After at most n reweights we can do an augmentation.
- A reweighting can be trivially performed in time $\mathcal{O}(n^2)$ (keeping track of the tight edges).
- An augmentation takes at most O(n) time.
- In total we obtain a running time of $\mathcal{O}(n^4)$.
- ▶ A more careful implementation of the algorithm obtains a running time of $\mathcal{O}(n^3)$.

Construct an alternating tree.



even nodes odd nodes

Case 4:

y is already contained in T as an even vertex

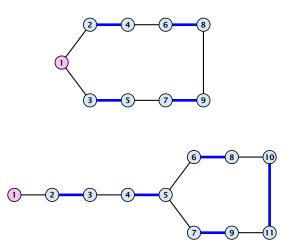
can't ignore y

The cycle $w \leftrightarrow y - x \leftrightarrow w$ is called a blossom. w is called the base of the blossom (even node!!!). The path u-w is called the stem of the blossom.

Definition 92

A flower in a graph G = (V, E) w.r.t. a matching M and a (free) root node r, is a subgraph with two components:

- A stem is an even length alternating path that starts at the root node r and terminates at some node w. We permit the possibility that r = w (empty stem).
- ▶ A blossom is an odd length alternating cycle that starts and terminates at the terminal node *w* of a stem and has no other node in common with the stem. *w* is called the base of the blossom.

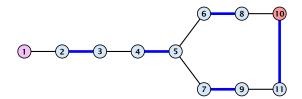


Properties:

- 1. A stem spans $2\ell+1$ nodes and contains ℓ matched edges for some integer $\ell \geq 0$.
- **2.** A blossom spans 2k + 1 nodes and contains k matched edges for some integer $k \ge 1$. The matched edges match all nodes of the blossom except the base.
- 3. The base of a blossom is an even node (if the stem is part of an alternating tree starting at r).

Properties:

- 4. Every node x in the blossom (except its base) is reachable from the root (or from the base of the blossom) through two distinct alternating paths; one with even and one with odd length.
- 5. The even alternating path to x terminates with a matched edge and the odd path with an unmatched edge.



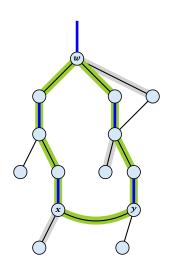
Shrinking Blossoms

When during the alternating tree construction we discover a blossom B we replace the graph G by G' = G/B, which is obtained from G by contracting the blossom B.

- Delete all vertices in B (and its incident edges) from G.
- Add a new (pseudo-)vertex b. The new vertex b is connected to all vertices in $V \setminus B$ that had at least one edge to a vertex from B.

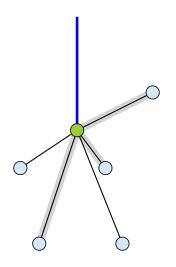
Shrinking Blossoms

- Edges of T that connect a node u not in B to a node in B become tree edges in T' connecting u to b.
- Matching edges (there is at most one) that connect a node u not in B to a node in B become matching edges in M'.
- Nodes that are connected in G to at least one node in B become connected to b in G'.



Shrinking Blossoms

- Edges of T that connect a node u not in B to a node in B become tree edges in T' connecting u to b.
- Matching edges (there is at most one) that connect a node u not in B to a node in B become matching edges in M'.
- Nodes that are connected in G to at least one node in B become connected to b in G'.



Example: Blossom Algorithm

Animation of Blossom Shrinking algorithm is only available in the lecture version of the slides.

Assume that in G we have a flower w.r.t. matching M. Let r be the root, B the blossom, and w the base. Let graph G' = G/B with pseudonode b. Let M' be the matching in the contracted graph.

Lemma 93

If G' contains an augmenting path P' starting at r (or the pseudo-node containing r) w.r.t. the matching M' then Gcontains an augmenting path starting at γ w.r.t. matching M.

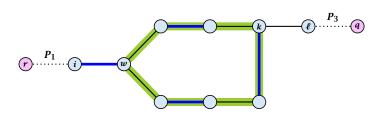
Proof.

If P' does not contain b it is also an augmenting path in G.

Case 1: non-empty stem

Next suppose that the stem is non-empty.



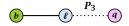


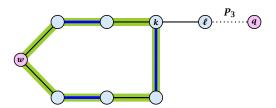
- After the expansion ℓ must be incident to some node in the blossom. Let this node be k.
- ▶ If $k \neq w$ there is an alternating path P_2 from w to k that ends in a matching edge.
- ▶ $P_1 \circ (i, w) \circ P_2 \circ (k, \ell) \circ P_3$ is an alternating path.
- ▶ If k = w then $P_1 \circ (i, w) \circ (w, \ell) \circ P_3$ is an alternating path.

Proof.

Case 2: empty stem

If the stem is empty then after expanding the blossom, w = r.





▶ The path $r \circ P_2 \circ (k, \ell) \circ P_3$ is an alternating path.

Lemma 94

If G contains an augmenting path P from r to q w.r.t. matching M then G' contains an augmenting path from r (or the pseudo-node containing r) to q w.r.t. M'.

Proof.

- ▶ If *P* does not contain a node from *B* there is nothing to prove.
- ▶ We can assume that r and q are the only free nodes in G.

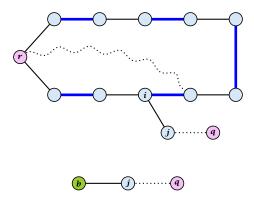
Case 1: empty stem

Let i be the last node on the path P that is part of the blossom.

P is of the form $P_1 \circ (i, j) \circ P_2$, for some node j and (i, j) is unmatched.

 $(b, j) \circ P_2$ is an augmenting path in the contracted network.

Illustration for Case 1:



Case 2: non-empty stem

Let P_3 be alternating path from r to w; this exists because r and w are root and base of a blossom. Define $M_+ = M \oplus P_3$.

In M_+ , γ is matched and w is unmatched.

G must contain an augmenting path w.r.t. matching M_+ , since M and M_+ have same cardinality.

This path must go between w and q as these are the only unmatched vertices w.r.t. M_{+} .

For M'_+ the blossom has an empty stem. Case 1 applies.

G' has an augmenting path w.r.t. M'_+ . It must also have an augmenting path w.r.t. M', as both matchings have the same cardinality.

This path must go between r and q.

	The lecture slides	
	contain a step by	
Algorithm 50 search(r , found)	step explanation.	
1: set $\bar{A}(i) \leftarrow A(i)$ for all nodes i		
2: found ← false		
3: unlabel all nodes;		

4: give an even label to r and initialize $list \leftarrow \{r\}$

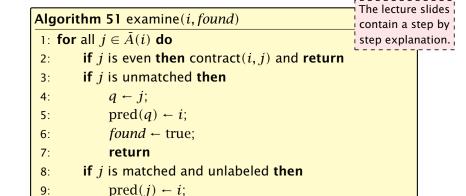
5: while $list \neq \emptyset$ do

6: delete a node i from list

7: examine(i, found)

8: **if** *found* = true **then return**

Search for an augmenting path starting at r.



Examine the neighbours of a node *i*

 $pred(mate(j)) \leftarrow j;$ add mate(j) to list

10:

11:

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
- 3: label b even and add to list
- 4: update $\bar{A}(j) \leftarrow \bar{A}(j) \cup \{b\}$ for each $j \in \bar{A}(b)$
- 5: form a circular double linked list of nodes in B
- 6: delete nodes in B from the graph

Contract blossom identified by nodes *i* and *j*

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
- 3: label b even and add to list
- 4: update $\bar{A}(j) \leftarrow \bar{A}(j) \cup \{b\}$ for each $j \in \bar{A}(b)$
- 5: form a circular double linked list of nodes in B
- 6: delete nodes in B from the graph

Get all nodes of the blossom.

Time: $\mathcal{O}(m)$

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
- 3: label b even and add to list
- 4: update $\bar{A}(j) \leftarrow \bar{A}(j) \cup \{b\}$ for each $j \in \bar{A}(b)$
- 5: form a circular double linked list of nodes in B
- 6: delete nodes in B from the graph

Identify all neighbours of b.

Time: $\mathcal{O}(m)$ (how?)

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
- 3: label b even and add to list
- 4: update $\bar{A}(j) \leftarrow \bar{A}(j) \cup \{b\}$ for each $j \in \bar{A}(b)$
- 5: form a circular double linked list of nodes in B
- 6: delete nodes in B from the graph

b will be an even node, and it has unexamined neighbours.

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
- 3: label *b* even and add to *list*
- 4: update $\bar{A}(j) \leftarrow \bar{A}(j) \cup \{b\}$ for each $j \in \bar{A}(b)$
- 5: form a circular double linked list of nodes in B
- 6: delete nodes in B from the graph

Every node that was adjacent to a node in B is now adjacent to b

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
- 3: label *b* even and add to *list*
- 4: update $\bar{A}(j) \leftarrow \bar{A}(j) \cup \{b\}$ for each $j \in \bar{A}(b)$
- 5: form a circular double linked list of nodes in B
- 6: delete nodes in B from the graph

Only for making a blossom expansion easier.

- 1: trace pred-indices of i and j to identify a blossom B
- 2: create new node b and set $\bar{A}(b) \leftarrow \bigcup_{x \in B} \bar{A}(x)$
- 3: label b even and add to list
- 4: update $\bar{A}(j) \leftarrow \bar{A}(j) \cup \{b\}$ for each $j \in \bar{A}(b)$
- 5: form a circular double linked list of nodes in B
- 6: delete nodes in *B* from the graph

Only delete links from nodes not in B to B.

When expanding the blossom again we can recreate these links in time O(m).

Analysis

- A contraction operation can be performed in time $\mathcal{O}(m)$. Note, that any graph created will have at most m edges.
- The time between two contraction-operation is basically a BFS/DFS on a graph. Hence takes time $\mathcal{O}(m)$.
- ► There are at most *n* contractions as each contraction reduces the number of vertices.
- The expansion can trivially be done in the same time as needed for all contractions.
- An augmentation requires time $\mathcal{O}(n)$. There are at most n of them.
- In total the running time is at most

$$n \cdot (\mathcal{O}(mn) + \mathcal{O}(n)) = \mathcal{O}(mn^2)$$
.



Example: Blossom Algorithm

Animation of Blossom Shrinking algorithm is only available in the lecture version of the slides.