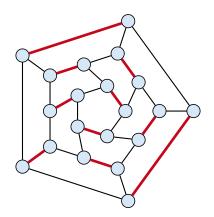
### Part V

# **Matchings**

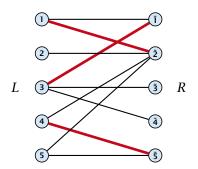
### **Matching**

- ▶ 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



### **Bipartite Matching**

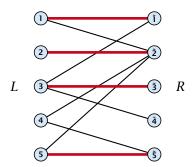
- ▶ Input: undirected, bipartite graph  $G = (L \uplus R, E)$ .
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### **Bipartite Matching**

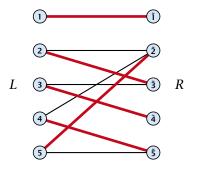
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### **Bipartite Matching**

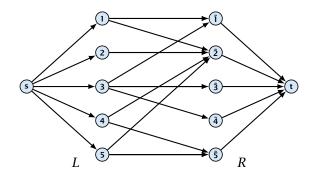
- ▶ A matching M is perfect if it is of cardinality |M| = |V|/2.
- For a bipartite graph  $G = (L \uplus R, E)$  this means |M| = |L| = |R| = n.



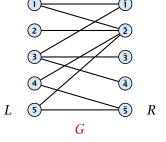


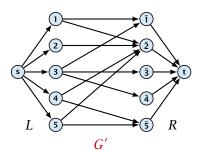
### 17 Bipartite Matching via Flows

- ▶ Input: undirected, bipartite graph  $G = (L \uplus R \uplus \{s, t\}, E')$ .
- ▶ Direct all edges from *L* to *R*.
- Add source s and connect it to all nodes on the left.
- Add t and connect all nodes on the right to t.
- All edges have unit capacity.



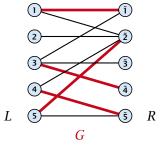
- Given a maximum matching M of cardinality k.
- ▶ Consider flow f that sends one unit along each of k paths.
- f is a flow and has cardinality k.

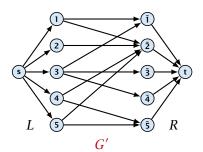






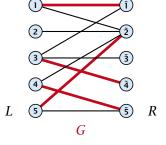
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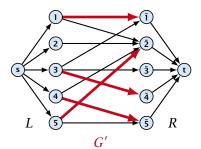






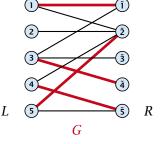
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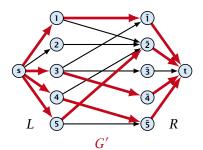






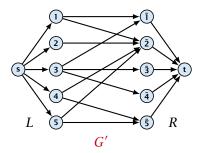
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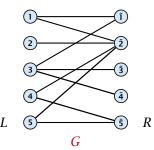






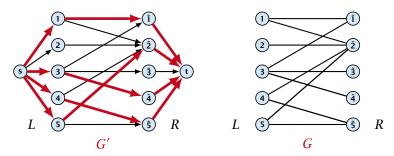
- Let f be a maxflow in G' of value k
- ▶ Integrality theorem  $\Rightarrow k$  integral; we can assume f is 0/1.
- Consider M= set of edges from L to R with f(e) = 1.
- ▶ Each node in *L* and *R* participates in at most one edge in *M*.
- |M| = k, as the flow must use at least k middle edges.



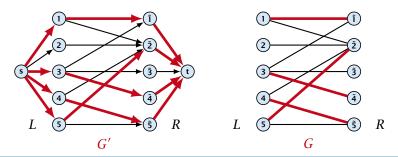




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### 17 Bipartite Matching via Flows

#### Which flow algorithm to use?

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



#### 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 1

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



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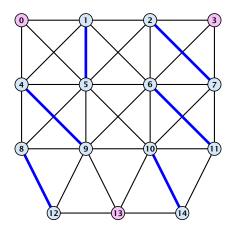
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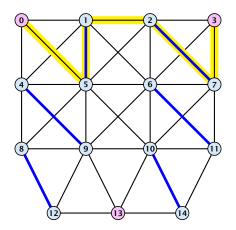
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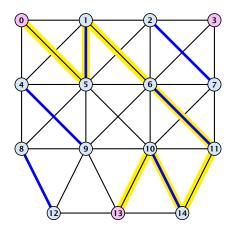




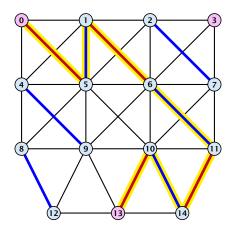




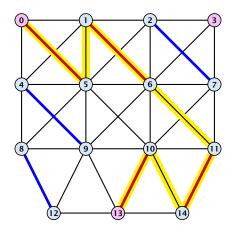




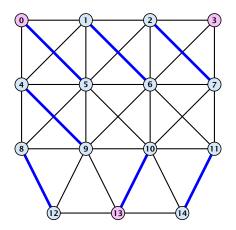














#### Proof.

- $\Rightarrow$  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.



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#### 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 2

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'.



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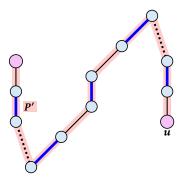






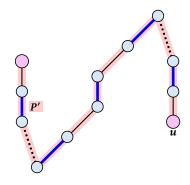
#### **Proof**

Assume there is an augmenting path P' w.r.t. M' starting at u.



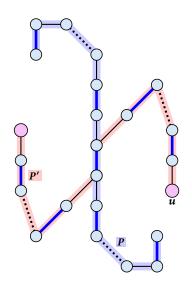


- Assume there is an augmenting path P' w.r.t. M' starting at u.
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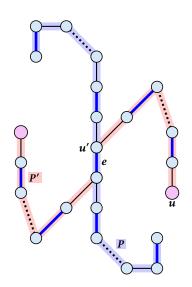


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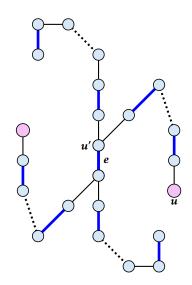


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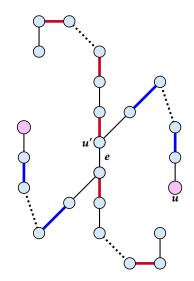


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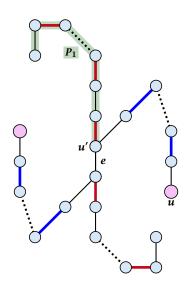


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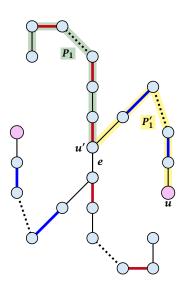


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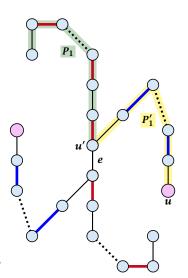






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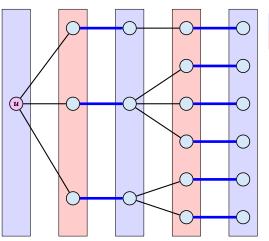
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- u' splits P into two parts one of which does not contain e. Call this part  $P_1$ . Denote the sub-path of P'from u to u' with  $P'_1$ .
- ▶  $P_1 \circ P_1'$  is augmenting path in M (§).





FADS

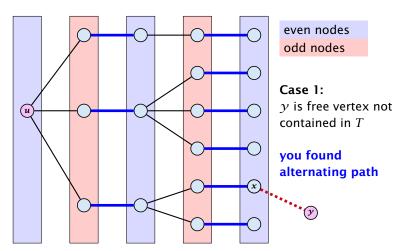
### Construct an alternating tree.



even nodes odd nodes

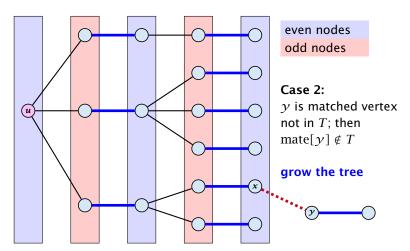


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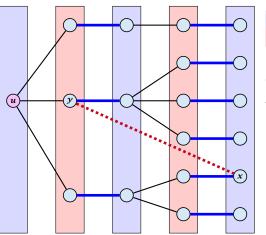
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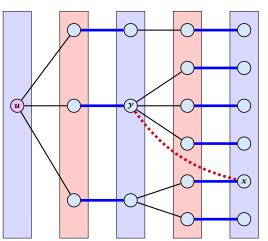
even nodes odd nodes

**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 52 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
6:
          for i = 1 to m do parent[i'] \leftarrow 0
7:
    Q \leftarrow \emptyset; Q. append(r); aug \leftarrow false;
          while aug = false and Q \neq \emptyset do
8:
```

 $x \leftarrow O.$  dequeue():

*aug* ← true;

 $free \leftarrow free - 1$ ;

for  $\gamma \in A_{\chi}$  do

else

9:

10:

11:

12:

13:

14.

15:

16:

17:

18:

graph  $G = (S \cup S', E)$  $S = \{1, ..., n\}$  $S' = \{1', \dots, n'\}$ 

if mate[y] = 0 then augm(mate, parent, y);if parent[v] = 0 then  $parent[y] \leftarrow x;$ Q. enqueue(mate[y]);

```
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- 3: while  $free \ge 1$  and r < n do
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- 5: **if** mate[r] = 0 **then**
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- 9:  $x \leftarrow O.$  dequeue():
- 10: for  $\gamma \in A_{\chi}$  do
- 11: if mate[y] = 0 then
- 12: augm(mate, parent, y);

else

13:

14.

15:

16:

17:

18:

**for** i = 1 **to** m **do**  $parent[i'] \leftarrow 0$ 

*aug* ← true;

 $free \leftarrow free - 1$ :

if parent[y] = 0 then

 $parent[y] \leftarrow x$ ; Q. enqueue(mate[y]);

- empty matching

start with an

```
Algorithm 52 BiMatch(G, match)
```

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- 12: augm(mate, parent, y);
- 13: *aug* ← true;
- 14.  $free \leftarrow free - 1$ ; else
- 15: 16:
- if parent[y] = 0 then 17:  $parent[y] \leftarrow x$ ; Q. enqueue(mate[y]); 18:

free: number of unmatched nodes in S r: root of current tree

# **Algorithm 52** BiMatch(*G*, *match*)

1: for  $x \in V$  do  $mate[x] \leftarrow 0$ : 2:  $r \leftarrow 0$ ; free  $\leftarrow n$ ;

6:

7:

8: 9:

10:

11:

12:

13:

14.

3: while  $free \ge 1$  and r < n do

4:  $r \leftarrow r + 1$ 

5: **if** mate[r] = 0 **then** 

for i = 1 to m do parent[i']  $\leftarrow 0$ 

 $Q \leftarrow \emptyset$ ; Q. append(r); aug  $\leftarrow$  false;

while aug = false and  $Q \neq \emptyset$  do

 $x \leftarrow O.$  dequeue():

for  $\gamma \in A_{\chi}$  do

if mate[y] = 0 then augm(mate, parent, y);

*aug* ← true;

 $free \leftarrow free - 1$ :

15: else 16: if parent[y] = 0 then 17:  $parent[y] \leftarrow x$ ; Q. enqueue(mate[y]); 18:

as long as there are unmatched nodes and we did not yet try to grow from all nodes we continue

```
Algorithm 52 BiMatch(G, match)
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 2: r \leftarrow 0; free \leftarrow n;
 3: while free \ge 1 and r < n do
 4: r \leftarrow r + 1
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    Q \leftarrow \emptyset; Q. append(r); aug \leftarrow false;
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10:

11.

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14.

15:

16:

17:

18:

while aug = false and  $Q \neq \emptyset$  do

*aug* ← true;

 $x \leftarrow O.$  dequeue():

for  $\gamma \in A_{\chi}$  do

else

 $\gamma$  is the new node that we grow from.

```
if mate[y] = 0 then
   augm(mate, parent, y);
   free \leftarrow free - 1:
   if parent[y] = 0 then
      parent[y] \leftarrow x;
```

Q. enqueue(mate[y]);

```
Algorithm 52 BiMatch(G, match)
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18:

If  $\gamma$  is free start tree construction

```
if mate[y] = 0 then
   augm(mate, parent, y);
   free \leftarrow free - 1;
   if parent[y] = 0 then
      parent[y] \leftarrow x;
```

Q. enqueue(mate[y]);

## **Algorithm 52** 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:  $r \leftarrow r + 1$
- if mate[r] = 0 then
- 6: for i = 1 to m do  $parent[i'] \leftarrow 0$  $Q \leftarrow \emptyset$ ; Q. append(r); aug  $\leftarrow$  false; 7:
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- 9:  $x \leftarrow O.$  dequeue():
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- 11: if mate[y] = 0 then
- 12: augm(mate, parent, y);
- 13: *aug* ← true; 14.  $free \leftarrow free - 1$ :
  - else if parent[y] = 0 then
- 15: 16: 17:  $parent[y] \leftarrow x$ ; Q. enqueue(mate[y]); 18:

Initialize an empty tree. Note that only nodes i'have parent pointers.

# **Algorithm 52** 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**
- 6: for i = 1 to m do parent[i']  $\leftarrow 0$
- $Q \leftarrow \emptyset$ ; Q. append(r); aug  $\leftarrow$  false; 7:
- while aug = false and  $Q \neq \emptyset$  do 8:
- 9:  $x \leftarrow O.$  dequeue(): 10: for  $\gamma \in A_{\kappa}$  do
- 11: if mate[y] = 0 then
- 12: augm(mate, parent, y);
- 13: *aug* ← true;
- 14.  $free \leftarrow free - 1$ : 15: else
- 16: if parent[y] = 0 then 17:  $parent[y] \leftarrow x$ ; Q. enqueue(mate[y]); 18:

Q is a queue (BFS!!!). aua is a Boolean that stores whether we already found an augmenting path.

## **Algorithm 52** BiMatch(*G*, *match*) 1: for $x \in V$ do $mate[x] \leftarrow 0$ :

2:  $r \leftarrow 0$ ; free  $\leftarrow n$ ;

6:

7:

8: 9:

10:

- 3: while  $free \ge 1$  and r < n do
- 4:  $r \leftarrow r + 1$
- 5: **if** mate[r] = 0 **then** 
  - **for** i = 1 **to** m **do**  $parent[i'] \leftarrow 0$
  - $Q \leftarrow \emptyset$ ; Q. append(r); aug  $\leftarrow$  false;
    - while aug = false and  $Q \neq \emptyset$  do
      - $x \leftarrow O.$  dequeue():
      - for  $\gamma \in A_{\chi}$  do
- 11: if mate[y] = 0 then
- 12: augm(mate, parent, y);
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- 14.  $free \leftarrow free - 1$ :
  - else
- 15: 16: if parent[y] = 0 then 17:  $parent[y] \leftarrow x$ ; Q. enqueue(mate[y]); 18:

as long as we did not augment and there are still unexamined leaves continue...

```
Algorithm 52 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
6:
           for i = 1 to m do parent[i'] \leftarrow 0
 7:
    Q \leftarrow \emptyset; Q. append(r); aug \leftarrow false;
           while aug = false and Q \neq \emptyset do
8:
               x \leftarrow Q. dequeue();
9:
10:
               for \gamma \in A_{\kappa} do
11:
                   if mate[y] = 0 then
```

else

augm(mate, parent, y);

if parent[y] = 0 then

 $parent[y] \leftarrow x;$ Q. enqueue(mate[y]);

*aug* ← true;

 $free \leftarrow free - 1$ :

12:

13:

14.

15:

16:

17:

18:

take next unexamined leaf

## **Algorithm 52** 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**
- 6: for i = 1 to m do parent[i']  $\leftarrow 0$
- 7:  $Q \leftarrow \emptyset$ ; Q. append(r); aug  $\leftarrow$  false;
- while aug = false and  $Q \neq \emptyset$  do 8:
- 9:  $x \leftarrow O.$  dequeue():
- 10: for  $\gamma \in A_{\kappa}$  do
- 11: if  $mate[\gamma] = 0$  then
- 12: augm(mate, parent, y);
- 13: *aug* ← true;
- 14.  $free \leftarrow free - 1$ : 15: else 16:
- if parent[y] = 0 then 17:  $parent[y] \leftarrow x$ ; Q. enqueue(mate[y]); 18:

if x has unmatched neighbour we found an augmenting path (note that  $y \neq r$  because we are in a bipartite graph)

```
Algorithm 52 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
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                      augm(mate, parent, y);
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                      aug ← true;
14.
                      free \leftarrow free - 1:
15:
                  else
16:
                      if parent[y] = 0 then
17:
                          parent[y] \leftarrow x;
```

18:

Q. enqueue(mate[y]);

do an augmentation...

## **Algorithm 52** 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**
- 6: for i = 1 to m do parent[i']  $\leftarrow 0$ 7:  $Q \leftarrow \emptyset$ ; Q. append(r); aug  $\leftarrow$  false;
- while aug = false and  $Q \neq \emptyset$  do 8:
- 9:  $x \leftarrow O.$  dequeue():
- 10: for  $\gamma \in A_{\chi}$  do
- 11: if mate[y] = 0 then

18:

- 12: augm(mate, parent, y);13: *aug* ← true;
- 14:  $free \leftarrow free - 1$ : 15: else
- 16: if parent[y] = 0 then 17:  $parent[y] \leftarrow x$ ; Q. enqueue(mate[y]);

setting aug = trueensures that the tree construction will not continue

```
Algorithm 52 BiMatch(G, match)

1: for x \in V do mate[x] \leftarrow 0;

2: r \leftarrow 0; free \leftarrow n;

3: while free \geq 1 and r < n do

4: r \leftarrow r + 1

5: if mate[r] = 0 then

6: for i = 1 to m do parent[i'] \leftarrow 0

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8: while aug = false and Q \neq \emptyset do
```

 $x \leftarrow O.$  dequeue():

if mate[y] = 0 then

 $free \leftarrow free - 1$ :

aug ← true;

augm(mate, parent, y);

if parent[y] = 0 then

Q. enqueue(mate[y]);

 $parent[y] \leftarrow x$ ;

for  $\gamma \in A_{\chi}$  do

else

9:

10:

11:

12:

13:

14:

15:

16:

17:

18:

reduce number of free nodes

```
Algorithm 52 BiMatch(G, match)
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              for \gamma \in A_{\chi} do
11:
                  if mate[y] = 0 then
12:
                      augm(mate, parent, y);
13:
                      aug ← true;
```

else

 $free \leftarrow free - 1$ :

if parent[y] = 0 then  $parent[y] \leftarrow x$ ;

Q. enqueue(mate[y]);

14.

15: 16:

17:

18:

if  $\boldsymbol{\mathcal{Y}}$  is not in the tree yet

```
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12:
                      augm(mate, parent, y);
13:
                      aug ← true;
14.
                      free \leftarrow free - 1:
```

else

if parent[v] = 0 then

Q. enqueue(mate[y]);

 $parent[y] \leftarrow x$ ;

15:

16:

17:

18:

...put it into the tree

## **Algorithm 52** BiMatch(G, match)

1: for  $x \in V$  do  $mate[x] \leftarrow 0$ : 2:  $r \leftarrow 0$ ; free  $\leftarrow n$ ;

6:

7:

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10:

11:

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$$i = 1$$
 to  $m$  do  $parent[i'] \leftarrow 0$ 

$$Q \leftarrow \emptyset$$
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while aug = false and  $Q \neq \emptyset$  do  $x \leftarrow O.$  dequeue():

# for $\gamma \in A_{\chi}$ do

# if mate[y] = 0 then

- augm(mate, parent, y);
- *aug* ← true;
- 14.  $free \leftarrow free - 1$ : else
- 15: 16: if parent[y] = 0 then  $parent[y] \leftarrow x$ ; 17: O. enqueue(mate[v]); 18:

add its buddy to the set of unexamined leaves

# 19 Weighted Bipartite Matching

## 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?]):

- assume that |L| = |R| = n
- assume that there is an edge between every pair of nodes  $(\ell,r) \in V \times V$





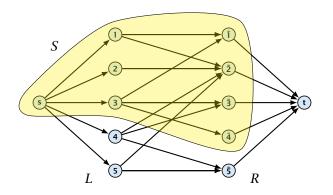
# **Weighted Bipartite Matching**

### Theorem 3 (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.



# 19 Weighted Bipartite Matching



- 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|.

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- $\Rightarrow$  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  $L_S \cong L \cap S$  and  $R_S \cong R \cap S$  denote the portion of S inside L and R, respectively.
  - Clearly, all neighbours of nodes in L<sub>S</sub> 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|.



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We introduce a node weighting  $\vec{x}$ . Let for a node  $v \in V$ ,  $x_v \ge 0$  denote the weight of node v.

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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.



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#### 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 matching M has

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



#### 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  $\Gamma$  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).



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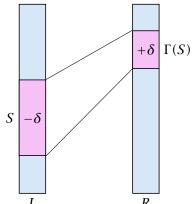
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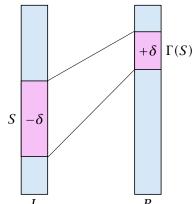


- ► Total node-weight decreases
- ▶ Only edges from S to  $R \Gamma(S)$  decrease in their weight.
- Since, none of these edges is tight (otw. the edge would be contained in  $H(\vec{x})$ , and hence would go between S and  $\Gamma(S)$ ) we can do this decrement for small enough  $\delta>0$  until a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



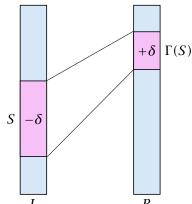


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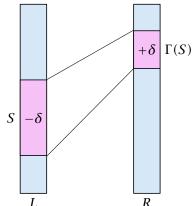


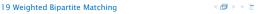


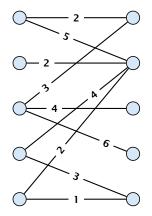
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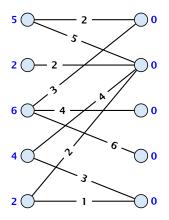
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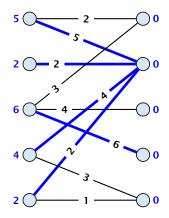




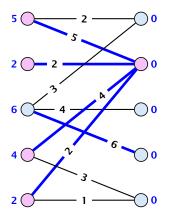






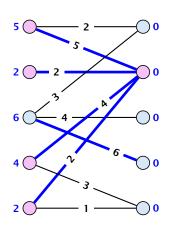


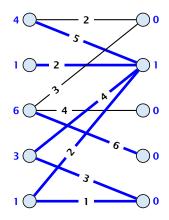




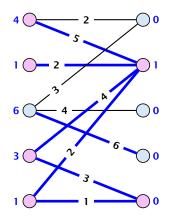


$$\delta = 1$$



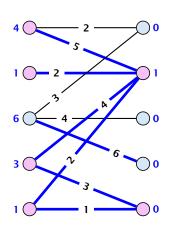


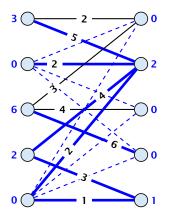




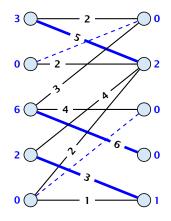


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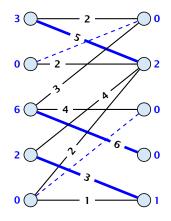














- One reweighting step increases the number of edges out of S by at least one.
- Assume that we have a maximum matching that saturates the set  $\Gamma(S)$ , in the sense that every node in  $\Gamma(S)$  is matched to a node in S (we will show that we can always find S and a matching such that this holds).
- ▶ 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.

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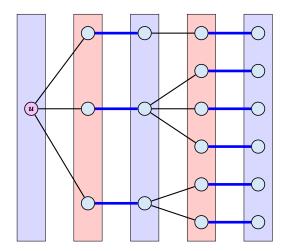
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- We will show that after at most n reweighting steps the size of the maximum matching can be increased by finding an augmenting path.
- This gives a polynomial running time.

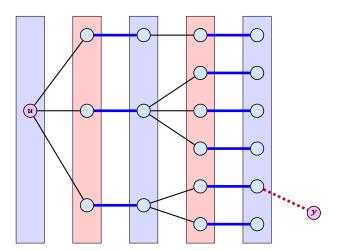
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#### Construct an alternating tree.



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- Start on the left and compute an alternating tree, starting at any free node u.
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- 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.



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- ▶ The current matching does not have any edges from  $V_{\rm odd}$  to outside of  $L \setminus V_{\rm even}$  (edges that may possibly be deleted by changing weights).
- After changing weights, there is at least one more edge connecting  $V_{\mathrm{even}}$  to a node outside of  $V_{\mathrm{odd}}$ . 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 otain a running time of  $O(n^4)$ .
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# A Fast Matching Algorithm

## **Algorithm 53** Bimatch-Hopcroft-Karp(G)

3: let  $\mathcal{P} = \{P_1, \dots, P_k\}$  be maximal set of 4: vertex-disjoint, shortest augmenting path w.r.t. M.

5:  $M \leftarrow M \oplus (P_1 \cup \cdots \cup P_k)$ 

6: until  $\mathcal{P} = \emptyset$ 

7: return M

We call one iteration of the repeat-loop a phase of the algorithm.

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- ▶ Consider the graph  $G = (V, M \oplus M^*)$ , and mark edges in this graph blue if they are in M and red if they are in  $M^*$ .
- ▶ The connected components of *G* are cycles and paths
- ▶ The graph contains  $k ext{ # } |M^*| |M|$  more red edges than blue edges.
- ► Hence, there are at least *k* components that form a path starting and ending with a blue edge. These are augmenting paths w.r.t. *M*.



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- Let  $P_1, ..., P_k$  be a maximal collection of vertex-disjoint, shortest augmenting paths w.r.t. M (let  $\ell = |P_i|$ ).
- $M' \stackrel{\text{def}}{=} M \oplus (P_1 \cup \cdots \cup P_k) = M \oplus P_1 \oplus \cdots \oplus P_k.$
- Let P be an augmenting path in M'.

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The set  $A \cong M \oplus (M' \oplus P) = (P_1 \cup \cdots \cup P_k) \oplus P$  contains at least  $(k+1)\ell$  edges.



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- ▶ This edge is not contained in *A*.
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#### Proof

The symmetric difference between M and  $M^*$  contains  $|M^*| - |M|$  vertex-disjoint augmenting paths. Each of these paths contains at least  $\ell+1$  vertices. Hence, there can be at most  $\frac{|V|}{\ell+1}$  of them.



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#### Lemma 7

The Hopcroft-Karp algorithm requires at most  $2\sqrt{|V|}$  phases.

- After iteration  $\lfloor \sqrt{|V|} \rfloor$  the length of a shortest augmenting path must be at least  $\lfloor \sqrt{|V|} \rfloor + 1 \geq \sqrt{|V|}$ .
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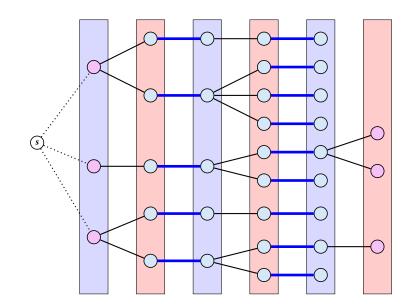


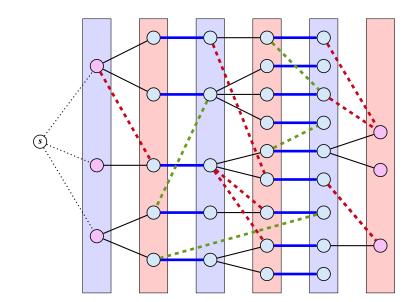
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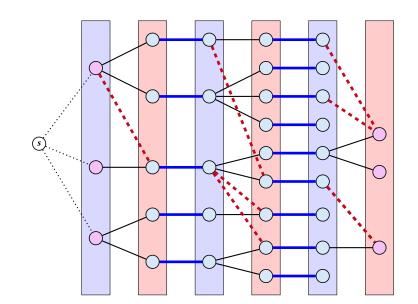
One phase of the Hopcroft-Karp algorithm can be implemented in time O(m).

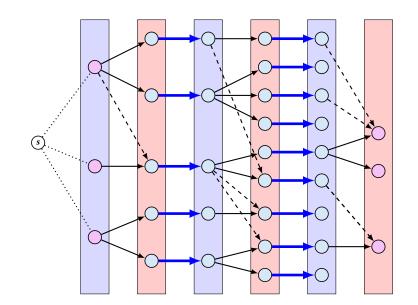
- ▶ Do a breadth first search starting at all free vertices in the left side L.
  - (alternatively add a super-startnode; connect it to all free vertices in L and start breadth first search from there)
- ► The search stops when reaching a free vertex. However, the current level of the BFS tree is still finished in order to find a set F of free vertices (on the right side) that can be reached via shortest augmenting paths.

- Then a maximal set of shortest path from the leftmost layer of the tree construction to nodes in F needs to be computed.
- Any such path must visit the layers of the BFS-tree from left to right.
- To go from an odd layer to an even layer it must use a matching edge.
- To go from an even layer to an odd layer edge it can use edges in the BFS-tree or edges that have been ignored during BFS-tree construction.
- We direct all edges btw. an even node in some layer  $\ell$  to an odd node in layer  $\ell+1$  from left to right.
- A DFS search in the resulting graph gives us a maximal set of vertex disjoint path from left to right in the resulting graph.



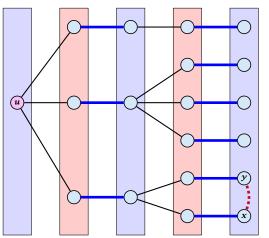






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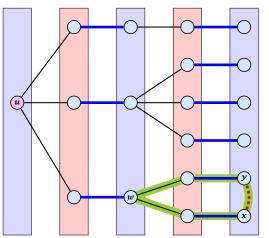
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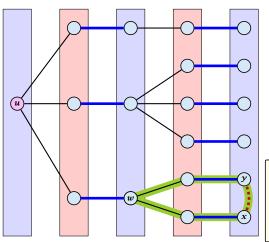
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The cycle  $w \leftrightarrow y - x \leftrightarrow w$ is called a blossom. w is called the base of the blossom (even node!!!). The path u-w path is called the stem of the blossom.





#### **Definition 9**

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.



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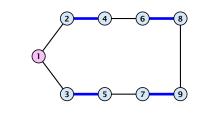
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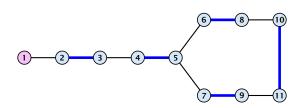


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- 1. A stem spans  $2\ell+1$  nodes and contains  $\ell$  matched edges for some integer  $\ell \geq 0$ .
- **2.** A blossom spans 2k + 1 nodes and contains k matched edges for some integer  $k \ge 1$ . The matched edges match all nodes of the blossom except the base.
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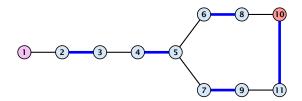
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- **4.** Every node x in the blossom (except its base) is reachable from the root (or from the base of the blossom) through two distinct alternating paths; one with even and one with odd length.
- **5.** The even alternating path to *x* terminates with a matched edge and the odd path with an unmatched edge.



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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*.
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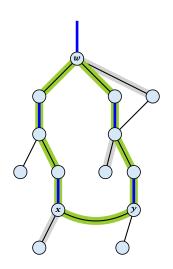
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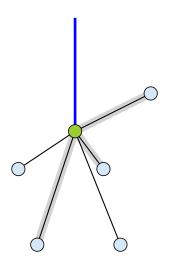
- Edges of T that connect a node u not in B to a node in B become tree edges in T' connecting u to h.
- Matching edges (there is at most one) that connect a node u not in B to a node in B become matching edges in M'.
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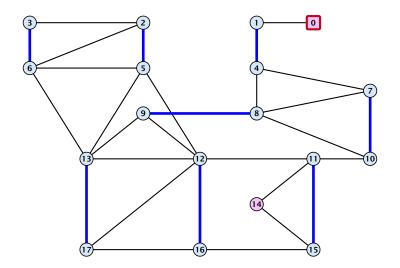
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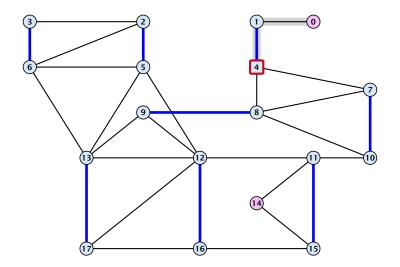


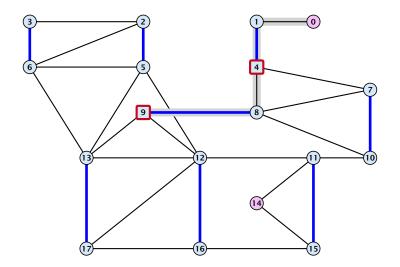


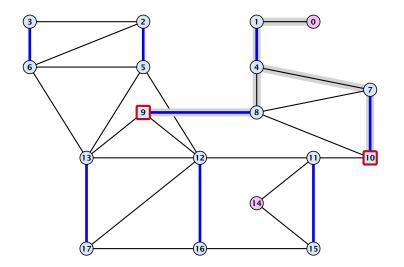
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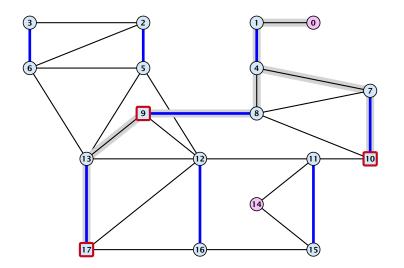




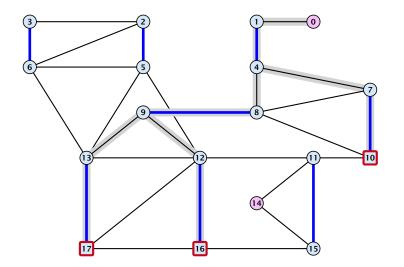




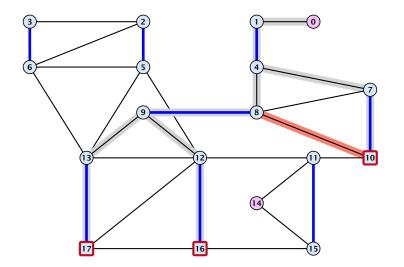




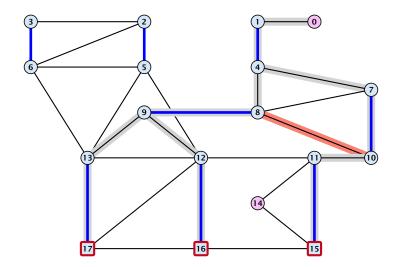




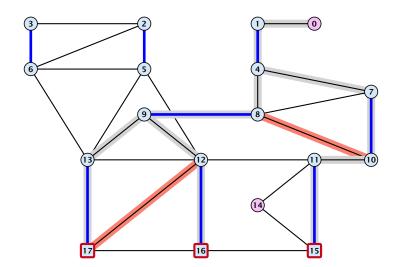




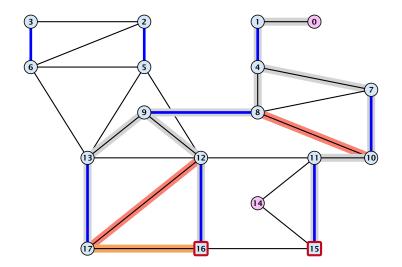




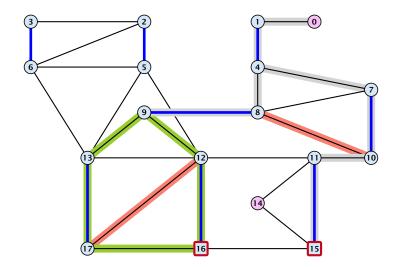




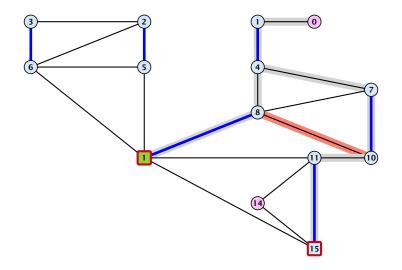




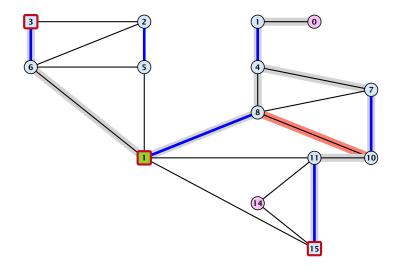




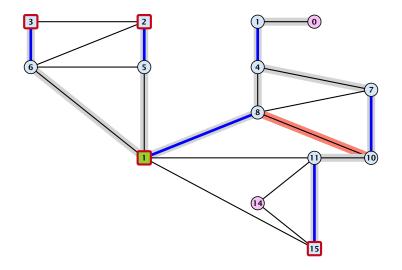




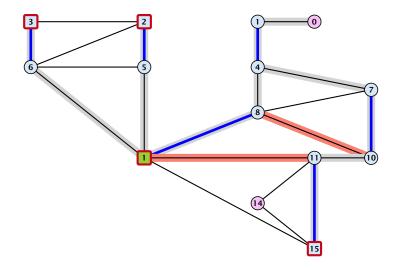




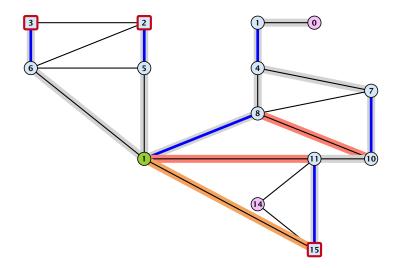




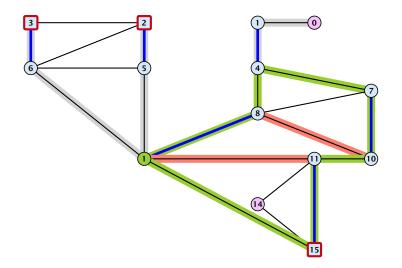




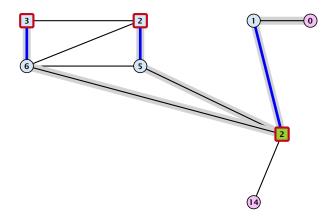


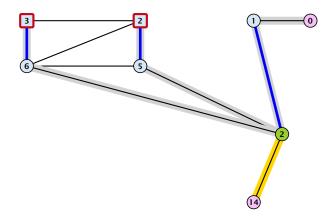


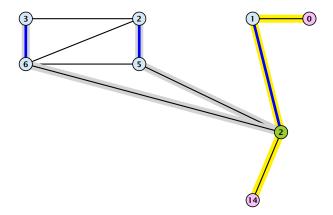


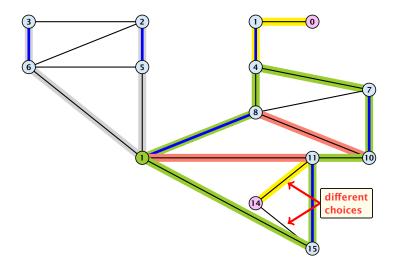




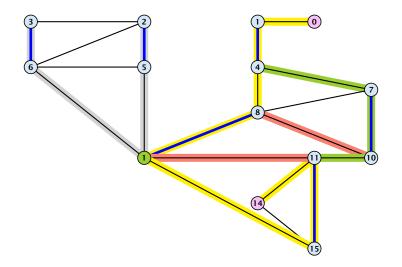


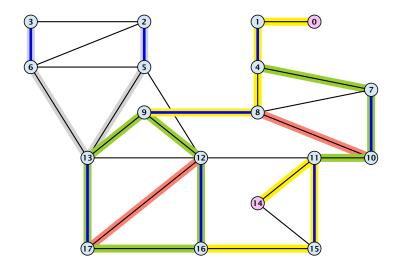




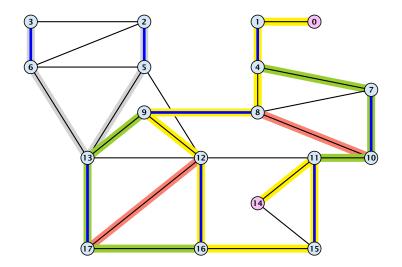




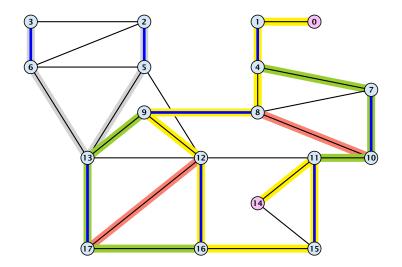














Assume that in G we have a flower w.r.t. matching M. Let r be the root, B the blossom, and W the base. Let graph G' = G/B with pseudonode b. Let M' be the matching in the contracted graph.

#### Lemma 10

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



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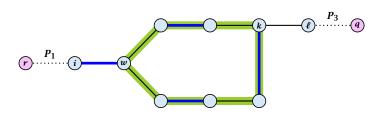
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Next suppose that the stem is non-empty.







- $\blacktriangleright$  After the expansion  $\ell$  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.



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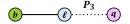
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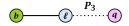
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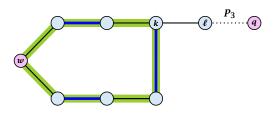


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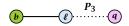


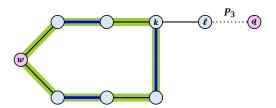


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▶ The path  $r \circ P_2 \circ (k, \ell) \circ P_3$  is an alternating path.



#### Lemma 11

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.



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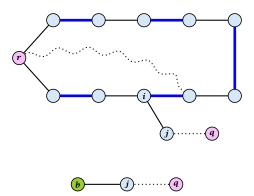
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#### Illustration for Case 1:





# 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  $M_+$ , r 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  $oldsymbol{w}$  and  $oldsymbol{q}$  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.





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- 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.

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A(i) contains neighbours of node i.

We create a copy  $\bar{A}(i)$  so that we later can shrink blossoms.

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found is just a Boolean that allows to abort the search process...

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In the beginning no node is in the tree.

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Put the root in the tree.

*list* could also be a set or a stack.

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As long as there are nodes with unexamined neighbours...

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...examine the next one

- 1: set  $\bar{A}(i) \leftarrow A(i)$  for all nodes i
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- 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**

If you found augmenting path abort and start from next root.

```
Algorithm 55 examine(i, found)

1: for all j \in \bar{A}(i) do

2: if j is even then contract(i, j) and return

3: if j is unmatched then

4: q \leftarrow j;

5: pred(q) \leftarrow i;

6: found \leftarrow true;

7: return
```

 $pred(j) \leftarrow i$ ;

8:

9:

10:

11:

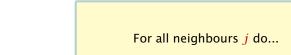
Examine the neighbours of a node  $\it i$ 

 $pred(mate(j)) \leftarrow j$ ;

add mate(j) to *list* 

if j is matched and unlabeled then

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You have found a blossom...

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10:
```

You have found a free node which gives you an augmenting path.

add mate(j) to *list* 

11:

```
Algorithm 55 examine(i, found)
1: for all j \in \bar{A}(i) do
        if j is even then contract(i, j) and return
2:
3:
      if j is unmatched then
4:
             q \leftarrow j;
             pred(q) \leftarrow i;
5:
             found ← true;
6:
7:
             return
        if i is matched and unlabeled then
8:
9:
             pred(j) \leftarrow i;
              pred(mate(j)) \leftarrow j;
10:
```

If you find a matched node that is not in the tree you grow...

add mate(j) to *list* 

11:

```
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             add mate(j) to list
11:
```

mate(j) is a new node from
which you can grow further.

- 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* 



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Get all nodes of the blossom.

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



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Identify all neighbours of b.

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



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



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



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Only for making a blossom expansion easier.



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Only delete links from nodes not in B to B.

When expanding the blossom again we can recreate these links in time  $\mathcal{O}(m)$ .



- A contraction operation can be performed in time 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

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



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