Implementation of Edmond's Blossom Algorithm for Maximum Matching in a Graph

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Abstract

The blossom algorithm is an algorithm in graph theory for constructing maximum matchings on graphs. The algorithm was developed by Jack Edmonds in 1961, and published in 1965. This polynomial time algorithm is used in several applications including the assignment problem, the marriage problem, and the Hamiltonian cycle and path problems (i.e., Traveling Salesman Problem). Through this project, we aim to implement the blossom algorithm and visualize the steps leading to the maximum matching by using a graphical interface. The user creates the graph on screen by virtue of mouse inputs and obtains the maximum matching for the given input graph. The interface also allows the user to control the steps which show the implementation of the algorithm. The inputs and outputs are also stored in text files for future reference and reuse.

1 | Blossom algorithm

The blossom algorithm is an algorithm in graph theory for constructing maximum matching on graphs. The algorithm was developed by Jack Edmond's in 1961, and published in 1965.

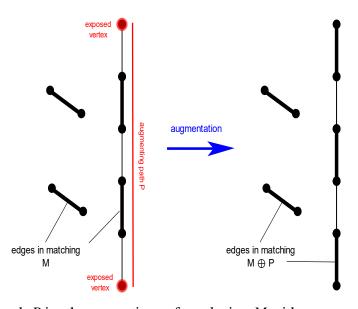
Given a general graph G = (V, E), the algorithm finds a matching M such that each vertex in V is incident with at most one edge in M and |M| is maximized. The matching is constructed by iteratively improving an initial empty matching along augmenting paths in the graph. Unlike bipartite matching, the key new idea is that an odd-length cycle in the graph (blossom) is contracted to a single vertex, with the search continuing iteratively in the contracted graph.

The algorithm runs in time $O(|E||V|^2)$, where |E| is the number of edges of the graph and |V| is its number of vertices. A better running time of $O(|E||V|^{1/2})$ for the same task can be achieved with the much more complex algorithm of Micali and Vazirani.

A major reason that the blossom algorithm is important is that it gave the first proof that a maximum-size matching could be found using a polynomial amount of computation time.

1. 1 Augmenting paths

Given G =(V, E)and a matching M of G, a vertex v is exposed if no edge of M is incident with v. A path in G is an alternating path, if its edges are alternately not in M and in M (or in M and not in M). An augmenting path P is an alternating path that starts and ends at two distinct exposed vertices. Note that the number of unmatched edges in an augmenting path is greater by one than the number of matched edges, and hence the total number of edges in an augmenting path is odd. A matching



augmentation along an augmenting path P is the operation of replacing M with a new $M_1 = M \oplus P = (M \setminus P) \ U \ (P \setminus M)$.

By Berge's Theorem, matching M is maximum if and only if there is no M-augmenting path in G. Hence, either a matching is maximum, or it can be augmented. Thus, starting from an initial matching, we can compute a maximum matching by augmenting the current matching with augmenting paths as long as we can find them, and return whenever no augmenting paths are left. We can formalize the algorithm as follows:

```
INPUT: Graph G, initial matching M on G
  OUTPUT: maximum matching M* on G
A1 function find_maximum_matching(G, M) : M*
A2   P ← find_augmenting_path(G, M)
A3   if P is non-empty then
A4       return find_maximum_matching(G, augment M along P)
A5   else
A6       return M
A7   end if
A8 end function
```

We still have to describe how augmenting paths can be found efficiently. The subroutine to find them uses blossoms and contractions.

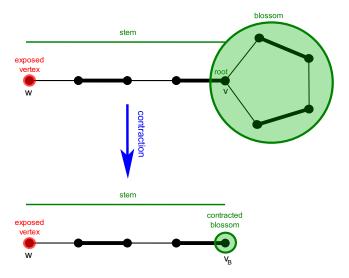
1.2 Blossoms and contractions

Given G = (V, E) and a matching M of G, a blossom B is a cycle in G consisting of 2k + l edges of which exactly k belongs to M, and where one of the vertices v of the cycle (the base) is such that there exists an alternating path of even length (the stem) from v to an exposed vertex w.

Finding Blossoms:

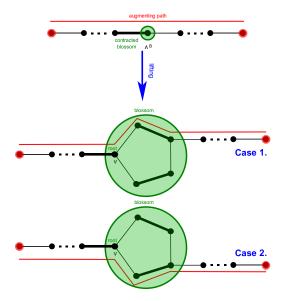
- Traverse the graph starting from an exposed vertex.
- Starting from that vertex, label it as an outer vertex "o".
- Alternate the labelling between vertices being inner "i" and outer "o" such that no two adjacent vertices have the same label.
- If we end up with two adjacent vertices labelled as outer "o" then we have an odd-length cycle and hence a blossom.

Define the *contracted graph G* as the graph obtained from G by contracting every edge of B, and define the *contracted matching M* as the matching of G corresponding to M.

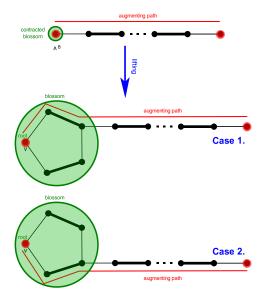


G' has an M'-augmenting path if and only if G has an M-augmenting path, and that any M'-augmenting path P' in G' can be *lifted* to an M-augmenting path in G by undoing the contraction by B so that the segment of P' (if any) traversing through v_B is replaced by an appropriate segment traversing through B. In more detail:

• if P' traverses through a segment $u \to v_B \to w$ in G', then this segment is replaced with the segment $u \to (u' \to ... \to w') \to w$ in G, where blossom vertices u' and w' and the side of B, $(u' \to ... \to w')$, going from u' to w' are chosen to ensure that the new path is still alternating (u') is exposed with respect to $M \cap B$ $\{w', w\} \in E \setminus M$.



• if P' has an endpoint v_B , then the path segment $u \to v_B$ in G' is replaced with the segment $u \to (u' \to ... \to v')$ in G, where blossom vertices u' and v' and the side of B, $(u' \to ... \to v')$, going from u' to v' are chosen to ensure that the path is alternating (v' is exposed, $\{u',u\} \in E \setminus M$).



Thus, blossoms can be contracted and search performed in the contracted graphs. This reduction is at the heart of Edmonds' algorithm.

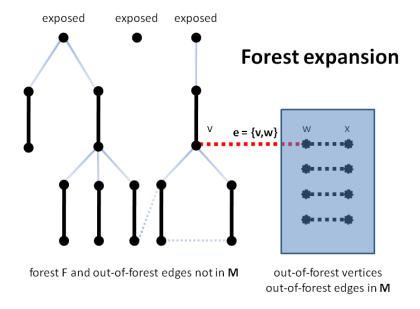
1.3 Finding an augmenting path

The search for an augmenting path uses an auxiliary data structure consisting of a forest F whose individual trees correspond to specific portions of the graph G. In fact, the forest F is the same that would be used to find maximum matchings in bipartite graph (without need for shrinking blossoms). In each iteration the algorithm either (1) finds an augmenting path, (2) finds a blossom and recurses onto the corresponding contracted graph, or (3) concludes there are no augmenting paths. The auxiliary structure is built by an incremental procedure discussed next. The construction procedure considers vertices v and edges e in G and incrementally updates F as appropriate. If v is in a tree T of the forest, we let root(v) denote the root of T. If both u and v are in the same tree T in F, we let distance(u,v) denote the length of the unique path from u to v in T.

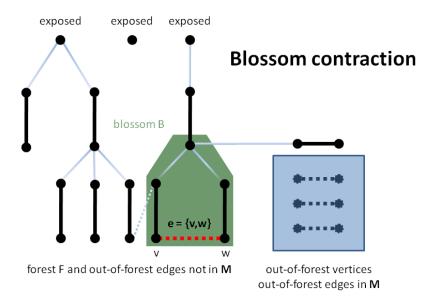
```
INPUT: Graph G, matching M on G
    OUTPUT: augmenting path P in G or empty path if none found
B01 function find augmenting path(G, M) : P
B02
        F \leftarrow \text{empty forest}
        unmark all vertices and edges in G, mark all edges of M
B05
        for each exposed vertex v do
B06
             create a singleton tree { v } and add the tree to F
        while there is an unmarked vertex v in F with distance(v, root(v)) even do
B08
B09
             while there exists an unmarked edge e = \{ v, w \} do
B10
                  if w is not in F then
                       // w is matched, so add e and w's matched edge to F
В11
                       x \leftarrow \text{vertex matched to } w \text{ in } M
В12
                       add edges { v, w } and { w, x } to the tree of v
B13
                       if distance(w, root(w)) is odd then
В14
                            // Do nothing.
B15
                       else
                            if root(v) \neq root(w) then
B16
                                 // Report an augmenting path in F U { e }.
В17
                                 P \leftarrow \text{path} (root(v) \rightarrow \ldots \rightarrow v) \rightarrow (w \rightarrow \ldots \rightarrow root(w))
B18
                                return P
B19
                            else
                                 // Contract a blossom in G and look for the path in the
contracted graph.
B20
                                B \leftarrow \text{blossom formed by } e \text{ and edges on the path } v \rightarrow w \text{ in } T
B21
                                 G', M' \leftarrow \text{contract } G \text{ and } M \text{ by } B
                                 P' \leftarrow find augmenting\_path(G', M')
B22
                                 P \leftarrow \text{lift } P' \text{ to } G
B23
B24
                                return P
B25
                            end if
B26
                       end if
B27
                  end if
B28
                 mark edge e
B29
             end while
B30
             mark vertex v
        end while
B31
B32
       return empty path
B33 end function
```

1.3.1 Examples

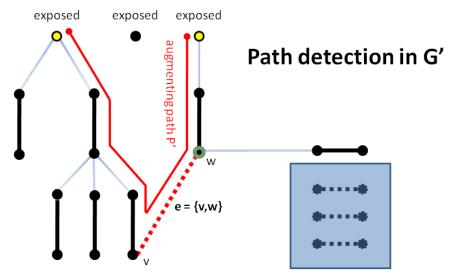
The following four figures illustrate the execution of the algorithm. Dashed lines indicate edges that are currently not present in the forest. First, the algorithm processes an out-of-forest edge that causes the expansion of the current forest (lines B10-B12).



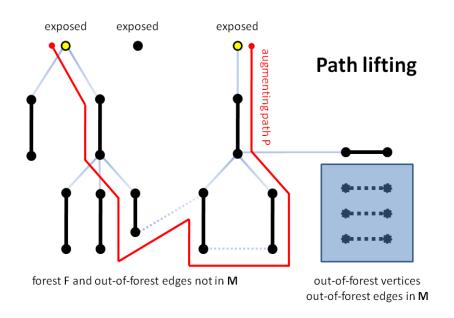
Next, it detects a blossom and contracts the graph (lines B20 – B21).



Finally, it locates an augmenting path P' in the contracted graph (line B22) and lifts it to the original graph (line B23). Note that the ability of the algorithm to contract blossoms is crucial here; the algorithm cannot find P in the original graph directly because only out-of-forest edges between vertices at even distances from the roots are considered on line B17 of the algorithm.



forest F' in G' and out-of-forest edges not in $\mathbf{M'}$ out-of-forest vertices out-of-forest edges in $\mathbf{M'}$



1.3.2 Analysis

The forest *F* constructed by the *find_augmenting_path()* function is an alternating forest.

- a tree T in G is an alternating tree with respect to M, if
 - \circ T contains exactly one exposed vertex r called the tree root
 - \circ every vertex at an odd distance from the root has exactly two incident edges in T, and
 - o all paths from *r* to leaves in *T* have even lengths, their odd edges are not in *M* and their even edges are in *M*.
- a forest F in G is an alternating forest with respect to M, if
 - o its connected components are alternating trees, and
 - \circ every exposed vertex in G is a root of an alternating tree in F.

Each iteration of the loop starting at line B09 either adds to a tree T in F (line B10) or finds an augmenting path (line B17) or finds a blossom (line B20). It is easy to see that the running time is $O(|E||V|^2)$.

1.3.3 Bipartite matching

When G is bipartite, there are no odd cycles in G. In that case, blossoms will never be found and one can simply remove lines B20 - B24 of the algorithm. The algorithm thus reduces to the standard algorithm to construct maximum cardinality matchings in bipartite graphs where we repeatedly search for an augmenting path by a simple graph traversal: this is for instance the case of the Ford-Fulkerson Algorithm.

1.3.4 Weighted matching

The matching problem can be generalized by assigning weights to edges in G and asking for a set M that produces a matching of maximum (minimum) total weight: this is the maximum weight matching problem. This problem can be solved by a combinatorial algorithm that uses the unweighted Edmonds's algorithm as a subroutine.

2 | Tools & Technologies used for the implementation

The following tools and technologies have been used in this project for the implementation of the blossom's algorithm in a graphical interface. VS Code was used as a code-editor, C++ was the language used for programming and the OpenGL API was used to design the graphical interface. Here is a brief introduction on each:

2.1 OpenGL

OpenGL (Open Graphics Library) is a cross-language, cross-platform Application Programming Interface (API) for rendering 2D and 3D vector graphics. The API is typically used to interact with a Graphics Processing Unit (GPU), to achieve hardware-accelerated rendering.



2.2 Visual Studio Code

Visual Studio Code, also commonly referred to as VS Code, is a source-code editor made by Microsoft for Windows, Linux and macOS. Features include support for debugging, syntax highlighting, intelligent code completion, snippets, code refactoring, and embedded Git. Users can change the theme, keyboard shortcuts, preferences, and install extensions that add additional functionality.



2.3 C++

C++ is a general-purpose programming language created by Danish computer scientist *Bjarne Stroustrup* as an extension of the C programming language, or "C with Classes". The language has expanded significantly over time, and modern C++ now has object-oriented, generic, and functional features in addition to facilities for low-level memory manipulation. It is almost always implemented as a compiled language, and many vendors provide C++ compilers, including the Free Software Foundation, LLVM, Microsoft, Intel, Oracle, and IBM so, it is available on many platforms.



3 | Implementation with a Graphical Interface

To visualize how the algorithm works to find the maximum matching we have used the OpenGL API to take inputs via mouse clicks to create the graph and then keyboard inputs to follow the steps as the algorithm proceeds.

- We use a single buffer RGBA colour profile.
- The Input window is set to a resolution of 1280 x 720 pixels, positioned at (0,0), i.e., at the top-left corner of the computer screen and titled "Edmond's Blossom Algorithm".
- The Background colour is set to white by glClearColor (1.0, 1.0, 1.0, 1.0).
- **gpMouse** is the callback for the *glutMouseFunc()* and manages the user inputs to create and operate on the graph.
- In the main function a file "*Input.txt*" is opened with write permissions to store the vertices and edges for future use.
- In the code snippet give below, we see the main function and the *gpInit()* function which initialises the plot session. The description of the various parameters and instructions have been mentioned above.

```
int main(int argc, char *argv[])
    fp = fopen("input.txt", "w+");
   glutInit(&argc, argv);
   glutInitDisplayMode(GLUT_SINGLE | GLUT_RGBA);
   glutInitWindowSize(WW, WH);
   glutInitWindowPosition(0, 0);
   glutCreateWindow("Edmond's Blossom Algorithm");
   glutMouseFunc(gpMouse);
   gpInit();
   glutMainLoop();
   return 0;
void gpInit()
   glClearColor(1.0, 1.0, 1.0, 1.0);
    glColor4f(0.0, 0.0, 0.0, 1.0);
   glMatrixMode(GL PROJECTION);
    glLoadIdentity();
   gluOrtho2D(0, WW, 0, WH);
   glClear(GL COLOR BUFFER BIT);
   glFlush();
```

3.1 Utility functions for plotting points and lines

Before we actually see how the mouse inputs are used to create vertices and edges, we need to understand certain utility functions which will later be used for drawing points and lines.

3.1.1 Drawing points

There are 3 types of points: regular, blossom and small points. They are discussed below.

3.1.1.1 Drawing regular points

The *drawpoint()* function takes in two integer parameters *x* and *y*, the respective *x* and *y* coordinates of the point where the user clicked on the screen and displays it in the colour as passed into the function as the string parameter 'col'.

- The 4 global variables [r, g, b, a] are respectively set to the values of red, green, blue and alpha that will be taken for a given colour as specified by the string "col". This is done by the multiple if-statement blocks some of which are shown collapsed in the code snippet below.
- The colours are set by glColor4f(r, g, b, a).
- The point size is set to "*radius*" which is also globally declared to be equal to 20 pixels, so that a vertex appears as a small circle or a thick point.
- The point is also set to be of smooth nature by glEnable(GL_POINT_SMOOTH).
- The actual point is plotted by glVertex2i(x, y) placed between $glBegin(GL_POINTS)$ and glEnd().

```
void drawPoint(int x, int y, string col)
    if (col == "light") ···
    if (col == "red") ...
    if (col == "black") ···
    if (col == "gray") ⋯
    if (col == "blue") ···
    if (col == "green")
        r = 0.0;
        g = 1.0;
        b = 0.0;
        a = 1.0;
    glColor4f(r, g, b, a);
    glPointSize((double)radius);
    glEnable(GL_POINT_SMOOTH);
    glBegin(GL_POINTS);
    glVertex2i(x, y);
    glEnd();
```

3.1.1.2 Drawing Blossom points

The drawblossompoint() function takes in two integer parameters x and y, the respective x and y coordinates of the vertex which is a part of a blossom which is to be contracted and highlights that vertex with a yellow semi-transparent colour [glColor4f(1.0, 0.5, 0.0, 0.5)]. The thickness of the point is determined by the argument "rad" which is passed to the function. Usually, this point is made larger than a regular-sized vertex and appears as a glow over a regular-sized vertex. The other properties of the point are similar to what we had in the function drawpoint().

3.1.1.3 Drawing Small points

The drawsmallpoint() function takes in two integer parameters x and y, the respective x and y coordinates of the vertex which is a father of the activeVertex and highlights that vertex with an orange semi-transparent colour [glColor4f(1.0, 1.0, 0.0, 0.5)]. The thickness of the point is determined by the argument "rad" which is passed to the function. Usually, this point is made smaller than a regular-sized vertex and appears as a dot inside a regular-sized vertex. The other properties of the point are similar to what we had in the function drawpoint().

3.1.2 Drawing Lines

The *drawLine()* function takes in four integer parameters (*xa*, *ya*) and (*xb*, *yb*), the x and y coordinates of the initial and final points respectively of the line to be drawn representing and edge between two vertices displays it in the colour as passed into the function as the string parameter 'col'.

- The 4 global variables [r, g, b, a] are respectively set to the values of red, green, blue and alpha that will be taken for a given colour as specified by the string "col". This is done by the multiple if-statement blocks some of which are shown collapsed in the code snippet below.
- The colours are set by glColor4f(r, g, b, a).
- The line width is set to be 5 pixels so that the edge appears visibly thick.
- The line is also set to be of smooth nature by *glEnable(GL_LINE_SMOOTH)*.
- Within glBegin(GL_LINES) and glEnd(), the starting and ending points of the line to be plotted are set by glVertex2i(xa, ya) and glVertex2i(xb, yb).

```
void drawLine(int xa, int ya, int xb, int yb, string col)
    if (col == "red") ···
    if (col == "black") ···
    if (col == "gray") ···
    if (col == "blue") ···
    if (col == "green")
        r = 0.0;
        g = 1.0;
        b = 0.0;
        a = 1.0;
    glColor4f(r, g, b, a);
    glLineWidth(5);
    glEnable(GL LINE SMOOTH);
    glBegin(GL LINES);
    glVertex2i(xa, ya);
    glVertex2i(xb, yb);
    glEnd();
```

3.2 | Mouse function for graph creation

Let us take a look at some of the important data structures that will be required to create the graph by mouse inputs.

- The map "vertices" is used to store every vertex input from the user and it is mapped to a pair of integers which are the x and y coordinates of the point where the vertex is located on the screen.
- The vector of pairs "edges" stores two integers in a pair where each is a vertex and a pair denotes an edge between those two vertices.

```
map<int, pair<int, int>> vertices;
vector<pair<int, int>> edges;
```

Some more variables to be observed are:

- bool inputphase: Initially set to *true*. Once the inputs from user are taken, it is set to *false* to avoid any more inputs by stray mouse clicks on the screen while the blossom algorithm is being implemented.
- *int click*: Initialised to zero, this is used to maintain the number of left clicks made by the user.

3.2.1 Utility function for an already existing vertex

The function $int\ existVertex(int\ xp,\ int\ yp)$ is used to check if the points $(xp,\ yp)$ passed to the function are the coordinates of an existing vertex already input by the user previously or not. If yes, then the number of that vertex is returned which $(xp,\ yp)$ points to. If no, then -1 is returned and this point $(xp,\ yp)$ is treated as a new point to be plotted in the mouse function.

We traverse the map "vertices" and check for every existing vertex whether (xp, yp) belongs inside the circle centred at xcentre and ycentre with radius equal to "radius", where xcentre and ycentre are the x and y coordinates of the centres of that vertex. If such a point exists inside the circle, that vertex is returned by the function. After the search for all vertices is completed, the control reaches the end of the function only when no vertex could be found to exist at the point (xp, yp). We then return -1, indicating a new vertex is found.

```
int existVertex(int xp, int yp)

for (auto q : vertices)

{
    for (auto q : vertices)

    {
        int xcentre = q.second.first;
        int ycentre = q.second.second;
        if (((xcentre - xp) * (xcentre - xp) + (ycentre - yp)) * (ycentre - yp)) < radius * radius)
        return q.first;
}

return -1;
}
</pre>
```

3.2.2 The gpMouse() function

The function has 4 parameters: *button, state, x, y.* The *button* parameter specifies the button pressed left or right. The *state* parameter specifies the state of the button whether pressed or released. The parameters *x* and *y* contain the coordinates of the point where the click occurred on the screen.

- When a left-click is encountered, the click value increments and if we are still in the input phase, we proceed to the next steps, else we do nothing.
- We follow a convention that in every odd-numbered click, we set a starting vertex for an edge and in an even-numbered click, we set the ending vertex for that edge.
- In both these cases, we check if the point where the click occurred was an existing vertex. If it was then, we reset the values of x and y to that of the existing vertex and we set the current vertex denoted by 'u' or 'v' to be this existing vertex.
- Otherwise, if it is not an existing vertex, we increment '*Vnum*' as a new vertex is to be added. We set 'u' or 'v' to be '*Vnum*' and we add the pair {x,y} to the map '*vertices*' at '*Vnum*'. Then, we plot this new point (x, y) on the screen in gray colour using drawpoint() function.

```
void gpMouse(int button, int state, int x, int y)
    y = glutGet(GLUT WINDOW HEIGHT) - y;
    if ((button == GLUT LEFT BUTTON) && (state == GLUT DOWN))
        click++;
        if (inputphase)
            if (click % 2 != 0)
                int val = existVertex(x, y);
                if (val == -1)
                    Vnum++;
                    u = Vnum;
                    vertices[Vnum] = \{x, y\};
                    drawPoint(x, y, "gray");
                else
                    u = val;
                    x = vertices[val].first;
                    y = vertices[val].second;
                xa = x;
                ya = y;
```

- In case of odd-numbered clicks, (xa, ya) is set to (x, y).
- In case of even-numbered clicks, (xb, yb) is set to (x, y) and then the edge is drawn from (xa, ya) to (xb, yb) in black colour using the drawLine() function. Also, the vertices are plotted again for neat presentation. Finally, the pair of vertices forming this edge {u,v} is added to the vector 'edges'.
- When a right-click is encountered, it marks the end of the input phase and hence the variable *inputphase* is set to false and the *display()* function is called which shall proceed with the execution of the Edmond's Blossom Algorithm for the graph input in the input phase.

```
else
            int val = existVertex(x, y);
            if (val == -1)
                Vnum++;
                v = Vnum;
                vertices[Vnum] = {x, y};
                drawPoint(x, y, "gray");
            else
                x = vertices[val].first;
                y = vertices[val].second;
                v = val;
            xb = x;
            yb = y;
            drawLine(xa, ya, xb, yb, "black");
            drawPoint(xa, ya, "gray");
            drawPoint(xb, yb, "gray");
            edges.push_back({u, v});
  ((button == GLUT_RIGHT_BUTTON) && (state == GLUT_DOWN))
    if (inputphase)
        inputphase = false;
        glFlush();
        display();
glFlush();
return;
```

3.3.1 The display() function

The *display* function performs two types of tasks. Firstly, it sorts the '*edges*' vector for a neat output into the "*input.txt*" file. The first line of "*input.txt*" contains the number of vertices and edges in the graph input. The following lines list all the edges as pairs of vertices in 1-based indexing separated by spaces. Secondly and finally, it calls the *edmondcall()* function.

3.3.2 The edmondcall() function

The *edmondcall* function opens "*input.txt*" in *read* mode and "*output.txt*" in *write* mode. It scans the number of vertices and edges and stores them in global variables 'V' and 'E'. It creates an object of class *Blossom* named 'bm' and initialises it with values for 'V' and 'E'. It then scans the next *E* lines from the input file and passes them to the *addEdge()* function in 0-based indexing. It then calls the *edmondsBlossomAlgorithm()* function to calculate the maximum matching and then prints it to "*output.txt*" using the *printMatching()* function.

```
void edmondcall()
{

fpin = fopen("input.txt", "r+");
    fpout = fopen("output.txt", "w+");
    int u, v;
    int V, E;
    rewind(fpin);
    fscanf(fpin, "%d %d", &V, &E);

Blossom bm(V, E);
    while (E--)

{
        fscanf(fpin, "%d %d", &u, &v);
        bm.addEdge(u - 1, v - 1);
    }

int res = bm.edmondsBlossomAlgorithm();

bm.printMatching(res);
    fclose(fpin);
    fclose(fpout);
    cout << "Press Enter thrice to quit\n";
    flag = getc(stdin); flag = getc(stdin);
    exit(0);
}
</pre>
```

3.4 | The Class Blossom

Some data structures and variables to understand before we explore the functions in this class are mentioned as follows:

• *struct StructEdge*: This struct stores all pairs of vertices for all edges. The integer variable *vertex* stores the number of the vertex and the pointer *next* stores the next sibling of the vertex. *Edge* is a type-defined pointer for *StructEdge*.

```
struct StructEdge

int vertex;

StructEdge *next;

y;

typedef StructEdge *Edge;
```

- A *pool* array is created of type *StructEdge* to store all the vertices from the input file. Pointers of *StructEdge* type are created: "top" which points to the top of the input stack in *pool*; and *adj[M]*, an array which stores the address of the edge of last child of the respective indices in *adj[]*.
- Integers *V*, *E* store the number of vertices and edges respectively.
- Integers q_front and q_rear store the index of the front and rear of the bfs_queue array which is used for BFS Traversal in finding augmenting paths.
- Integer array *match[]* stores the matching for the edges. If a matching exists between *u* and *v*, *match[u]* will be equal to *v*, and *match[v]* will be equal to *u*. It is initialised to 1 and remains so for vertices which are unmatched.
- Integer array *father[]* stores the parent vertex from which we reached the current vertex during the BFS traversal.
- Integer array *base[]* stores the base for every vertex. Initially, it is set to its index value for every vertex, but in case a blossom is encountered, all vertices in the blossom will have their base values equal to one vertex in the blossom at which it shall be assumed to be contracted.
- Boolean array ed[][] is an adjacency matrix used to avoid duplicate entries in graph.
 Boolean arrays inq[] and inb[] are used to check if a particular vertex is in the BFS queue or if it is in a blossom respectively.

```
class Blossom

class Blossom

{

StructEdge pool[M * M * 2];

Edge top = pool, adj[M];

int V, E, q_front, q_rear;

int match[M], bfs_queue[M], father[M], base[M];

bool inq[M], inb[M], ed[M][M];

public:

Blossom(int V, int E) : V(V), E(E) {}
```

3.4.1 The initializer list

The initializer list in blossom class initialises the number of vertices V and number of edges E to the integers V and E passed to the constructor.

```
public:

Blossom(int V, int E) : V(V), E(E) {}
```

3.4.2 The addEdge() function

This function takes in two vertices u and v in 0-based indexing. The if-condition checks for duplicates and self-loops in inputs. If no such condition arises the vertices v and u are added to the *pool* and the respective next values for each are assigned the other's adj array value, i.e., for the StructEdge element for vertex v, next stores the pointer for adj[u] and vice-versa. Lastly, the vertices u and v are added to the adjacency matrix ed[j][].

```
void addEdge(int u, int v)

{
    if (!ed[u][v] && u != v)

    {
        top->vertex = v;
        top->next = adj[u];
        adj[u] = top++;

        top->vertex = u;
        top->next = adj[v];
        adj[v] = top++;

ed[u][v] = ed[v][u] = true;
}
```

3.4.3 The pausefunc() function

This function is paired with the *updateGraph()* function to display the updated graph and then halt the program execution till the user allows it to proceed to observe every intermediate step leading to the maximum matching.

```
void pausefunc()

{

| flag = getc(stdin);

| }
```

3.4.4 The updateGraph() function

- First, the screen is cleared and the background colour is set to white.
- Next for all edges, we check whether it is a matched edge or not. We represent matched
 edge with red colour and unmatched edges with black. This is done with the help of
 drawLine() function.
- Next for every vertex, we check for the following:
 - o If it is the *root* of a blossom, it is highlighted with a *large yellow* circle.
 - o If it is not a root but a part of the blossom it is highlighted in *yellow* with a *large* but smaller than the previous circle. Both these tasks are accomplished by drawblossompoint() function.
 - o If the vertex is the *originalVertex* i.e., the vertex for which we are trying to improve the matching, then it is displayed in *green* colour.
 - o If the vertex is the *startVertex* i.e., the vertex at the front of the BFS queue, whose children will be accessed, then it is displayed in *blue* colour.
 - o If the vertex is the *activeVertex* i.e., the current vertex we are exploring, it is then displayed in *red* colour.
 - Otherwise, if it is none of the above, it is a regular vertex and we denote it by *gray* colour. All these steps from *originalVertex* to regular vertex are accomplished by *drawpoint()* function.
- Lastly, the *father* of the *activeVertex* is denoted by a *smaller orange* point drawn using *drawsmallpoint()* function.

```
void updateGraph()
{
   glClear(GL_COLOR_BUFFER_BIT | GL_DEPTH_BUFFER_BIT);
   glClearColor(1.0f, 1.0f, 1.0f, 1.0f);
   for (auto checkedge : edges)
{
      int v1 = checkedge.first;
      int v2 = checkedge.second;
      xa = vertices[v1].first;
      ya = vertices[v2].first;
      yb = vertices[v2].second;
      if (match[v1 - 1] == (v2 - 1) && match[v2 - 1] == (v1 - 1))
            drawLine(xa, ya, xb, yb, "red");
      else
            drawLine(xa, ya, xb, yb, "red");
      else
            drawLine(xa, ya, xb, yb, "black");
   }
   for (auto checkv : vertices)
{
      int ve = checkv.first;
      if (bmnode != -1 && bmnode == (ve - 1))
            drawBolosocompoint(vertices[ve].first, vertices[ve].second, radius + 20);
      if (inb[ve - 1])
            drawBolosocompoint(vertices[ve].first, vertices[ve].second, radius + 10);
      if (originalVertex != -1 && (ve - 1) == originalVertex)
            drawBolosocompoint(vertices[ve].first, vertices[ve].second, "green");
      if (startVertex != -1 && (ve - 1) == startVertex)
            drawPoint(vertices[ve].first, vertices[ve].second, "blue");
      if (activeVertex != -1 && (ve - 1) == startVertex)
            drawPoint(vertices[ve].first, vertices[ve].second, "red");
      if (((ve - 1) != originalVertex) && ((ve - 1) != startVertex) && (newPoint(vertices[ve].first, vertices[ve].second, "gray");
      if (activeVertex != -1 && father[activeVertex] != -1)
            drawSmallpoint(vertices[father[activeVertex] != -1)
            drawSmallpoint(ver
```

3.4.5 Finding and displaying maximum matching

The *edmondsBlossomAlgorithm* function calculates the maximum number of matched edges and stores it in *match_counts* and returns this value.

- It initialises the *match[]* array to have all values -1.
- For every vertex which is unmatched, we find an augmenting path for that vertex u and we augment the path from u to v, where v is the vertex returned by $find_augmenting_path(u)$. If the path could be augmented, the $match_counts$ is incremented by 1.
- Finally, after maximum matching has been found all variables *activeVertex*, *startVertex*, *originalVertex*, *bmnode* and the *inb[]* array is reset to their initial values so that only the final matching is displayed on the screen.

The *printMatching* function takes in the maximum number of edges matched and prints it to the output file, followed by all the matched edges in 1-based indexing. For all vertices, we check if the value in *match[]* array for that vertex exists and is greater than that vertex, to avoid printing same edge twice. If it is so, we print the vertex *i* and the *match[i]*.

```
int edmondsBlossomAlgorithm()
    int match_counts = 0;
    memset(match, -1, sizeof(match));
    for (int u = 0; u < V; u++)</pre>
        if (match[u] == -1)
            match_counts += augment_path(u, find_augmenting_path(u));
    activeVertex = originalVertex = startVertex = -1, bmnode = -1;
    memset(inb, 0, sizeof(inb));
    updateGraph();
    return match counts;
void printMatching(int maxMatch)
    rewind(fpout);
    fprintf(fpout, "%d\n", maxMatch);
    for (int i = 0; i < V; i++)</pre>
        if (i < match[i])</pre>
            fprintf(fpout, "%d %d\n", i + 1, match[i] + 1);
```

3.4.6 Finding Augmenting Paths

The find_augmenting_path function accepts a vertex s and returns the last vertex in the augmenting path beginning from s. It returns -1 if no augmenting path was found.

- Initially the *inq[]* and *father[]* arrays are initialised to *false* and -1 respectively and *base* of every vertex is set to itself.
- We begin by inserting *s* into *bfs_queue* and while the queue is not empty, we perform BFS on the graph.
- We traverse all children v of u, where u is the q_front. Only if the base of both u and v are different and there is no matching existing between u and v, we proceed.
- If *v* becomes equal to *s* or a matching exists for *v* with a *father* present, we have found an *odd-length cycle* and we *contract the blossom*.
- Otherwise, if v doesn't have a *father*, we set *father* of v equal to u.
- If *v* doesn't have a matching we return this value to *augment_path()* function as we have found an augmenting path.
- If matching exists for v, but the matched vertex is absent in current BFS, we add it to the *rear* of bfs_queue .

```
int find_augmenting_path(int s)
   originalVertex = s;startVertex = -1;activeVertex = -1;
   updateGraph(); pausefunc();
   memset(inq, 0, sizeof(inq));
   memset(father, -1, sizeof(father));
   for (int i = 0; i < V; i++)</pre>
   inq[bfs_queue[q_front = q_rear = 0] = s] = true;
   while (q_front <= q_rear)</pre>
       int u = bfs_queue[q_front++];
       startVertex = u; activeVertex = -1;
       updateGraph(); pausefunc();
       for (Edge e = adj[u]; e; e = e->next)
            int v = e->vertex;
            updateGraph(); pausefunc();
            if (base[u] != base[v] && match[u] != v)
                if ((v == s) || (match[v] != -1 && father[match[v]] != -1))
                    updateGraph(); pausefunc();
                else if (father[v] == -1)
                    updateGraph(); pausefunc();
                    if (match[v] == -1)
                        return v;
                    else if (!inq[match[v]])
                        inq[bfs_queue[++q_rear] = match[v]] = true;
```

3.4.7 Augmenting a path

The *augment_path* function takes in two parameters *s* and *t* where *t* is the exposed vertex returned by the *find_augmenting_path* function. If there was an augmenting path found, i.e., *t* was not -1, *augment_path()* returns 1 as the matching will be improved by 1.

- The algorithm begins from the exposed vertex u=t finds its *father* v and makes a matching between u and v and then u moves to the previous matching of v.
- At any time if there ceases to be a matching for v, then u will become -1 and hence loop will terminate thus having created the maximum matching for that augmenting path starting from s and ending at t.

```
int augment_path(int s, int t)
{
    int u = t, v, w;
    while (u != -1)
    {
        v = father[u];
        w = match[v];
        match[v] = u;
        match[u] = v;
        u = w;
        updateGraph(); pausefunc();
    }
    return t != -1;
}
```

3.4.8 Finding lowest common ancestor for blossoms

This function is used to find the lowest common ancestor for two vertices u and v in a blossom with respect to a root vertex. Firstly, we find a path from u to the root and then we start from v and return the first vertex that was present in the previous path.

```
int LCA(int root, int u, int v)
{
    static bool inp[M];
    memset(inp, 0, sizeof(inp));
    while (1)
    {
        inp[u = base[u]] = true;
        if (u == root)
            break;
        u = father[match[u]];
    }
    while (1)
    {
        if (inp[v = base[v]])
            return v;
        else
            v = father[match[v]];
    }
}
```

3.4.9 Marking and Contracting Blossoms

The *mark_blossom* function is used to mark and add to the blossom all vertices from vertex *u* to the vertex *lca* which are accepted as parameters.

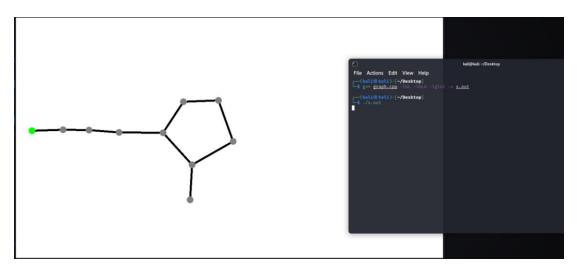
The *blossom_contraction* function finds the *LCA* of *u* and *v* with root *s*. Initialises *inb[]* array to all *false* values, then *marks* all vertices in blossom from *u* to *lca* and then from *v* to *lca* using *mark_blossom()*. Finally, for all vertices in the blossom, it changes their *base* to the *lca* value found and adds all vertices to the *rear* of *bfs_queue* if not previously present. This is done so that, all vertices in the blossom can be treated to be contracted to the *LCA* vertex and when we try to find an augmenting path in the graph with a *contracted blossom*, we treat every matching to be created with a blossom vertex to be supposedly with the *LCA* vertex.

```
void mark blossom(int lca, int u)
    while (base[u] != lca)
        int v = match[u];
        inb[base[u]] = inb[base[v]] = true;
        u = father[v];
        if (base[u] != lca)
            father[u] = v;
        updateGraph(); pausefunc();
void blossom_contraction(int s, int u, int v)
    int lca = LCA(s, u, v);
    bmnode = 1ca;
    updateGraph(); pausefunc();
    memset(inb, 0, sizeof(inb));
   mark blossom(lca, u);
    mark_blossom(lca, v);
    if (base[u] != lca)
        father[u] = v;
    if (base[v] != lca)
        father[v] = u;
    for (int u = 0; u < V; u++)
        if (inb[base[u]])
            base[u] = lca;
            if (!inq[u])
                inq[bfs queue[++q rear] = u] = true;
```

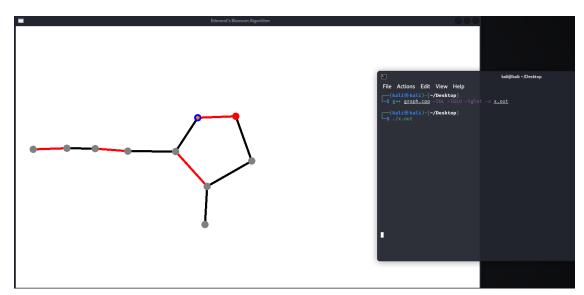
4 | Executing the code

The code works in both Linux and Windows environments but the system must have OpenGL and GLUT libraries installed in it. For this sample run, we will be operating in a Linux environment.

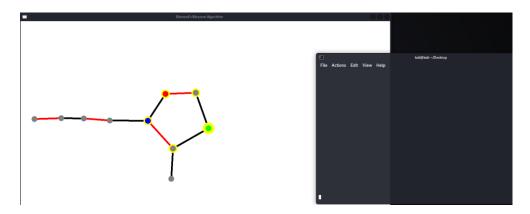
- First, we compile the code with the following headers in the terminal: g++ graph.cpp -lGL -lGLU -lglut -o x.out
- Now run the binary file x.out by typing "./x.out
- Plot the graph in the window by clicking with left-mouse button and once all inputs are done, right-click the mouse once to terminate the input phase. To make an edge the first left-click sets the starting vertex for edge and the next left-click completes the edge by setting the other vertex for that edge.



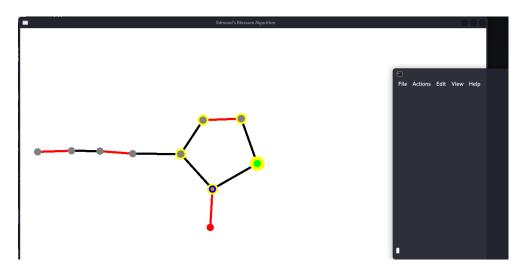
• Now move over to the terminal and keep pressing "Enter" to proceed with the algorithm. As the algorithm proceeds, we see the current vertex denoted by red finding a father denoted by an orange dot and then forming matchings denoted by red lines.



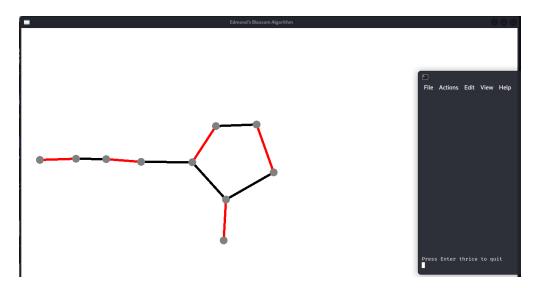
• The algorithm continues and finds a blossom some steps later. Now, it finds the LCA of the red and blue vertex with respect to green root vertex and also marks the blossom vertices. Thus, the blossom has now been contracted.



• The algorithm finds an external node to the blossom and the augmented path is adjusted with respect to that.



• Finally, the augmented path inside the blossom is completed and the maximum matching is obtained.



5 | A simple implementation in a real-life problem

Problem Statement

Suppose we are given a large collection of people and we are asked to match them into the pairs such that people who are known to each other are preferred to form a pair between them. We have to find the maximum matching number of pairs possible. Also, it is to be noted that every person can belong only to a single pair.

Solution approach

Let us assume each person to be a vertex in a graph and the acquaintances to be an edge in a graph between the vertices representing those people. Thus, we create an unweighted, undirected graph G = (V, E) on which we can apply the Blossom Algorithm to find the maximum matching. We can map each person to a vertex and once the matching is maximised, we can again map the results back to the names of the people and display which are the pairs which will result in maximum matching. We use blossom algorithm for this because it finds the maximum matching in general graphs in polynomial-time. In addition to finding maximum matching, at last we form the unmatched vertices amongst themselves in pairs in random fashion as there is no edge between any of them. Thus, the problem is solved.

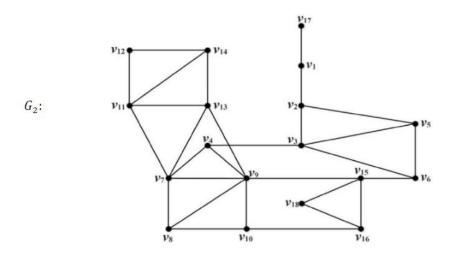
6 | Implementation in the Battle of Britain (1940)

In the Battle of Britain (1940), The Royal Air Force provided several aircrafts to thwart the German air force attack. The aircrafts provided by the Royal Air Force could be flown only by two people as a pilot and co-pilot. To get the best results in the mission to thwart German air force attack, the Royal Air Force brought some of the best pilots from all regions in Britain. However, there was a problem to solve by the Royal Air Force: among all the pilots that were called, some of them could not fly together in one aircraft because there were differences in terms of language, the ability of flying techniques, as well as in terms of flying experience. Suppose the Royal Air Force obtained 18 best pilots from all regions in Britain that were ready to fly the aircraft. Based on the problem, the Royal Air Force wished to pair the 18 pilots where each pair consisted of two pilots in such a way that the number of all the possible pairs is the highest number. In order to solve such problem, the Royal Air Force conducted a series of tests to the 18 pilots including language test, flying technique test, and psychological test to determine the suitability of the 18 pilots to fly together in one aircraft as a pilot and co-pilot.

Table of pairing between each aircraft pilot

The pilot's number (i=1,2, 3,,18)	The result of pairing between each aircraft pilot
1 st Pilot	Pilot: 2 nd and 17 th
2 nd Pilot	Pilot: 1 st , 3 rd , and 5 th
3 rd Pilot	Pilot: 2 nd , 4 th , 5 th , and 6 th
4 th Pilot	Pilot: 3 rd , 7 th , and 9 th
5 th Pilot	Pilot: 2 nd , 3 rd , and 6 th
6 th Pilot	Pilot: 3 rd , 5 th , and 15 th
7 th Pilot	Pilot: 4 th , 8 th , 9 th , 11 th , and 13 th
8 th Pilot	Pilot: 7 th , 9 th , and 10 th
9 th Pilot	Pilot: 4 th , 7 th , 8 th , 10 th , 13 th , and 15 th
10 th Pilot	Pilot: 8 th , 9 th , and 16 th
11 th Pilot	Pilot: 7 th , 12 th , 13 th , and 14 th
12 th Pilot	Pilot: 11 th and 14 th
13 th Pilot	Pilot: 7 th , 9 th , 11 th , and 14 th
14 th Pilot	Pilot: 11 th , 12 th , and 13 th
15 th Pilot	Pilot: 6 th , 9 th , 16 th , and 18 th
16 th Pilot	Pilot: 10 th , 15 th , and 18 th
17 th Pilot	Pilot 1 th
18 th Pilot	Pilot: 15 th and 16 th

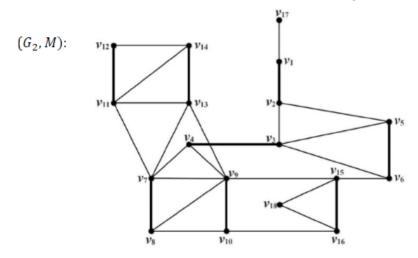
By transforming the problem into the form of graph, where the pilots are represented as a vertex and the corresponding relationship between each pilot is represented as an edge, the following graph is obtained:



By following the steps from Edmonds' cardinality matching algorithm, we obtain maximum matching on graph G_2 as follows:

Step 1 (initialization):

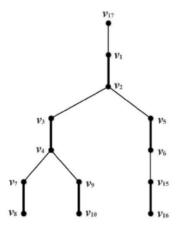
By doing the process of matching initialization to graph G_2 , we obtain graph (G_2, M) with $M = \{v_1v_2, v_3v_4, v_5v_6, v_7v_8, v_9v_{10}, v_{11}v_{12}, v_{13}v_{14}, v_{15}v_{16}\}$ as follows:



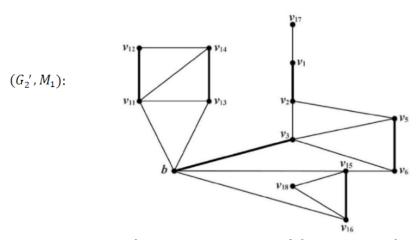
Since there are still two exposed vertices left that are v_{17} and v_{18} , then go to step 2.3 Step 2.3 :

Vertex v_{17} is selected as the root, by using BFS we obtain an alternating tree T_1 that corresponds to graph (G_2, M) as follows:

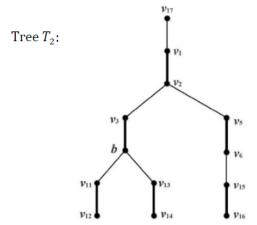
Tree T_1 :



Since blossom $B = \{v_4, v_8, v_9, v_{10}\}$ has been encountered, then go to step 4. Step 4: Shrink blossom $B = \{v_4, v_8, v_9, v_{10}\}$ to obtain a graph as follows:

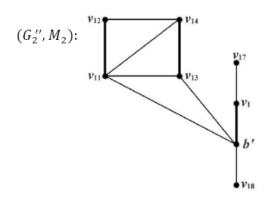


By continuing the constructing process of alternating tree that corresponds to graph (G_2', M_1) , we obtain:

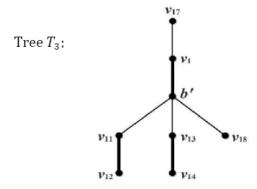


Since blossom $B'=\{b,v_2,v_3,v_5,v_6,v_{15},v_{16}\}$ has been encountered, then go to Step 4:

Shrink blossom $B' = \{b, v_2, v_3, v_5, v_6, v_{15}, v_{16}\}$ to obtain a graph as follows:



By continuing the constructing process of alternating tree that corresponds to graph (G_2'', M_2) , we obtain:



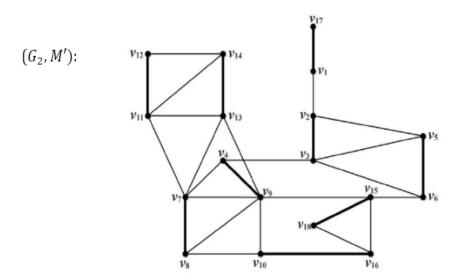
Since an augmenting path P'': $v_{18}-b'-v_1-v_{17}$ that contains *pseudovertex b'* has been obtained, then go to step 3.1

Step 3.1:

By unshrinking the pseudovertex b', we obtain augmenting path P': $v_{18} - v_{15} - v_{16} - b - v_3 - v_2 - v_1 - v_{17}$ that contains pseudovertex b, then go to step 3.1 Step 3.1:

By usnhrinking the pseudovertex b, we obtain augmenting path P: $v_{18}-v_{15}-v_{16}-v_{10}-v_{9}-v_{4}-v_{3}-v_{2}-v_{1}-v_{17}$ which does not contain any pseudovertex , then go to step 3.2 Step 3.2:

Augment the initial matching M using augmenting path P, we obtain a new matching in (G_2, M) as follows:



Since all vertices in (G_2, M') are incident with a matching edge, then go to step 5. Step 5:

Matching M' in graph G_2 is maximum, therefore the finding process is discontinued.

From the steps that have been done, we obtain a maximum matching M' in G_2 with cardinality of matching M' that is |M'| = 9. Matching M' that has been obtained is the solution of the problem of pairing the 18 pilots as described above with details as follows:

- 1. Pilot number 1 pairs up with pilot number 17
- 2. Pilot number 2 pairs up with pilot number 3
- 3. Pilot number 4 pairs up with pilot number 9
- 4. Pilot number 5 pairs up with pilot number 6
- 5. Pilot number 7 pairs up with pilot number 8
- 6. Pilot number 10 pairs up with pilot number 16
- 7. Pilot number 11 pairs up with pilot number 12
- 8. Pilot number 13 pairs up with pilot number 14
- 9. Pilot number 15 pairs up with pilot number 18

And the maximum number of aircrafts that can be flown is as many as 9 aircrafts.

7 | Conclusion

Matching M in graph G is a subset of E(G) edge set where there are no two edges in Matching M that meet each other at the same vertex in graph G. Generally, Edmonds' maximum matching algorithm is divided into three steps where the first step is the process of matching initialization. The second step is shrinking and unshrinking of blossom B_i . The third step is to execute augmenting-M if the augmenting P path has been found. Edmonds' blossom algorithm can be applied to find for maximum matching especially towards non-bipartite graph. The process of matching initialization is the effort to find the first matching to fasten the finding of maximum matching. Alternating T tree is a T graph concept that is used to find an augmenting path to obtain a new matching that has bigger cardinality than the first matching from the process of matching initialization.

8 | References

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