

# An Enumerative Algorithm for the Maximum Clique Problem

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## Abstract

Partial enumerative algorithms are presented for maximum clique problems which are very easy to implement. Search results for efficient implementation of any graph for randomly generated graphs of up to 10 vertices and more than ten sides. Appropriate specifications are also provided for testing problems to facilitate future comparisons. In addition, to understand the maximum clique search without using any software. Search results are presented in the form of examples.

**Keyword:** maximum clique, enumerative algorithm, current best clique.

## 1. Introduction

Given a graph  $G(V, E)$ , where  $V$  is the set of vertices and  $E$  is the set of edges, a complete subgraph of  $G$  is one whose vertices are pairwise adjacent. The *maximum clique problem* is to find the maximum complete subgraph of  $G$ . An independent set is a set of vertices whose elements are pairwise nonadjacent. The *maximum independent set problem* is to find the largest such independent set. A vertex cover is a set of vertices that covers all the edges of  $G$ . The *minimum vertex cover problem* is to find the minimum of such set. These three problems are computationally equivalent. They are also known to be NP-complete (P. Berman and G. Schnitger, 1989[2]).

The maximum clique problem has numerous applications in science and engineering (E. Balas and C.S. Yu, 1986) [1]. One way to cope with an NP-complete problem is to design an algorithm that performs fast on the average with respect to a natural probability distribution of inputs. Most of the algorithms for the maximum clique problem degenerate to an exhaustive search. An obvious enumerative algorithm is to consider all subsets of  $V$  in decreasing order of cardinality and check if each one forms a clique. Stop the checking when such a set has been found. Algorithms for the maximum clique problem have been studied by many authors (E. Balas and C.S. Yu, 1986), (C. Bron and J. Kerbosch, 1973) and (R.E. Tarjan and A.E. Trojanowski, 1977). Most of the proposed algorithms have been implemented and tested on medium size problems using random graphs. Recently, using a

branch and bound method based on quadratic 0-1 programming (P.M. Pardalos and G.P. Rodgers, 1990), problems with 1000 vertices and as many as 150000 edges were solved.

In this paper, we propose a simple and very efficient algorithm for finding the maximum clique in a graph  $G$ . The algorithm uses a partial enumeration and is very simple to implement. Computational experience with randomly generated graphs shows that the algorithm is faster than any previous known method.

## 2. Algorithm description

Initially, the algorithm considers an ordering of the vertices of  $G$ , say  $v_1, v_2, \dots, v_n$ , where  $v_1$  is a vertex of smallest degree in  $G$ ,  $v_2$  is a vertex of smallest degree in  $G - (v_1)$ , and generally  $v_k$  is a vertex of smallest degree in  $G - \{v_1, v_2, \dots, v_{k-1}\}$ ,  $k \leq n - 2$ .

It has been observed that this vertex ordering reduces the computational time in problems with dense graphs. In addition, this ordering can be reapplied at depths  $> 1$ . For lower density problems the algorithm is faster when no ordering of the vertices is performed.

Without any pruning, the algorithm produces a most  $n$  cliques of increasing size. Initially, it finds the largest clique  $C_1$  that contains the vertex  $v_1$ . Then it finds  $C_2$ , the largest clique in  $G - (v_1)$ , that contains  $v_2$  and so on. Applying heuristics and pruning techniques, we can reduce the search space dramatically.

Crucial to the understanding of the algorithm is the notion of the *depth*. Suppose we consider vertex  $v_1$ . At depth 2, we consider all vertices adjacent to  $v_1$ . At depth 3, we consider all vertices (that are in depth 2) adjacent to the first vertex in depth 2 (the example below makes this clear). Let  $v_{di}$  be the vertex we are currently expanding at depth  $d$  and step  $i$ . That is,

$$\text{Depth } d: v_{d1} \dots v_{di} \dots v_{dm}.$$

If  $d + (m - i) \leq \text{size of current best clique (CBC)}$ , then we prune, since the size of the largest possible clique (formed by expanding  $v_{di}$ ) would be less or equal to the size of CBC. If we are at depth 1 and this inequality holds then we stop; we have found the maximum clique (R. Carraghan and P. M. Pardalos, 1990).

If it is known that the maximum clique has size  $\geq \alpha$ , then we may use as pruning (stopping) criterion the following:

$$d + (m - i) \leq \alpha.$$

If  $\alpha$  is close to the size of the actual maximum clique, the computational time can be reduced dramatically for graphs with high density. In (P. Turan, 1954) proved that every graph with  $n$  vertices and  $m$  edges contains a clique of size  $\geq n^2 / (n^2 - 2m)$ .

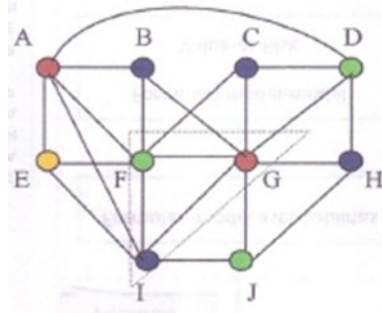
Hence, if the graph has density  $D$  then we may take

$$\alpha = \frac{n}{(1 - D)n + D}.$$

Unfortunately, the value of  $\alpha$  is helpful only if it is very close to the actual maximum clique size. However, polynomial time algorithms can be used to find very good values for  $\alpha$  (E. Balas and C.S. Yu, 1986).

## 3. Result

In this section we present results to show the efficiency of our algorithm. Consider the graph in Figure 1, with the steps of the algorithm shown in Table 1. We have found the maximum clique for this graph  $\{A, E, F, I\}$ , with size 4.



Figur 1. Vertex Coloring Graph.

Table 1. Vertex Coloring

1	Initialization orders the vertices	H J B C E D A I F G	
2	Depth 1:	<b>H</b> J B C E D A I F G	
	Depth 2:	<b>J D G</b>	
	Depth 3:	<b>G</b>	Cannot expand, so CBC is {H, J, G}, size 3
	Depth 2:	<b>J D G</b>	Since $d + (m - i) = 2 + (3 - 2) = 3 \leq 3$ , we prune.
	We are now finished with vertex H		
	The largest clique containing this node is {H, J, G}		
3	Depth 1:	H <b>J</b> B C E D A I F G	
	Depth 2:	I <b>G</b>	$2+(2-1) \leq \text{CBC}$ , so prune
4	Depth 1:	H J <b>B</b> C E D A I F G	
	Depth 2:	A <b>F</b> G	
	Depth 3:	<b>F</b>	Cannot expand, so CBC is {B, A, F}, size 3
	Depth 2:	A <b>F</b> G	$2+(3-2) \leq \text{CBC}$ , so prune
5	Depth 1:	H J B <b>C</b> E D A I F G	
	Depth 2:	D <b>F</b> G	
	Depth 3:	<b>G</b>	Cannot expand, so CBC is {C, D, G}, size 3
	Depth 2:	D <b>F</b> G	$2+(3-2) \leq \text{CBC}$ , so prune
6	Depth 1:	H J B C E <b>D</b> A I F G	
	Depth 2:	A <b>F</b> I	
	Depth 3:	<b>F</b> I	
	Depth 4:	<b>I</b>	{E, A, F, I} becomes our new CBC of size 4
	Depth 3:	<b>F</b> I	$3+(2-1) \leq 4$ , so prune
	Depth 2:	A <b>F</b> I	$2+(3-2) \leq \text{CBC}$ , so prune
7	Depth 1:	H J B C E D <b>A</b> I F G	$1+(10-4) \geq \text{CBC}$ and depth 1, so stop

We have the maximum clique for this graph {A, E, F, I}, with size 4 of step 6.

#### 4. Concluding remarks

We have shown that a simple algorithm implemented very efficiently can solve very large maximum clique problems. Since this algorithm is very simple to implement, it can be used as a basis for computational comparison when different algorithms are used. The proposed algorithm can be easily parallelized, since vertex  $v_{1k}$  (depth 1) can be processed independently of vertices  $v_{11}, \dots, v_{1(k-1)}$ . The performance of the algorithm for dense problems can be improved by reapplication of vertex ordering at depths  $> 1$ . In addition, the algorithm will be faster when a good heuristic is used to determine  $\alpha$ .

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