

COMMUNICATION

PSEUDODOMISHOLD GRAPHS

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A graph G is called pseudodomishold if there are nonnegative real numbers c_1, \dots, c_n, t (not all of them zero) such that for every subset U of VG and for its characteristic zero-one vector (e_1, \dots, e_n) the implications

$$(t < c_1 e_1 + \dots + c_n e_n) \Rightarrow (U \text{ is a dominating set of } G),$$

$$(t > c_1 e_1 + \dots + c_n e_n) \Rightarrow (U \text{ is not a dominating set of } G)$$

are true.

A pseudodomishold graph is called a hereditary one (briefly, HP-graph) if every its induced subgraph is also pseudodomishold.

In this note we describe the structure of HP-graphs in terms of the composition of simplest components and minimal forbidden induced subgraphs. This result gives rise to a simple recurrence formula and estimates for the number u_n of n -vertex HP-graphs. It also implies the linear-time algorithms recognizing HP-graphs and their degree sequences.

Up to now the various classes of P-threshold graphs, namely threshold, pseudothreshold and boxthreshold, have been thorough studied. A number of structure characterizations, recognition algorithms, classification and enumeration aspects concerning P-threshold graphs can be found in [1–9]. There has been little knowledge of pseudodomishold graphs. Unlike thresholdness, pseudothresholdness and domisholdness in definition of which a separating hyperplane plays a principal role, pseudodomisholdness is not a hereditary property, i.e. it is not preserved in proceeding to some induced subgraph. The aim of this communication is to give description and enumeration of hereditary pseudodomishold graphs (briefly, HP-graphs). It is a rather wide class containing the ones of domishold and threshold graphs [1–2].

The main result is Theorem 1 characterizing HP-graphs in terms of the composition of the simplest components and minimal forbidden induced subgraphs. It, in turn, gives rise to a simple recurrence formula and estimates for the number u_n of n -vertex HP-graphs (Corollaries 1–2). Theorem 1 also implies an easy guessed linear-time algorithms recognizing HP-graphs and their degree sequences. It is interesting to compare the estimates for u_n with the asymptotics for the number d_n of n -vertex domishold graphs given in Corollary 3.

All graphs considered are finite, undirected, without loops and multiple edges. VG is the vertex set of a graph G and EG is its edge set. A n -vertex graph G is called pseudodomishold if there are nonnegative real numbers c_1, \dots, c_n, t , not all of them equal zero, such that for every subset U of VG we have

$$\begin{aligned} (c_1 e_1 + \dots + c_n e_n > t) &\Rightarrow (U \text{ is dominating}), \\ (c_1 e_1 + \dots + c_n e_n < t) &\Rightarrow (U \text{ is not dominating}) \end{aligned} \quad (1)$$

where (e_1, \dots, e_n) is the characteristic vector of U .

If the first inequality in (1) were substituted by the nonstrict inequality $c_1 e_1 + \dots + c_n e_n \geq t$ we would have the definition of a domishold graph [1].

Notation. \bar{G} is a graph being complementary to G . K_n is a n -vertex complete graph. M is the set of graphs H such that either any vertex degree in H exceeds $|VH| - 4$ (i.e. all connected components of \bar{H} are only chains or cycles) or H is a 5-vertex chain. t_n is the number of n -vertex graphs from M not containing dominating vertices, isolated vertices and 2-vertex edgeless dominating sets.

So called *Forbidden* configurations are shown in Fig. 1: broken lines indicate non-edges.

We will write $G(A, B)$ if the fixed partition $VG = A \cup B$ is meant ($A \cap B = \emptyset$). One of the parts A, B can be empty. If $G = \bar{K}_n$ then Q_n will denote $G(VG, \emptyset)$. If $G = K_1$ then O_1 will denote $G(\emptyset, VG)$. If G is a 4-vertex chain then P_4 will denote the graph $F(A, B)$ where $|A| = |B| = 2$ and B consists of vertices with degrees being equal to 1.

The composition \circ is defined as follows [8]: given $G(A, B)$ and H , $VH \cap VG = \emptyset$ then

$$G(A, B) \circ H = G \cup H \cup K_{A, VH}$$

where \cup is the sign of the union of graphs, $K_{A, VH}$ is the complete bipartite graph with parts A and VH .

Theorem 1. The following assertions are equivalent:

- (i) G is a HP-graph.

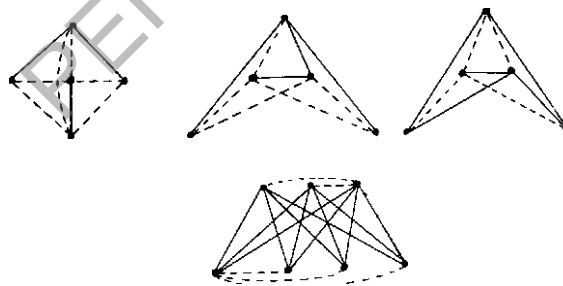


Fig. 1.

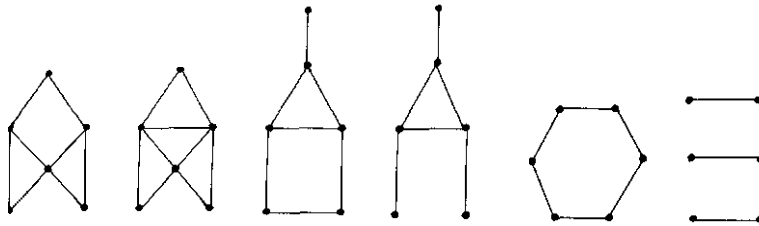


Fig. 2.

(ii) G contains neither forbidden configurations shown in Fig. 1 nor induced subgraphs shown in Fig. 2.

(iii) G can be represented in the form

$$G = X_1 \circ \dots \circ X_m \circ H, m \geq 0,$$

where $H \in M$, $X_i \in \{Q_1, Q_2, O_1, P_4\}$.

Corollary 1.

$$u(x) = \frac{t(x) + x - x^3}{-x^4 + x^3 - x^2 - 2x + 1}$$

where $u(x) = \sum_{n=0}^{\infty} u_n x^n$, $t(x) = \sum_{n=1}^{\infty} t_n x^n$.

Corollary 2. For $n \geq 6$

$$(2, 32)^{n-1} < u_n < (2, 37)^{n-1}.$$

Corollary 3. If d_n is the number of n -vertex domishold graphs then

$$d(x) = \sum_{n=1}^{\infty} d_n x^n = \frac{-x^3 + x}{x^3 - x^2 - 2x + 1}$$

and

$$d_n \sim cr^n$$

where r is the maximum (in absolute value) root of the equation $x^3 - 2x^2 - x + 1 = 0$ ($r \approx 2, 24$).

Problem. What is the complexity of recognizing pseudodomishold (not necessary hereditary) graphs?

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