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Block allocation of a sequential resource

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Abstract

An H -packing of G is a collection of vertex-disjoint subgraphs of G such that each component is isomorphic to H . An H -packing of G is maximal if it cannot be extended to a larger H -packing of G . In this paper we consider problem of random allocation of a sequential resource into blocks of m consecutive units and show how it can be successfully modeled in terms of maximal P_m -packings. We enumerate maximal P_m -packings of P_n of a given cardinality and determine the asymptotic behavior of the enumerating sequences. We also compute the expected size of m -packings and provide a lower bound on the efficiency of block-allocation.

Keywords: Maximal matching, maximal packing.

Math. Subj. Class.: 05C70, 05A15, 05A16

1 Matchings and packings

A *matching* M in a graph G is a collection of edges of G such that no two edges from M have a vertex in common. The number of edges of M is called the *size* of the matching. Small matchings are not interesting – they are easy to find and enumerate. Hence, we are mostly interested in matchings that are as large as possible. There are two ways to quantify the idea of “large” matchings, one of them based on their cardinality, the other based on the set inclusion.

A matching M is *maximum* if there is no matching in G with more edges than M . The cardinality of any maximum matching in G is called the *matching number* of G and denoted by $\nu(G)$. The matching number of a graph on n vertices, obviously, cannot exceed $\lfloor n/2 \rfloor$, since each edge saturates two vertices. A matching that saturates all vertices of G is called a *perfect matching*.

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A matching M in G is *maximal* if it cannot be extended to a larger matching in G , i.e., if no other matching in G contains it as a proper subset. Obviously, every maximum matching is also maximal, but the opposite is generally not true. The cardinality of any smallest maximal matching in G , denoted by $s(G)$, is the *saturation number* of G ; the largest size of a maximal matching is, of course, $\nu(G)$.

Matchings are natural models for many problems in natural, technical and social sciences. Worth mentioning are applications of perfect matchings in organic chemistry and solid state physics. For a general background on matching theory and terminology we refer the reader to the classical monograph by Lovász and Plummer [14]. For graph theory terms not defined here we also recommend [3, 19].

A closely related concept of packing is a generalization of matching. There are several varieties of packing; we consider here only the simplest case. An H -packing of G is a collection of vertex-disjoint subgraphs of G such that each component is isomorphic to H [3]. Hence, a matching of G is a P_2 -packing in G , where P_2 denotes a path on 2 vertices. Again, we are interested only in large packings. If a packing is a spanning subgraph, we say that the packing is *perfect*; if no other H -packing has more components, the packing is *maximum*; finally, if an H -packing cannot be extended to a valid H -packing, we say that it is a *maximal H -packing*. The *H -packing number* and *H -saturation number* are defined in the same way as for matchings. When $H = P_m$ we denote these two quantities by $\nu_m(G)$ and $s_m(G)$ and call them the *m -packing number* and *m -saturation number*, respectively. We refer the reader to [12, 13] for some aspects of P_3 -packings in claw-free and in subcubic graphs and to [15] for similar problems in directed graphs.

Maximal matchings and packings can serve as models of several physical and technical problems such as the block-allocation of a sequential resource or adsorption of dimers and/or polymers on a structured substrate or a molecule. When that process is random, it is clear that the substrate can become saturated by a number of units much smaller than the theoretical maximum. The respective saturation numbers provide an information on the worst possible case of clogging; they measure how inefficient the adsorption or the allocation process can be. However, in order to assess its efficiency, we also need to know how likely it is that a given number of units will saturate the substrate. Hence, we must study the enumerative aspects of the problem.

For the matching case, the question has been answered in [7]. The main goal of this paper is to contribute to the corpus of knowledge about the enumerative aspects of maximal P_m -packings in paths and cycles. Specifically, we compute the efficiency of block-allocation of length m of a sequential linear or cyclic resource. In some cases we provide explicit formulas for the number of maximal m -packings of a given cardinality, while in other cases we establish the recurrences for the enumerating sequences and then use their uni- and bivariate generating functions to determine their asymptotic behavior.

Finally, in the concluding section we discuss some open problems and indicate some directions of possible future research.

2 Paths and cycles

2.1 Paths

We remind the reader that throughout this paper P_n denotes the path on n vertices, hence of length $n - 1$. As a motivation, we consider a parking lot made of n parallel concrete strips such that a car can be parked on any two neighboring strips. In ideal situation, when all

drivers take care and park responsibly, the lot can accommodate $\lfloor n/2 \rfloor$ cars. However, if the drivers are careless, the lot can become saturated by a smaller number of cars, as shown in Figure 1. In the worst possible case, it can become saturated by as few as $\lfloor (n+1)/3 \rfloor$

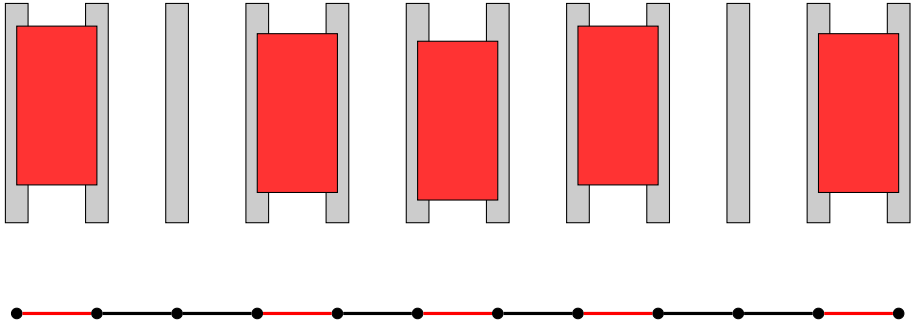


Figure 1: A saturated parking lot and the corresponding maximal matching.

cars. Hence, it is of interest to find out how likely is this to happen, and what is the expected number of cars under the random regime.

In the continuous setting, this problem is known as the random car-parking problem of Rényi [16, 17], while in discrete setting it has a natural representation as a problem of maximal matching in P_n , as shown in Figure 1; it was considered in detail in [7], where its full solution was obtained, including the explicit formulas for the number of different configurations accommodating a given number of cars. Also, the expected number of cars under the random regime was computed, and the asymptotic behavior of the sequence enumerating all possible parking arrangement was determined.

But what happens if we wish to park trucks such that each of them is twice as wide as a car? Each truck will then consume three consecutive strips, as shown in Figure 2, and the corresponding graph-theoretical model will not be a matching, but a packing of copies of

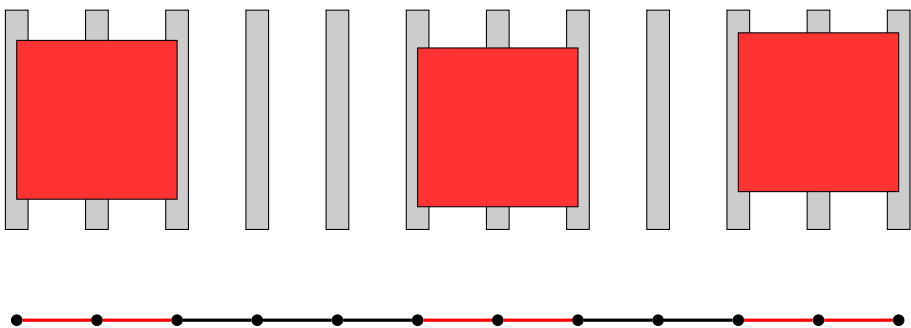


Figure 2: A parking lot saturated with trucks.

P_3 in P_n . Obviously, the structure of the problem remains the same if instead of parking lots and cars and trucks we consider any sequential resource of length n which is allocated in blocks of $m \geq 2$ consecutive units. All such situations can be studied as problems of packing copies of P_m in P_n . We call such a packing an m -packing. In this subsection we consider the enumerative aspects of m -packings in paths. Before counting them, we state

(without proof) two results about the smallest and the largest possible size of m -packings in P_n .

Proposition 2.1. *Let P_n be a path on n vertices. Then*

$$s_m(P_n) = \left\lfloor \frac{n + m - 1}{2m - 1} \right\rfloor \quad \text{and} \quad \nu_m(P_n) = \left\lfloor \frac{n}{m} \right\rfloor.$$

We now start counting all maximal m -packings in P_n . Let $\psi_{n,k}^{(m)}$ denote the total number of maximal m -packings in P_n with exactly k copies of P_m .

Proposition 2.2. *The sequence $\psi_{n,k}^{(m)}$ is given by the recurrence*

$$\psi_{n,k}^{(m)} = \sum_{l=m}^{2m-1} \psi_{n-l,k-1}^{(m)}$$

for $n \geq 2m - 1$ and with the initial conditions

$$\psi_{0,0}^{(m)} = \psi_{1,0}^{(m)} = \dots = \psi_{m-1,0}^{(m)} = 1$$

and $\psi_{l,0}^{(m)} = 0$ for all other values of l .

Proof. Let us label the vertices of P_n by v_1, \dots, v_n . Let v_l be the vertex with the highest label that is covered by a copy of P_m in a maximal m -packing of size k . Clearly, $v_l \in \{v_{n-m+1}, \dots, v_n\}$ (otherwise there would be enough place to pack one more copy of P_m , contrary to the assumption of maximality), and the remaining $k - 1$ copies of P_m must form a valid maximal packing of P_m of size $k - 1$ in the remaining portion of P_n , i.e., in P_{l-m+1} . The initial conditions count trivial packings of size zero. \square

From the above recurrence one can immediately compute the bivariate generating function for the numbers $\psi_{n,k}^{(m)}$ by multiplying them throughout by $x^n y^k$ and summing over all $n \geq 2m - 1, k \geq 1$. We state the result omitting the computational details.

Theorem 2.3. *Let $F_m(x, y) = \sum_{n,k \geq 0} \psi_{n,k}^{(m)} x^n y^k$ be the bivariate generating function of $\psi_{n,k}^{(m)}$. Then*

$$F_m(x, y) = \frac{p_m(x)}{1 - yq_m(x)},$$

where $p_m(x) = \frac{1-x^m}{1-x}$ and $q_m(x) = x^m p_m(x)$.

Corollary 2.4. *The bivariate generating function of $\psi_{n,k}^{(m)}$ is given by*

$$F_m(x, y) = \frac{1 - x^m}{1 - x - x^m(1 - x^m)y}.$$

The generating function $F_m(x) = \sum_{n \geq 0} \psi_n^{(m)} x^n$ for the sequence enumerating the total number of m -packings in P_n is now obtained by substituting $y = 1$ into the expression for $F_m(x, y)$.

Corollary 2.5. *The generating function of the sequence enumerating the total number of maximal m -packings in P_n is given by*

$$F_m(x) = \frac{1 - x^m}{1 - x - x^m + x^{2m}}.$$

From the above result we can deduce the recurrence satisfied by $\psi_n^{(m)}$.

Corollary 2.6. *The numbers $\psi_n^{(m)}$ satisfy the recurrence*

$$\psi_n^{(m)} = \psi_{n-m}^{(m)} + \dots + \psi_{n-2m+1}^{(m)}$$

for $n \geq 2m - 1$ with the initial conditions $\psi_0^{(m)} = \dots = \psi_m^{(m)} = 1$ and $\psi_{m+i}^{(m)} = i + 1$ for $1 \leq i \leq m - 2$.

The numbers $\psi_{n,k}^{(m)}$ form a triangular array with rows indexed by n and columns indexed by k . It can be deduced from the form of the bivariate generating function that the columns are, in fact, shifted rows of the triangle of multinomial (m -nomial) coefficients. Recall that the (p, q) -th m -nomial coefficient

$$t_{p,q}^{(m)} = \sum_{i=0}^{\lfloor q/m \rfloor} (-1)^i \binom{p}{i} \binom{p+q-1-im}{p-1}$$

is the coefficient of x^q in $(1 + x + \dots + x^{m-1})^p$. (See, for example, sequence [A035343](#) in [18] for $m = 5$.) The observation can be formally stated in the following way.

Corollary 2.7.

$$\psi_{n,k}^{(m)} = t_{k+1, n-mk}^{(m)}$$

As a consequence, we can obtain formulas for $\psi_{n,k}^{(m)}$ and $\psi_n^{(m)}$. We refer the reader to the *On-Line Encyclopedia of Integer Sequences* for more details on multinomial coefficients [18].

Corollary 2.8.

$$\psi_{n,k}^{(m)} = \sum_{i=0}^{\lfloor \frac{n-k}{m} \rfloor} (-1)^i \binom{k+1}{i} \binom{n+k-m(i+k)}{k};$$

$$\psi_n^{(m)} = \sum_{k=0}^{\lfloor \frac{n}{m} \rfloor} \sum_{i=0}^{\lfloor \frac{n-k}{m} \rfloor} (-1)^i \binom{k+1}{i} \binom{n+k-m(i+k)}{k}.$$

When $m = 2$, the above formulas reduce to known results about the number of maximal matchings [7].

As a further consequence, we note that the number of all maximal m -packings of size k in all paths is given by m^{k+1} .

Our next goal is to determine the asymptotic behavior of the enumerating sequences and then use it to compute the expected size of a maximal m -packing in P_n . We rely on the following version of Darboux's theorem [2].

Theorem A. *If the generating function $f(x) = \sum_{n \geq 0} a_n x^n$ of a sequence (a_n) can be written in the form $f(x) = (1 - \frac{x}{w})^\alpha h(x)$, where w is the smallest modulus singularity of f and h is analytic in w , then $a_n \sim \frac{h(w)n^{-\alpha-1}}{\Gamma(-\alpha)w^n}$, where Γ denotes the gamma function.*

As a consequence, the expected size of a maximal m -packing in P_n , $\pi_m(P_n)$, can be computed as

$$\pi_m(P_n) = \frac{[x^n] \frac{\partial F_m(x,y)}{\partial y} \Big|_{y=1}}{[x^n] F_m(x,y) \Big|_{y=1}},$$

where $[x^n]F(x)$ denotes the coefficient of x^n in the expansion of $F(x)$.

We refer the reader to [2, 20] for more information on obtaining the asymptotics of a sequence from its generating function.

We start by observing that $F_m(x) = F_m(x,y) \Big|_{y=1}$ and $\frac{\partial F_m(x,y)}{\partial y} \Big|_{y=1}$ can be represented as

$$F_m(x) = \left(1 - \frac{x}{w_m}\right)^{-1} \frac{p_m(x)}{w_m \frac{1-q_m(x)}{w_m-x}} = \left(1 - \frac{x}{w_m}\right)^{-1} g_m(x)$$

and

$$\frac{\partial F_m(x,y)}{\partial y} \Big|_{y=1} = \left(1 - \frac{x}{w_m}\right)^{-2} \frac{p_m(x)q_m(x)}{\left[w_m \frac{1-q_m(x)}{w_m-x}\right]^2} = \left(1 - \frac{x}{w_m}\right)^{-2} h_m(x).$$

Here w_m denotes the smallest (and the only) real solution of the equation $q_m(x) = 1$. By plugging this into Theorem A we obtain following results.

Theorem 2.9. *The asymptotics of the number of m -packings in P_n is given by*

$$\psi_n^{(m)} \sim g_m(w_m) \cdot w_m^{-n}.$$

Theorem 2.10. *The expected size of a maximal m -packing in P_n is given by*

$$\pi_m(P_n) = \frac{1}{w_m q'_m(w_m)} n,$$

where w_m is the only real solution of $q_m(x) = 1$.

Now we can define the *efficiency* of random m -packing in P_n as the quotient of the expected and the optimal size of an m -packing. Since the size of any largest possible m -packing in P_n is $\lfloor n/m \rfloor$, the efficiency is given by

$$\varepsilon(m) = \frac{m}{w_m q'_m(w_m)}.$$

It is, hence, of interest to investigate the behavior of the above quotient for large values of n and m . (We will assume that $n \gg m$, since the opposite case is not very interesting.) Numerical computations indicate that it initially decreases from 0.823 for $m = 2$ and achieves the minimum value of 0.758317 for $m = 9$, and then increases slowly (apparently monotonously) so that for $m = 100$ it has the value of approximately 0.796. In the rest of this subsection we show that $\varepsilon(m)$ remains bounded from below for all values of m .

For the beginning, we transform the expression for $q'_m(x)$ as follows:

$$\begin{aligned} q'_m(x) &= \frac{mx^{m-1}}{1-x}(1-2x^m) + \frac{x^m(1-x^m)}{1-x} \\ &= \frac{x^m(1-x^m)}{1-x} \left[\frac{2m}{x} + \frac{1}{1-x} - \frac{m}{x} \frac{1}{1-x^m} \right] \end{aligned}$$

By plugging in $x = w_m$, the first term on the right-hand side becomes 1, and by multiplying the resulting equation through by w_m , we obtain

$$w_m q'_m(w_m) = \left(2 - \frac{1}{1-w_m^m} \right) m + \frac{w_m}{1-w_m}.$$

We would like to estimate the right-hand side and give some upper bound. The first term never exceeds m ; it is enough to note that $w_m > 1/2$ for all $m \geq 2$, and from there it follows $2 - \frac{1}{1-w_m^m} < 2 - \frac{1}{1-2^{-m}} < 1$. In order to bound the second term, we notice that for large enough values of m we must have $w_m < 1 - \frac{3}{m}$. Indeed, this is equivalent to

$$\left(1 - \frac{3}{m} \right)^m - \left(1 - \frac{3}{m} \right)^{2m} > \frac{3}{m},$$

and this is true, since the left-hand side tends to $e^{-3} - e^{-6} \approx 0.047308$, while the right-hand side tends to zero. Numerical computations show that “large enough” here means $m = 68$. By plugging in the upper bound $w_m < 1 - \frac{3}{m}$ into the second term, we obtain $\frac{w_m}{1-w_m} < \frac{m}{3}$. Now the right-hand side can be bounded from above by $\frac{4m}{3}$. This gives us a lower bound on the efficiency.

Proposition 2.11. *The efficiency of m -packings is bounded from below. For all $m \geq 2$,*

$$\varepsilon(m) > \frac{3}{4}.$$

The same argument as above could be used to show that for large enough values of m and for any real $a > 0$, an expression of the type $1 - \frac{a}{m}$ will be an upper bound on w_m . This implies that the right-hand side of the expression for $w_m q'_m(w_m)$ can be bounded from above by $\frac{a+1}{a} m$, and consequently, that $\lim_{m \rightarrow \infty} \varepsilon(m) = 1$.

Our results indicate that longer blocks achieve better efficiency of random block allocation of a sequential resource. The dependency is rather mild, and the growth is slow. For example, a hundredfold increase of the block length from $m = 1000$ to $m = 100\,000$ results in the moderate increase of efficiency from $\varepsilon(1000) = 0.844$ to $\varepsilon(100\,000) = 0.903$. Still, the block length of nine seems to be a bad choice.

Before we move to the cycles, we mention that our analysis assumes that all packings are equally probable. It is known for maximal matchings that the efficiency is slightly better if instead one considers dynamics, i.e., the situation where the dimers arrive sequentially and try to bind to the substrate [9]. It would be interesting to see how such approach would affect the efficiency here.

2.2 Cycles

Let us now consider the number of maximal m -packings in a cycle C_n of length $n \geq 3$, $n \geq m$. We denote it by $\varphi_n^{(m)}$, and the number of maximal m -packings in C_n of size k by $\varphi_{n,k}^{(m)}$.

Proposition 2.12. *The numbers $\varphi_{n,k}^{(m)}$ are given by*

$$\varphi_{n,k}^{(m)} = m\psi_{n-m,k-1}^{(m)} + \sum_{i=1}^{m-1} i\psi_{n-2m-1,k-2}^{(m)}$$

for $n \geq 3, k \geq 2$, where $\psi_{n,k}^{(m)}$ count maximal m -packings of size k in P_n .

Proof. Let us consider vertex v_n in C_n . If it is not covered by a copy of P_m in an m -packing, then it must be in a “hole” of size i for some $1 \leq i \leq m - 1$. At each side of the hole there must be a copy of P_m . Hence the remaining $k - 2$ copies of P_m must form a valid m -packing in P_{n-2m-1} , and those are counted by $\psi_{n-2m-1,k-2}^{(m)}$. As there are i holes of size i containing vertex v_n , the second term in the right-hand side of the above expression counts all of them. The first term counts the m -packings in C_n that cover v_n by a copy of P_m . \square

Proposition 2.13. *The numbers $\varphi_n^{(m)}$ satisfy the same recurrence as the numbers $\psi_n^{(m)}$, i.e.,*

$$\varphi_n^{(m)} = \varphi_{n-m}^{(m)} + \dots + \varphi_{n-2m+1}^{(m)}$$

with the initial conditions

$$\varphi_3^{(m)} = \dots = \varphi_{m-1}^{(m)} = 1$$

and $\varphi_{m+i}^{(m)} = m + i$ for $0 \leq i \leq m - 1$.

Hence, the asymptotic behavior, the expected size and the efficiency of m -packings in C_n are the same as in P_n .

3 Future developments

This manuscript presents a systematic attempt to address enumerative aspects of maximal P_m -packings in some classes of graphs with simple connectivity patterns. It continues the line of research of a recent paper concerned with maximal matchings [7]. As this is, to the best of my knowledge, the first paper of this type, it leaves unanswered many questions that arise in the course of research. In this last section we outline some of the open problems and suggest some possible directions for future research.

The most natural thing would be to count m -packings in some other families of graphs with repetitive structure that have low connectivity. Examples of such graphs are cactus chains, such as those considered in [5, 6, 7]. Due to their simple structure, it is reasonable to expect that the enumerating sequences will satisfy (rather short) linear recurrences with constant coefficients, yielding thus to the same type of asymptotic analysis as obtained here. Besides finding the asymptotics, an interesting problem would be to find the extremal chains. For maximal matchings ($m = 2$) the problem is solved for hexagonal cacti and it would be interesting to see if the pattern persists for larger values of m .

Another promising class could be the so-called thorny graphs. From a given graph G one obtains the t -thorny graph $T_t(G)$ by appending t pendent vertices to every vertex of G . When G has a simple structure, the methods of this paper could be employed to obtain the recurrences for the number of m -packings in $T_t(G)$. As an example, we consider 3-packings in $T_t(P_n)$.

Proposition 3.1. Let $p_n^{(3)}$ denote the number of 3-packings in $T_t(P_n)$. Then

$$p_n^{(3)} = \binom{t}{2} p_{n-1}^{(3)} + 2t p_{n-2}^{(3)} + p_{n-3}^{(3)}$$

for $n \geq 3$ with the initial conditions that can be verified by direct computation.

The next step could be to consider linear polymers of connectivity 2. Among them, the most interesting are without doubt the benzenoid chains. Again, there are some results for maximal matchings [6, 7] for benzenoid and polyomino chains, but for other classes of fascia- and rota-graphs [11] not even that case is investigated.

Another direction could be to consider structural and enumerative problems of m -packings in composite graphs, i.e., in graphs that arise from simpler building blocks via various binary operations known as graph products. We have considered here one such example of low connectivity (the thorny graph, that could be thought of as the corona product of G and $\overline{K_t}$). However, many interesting operations such as, e.g., the Cartesian product, actually increase the connectivity. It would be too optimistic to expect that complete results of the type presented here could be obtained in general cases, but we believe that the cases when one component is a path or a cycle should be feasible. Another interesting problem would be to determine the m -saturation number of such graphs, in particular for the finite portions of grids and lattices. Also, nanostructures and fullerenes are natural candidates for investigation of structural properties related to m -packings. The results would generalize those for maximal matchings [1, 4].

A graph G is *equimatchable* [10, 14] if every maximal matching in G is also maximum, i.e., if all maximal matchings are of the same size. What can be said about *equipackable* graphs in which every maximal m -packing is also maximum m -packing?

Finally, it would be interesting to see if packing polynomials and maximal packing polynomials, modelled after their matching counterparts [7, 8, 14], would be useful in the study of packing enumeration.

References

- [1] V. Andova, F. Kardoš and R. Škrekovski, Sandwiching saturation number of fullerene graphs, *MATCH Commun. Math. Comput. Chem.* **73** (2015), 501–518, http://match.pmf.kg.ac.rs/electronic_versions/Match73/n2/match73n2_501-518.pdf.
- [2] E. A. Bender and S. G. Williamson, *Foundations of Combinatorics with Applications*, Dover, 2006, <http://www.math.ucsd.edu/~ebender/CombText/>.
- [3] R. Diestel, *Graph Theory*, volume 173 of *Graduate Texts in Mathematics*, Springer, Heidelberg, 4th edition, 2010, doi:10.1007/978-3-642-14279-6.
- [4] T. Došlić, Saturation number of fullerene graphs, *J. Math. Chem.* **43** (2008), 647–657, doi:10.1007/s10910-006-9217-3.
- [5] T. Došlić and F. Måløy, Chain hexagonal cacti: matchings and independent sets, *Discrete Math.* **310** (2010), 1676–1690, doi:10.1016/j.disc.2009.11.026.
- [6] T. Došlić and T. Short, Maximal matchings in polyspiro and benzenoid chains, submitted, [arXiv:1511.00590](https://arxiv.org/abs/1511.00590) [math.CO].
- [7] T. Došlić and I. Zubac, Counting maximal matchings in linear polymers, *Ars Math. Contemp.* **11** (2016), 255–276, doi:10.26493/1855-3974.851.167.

- [8] E. J. Farrell, An introduction to matching polynomials, *J. Comb. Theory Ser. B* **27** (1979), 75–86, doi:10.1016/0095-8956(79)90070-4.
- [9] P. J. Flory, Intramolecular reaction between neighboring substituents of vinyl polymers, *J. Am. Chem. Soc.* **61** (1939), 1518–1521, doi:10.1021/ja01875a053.
- [10] A. Frendrup, B. Hartnell and P. D. Vestergaard, A note on equimatchable graphs, *Australas. J. Combin.* **46** (2010), 185–190, https://ajc.maths.uq.edu.au/pdf/46/ajc_v46_p185.pdf.
- [11] M. Juvan, B. Mohar, A. Graovac, S. Klavžar and J. Žerovnik, Fast computation of the Wiener index of fasciagraphs and rotagraphs, *J. Chem. Inf. Comput. Sci.* **35** (1995), 834–840, doi:10.1021/ci00027a007.
- [12] A. Kelmans, Packing 3-vertex paths in claw-free graphs and related topics, *Discrete Appl. Math.* **159** (2011), 112–127, doi:10.1016/j.dam.2010.05.001.
- [13] A. Kosowski, M. Małafiejski and P. Żyliński, Packing three-vertex paths in a subcubic graph, in: S. Felsner (ed.), *2005 European Conference on Combinatorics, Graph Theory and Applications (EuroComb '05)*, DMTCS, Nancy, France, volume AE of *DMTCS Proceedings Series*, 2005 pp. 213–218, proceedings of a conference held at Technische Universität, Berlin, September 5–9, 2005, <https://dmtcs.episciences.org/3413>.
- [14] L. Lovász and M. D. Plummer, *Matching Theory*, volume 121 of *North-Holland Mathematics Studies*, North-Holland, Amsterdam, 1986.
- [15] S. Pantel, *Graph Packing Problems*, Master's thesis, Simon Fraser University, Canada, 1993.
- [16] M. D. Penrose, Random parking, sequential adsorption, and the jamming limit, *Comm. Math. Phys.* **218** (2001), 153–176, doi:10.1007/s002200100387.
- [17] A. Rényi, On a one-dimensional problem concerning random space filling, *Magyar Tud. Akad. Mat. Kutató Int. Közl.* **3** (1958), 109–127.
- [18] N. J. A. Sloane (ed.), *The On-Line Encyclopedia of Integer Sequences*, published electronically at <https://oeis.org>.
- [19] D. B. West, *Introduction to Graph Theory*, Prentice Hall, Upper Saddle River, New Jersey, 1996.
- [20] H. S. Wilf, *generatingfunctionology*, Academic Press, Boston, Massachusetts, 1990.