

Comparing mean field and Euclidean matching problems

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Received: 1st December 1997 / Revised: 6 May 1998 / Accepted: 30 June 1998

Abstract. Combinatorial optimization is a fertile testing ground for statistical physics methods developed in the context of disordered systems, allowing one to confront theoretical mean field predictions with actual properties of finite dimensional systems. Our focus here is on minimum matching problems, because they are computationally tractable while both frustrated and disordered. We first study a mean field model taking the link lengths between points to be independent random variables. For this model we find perfect agreement with the results of a replica calculation, and give a conjecture. Then we study the case where the points to be matched are placed at random in a d -dimensional Euclidean space. Using the mean field model as an approximation to the Euclidean case, we show numerically that the mean field predictions are very accurate even at low dimension, and that the error due to the approximation is $O(1/d^2)$. Furthermore, it is possible to improve upon this approximation by including the effects of Euclidean correlations among k link lengths. Using $k = 3$ (3-link correlations such as the triangle inequality), the resulting errors in the energy density are already less than 0.5% at $d \geq 2$. However, we argue that the dimensional dependence of the Euclidean model's energy density is non-perturbative, *i.e.*, it is beyond all orders in k of the expansion in k -link correlations.

PACS. 75.10.Nr Spin-glass and other random models – 02.60.Pn Numerical optimization

1 Introduction

1.1 Background

The study of disordered and frustrated systems, and in particular spin-glasses, has long been a major issue in condensed matter physics (for reviews see [1–3]). Most efforts have been based on replicas, in part because that method has led to the exact solution [4,5] of the Sherrington-Kirkpatrick (SK) model [6]. However, since the SK model is of infinite range, it is not clear [7–9] how relevant its solution is for understanding finite dimensional spin-glasses such as the Edwards-Anderson (EA) model [10]. The application of the replica formalism to finite dimensional systems, on the other hand, is hampered by two major difficulties [11,12]. First, in the saddle point equations, finite connectivities lead to an infinite number of order parameters: one has to deal with order parameters $q_{\alpha\beta\gamma\dots}$ having an arbitrarily large number of indices [13]. In contrast, the SK model requires only the order parameter $q_{\alpha\beta}$, with two indices. The second difficulty comes from the Euclidean nature of space: the metric structure introduces constraints on the possible values of the quenched disorder variables between points i and j . In the infinite range model, these variables are independent, but in the

short range models, the values allowed depend on the distance between points i and j , thereby introducing strong correlations. These correlations make the replica analysis much more difficult.

In this article we study matching problems; these are disordered and frustrated short range models arising in combinatorial optimization. They are simpler than spin-glasses and the difficulties just stressed have to a large extent been overcome. In particular, for the minimum matching problem based on *independent random* link lengths, Mézard and Parisi (M & P) have worked out the order parameters $q_{\alpha\beta\gamma\dots}$ exactly [14] and have introduced a link correlation expansion [15] to take into account correlations among link variables. Using extensive numerical simulations of the ground states, we study the following: (i) the probability distribution of ground state energies in the disorder ensembles; (ii) the distribution of “local” energies in the ground state configurations; (iii) the validity of their replica approach in the case without correlations; (iv) the accuracy of this random-link length model as a mean-field approximation to the Euclidean model; (v) the accuracy of the link correlation expansion.

1.2 Models

Consider N points (N even), and a set of links with lengths $l_{ij} = l_{ji}$ connecting points i and j , ($i, j = 1, \dots, N$).

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We call an *instance* a specification of these link lengths. A *matching* of the points is a set of bonds in which each of the N points is the extremity of one and only one bond. In effect this is a dimerization: points are linked in pairs. The length or cost of a matching is then the sum of the lengths of its bonds. (Since the points will often belong to a Euclidean space, we will use the term length rather than cost hereafter.) The minimum matching problem (MMP) [16] can be defined as the problem of finding a matching of minimum length, given the l_{ij} 's. A variant of this problem is the minimum bipartite matching problem (MBMP) [16]: we now have *two* sets of N points instead of one (N is no longer necessarily even) and a bipartite matching consists of a dimerization where each pair contains one point from each set. The length and the minimization of the bipartite matching are defined as for the MMP. From an algorithmic point of view, both problems belong to the class P , meaning that the exact solution of any instance of size N can be found in a time at most polynomial in N . In fact, the MMP can be solved in a time of $O(N^3)$ and the MBMP in a time of $O(N^2 \ln N)$. This property is important for a numerical study since it allows extensive exact computations. We have implemented the $O(NE \ln(E))$ matching algorithm as exposed in [17] where E is the number of edges in the graph. The resulting program solves a typical $N = 200$ random point instance in less than a second on a Dec-Alpha machine. The total computation time spent generating the data summarized in the tables of this paper amounts to about 100 days of machine time.

These combinatorial optimization problems may be mapped onto physical systems, where each matching is one state of the system with an energy equal to its length. Then the minimum matching problem is equivalent to finding the ground state of that system. The physical systems built in this way are frustrated since in general not all points may be matched with their nearest neighbor. The thermodynamics of these systems can be studied by introducing Boltzmann weights for the states as was proposed by Kirkpatrick *et al.* [18]. Since this article focuses on ground states, however, we restrict our discussion to the zero temperature properties of these systems.

In any disordered system, be it a spin-glass or a matching problem, one is not particularly interested in the properties of a given instance of the problem. More relevant physically are typical properties, or averages over an ensemble of instances. The l_{ij} 's then become quenched random variables and one speaks of the stochastic M(B)MP, problems that are both disordered and frustrated. Moreover, we are interested in the infinite "volume" limit, meaning in this case the limit $N \rightarrow \infty$. These systems can then be studied using the replica or the cavity method (for a review see [1]) as developed by M & P [14,19].

Let us now describe the different ensembles of l_{ij} 's we consider. The first is the Euclidean ensemble (for the MMP as well as for the MBMP) where we have N (or $2N$) random and independent points chosen uniformly in a d -dimensional volume (*e.g.*, a unit hypercube), and the lengths l_{ij} are given by the Euclidean distances between the points. This is a short-range model in that the

points lie in a Euclidean space and only the first few nearest neighbors are relevant for the minimum matching. (In the formulation based on spins [14], the coupling between spins decreases exponentially with distance.) Using a spin-glass analogy, this Euclidean model is like the EA model (both models are short range, leading to important Euclidean correlations in the disorder variables). Not surprisingly, the Euclidean MMP – like the EA model – is not solvable analytically. This suggests that one should consider a different ensemble for the disorder variables in order to render the problem more tractable [20]. Indeed, this may be accomplished by taking the lengths l_{ij} to be independent, identically distributed random variables. The corresponding model is called the "random-link" model. (Note that the points do not lie in a metric space and the l_{ij} 's no longer have correlations; in particular the triangle inequality does not hold.) Pursuing our spin-glass analogy, the random-link MMP is like the model of Viana and Bray [21] for spin-glasses. Both models are "infinite dimensional" in the sense that there is no underlying geometry and thus there are no Euclidean correlations. Furthermore, in both cases, the effective connectivity at each site stays fixed as the size of the system grows. For the Viana and Bray model, this is enforced by having the number of non-zero couplings to a spin be size independent; for the random-link MMP, this occurs because only the first few nearest neighbors of a site effectively contribute to the minimum matching. These models thus interpolate between the infinite range/infinite connectivity case and the finite range/finite connectivity case. Thus, since it is expected that the Viana and Bray model provides a better approximation than the SK model to the EA model, the fact that its analogue here (the random-link MMP) is exactly solvable is of major interest.

A connection between the Euclidean and the random-link models was first given by M & P [19]: they pointed out that the one- and two-link distributions could be made identical in both problems, leading to a particular family of random-link models parametrized by d , where d is the spatial dimension for the corresponding Euclidean model. With such a choice, both models have the same Cayley tree approximation. One may then consider the random-link MMP to be a mean-field model for the Euclidean MMP in which correlations between link lengths (the quenched disorder variables) have been neglected. What we call from now on the "random-link approximation" [22] consists of using the thermodynamic functions of the random-link models as estimators for those of the Euclidean models.

1.3 Outline

This paper expands upon previous work [23], and provides an in depth study of ground state properties (disorder induced distributions of the energy, length of dimers, dimensional dependence, *etc.*) in the MMP and the MBMP. The outline is as follows. In Section 2, we examine the random-link models. First, we discuss self-averaging and

large N scaling properties of the ground state energy. Second, we recall results derived assuming replica symmetry and check them numerically. Third, we propose a conjecture for the frequency with which a point connects to its k th nearest neighbor in the ground state of the M(B)MP. In Section 3, we study the Euclidean models. First, we discuss self-averaging of the ground state energy. Second, we quantify numerically the precision of the random-link approximation. Third, we consider the Euclidean corrections to this approximation in the case of the MMP. Finally, Section 4 discusses our results and provides an outlook on possible generalizations to other systems.

In Appendix A, we present an N -independent upper bound for the length of the random-link MMP when $d = 1$. In Appendix B, we prove a self-averaging property for the Euclidean MBMP in dimensions greater or equal to 3.

2 Random-link models

In order to use the random-link models as an approximation for the Euclidean ones, we must, as previously mentioned, set the random-link distribution to match the one-link Euclidean distribution. In the Euclidean model, one possible approach would be to take the large N limit at fixed density of points, in which case the volume would scale linearly with N . However, for historical reasons, the standard practice is to take $N \rightarrow \infty$ in a fixed volume. These two pictures are equivalent, and are mapped onto one another by rescaling all the lengths by a factor $N^{1/d}$. As a consequence, in the units we shall use in the rest of this paper, the mean length between neighboring points scales as $N^{-1/d}$.

Consider two points i and j randomly chosen in the d -dimensional unit hypercube. In the absence of edge effects, the distribution of their distance l_{ij} is given by $\rho_d(l_{ij} = r) = dB_d r^{d-1}$, where $B_d = \pi^{d/2}/(d/2)!$ is the volume of the d -dimensional unit ball. We thus take from now on $\rho_d(l)$ as the individual distribution for the link lengths in the random-link model. The random-link M(B)MP models are then described by a single parameter d . Two comments are in order. First, ρ_d is not normalized, and so must be cut off; this can be done arbitrarily as the large N scaling of the minimum matching depends only on the behavior of $\rho_d(l)$ at small l . Second, neglecting edge effects, any two lengths are uncorrelated in Euclidean space, so the prescription just given also matches the two-link Euclidean distribution.

2.1 Large N limit

We denote by L_{MM}^{RL} the length (or energy) of the minimum matching in the MMP; L_{MM}^{RL} is a random variable depending on the instance (*i.e.*, on the l_{ij} 's). It is a sum of $N/2$ terms, each of which is typically the length between near neighbors. We have seen that these lengths scale as $N^{-1/d}$, so L_{MM}^{RL} is expected to scale as $N^{1-1/d}$. For instance, consider the length d_i between point i and

Table 1. Numerical measurements of the relative standard deviation σ and skewness s of the distribution of L_{MM}^{RL} and L_{MBM}^{RL} at $d = 1$ and $d = 2$.

N	MMP $d = 1$		MMP $d = 2$	
	$\sigma\sqrt{N/2}$	$s\sqrt{N/2}$	$\sigma\sqrt{N/2}$	$s\sqrt{N/2}$
50	0.784	1.39	0.421	0.33
100	0.798	1.50	0.423	0.32
200	0.798	1.39	0.427	0.36
400	0.808	1.56	0.424	0.24

N	MBMP $d = 1$		MBMP $d = 2$	
	$\sigma\sqrt{N}$	$s\sqrt{N}$	$\sigma\sqrt{N}$	$s\sqrt{N}$
25	0.780	1.29	0.419	0.25
50	0.795	1.36	0.423	0.29
100	0.802	1.36	0.427	0.35
200	0.811	1.57	0.428	0.45

its nearest neighbor. It is easy to show that in the large N limit, $\langle d_i \rangle \sim D_1(d)/N^{1/d}$ where

$$D_1(d) = (1/d)B_d^{-1/d} \tag{1}$$

is the rescaled average nearest neighbor length at large N . (In the units where the density of points is equal to one, $D_1(d)$ is exactly the mean nearest neighbor length.) In the case of the minimum matching length, it can be proven [24] that $L_{MM}^{RL}/N^{1-1/d}$ becomes peaked around its mean value as $N \rightarrow \infty$. A stronger property, called self-averaging, would be that $L_{MM}^{RL}/N^{1-1/d}$ tends with probability one to a (non-random, N -independent) constant $\beta_{MM}^{RL}(d)$, as $N \rightarrow \infty$. Although this second property has not yet been proven, it is strongly supported by previous numerical studies [25] as well as the simulations described in Section 2.3. Moreover, bounds can be found for $\beta_{MM}^{RL}(d)$ (see Appendix A) that reinforce this hypothesis. In the following, we assume the existence of the $\beta_{MM}^{RL}(d)$. The analogous discussion applies to MBMP as well, so we will assume $\beta_{MBM}^{RL}(d) = \lim_{N \rightarrow \infty} L_{MBM}^{RL}/N^{1-1/d}$.

It is of interest to understand the *distribution* of L_{MM}^{RL} in the large N limit. In particular, one may wonder whether the self-averaging property comes from some kind of central limit theorem which would lead to a Gaussian (normal) limit distribution for L_{MM}^{RL} . The central limit theorem (CLT) states that for a sum S of N independent identically distributed random variables the relative standard deviation $\sigma = \sqrt{\text{Var}(S)}/\langle S \rangle$ and the skewness $s = \langle (S - \langle S \rangle)^3 \rangle / \sigma^3$ decrease as $1/\sqrt{N}$. Although in fact the terms entering the sum L_{MM}^{RL} are correlated, if the correlations are not too strong, we can expect the same CLT-type scaling to hold for L_{MM}^{RL} . We have computed σ and s numerically for L_{MM}^{RL} and find a behavior that is in excellent agreement with the expected CLT scaling laws (see Tab. 1). (Since the finite size effects and the statistical noise are significant, we have not extrapolated our data to the $N \rightarrow \infty$ limit.)

Similarly, we have computed σ and s for the MBMP, and find nearly identical results (see again Tab. 1). Clearly, the large N behavior follows the CLT scaling laws for both the MMP and the MBMP. Furthermore, our data suggest that the amplitudes associated with these scalings are equal in the two models. We conjecture that the two are equal for *any* of the connected moments, and provide theoretical support for this in Section 2.2. We will also derive a theoretical estimate for the values of σ and s in Section 2.3.

2.2 Survey of analytical results

The random-link MMP was first solved by M & P [14] using the replica method, with a replica symmetric ansatz. They verified [26] the stability of the replica symmetric solution (at least for $d = 1$), suggesting that most likely the ansatz is exact. They also confirmed their results by the cavity method [19]. In our notation their results may be written

$$\beta_{MM}^{RL}(d) = \frac{dD_1(d)}{2(1/d)!} \int_{-\infty}^{+\infty} G_d(x) e^{-G_d(x)} dx, \quad (2)$$

where $G_d(x)$ satisfies the integral equation

$$G_d(x) = d \int_{-x}^{+\infty} (x+y)^{d-1} e^{-G_d(y)} dy \quad (3)$$

and $D_1(d)$ is as defined in equation (1). Equation (3) can be solved at $d = 1$, leading to $G_1(x) = \ln(1 + e^x)$ and $\beta_{MM}^{RL}(1) = \pi^2/24$. Furthermore, M & P calculated the distribution $P_d(l)$ of the rescaled bond length $N^{1/d}l_{ij}$ in the minimum matching in the limit $N \rightarrow \infty$, and found

$$P_d(l) = d l^{d-1} \int_{-\infty}^{+\infty} \frac{dG_d}{dx}(x) e^{-G_d(x) - G_d(l-x)} dx. \quad (4)$$

In the case $d = 1$, they found $P_1(l) = 2(2l - e^{-2l} \sinh(2l)) / \sinh^2(2l)$. Finally, since replica symmetry is not broken, one expects the mean length to have a $1/N$ expansion:

$$\frac{\langle L_{MM}^{RL} \rangle}{N^{1-1/d}} = \beta_{MM}^{RL}(d) \left(1 + \frac{A(d)}{N} + \frac{B(d)}{N^2} + \dots \right). \quad (5)$$

M & P have calculated [26] the first subleading term at $d = 1$ and have found $A(1) \approx -0.13$.

Similar calculations were performed for the MBMP (Orland [27], M & P [14, 19, 26]). In our units, they find:

$$\beta_{MBM}^{RL}(d) = 2\beta_{MM}^{RL}(d). \quad (6)$$

At $d = 1$, they obtain $\beta_{MBM}^{RL}(1) = \pi^2/12$ and $A(1) \approx -1.61$.

The MBMP and the MMP are very closely related. This is best seen within the convention we have used where the MMP matches N points, the MBMP matches $2N$ points and both models have the same individual link

length distribution ρ_d . One then sees that the saddle point equations for the partition functions (in the limit of large N) become identical in the two models. Thus $P_d(l)$ is the same in the two models, and in fact any reasonable observable will be the same in both models at large N . This remarkable property has apparently gone unnoticed so far. (Given this property and our conventions, the factor 2 in equation (6) simply follows from the fact that a bipartite matching has twice as many bonds as a simple matching.) A consequence of this correspondence is that the moments of L_{MM}^{RL} and L_{MBM}^{RL} should be the same at large N . We are thus able to provide theoretical support for the conjecture, given in the previous section, and based on our numerical results.

It is also of interest to study the limit of large d . We have derived a $1/d$ expansion of $\beta_{MM}^{RL}(d)$. One way to do this is to set $\tilde{G}_d(x) = G_d(\tilde{x} = x/d + 1/2)$ and then write $\tilde{G}_d(x)$ as a power series in $1/d$. From this we find [24]

$$\beta_{MM}^{RL}(d) = \frac{D_1(d)}{2} \times \left[1 + \frac{1-\gamma}{d} + \frac{\pi^2/12 + \gamma^2/2 - \gamma}{d^2} + O\left(\frac{1}{d^3}\right) \right] \quad (7)$$

where $\gamma = 0.5772\dots$ is Euler's constant.

2.3 Numerical verifications and a new conjecture

Brunetti *et al.* [25] have used numerical simulations of the random-link models to confirm the predictions of β^{RL} to the level of 0.2% for the MMP and 0.7% for the MBMP at $d = 1$ and $d = 2$. They have also checked the $O(1/N)$ corrections to $\langle L^{RL} \rangle$ at $d = 1$ and find relatively good agreement with the theory.

To obtain further confirmation we have estimated $\beta_{MM}^{RL}(d)$ and $\beta_{MBM}^{RL}(d)$ numerically for $1 \leq d \leq 10$, and have found accordance with the replica symmetric predictions to the level of 0.03% for the MMP and to the level of 0.1% for the MBMP. In order to reduce the statistical fluctuations and get quantitative errors on our estimates for β , we use the following procedure. First we compute the ensemble average $\langle L_{MM}^{RL} \rangle / N^{1-1/d}$ using a variance reduction trick [22, 28]. (The values of N used here and for the other calculations of β given later on are $N = 50, 70, 100, 150, 200, 260$, and 400 ; we performed averages over 5×10^5 instances at $N = 50$ down to $7,500$ instances for $N = 400$.) Then, to get the large N limit, we fit our data using a $1/N$ series (truncated after the second order) as indicated by the theory (Eq. (5)). The fits are good, with χ^2 values confirming the finite-size scaling law. The statistical error bar on $\beta_{MM}^{RL}(d)$ is then obtained by the standard method [29], whereby fixing the fit's leading coefficient at $\beta \pm \sigma$ makes χ^2 increase by one from its minimum value. Our results are summarized in Table 2.

We have also checked the prediction for the distribution $P_d(l)$ of the bond lengths in the minimum (bipartite) matching. We find such good agreement with theory that the numerical data is indistinguishable from the replica

Table 2. Comparison of theoretical and numerical values of $\beta_{MM}^{RL}(d)$ and $\beta_{MBM}^{RL}(d)$. Numbers in parentheses represent the statistical error bar on the last digit(s).

d	β_{th}	MMP	MBMP
		β_{num}	$\beta_{num}/2$
1	0.411234	0.41142(12)	0.41134(9)
2	0.322580	0.32257(5)	0.32262(4)
3	0.326839	0.32684(4)	0.32691(4)
4	0.343227	0.34323(3)	0.34327(4)
5	0.362175	0.36210(3)	0.36222(3)
6	0.381417	0.38143(3)	0.38143(3)
7	0.400277	0.40026(5)	0.4002(5)
8	0.418548	0.41852(5)	0.4180(5)
9	0.436185	0.43612(5)	0.4362(5)
10	0.453200	0.45310(5)	0.4531(4)

predictions (in a figure one would not be able to tell the two curves apart).

We will now use $P_d(l)$ to obtain an estimate of the quantities computed numerically in Table 1. Neglecting the correlations among the bond lengths in the optimal matching, we can use equation (4) to predict σ and s as a function of N . At $d = 1$, we find $\sigma\sqrt{N/2} \approx 0.73$ and $s\sqrt{N/2} \approx 1.85$ for the MMP. For the MBMP, the same formulae apply if $N/2$ is replaced by N . These theoretical predictions (to be compared with Tab. 1) are about 10% too small for σ and about 30% too large for s , showing that correlations cannot be neglected, but are nevertheless relatively small.

Another interesting quantity is the mean fraction of points connected to their k th nearest neighbor in the optimal matching. Call this fraction $p_d(k)$. In view of our numerical data at $d = 1$ (see Tab. 3), we conjecture for both the MMP and the MBMP that in the limit $N \rightarrow \infty$,

$$p_1(k) = 2^{-k}. \quad (8)$$

In addition the N -dependence of these fractions seems to be linear in $1/N$ as one could expect from equation (5). It seems likely that our conjecture may be confirmed by using the replica method. There may also be analogous relations in higher dimensions, but unfortunately we have not found any convincing formulae. At best, our data are approximately fitted by stretched exponentials (see also [28]). This same kind of behavior also arises in other combinatorial optimization problems such as the traveling salesman problem [30]. It is also interesting to note that another conjecture has been proposed for a random-link MBMP by Parisi [31]. His conjectured relation gives the mean length of the matching at all values of N , not just in the limit $N \rightarrow \infty$.

3 Euclidean models

3.1 Large N limit

Let L_{MM}^E be the length of the minimum matching in the Euclidean MMP. Hereafter, we take the points to be distributed randomly in the d -dimensional unit hypercube. Following the same argument as for the random-link MMP, one expects L_{MM}^E to scale as $N^{1-1/d}$. In fact, it has been proven [32] that the Euclidean MMP has the self-averaging property in any dimension, so $L_{MM}^E/N^{1-1/d}$ tends to a constant $\beta_{MM}^E(d)$ as $N \rightarrow \infty$ with probability one.

For the Euclidean MBMP, the situation is more complex. There are two sets of points, so local density differences have a large effect. In particular, the $N^{1-1/d}$ scaling law is *not* valid for $d \leq 2$. At $d = 1$, it is easy to see that the optimum corresponds to matching the points left to right. Then a quick estimate shows that L_{MBM}^E scales as \sqrt{N} instead of as N^0 . Furthermore L_{MBM}^E/\sqrt{N} does not become peaked, so there is no self-averaging. At $d = 2$, the situation is more interesting: L_{MBM}^E scales [33] as $\sqrt{N \ln N}$ (instead of as \sqrt{N} for L_{MBM}^{RL}). The question of self-averaging has not yet been settled but numerical simulations indicate that the property does hold, and we will assume this is the case hereafter. At higher dimensions, we have proven self-averaging (see Appendix B), so the quantity $\beta_{MBM}^E(d) = \lim_{N \rightarrow \infty} L_{MBM}^E/N^{1-1/d}$ exists for $d \geq 3$.

As with the random-link models, one may wonder whether the central limit theorem is at work. Following what was presented in Section 2.1, it is natural to investigate the limiting distribution of the optimum length. It is convenient in the numerical study to avoid boundary effects; to do so, we work in the unit hypercube with periodic boundary conditions. Using numerical measurements of the moments of the distributions we find that the MMP obeys the CLT scaling laws (see Tab. 4). This is not surprising: for the Euclidean MMP, we can divide the hypercube into subvolumes. Then the length of the minimum matching is close to the sum of the minimum matching length in these subvolumes, and the CLT scalings should hold. On the other hand, the CLT scalings do *not* hold for the MBMP, and furthermore the limit distribution is *not* Gaussian. In particular, we find that s does not tend to zero, rather it *grows* with N . The CLT argument used for the MMP does not apply to this problem because the subvolumes just mentioned will not in general contain equal number of points from each set. Nevertheless, as the dimension increases, the density fluctuations decrease, explaining why at fixed N the MBMP values of σ and s get closer to the MMP ones as d increases. (For a formal application of this argument, see Appendix B.)

3.2 Numerical results and the random-link approximation

To date, little has been done to compute the ground-state energy densities $\beta_{MM}^E(d)$ and $\beta_{MBM}^E(d)$. The best

Table 3. Numerical results of $10^5(p_d(k) - 2^{-k})$ for the random-link MMP and MBMP in the case $d = 1$. Numbers in parentheses represent the statistical error on the last digit(s).

k	MMP			MBMP		
	$N = 50$	$N = 100$	$N = 200$	$N = 50$	$N = 100$	$N = 200$
1	422(22)	181(16)	86(16)	1185(16)	608(11)	305(11)
2	174(19)	96(14)	47(14)	113(14)	47(10)	17(10)
3	-26(15)	-12(10)	1(10)	-238(10)	-108(7)	-52(7)
4	-81(11)	-30(8)	-18(8)	-289(7)	-157(5)	-77(5)
5	-112(8)	-49(5)	-31(5)	-248(5)	-123(4)	-56(4)

Table 4. Numerical results for the relative standard deviation σ and skewness s of the distribution of L_{MM}^E and L_{MBM}^E at $d = 2, 3$ and 4 .

N	MMP $d = 2$		MMP $d = 3$		MMP $d = 4$	
	$\sigma\sqrt{N/2}$	$s\sqrt{N/2}$	$\sigma\sqrt{N/2}$	$s\sqrt{N/2}$	$\sigma\sqrt{N/2}$	$s\sqrt{N/2}$
50	0.302	-0.90	0.244	-0.65	0.206	-0.68
100	0.299	-0.93	0.244	-0.66	0.205	-0.64
200	0.290	-0.87	0.243	-0.52	0.204	-0.52
400	0.295	-0.85	0.243	-0.88	0.203	-0.45

N	MBMP $d = 2$		MBMP $d = 3$		MBMP $d = 4$	
	$\sigma\sqrt{N}$	$s\sqrt{N}$	$\sigma\sqrt{N}$	$s\sqrt{N}$	$\sigma\sqrt{N}$	$s\sqrt{N}$
25	0.603	2.61	0.350	1.16	0.258	0.42
50	0.744	4.20	0.387	1.94	0.274	1.12
100	0.938	6.49	0.431	3.26	0.289	1.88
200	1.197	9.97	0.481	4.71	0.305	2.53

Table 5. Comparison of MMP ground state energies for the three models: Euclidean, random-link, and random-link including 3-link Euclidean corrections ($1 \leq d \leq 10$). For $\beta^E(d)$ the numbers in parentheses are statistical errors on the last digit.

d	$\beta^E(d)$	$\beta^{RL}(d)$	$\frac{\beta^{RL} - \beta^E}{\beta^E}$	$d\frac{\beta^{RL} - \beta^E}{\beta^E}$	$\beta^{EC}(d)$	$\frac{\beta^{EC} - \beta^E}{\beta^E}$
1	0.5	0.411234	-17.75%	-0.178		
2	0.3104(2)	0.322580	3.92%	0.078	0.30915	-0.40%
3	0.3172(2)	0.326839	3.04%	0.091	0.31826	0.33%
4	0.3365(3)	0.343227	2.01%	0.080	0.33756	0.30%
5	0.3572(2)	0.362175	1.39%	0.070	0.35818	0.27%
6	0.3777(1)	0.381417	0.98%	0.059	0.37849	0.21%
7	0.3972(1)	0.400277	0.77%	0.054	0.39807	0.22%
8	0.4162(1)	0.418548	0.56%	0.045	0.41685	0.17%
9	0.4341(1)	0.436185	0.48%	0.042	0.43485	0.17%
10	0.4515(1)	0.453200	0.38%	0.038	0.45214	0.14%

estimates prior to our recent work [23] were [34,28] $\beta_{MM}^E(2) \approx 0.312$ and $\beta_{MM}^E(3) \approx 0.318$; for $\beta_{MBM}^E(d)$, no valid estimates have yet been published. Expanding upon the work in [23], we now provide very accurate measurements of these quantities, using the same procedure as in the random-link case. Again we find that the χ^2 values justify the use of a truncated $1/N$ series.

As previously remarked, our random-link distributions were established in order to match the one- and two-link distributions of the Euclidean model. So, if the effects coming from the Euclidean correlations among three or more link lengths are small, then the properties of the

random-link and Euclidean M(B)MP should be quantitatively close. In fact, replacing $\beta_{MM}^E(d)$ and $\beta_{MBM}^E(d)$ by $\beta_{MM}^{RL}(d)$ and $\beta_{MBM}^{RL}(d)$ leads to a very precise approximation. As shown in Table 5, $\beta_{MM}^{RL}(d)$ differs from $\beta_{MM}^E(d)$ by 17.8% at $d = 1$ and by 3.9% at $d = 2$, and this difference decreases quickly as the dimension increases (the quantity β^{EC} given in the table will be discussed in Sect. 3.3). Likewise for the MBMP, shown in Table 6: $\beta_{MBM}^{RL}(d)$ differs from $\beta_{MBM}^E(d)$ by only 7.7% at $d = 3$. Note that comparing $\beta_{MBM}^{RL}(d)$ and $\beta_{MBM}^E(d)$ at $d \leq 2$ is meaningless since the scaling laws are different as we mentioned in the beginning of Section 3.1. Nevertheless,

Table 6. Comparison of MBMP ground state energies for the two models: Euclidean and random-link ($3 \leq d \leq 10$). The rightmost column compares the Euclidean MBMP and MMP models. For $\beta^E(d)$ the numbers in parentheses are statistical errors on the last digit(s). In the last column $\Delta_E = (\beta_{MBM}^E - 2\beta_{MM}^E)/\beta_{MBM}^E$.

d	$\beta^E(d)$	$\beta^{RL}(d)$	$\frac{\beta^{RL} - \beta^E}{\beta^E}$	$d^2 \Delta_E$
3	0.7080(2)	0.653679	-7.68%	0.664
4	0.7081(2)	0.686455	-3.06%	0.597
5	0.7349(1)	0.724350	-1.44%	0.514
6	0.7688(3)	0.762834	-0.77%	0.482
7	0.8039(2)	0.800554	-0.41%	0.461
8	0.8391(2)	0.837097	-0.24%	0.445
9	0.8736(2)	0.872370	-0.14%	0.429
10	0.9076(2)	0.906400	-0.13%	0.436

we have also computed $\beta_{MBM}^E(2)$; empirically, we found that using a $1/N$ fit could not do, but that good χ^2 values were obtained using a linear fit in $1/\ln(N)$. We find $\beta_{MBM}^E(2) = 0.340(1)$.

Since the random-link approximation was also found to be very good for the traveling salesman problem [22], our results suggest that this approximation should be widely applicable to link-based optimization problems. Furthermore, we can understand how the size of the error inherent to this approximation decreases as $d \rightarrow \infty$. Consider for instance the bond occupation probabilities for links connected to a given site. One expects that these probabilities have a large d limit, and that Euclidean correlations introduce $1/d$ corrections compared to the random-link case. (To be precise, one expects the corrections to be given by a $1/d$ expansion, with the leading term naturally being of order $1/d$.) In support of this argument, we have checked that the relative difference between the Euclidean value and the random-link value of $p_d(k)$ is indeed of order $1/d$. We can then see how quantities such as $\beta(d)$ depend on the link lengths vary when one goes from the random-link model to the Euclidean model. At large d , a simple calculation shows [22] that the mean link lengths between k th nearest neighbors have for different values of k a relative difference of $O(1/d)$ (regardless of whether we use the random-link or the Euclidean ensemble). Since the *occupation probabilities* $p_d(k)$ have Euclidean corrections of order $1/d$, the random-link approximation leads to an error of order $1/d^2$ for $\beta(d)$ and $P_d(l)$. We have tested this behavior numerically by considering the quantity $d(\beta_{MM}^{RL} - \beta_{MM}^E)/\beta_{MM}^E$, and the data scale as $1/d$ as expected. Thus the random-link approximation gives both the leading and subleading $1/d$ dependence of $\beta_{MM}^E(d)$. Then, from equation (7), we find

$$\beta_{MM}^E(d) = \frac{D_1(d)}{2} \left[1 + \frac{1-\gamma}{d} + O\left(\frac{1}{d^2}\right) \right]. \quad (9)$$

Furthermore we have directly confirmed this dependence by performing a fit of our $\beta_{MM}^E(d)$ values and we find

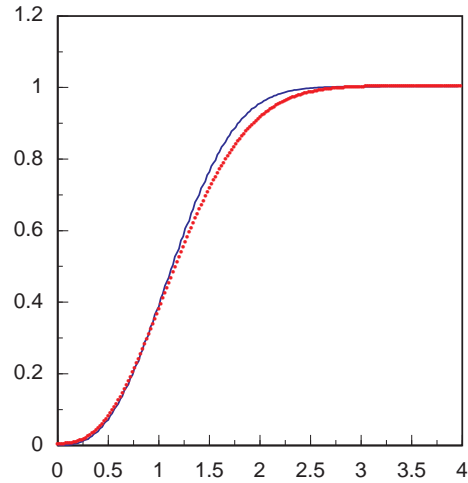


Fig. 1. Comparison of the random-link (continuous line) and Euclidean (points) integrated distributions of the rescaled bond lengths in the minimum bipartite matching for the MBMP at $d = 3$ and $N = 60$.

0.424 ± 0.008 for the coefficient of the $1/d$ term; this is to be compared to the theoretical value $1 - \gamma = 0.42278\dots$

Performing the same analysis for the MBMP, we find that $\beta_{MBM}^E(d)$ also satisfies equation (9) (omitting the factor $1/2$) or equivalently that $(\beta_{MBM}^E - 2\beta_{MM}^E)/\beta_{MBM}^E = O(1/d^2)$ (see Tab. 6). A direct fit to the $1/d$ term of equation (9) for β_{MBM}^E gives 0.42 ± 0.05 in agreement with the theoretical value $1 - \gamma$.

Finally, we have also applied the random-link approximation to compare $P_d(l)$ in the random-link and Euclidean models and found very good agreement (see Fig. 1 for the MBMP at $d = 3$, which has the largest discrepancy). The agreement becomes better as the dimension increases, with an error of order $1/d^2$.

3.3 Euclidean corrections to the random-link approximation

The mean field model (*i.e.*, the random-link model) provides a very good approximation to the finite dimensional Euclidean model. It is of major interest to push the approximation further and to derive, for instance, a large dimensional expansion. For the MMP, M & P have calculated [15] a correction to the random-link approximation by considering the effects of the three-link Euclidean correlations. Such correlations arise only when three links form a triangle. (For the MBMP, one would have to go to four-link correlation effects, and this has not yet been attempted). M & P's result is given in terms of a function G_d , but where G_d now satisfies a much more complicated integral equation (Eq. (34) in their paper). From this, one obtains a new estimate for $\beta_{MM}^E(d)$, which we denote here by $\beta_{MM}^{EC}(d)$ (EC stands for Euclidean corrections).

We solved numerically this modified integral equation for G_d and computed $\beta_{MM}^{EC}(d)$ for $2 \leq d \leq 10$ (see Tab. 5). Comparing with $\beta_{MM}^E(d)$, we see that these estimates are considerably more accurate than when using $\beta_{MM}^{RL}(d)$.

At $d = 2$, the random-link approximation leads to an error of 3.9%; this error is reduced by nearly a factor of 10 by including the corrections due to 3-link correlations. At $d = 3$, the error is reduced from 3.0% to 0.4%. At larger d , the error continues to decrease, though the effect is less significant.

To understand the dependence of these corrections on dimension, it is useful to consider how the difference $\beta_{MM}^{EC} - \beta_{MM}^{RL}$ scales with d . This difference is associated with a 3-link correction term which gives the probability of finding nearly equilateral triangles as $d \rightarrow \infty$. It is not difficult to see that this probability goes to zero exponentially with d . Thus the 3-link correlations give tiny corrections at large d (as confirmed by the numerics), and the power series expansion in $1/d$ of β_{MM}^{EC} is *identical* to that of β_{MM}^{RL} . In fact the $1/d$ series of β_{MM}^{EC} is not modified if one includes 4, 5, or *any* finite number of multi-link correlations. This follows from the fact that all non-zero multi-link correlations come from sets of links forming at least one loop, and that fixed sized (N -independent) loops connecting near neighbors become exponentially rare at large d . (Of course this behavior depends on having randomly placed points; the situation is very different on lattices, for instance, where small loops are important.) The main consequence of the nature of these multi-link correlations is that the M & P k -link correlation expansion will not converge towards $\beta_{MM}^E(d)$, and in particular it does not even allow one to compute the $O(1/d^2)$ term in the $1/d$ expansion of $\beta_{MM}^E(d)$. Formally, the Euclidean dimensional dependence is beyond all orders of the k -link correlation expansion. This non-perturbative behavior is quite remarkable, and indicates that the large N limit and the k -link correlation expansion do not commute. This property may have its analogue in other disordered systems.

4 Summary and discussion

In this article we have studied two versions of the stochastic minimum matching and minimum bipartite matching problems: the random-point Euclidean ensemble, and its correlation-free approximation, the random-link ensemble. For both ensembles, we have given evidence that the ground state energy is self-averaging and obeys (except for the Euclidean MBMP) the central limit theorem scaling laws. For both ensembles we have performed extensive numerical simulations in order to measure the ground state energy density $\beta(d)$ and the distribution $P_d(l)$ of bond lengths in the ground-state. For the random-link model we have checked to high precision the replica symmetric prediction of Mézard and Parisi and find excellent agreement. Furthermore, we have proposed a new conjecture at $d = 1$ that suggests further analytical calculations. For the Euclidean model, we have studied the accuracy of the random-link approximation and find the error to be small even at low dimensions. For example at $d = 2$ the error introduced by this approximation is 3.9% for β_{MM}^E . We have also been able to go beyond the random-link approximation by applying the formalism of Mézard and Parisi

in order to include 3-link correlations associated with the triangle inequality. We find that this improved estimate reduces the error by nearly a factor of 10 at low dimensions. In particular, the resulting $d = 2$ prediction for β_{MM}^E has an error of 0.4%.

The limit of high dimensions is also of major interest. Based on our simulations up to $d = 10$, we have given strong evidence that the first two terms of the $1/d$ expansion of the random-link model are the same as for the Euclidean model but that these two models differ at order $1/d^2$. Furthermore we have argued that for any fixed k , k -link correlations in Euclidean space only give rise to exponentially small contributions in d and thus do not modify the $1/d$ expansion. As a consequence, the Euclidean dimensional dependence is beyond all orders of the k -link correlation expansion.

Although our study was performed in the context of the MMP, our reasoning applies also to other link-based problems associated with random points, and leads us to the following picture. For these random-point systems, whenever the thermodynamic functions depend only on the *local* properties of the (short) link graph, we expect the error in the random-link approximation to be exponentially small in d . This will always be the case in the high temperature phase where correlations are weak. However, in the low temperature phase, the correlations may be such that the $N \rightarrow \infty$ limit and the k -link correlation expansion do not commute. We expect this to be the case in many combinatorial optimization problems such as the assignment problem and the traveling salesman problem, where k -link correlations with k growing in N remain important as $N \rightarrow \infty$. Arbitrarily large loops matter in these systems, and contribute to the thermodynamics at order $1/d^2$. This change in behavior can be illustrated using more physical language by considering a polymer on a random-point lattice. Then the arguments given previously indicate that the random-link approximation is exponentially good in the “dilute” phase, while it leads to an error in powers of $1/d$ in the collapsed phase. This non-perturbative behavior of the $1/d$ expansion is reminiscent of what occurs in lattice systems such as the Eden model, where the $1/d$ expansion does not commute with the $N \rightarrow \infty$ limit [35]. We would not be surprised if similar phenomena occurred in other disordered systems, whether they be based on links or spins; the calculation of their $1/d$ series would then be particularly difficult. Further insight into these issues may be obtained by looking at excited states of the M(B)MP to see whether they consist of arbitrary large loops (*i.e.*, whose sizes diverge as $N \rightarrow \infty$). Such a study could also allow a direct investigation of possible replica symmetry breaking in the Euclidean case.

It is tempting to speculate on how all of this might carry over to spin-glasses. There, one analogue of the random-point MMP is the Edwards-Anderson model with nearest neighbor interactions on a d -dimensional hypercubic lattice of connectivity $2d$. The mean field model for spin-glasses is usually taken to be the Sherrington-Kirkpatrick infinite connectivity model. However, a more appropriate mean field model in this context is that of

Viana and Bray [21] in which the connectivity is $2d$ and for which the couplings J_{ij} have the same *individual* distribution as in the Edwards-Anderson model. Just as in the matching problem, one thereby obtains a mean field model at any given dimension, and the mean field approximation then consists of using this model to estimate the thermodynamic quantities of the d -dimensional Edwards-Anderson model. We expect this approach to lead to errors of a few percent at low dimensions, and to $O(1/d)$ errors at large d . Unlike the random-link MMP, the Viana-Bray model has not yet been solved analytically, so that the mean field values would have to be calculated approximately. Nevertheless, we view these mean field models as providing a very promising approach to computing quantities in the Edwards-Anderson model. We hope that the potential reward will encourage new attempts to solve the Viana and Bray model, and that the challenge of determining the first $O(1/d)$ correction to “mean field” will be taken up.

We are grateful to D. Dean, W. Krauth, M. Mézard, H. Orland, N. Sourlas and J. Vannimenus for comments, and to O. Bohigas for his constant interest. Special thanks go to A. Percus for constructive criticisms. J.H. and J.H.B.dM. acknowledge a fellowship from the MENESR, and O.C.M. acknowledges support from the Institut Universitaire de France. The Division de Physique Théorique is an “Unité de Recherche des Universités Paris XI et Paris VI associée au CNRS”.

Appendix A: Bounds for the MMP

Here we review exact lower and upper bounds for $\beta_{MM}^{RL}(d)$, and provide to our knowledge the first finite upper bound at $d = 1$. First there is a trivial lower bound: each point is at best linked to its nearest neighbor, so

$$\beta_{MM}^{RL}(d) \geq D_1(d)/2. \quad (\text{A.1})$$

To get an upper bound, we use another optimization problem: the traveling salesman problem (TSP), which consists of finding a minimum length tour visiting all the N points. Indeed we can obtain a matching by removing every second bond in a tour (note that this argument fails for the MBMP), so we get

$$\beta_{MM}^{RL}(d) \leq \beta_{TSP}^{RL}(d)/2, \quad (\text{A.2})$$

where $\beta_{TSP}^{RL}(d)$ is the analog of $\beta_{MM}^{RL}(d)$ for the TSP. A bound for $\beta_{TSP}^{RL}(d)$ is already known for $d \geq 2$ (see in [20]). This bound comes from the greedy algorithm and is

$$\beta_{TSP}^{RL}(d) \leq \frac{D_1(d)}{1 - 1/d}. \quad (\text{A.3})$$

In the case $d = 1$, the greedy construction leads to a bound which grows logarithmically in N and thus is useless. We have obtained a finite bound for the case $d = 1$ using a different approach. The idea is that a bound is known [36, 37] for the *asymmetric* TSP (where l_{ij} and l_{ji} are independent). Let us call λ_{ij} the link lengths of the asymmetric

TSP. We denote by $\rho_{Asym}(l)$ the distribution density of the λ_{ij} , and associate a symmetric TSP to any asymmetric TSP by setting $l_{ij} = \min(\lambda_{ij}, \lambda_{ji})$. Then, in the limit of short link lengths, the distribution in this symmetric TSP is $\rho_{Sym}(l) = 2\rho_{Asym}(l)$. This gives

$$\beta_{TSP}^{Sym.}(d) \leq 2^{1/d} \beta_{TSP}^{Asym.}(d). \quad (\text{A.4})$$

And we thus obtain our bound at $d = 1$.

Appendix B: Self-averaging for the Euclidean MBMP

We shall denote by $L_{MBM}(X_1, \dots, X_N, Y_1, \dots, Y_N)$ the length of the minimum bipartite matching between the X_i 's and the Y_i 's. We prove here that for any $d \geq 3$, L_{MBM} satisfies a theorem analogous to the one Beardwood, Halton and Hammersley have proven for the TSP [38]. Specifically, we prove the following:

Let X_1, \dots, X_N, \dots and Y_1, \dots, Y_N, \dots be two sequences of random points independently and uniformly distributed in $[0, 1]^d$, where $d \geq 3$, and let $L_N = L_{MBM}(X_1, \dots, X_N; Y_1, \dots, Y_N)$. There exists a constant $\beta_{MBM}(d) > 0$ such that with probability one,

$$\lim_{N \rightarrow \infty} \frac{L_N}{N^{1-1/d}} = \beta_{MBM}(d). \quad (\text{B.1})$$

To begin with, we remark that to prove this theorem, it is sufficient to establish that $L_N/N^{1-1/d}$ converges in mean value to a constant $\beta_{MBM}(d)$. This is a consequence of the following lemma [39]: for any $t > 0$,

$$P\left(\left|\frac{L_N}{N^{1-1/d}} - \langle \frac{L_N}{N^{1-1/d}} \rangle\right| > t\right) \leq 2 \exp\left(-\frac{N^{1-2/d}t^2}{8d}\right). \quad (\text{B.2})$$

The theorem then follows easily from the convergence of $\langle L_N \rangle / N^{1-1/d}$ as $N \rightarrow \infty$, by applying the Borel-Cantelli lemma. We now wish to establish that for $d \geq 3$ the quantity $\langle L_N \rangle / N^{1-1/d}$ indeed converges to a constant $\beta_{MBM}(d) > 0$. To do this, we exploit the subadditivity properties of L_{MBM} (see [40]).

First we need to generalize L_{MBM} to matchings between two sets of different cardinalities. We shall define $L_{MBM}(X_1, \dots, X_{N_1}; Y_1, \dots, Y_{N_2})$ by requiring that the matchings contain as few unmatched points as possible, that is we leave $|N_1 - N_2|$ points unmatched.

Suppose the points $X_1, \dots, X_{N_1}, Y_1, \dots, Y_{N_2}$ belong to an arbitrary cube Q whose edges have length a , and divide Q into disjoint cubes Q_p , $p = 1, \dots, 2^d$ by splitting each edge in two halves. Construct in each Q_p an optimal matching in the sense just defined, between the $n_{1,p}$ points X_i and the $n_{2,p}$ points Y_i in Q_p , and denote its length by L_p . There will be $|n_{1,p} - n_{2,p}|$ points left unpaired in each Q_p , so if L_0 denotes the length of an optimal matching for

these points, one has

$$\begin{aligned} L_{MBM}(X_1, \dots, X_{N_1}; Y_1, \dots, Y_{N_2}) &\leq \sum_{p=1}^{2^d} L_p + L_0 \\ &\leq \sum_{p=1}^{2^d} L_p + \frac{1}{2} a \sqrt{d} \sum_{p=1}^{2^d} |n_{1,p} - n_{2,p}|. \end{aligned} \quad (\text{B.3})$$

Now we apply this to $Q = [0, 1]^d$. Let $Q_{p_1, p_1 = 1, \dots, 2^d}$ be the cubes obtained in the last subdivision, let $Q_{p_1 p_2}$ be the cubes obtained by splitting in two halves the edges of each Q_{p_1} , and so on. By repeating this operation K times, we get a subdivision with cubes $Q_{p_1 \dots p_K}$ whose edges are of length $1/2^K$. Let $n_{1, p_1 \dots p_K}$ and $n_{2, p_1 \dots p_K}$ be respectively the number of points X_i and Y_i in $Q_{p_1 \dots p_K}$. Apply (B.3) first to the $Q_{p_1, \dots, p_{K-1}}$'s, then to the $Q_{p_1 \dots p_{K-2}}$'s, etc., keeping at each step only those points that are still unpaired. It is easy to convince oneself that the number of unpaired points in each $Q_{p_1, \dots, p_{K-k}}$ just after step k is given by $|n_{1, p_1, \dots, p_{K-k}} - n_{2, p_1, \dots, p_{K-k}}|$. After step $k = K$ one obtains a matching between X_1, \dots, X_{N_1} and Y_1, \dots, Y_{N_2} where all but $|N_1 - N_2|$ of the points are matched. One is thus led to the following inequality:

$$\begin{aligned} L_{MBM}(X_1, \dots, X_{N_1}; Y_1, \dots, Y_{N_2}) &\leq \sum_{p_1 \dots p_K} L_{p_1 \dots p_K} \\ &+ \sum_{k=1}^K \frac{\sqrt{d}}{2^k} \sum_{p_1 \dots p_k} |n_{1, p_1 \dots p_k} - n_{2, p_1 \dots p_k}|. \end{aligned} \quad (\text{B.4})$$

We now proceed to derive a subadditivity property for the average value of L_{MBM} . To do this, it is useful to consider the case where N_1 and N_2 are not fixed integers but are independent Poisson random variables with the same fixed parameter N . For a given k , the numbers n_{1, p_1, \dots, p_k} and n_{2, p_1, \dots, p_k} are then also independent Poisson random variables, with parameter $N/2^{kd}$. Let $M(N) = \langle L_{MBM}(X_1, \dots, X_{N_1}; Y_1, \dots, Y_{N_2}) \rangle$. It is easy to see that

$$\langle L_{p_1 \dots p_K} \rangle = 2^{-K} M(N/2^{Kd}). \quad (\text{B.5})$$

Moreover, well-known properties of Poisson variables allow us to write

$$\langle |n_{1, p_1 \dots p_k} - n_{2, p_1 \dots p_k}| \rangle \leq \sqrt{2} \left(\frac{N}{2^{kd}} \right)^{1/2}. \quad (\text{B.6})$$

By taking mean values in (B.4), we are thus led to

$$M(N) \leq 2^{K(d-1)} M(N/2^{Kd}) + \sqrt{2dN} \sum_{k=1}^K 2^{k(d/2-1)}. \quad (\text{B.7})$$

Extending this construction, one may easily prove [24] that for $2^K \leq m < 2^{K+1}$ we have

$$M(N) \leq m^{d-1} M(N/m^d) + 2^d \sqrt{2dN} \sum_{k=0}^K 2^{k(d/2-1)}. \quad (\text{B.8})$$

Dividing this last inequality by $N^{1-1/d}$ and then replacing N by $m^d N$,

$$\frac{M(m^d N)}{(m^d N)^{1-1/d}} \leq \frac{M(N)}{N^{1-1/d}} + \frac{2^d \sqrt{2d}}{N^{1/2-1/d}} \sum_{k=0}^K 2^{-k(d/2-1)}. \quad (\text{B.9})$$

Standard arguments may now be used to show that the ratio $M(N)/N^{1-1/d}$ necessarily converges to a limit $\beta_{MBM}(d)$ as $N \rightarrow \infty$. Indeed, let $f(t) = M(t^d)/t^{d-1}$. One verifies at once that $f(t)$ satisfies

$$f(mt) \leq f(t) + C_d/t^{d/2-1} \quad (\text{B.10})$$

for all $t > 0$ and all integer m ; $f(t)$ is continuous, since $M(N)$ is a continuous function of N . So the expression $f(t) + C_d/t^{d/2-1}$ is bounded in $[1, 2]$ and since $[1, \infty)$ is the union of the intervals $m[1, 2]$, $m \geq 1$, it follows from (B.10) that $f(t)$ remains bounded as $t \rightarrow \infty$, thus $\lim^* f(t) < \infty$. Now define $\beta = \lim_* f(t)$. For any $\epsilon > 0$, choose $t_0 \gg 1$ and $\eta > 0$ such that $f(t) + C_d/t^{d/2-1} < \beta + \epsilon$ for t in the interval $I = [t_0 - \eta, t_0 + \eta]$. Since the intervals mI , $m \geq 1$ span a whole interval $[A, \infty)$ for A sufficiently large, it follows again from (B.10) that $\lim^* f(t) \leq \beta + \epsilon$. Since ϵ is arbitrary, we have $\lim^* f(t) = \beta$, hence $f(t) \rightarrow \beta$ as $t \rightarrow \infty$, from which it follows that

$$\lim_{N \rightarrow \infty} M(N)/N^{1-1/d} = \beta.$$

We have thus shown that for $d \geq 3$,

$$\begin{aligned} \langle L_{MBM}(X_1, \dots, X_{N_1}; Y_1, \dots, Y_{N_2}) \rangle \\ \sim \beta_{MBM}^E(d) N^{1-1/d}, \quad N \rightarrow \infty \end{aligned} \quad (\text{B.11})$$

when N_1 and N_2 are independent Poisson variables with parameter N . The same result for the mean value $\langle L_N \rangle$, where N is a fixed integer, then follows easily. Indeed, one has the obvious bound

$$\begin{aligned} |L_{MBM}(X_1, X_N; Y_1, Y_N) - L_{MBM}(X_1, X_{N_1}; Y_1, Y_{N_2})| \\ \leq \sqrt{d} (|N_1 - N| + |N_2 - N|), \end{aligned} \quad (\text{B.12})$$

and taking mean values and dividing by $N^{1-1/d}$, we deduce that

$$\lim_{N \rightarrow \infty} \frac{\langle L_N \rangle}{N^{1-1/d}} = \beta_{MBM}(d). \quad (\text{B.13})$$

For further discussion of self-averaging proofs see [32, 41].

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