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Activated Aging Dynamics and Effective Trap Model Description in the Random Energy Model

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Abstract

We study the out-of-equilibrium aging dynamics of the Random Energy Model (REM) ruled by a single spin-flip Metropolis dynamics. We focus on the dynamical evolution taking place on time-scales diverging with the system size. Our aim is to show to what extent the activated dynamics displayed by the REM can be described in terms of an effective trap model. We identify two time regimes: the first one corresponds to the process of escaping from a basin in the energy landscape and to the subsequent exploration of high energy configurations, whereas the second one corresponds to the evolution from a deep basin to the other. By combining numerical simulations with analytical arguments we show why the trap model description does not hold in the former but becomes exact in the second.

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1 Introduction

A large variety of liquids, if kept in the super-cooled metastable phase below the melting point, exhibit a spectacular steep increase of the relaxation time as the temperature decreases. Despite the inherently disordered microscopic structure, this slowing down of the dynamics results in an effectively solid behaviour of the sample, which is commonly denominated glass [1]. At high temperatures, the system behaves as a simple liquid characterised by a fast dynamics [2]. When the temperature T is lowered, motion both in real and configuration space becomes gradually slower and the relaxation time extracted from dynamical correlation functions increases faster than Arrhenius in the so-called fragile liquids. Such behaviour is characteristic of activated dynamics in which a system in order to relax (and flow) has to jump over larger and larger barriers as the temperature is lowered, thus leading to a time-scale

$$\tau(\Delta E) = \tau_0 \exp(\Delta E/k_B T), \quad (1)$$

where ΔE increases when T decreases (τ_0 sets the microscopic timescale, k_B is the Boltzmann constant which in this paper we will set to one).

A general theoretical understanding of activated dynamics, in particular approaching the glass transition, remains an open challenge although it is arguably the central problem of glassy dynamics, and it has implications and ramifications in many other fields from computer science to biology. There has been actually a lot of progress in the last decades in the study of infinite-dimensional or mean field (MF) glassy models [3, 4] both regarding thermodynamical and dynamical properties. However, since in MF models the energy barriers are extensively high, activated dynamics takes place on time-scales diverging exponentially with N and is very difficult to analyze; indeed so far analytical results have been obtained on the regime of times of order one [3] and there have been only few works which addressed the role and the determination of energy barriers [5, 6, 7, 8]. New information has been recently gained from rigorous analysis of trap models and generalizations. As shown by Bouchaud [9], the Trap model (TM) [10] provides a simple framework to study activated dynamics. The TM describes a motion in the space of configurations in which the only way to reach a new configuration is to attain a fixed energy threshold. From these high-energy threshold, it is possible to reach any other point of the phase space. Time scales in the TM are given by the Arrhenius law, and a detailed analysis of the dynamics can be worked out [9, 11, 12, 13] (see [14, 15]

for comparison and applications to glassy liquids). In recent years these results obtained by physicists were shown to hold rigorously, and more than that, they were generalized in many other contexts [16, 17]. This line of research culminated in the proof that the Random Energy model (REM) [18] (the simplest mean-field glassy model) endowed with single spin flip Metropolis dynamics has an aging dynamics on exponentially diverging time-scales effectively equivalent to the one of TM [19, 20, 21]¹. Yet, it remained unclear *how* TM dynamics emerges in REM. In this work we answer this open question by addressing the following points: (1) What happens at shorter, subexponential, time scales, (2) What is the mechanism that allows for a TM description at exponentially large times, (3) How are observables influenced in the different time regimes, (4) The role played by the energy landscape, and (5) The importance of finite size effects. All this additional information will be instrumental in the task of further seeking for evidences of a trap-like dynamics in other disordered systems.

The two key features that make the REM different from the TM are (i) that the states of the REM are defined on an N -dimensional hypercube, instead of a fully connected space, and (ii) that the system follows physical dynamics, such as single spin flip Metropolis Monte Carlo (MC), instead of the purely barrier-passing trap dynamics.

These differences make REM dynamics more difficult and interesting than the one of TMs. In particular, it retains some of the essential ingredients of activated dynamics present in more realistic systems. Its study is therefore a first step to develop a general theory. Developing a general understanding of how and to what extent TM-like dynamics emerge at long times in the REM is the aim of our work. In order to do this, we study dynamics on exponentially growing time-scales by performing numerical simulations of finite systems. This approach was pioneered by Crisanti and Ritort [22] to understand properties of activated glassy dynamics. Previous results have been obtained for a modified (exponential) REM using a microscopic Hamiltonian based on the number partitioning problem [23]. It was shown that many relevant observables such as the distribution of trapping times and the aging functions (autocorrelation functions), are indeed trap-like.

In the following, after setting up the notations and recalling the main models and results used in this work, we shall first show to what extent and on which time-scales REM dynamics is different from TM one. We will then explain and show how on very large time-scales the TM dynamics emerge.

2 Models and simulations

This section is devoted to recall the definition of the models we will refer to, list the results useful for our work, and set the notation.

2.1 Trap Models

Trap Models (TM) were proposed to give a first simplified framework for activated aging dynamics [9, 12]. They are defined in terms of a collection of M

¹The proof of [20] was made public recently, while this work was already at an advanced stage.

configurations whose energies E_i are independent identically distributed (i.i.d.) random variables extracted from an exponential distribution

$$\rho_{\text{exp}}(E) = \frac{1}{\alpha} \exp(E/\alpha) \Theta(-E) , \quad (2)$$

where $\alpha > 0$ is the mean of the exponential distribution, and $\Theta(x)$ is the Heaviside step function; when $M = 2^N$, these configurations may represent all the possible configurations of a system with N binary degrees of freedom, e.g. Ising spins.

Their continuous time dynamics is defined as follows: at each time step a configuration i is randomly chosen with flat probability $1/M$ among all the configurations at disposal, the system remains trapped in the configuration i for a *trapping time* t which is a random variable extracted from an exponential distribution with mean

$$\tau_i = \tau_0 \exp(-\beta E_i) , \quad (3)$$

where $\beta = 1/T$ is the inverse temperature, and τ_0 sets the time unit. Once it moves, the system jumps to a configuration drawn uniformly from the M available ones and the process described above repeats.

Note that the inverse of the average trapping time can be interpreted as the (Arrhenius-like) probability for the system to escape from its trapping configuration i

$$P_{Arr} = \exp(-\beta \Delta_i) , \quad (4)$$

once the configuration is surrounded in all the directions by barriers of height $\Delta_i = E_{top} - E_i$ with $E_{top} = 0$. In other words, the energy landscape is like a golf-course: during the dynamics the system needs to jump up to a fixed top level of energy E_{top} in order to access the rest of the energy landscape. Once this is achieved, the dynamics loses completely memory of the past because any configuration is accessible from any other irrespectively from the past dynamical history, and any new configuration is equally acceptable. Due to these two features TM dynamics is a *renewal process* [24] because after every jump the process starts anew. Despite this strong simplification, the dynamics retains a number of features that lead to a non trivial aging phenomenology, as it will be evident in what follows.

It can be easily shown that the mean trapping times τ obtained from trapping configurations with random energies E are random variables, power-law distributed [9]

$$\psi_{\text{exp}}(\tau) = \frac{T}{\alpha \tau} \left(\frac{\tau}{\tau_0} \right)^{-T/\alpha} \Theta(\tau - 1) \propto \frac{1}{\tau^{1+T/\alpha}} , \tau \gg 1. \quad (5)$$

The regime $T/\alpha < 1$ (or $T < \alpha = T_c$) corresponds to a dynamics with several interesting properties [9]. Among these, the configurational average of the mean trapping times τ_i is infinite (correspondingly the thermodynamic limit is not defined: the partition function is divergent). Moreover taking a collection of n of these random variables $\{\tau_i\}$ their sum is dominated by the maximum among them: $\sum_i^n \tau_i \sim \tau_{\text{max}}$. Equivalently, at a finite time from the beginning of the dynamics, $t_w = \sum_i^n \tau_i$, the system is trapped with finite probability in the lowest trap among the ones already explored (the initial condition is drawn uniformly from the M configurations). The energy E_{min} of the lowest trap can be therefore

evaluated equating its trapping time $\tau_{\max} \simeq \tau_0 \exp(-\beta E_{\min})$ to t_w and inverting the relation. Given that a great fraction of time is spent in the lowest trap, the average energy also scales similarly with t_w :

$$E(t_w) = -T \log \left(\frac{t_w}{\tau_0} \right). \quad (6)$$

This is an example of aging behaviour in which the relaxation time of the system is set by its age [25, 26].

TM are so simple that it is possible to explicitly evaluate average correlation functions along the dynamics. The dynamical observable considered in the literature is the persistence function $C(t_w, t_w + t)$: it is non zero (and equal to 1) as long as the system remains trapped in the same configuration between t_w and $t_w + t$. As soon as the system escapes from the trapping configuration it occupied at t_w , the persistence drops to zero. The average value of the persistence is given by the probability of not changing configuration between the two observed times t_w and $t_w + t$, which is traditionally defined as $\Pi(t_w, t_w + t)$. This quantity has been found [9] in the large time limit to only depend on the ratio $w = t/t_w$ according to the so-called Arcsin law [11, 16]:

$$\lim_{t_w \rightarrow \infty, t/t_w = w} \Pi(t_w, t_w + t) = \frac{\sin(\pi x)}{\pi} \int_w^\infty \frac{du}{u^x(1+u)} \equiv H_x(w), \quad (7)$$

where the parameter x depends on the mean α of the exponential distribution $\rho_{\text{exp}}(E)$, and on the temperature T of the dynamics as $x = T/\alpha < 1$.

Gaussian Trap Model We will want to compare the TM to the REM, defined in the next section. With this objective, as a first minor step towards a more realistic model for activated aging dynamics, we also take into account the Gaussian Trap model (GTM), in which the distribution of the energy is Gaussian,

$$\rho_{\text{gauss}}(E) = \frac{1}{\sqrt{2\pi N}} \exp \left(-\frac{E^2}{2N} \right), \quad (8)$$

and the distribution of trapping times is therefore

$$\psi_{\text{gauss}}(\tau) = \frac{T}{\tau \sqrt{2\pi N}} \exp \left(-\frac{T^2}{2N} \log(\tau/\tau_0)^2 \right). \quad (9)$$

Note that $\psi(\tau)$ is in this case lognormal.

This last model is directly inspired by the REM, introduced in [18]: the equilibrium properties of GTM (with $M = 2^N$) and REM coincide, but the dynamical rules are different. In the case of the GTM, these are equivalent to the ones defined before (for the REM see next section). From equilibrium studies, it is known that this model shows a thermodynamic entropy-vanishing transition at a critical temperature $T_c \equiv 1/\beta_c = \frac{1}{\sqrt{2 \log(2)}} \simeq 0.849$ [18]. For what concerns the dynamics, the GTM presents a dynamics with an infinite number of time-scales. On each one of them the dynamics is effectively identical to the one of an exponential trap model with a time-scale dependent α [27]. More precisely, the continuous time dynamics of the GTM behaves like the one of a TM

with a parameter $\alpha(t_w) = T/x(t_w) = N/(T \log t_w) < 1$ dependent on the age of the system [13] if the times at which the dynamics is observed are properly rescaled [11, 13, 28].²

2.2 Random Energy Model

As anticipated in the previous section, following its classical definition [18] we call REM a model that is defined as a collection of $M = 2^N$ configurations representing all the possible microstates of a system with N binary variables, that we call spins. To each configuration is associated an i.i.d. quenched random variable extracted from the Gaussian probability distribution $\rho_{\text{gauss}}(E)$, previously shown in equation (8).³

The most natural choice of dynamics to be considered for such a system is a single spin flip MC Metropolis dynamics [29], the study of which has been the motivation of a number of previous rigorous studies [21, 20] and will be the focus of this work.

Beside the classical REM, to fill the gap between the REM and TM, we will also consider a slight modification of this model that we will call Exponential Random Energy model (EREM) [30]. In this model the distribution of energies is exponential $\rho(E) = \rho_{\text{exp}}(E)$; also for this model we consider a single spin flip Metropolis dynamics as we do for the REM.

We underline again that from the *equilibrium* point of view REM and EREM are alike GTM and TM, respectively. Dynamically, two major elements of difference are introduced: (1) the system is not in general expected at every step to go back to the highest possible energy level at disposal, E_{top} ; (2) since only single spin flip moves are considered the number of neighbouring configuration is N (instead of 2^N), hence the number of dynamical paths at disposal is considerably reduced. In this case, the lattice representing the dynamical connections between configurations separated by a single move is not fully connected. It is instead a hypercube in dimensions N with edges corresponding to the moves induced by the spin-flips. These two new ingredients, despite the absence of static correlation between the energies of any pair of configurations, might introduce correlations in the energies sampled during the dynamics, and considerably change the aging behaviour.

Note that these two ingredients must be taken into account if we aim at develop the description of an out-of-equilibrium dynamics for more realistic systems where dynamics is always controlled by local moves and typical dynamical paths may allow relaxation without stepping through high energy levels. The additional introduction of static correlations between the energies of neighbouring configurations is a further element to consider when developing a theory of activated dynamics of realistic systems, but it will not be the argument of this work.

²This becomes evident if we rewrite (9) in the form of an exponential distribution in analogy with (5) and define an effective $T/\alpha \sim (T^2 \log \tau)/N$ where we know that the relevant time scale of the system τ is given by its age.

³The variance of $\rho_{\text{gauss}}(E)$ is slightly different from Derrida's original definition [18], since we have $\text{var}E = N$ instead of $N/2$. This conveys that the critical temperature is rescaled by a factor $\sqrt{2}$.

2.2.1 Basic Features of the Energy Landscape

Before focusing on the out of equilibrium dynamics of these two models we first list a number of features that characterise the typical local arrangement of energy levels around a given (say deep) configuration C_0 .

In order to do this, we shall use results on extreme value statistics [31] that we recall by using simple scaling arguments. We evaluate the typical energy level of the lowest configuration, C_1 , among the N configurations surrounding a given C_0 , of the second lowest configuration, C_2 , and so on and so forth, as it is represented in Fig. 1. Since the energies of configurations are uncorrelated,

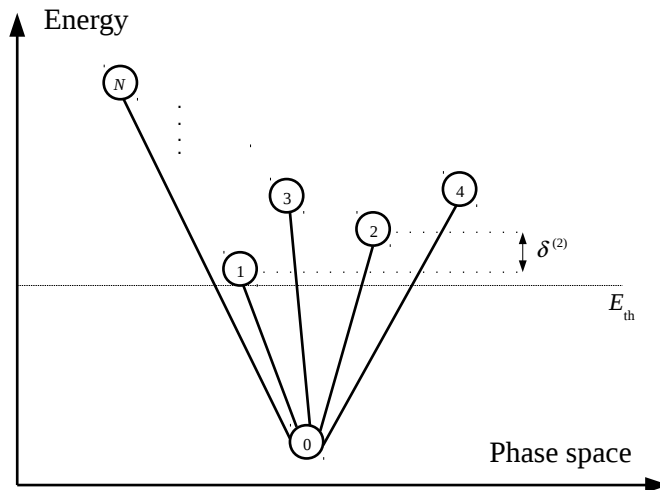


Figure 1: Schematic diagram of the connectivity of a low-lying configuration C_0 , with its neighbours C_1, C_2, \dots, C_N . E_{th} is the threshold energy, below which we say that the system is in a basin. We define E_{th} as the typical value of the smallest neighbour of any configuration (i.e. $E_{\text{th}} = E^{(1)}$). The energy difference $\delta^{(2)}$ is defined in Eq. (14).

these typical energy levels will not depend on C_0 . The energy $E^{(1)}$ of the lowest configuration among the nearest neighbours of C_0 is the minimum of N Gaussian variables, whose scaling can be obtained by evaluating the probability that out of the N energies at disposal one has an energy contained in the interval $[-\infty, E^{(1)}]$. The energy scale where this probability starts to be of the order of one sets the scale of $E^{(1)}$, leading to the equation:

$$O(1) \sim N \int_{-\infty}^{E^{(1)}} dE \rho(E) . \quad (10)$$

In the REM this leads to

$$E^{(1)} \sim -\sqrt{N \log N} + x_1 \sqrt{\frac{N}{\log N}} , \quad (11)$$

where the probability law of the order one variable x_1 can be obtained from extreme value statistics (it is the derivative of a Gumbel law) [31]. The result

for the energy $E^{(k)}$ of the k^{th} lowest configuration among the first dynamical neighbours of C_0 is analogous:

$$E^{(k)} \sim -\sqrt{N \log N} + x_k \sqrt{\frac{N}{\log N}} . \quad (12)$$

where of course $x_1 < x_k$ with probability one. The same arguments in the EREM lead to

$$E^{(k)} \sim -\alpha \log N + x_k . \quad (13)$$

Note that the difference in energy,

$$\delta^{(k)} \sim E^{(k)} - E^{(1)} , \quad (14)$$

between the 1st and the k^{th} deepest levels among N configurations at disposal is diverging with $N \rightarrow \infty$ in the REM,

$$\delta^{(k)} \sim \sqrt{\frac{N}{\log N}} , \quad (15)$$

and it is of order one in the EREM.

A final fundamental remark is that the ground state, or the configuration with lowest energy, E_{GS} , among all the M at disposal can be obtained from the equation (10) by substituting N with M . It gives in the REM

$$E_{GS} \sim -\sqrt{2N \log M} \quad (16)$$

and in the EREM

$$E_{GS} \sim -\alpha \log M . \quad (17)$$

When we consider a number of configuration $M = 2^N$, the typical ground state energy is in both cases extensive and corresponds to $E_{GS} \sim -N\sqrt{2 \log 2}$ in the first case, and to $E_{GS} \sim \alpha N \log 2$ in the second case.

Low temperature dynamics of REM and EREM (with $T < 1/\sqrt{2 \log 2}$ and $T < \alpha$, respectively) will tend to reach this ground state energy in the long time limit, leading the system to be trapped for long time intervals in very deep configurations C_0 surrounded by N configurations which are well above in the energy landscape (leading to a golf-course like landscape as stressed previously). The system has to progressively decrease its intensive energy; however any configuration with a negative intensive energy is surrounded by neighbours that have typically zero intensive energy. This leads to an extremely slow and activated dynamics which proceeds on time-scales diverging exponentially with N .

An important difference between the two models is that, for the REM, dynamics are activated and aging is observed until the system reaches equilibrium, for T both over and under T_c . In the EREM, instead, only the low-temperature phase has interesting dynamics. For $T > T_c$ the dynamics of the EREM is fast and not interesting for our purposes since the system remains floating at zero intensive energy, and does not display aging.

2.2.2 Energy Basin Description

In TMs, configurations are, by definition, schematic approximations of the deep basins, that in more realistic models are made up of several configurations with correlated energies. At first sight also for the REM and EREM basins consist in single configurations as we have seen in the previous section. However basins containing more than one configuration do exist (despite they are much rarer than basins with single configurations, as a matter of fact they are still exponentially numerous in N [21]) and as we shall show they can be relevant for the dynamical behaviour, at least for certain observables. In the following, in order to test if a TM description holds for the dynamics of the REM and EREM we need to map the MC Metropolis dynamics into a sequence of basin jumps, and hence give a pragmatical definition of basins. This mapping can be used also in more realistic models; by studying it for the REM and EREM we shall unveil subtleties and properties related to this mapping that can be useful when it is applied to more realistic cases.

Given the high dimensionality of the energy landscape, it is not straightforward to partition the system in a set of distinct basins. Therefore, inspired by a similar analysis proposed recently in [32], we use a dynamical definition of basin, which focuses on the one-dimensional dynamical path described by the energy evolution as a function of time, $E(t_w)$. We choose a threshold energy E_{th} and say that each time E becomes smaller than E_{th} the system is *in a basin*. When finally $E > E_{\text{th}}$ again, the basin is abandoned, and we say we are *on the barrier* between two basins. A schematic representation is provided in Fig. 2. A natural choice for E_{th} is the average minimum energy $E^{(1)}$ of the N configurations that can be reached from any configuration C_0 in a single dynamical step. With this choice, almost all the neighbours of any configuration are typically at $E \geq E_{\text{th}}$. This energy level does not depend on the configuration C_0 visited by the dynamics, whose energy E_0 is slowly decreasing towards E_{GS} . Hence the distance between E_{th} and the energy of the configuration typically explored by the dynamics will become larger and larger in the long time limit in a similar way as in TM dynamics.

Note that for short times, the visited basins are typically very close to E_{th} , implying that the system has to age enough time before the energy basins become similar to traps, where the system remains stuck for long times. In full generality, since within each basin B the system overwhelmingly spends the most of the time in its lowest-energy configuration, we associate to B the energy of the deepest configuration that belongs to that basin:

$$E_{\text{basin}} = \min_{C_i \in B} E(C_i). \quad (18)$$

The same applies to any static property we mention of a basin: it refers to the lowest-energy configuration within the considered basin.

Finally, to each basin we associate a trapping time as shown in figure 2.

3 Absence of trap-like behaviour on intermediate time scales

In this section we shall show that the dynamics of REM and EREM is not trap-like on time-scales corresponding to the exit from a basin and the exploration

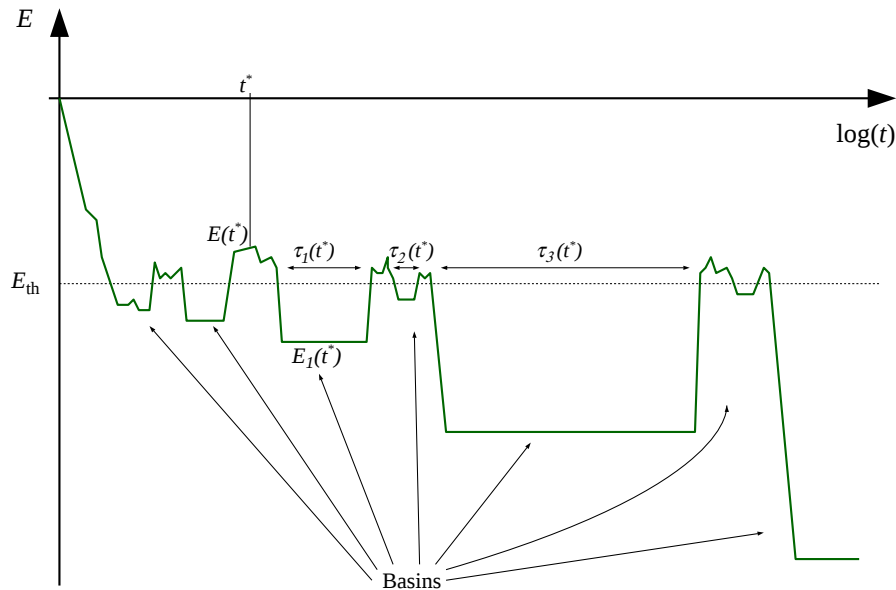


Figure 2: A sketch of the energy evolution in either the EREM or the REM. At a given time t^* , the instantaneous energy is $E(t)$. Independently of whether at time t^* the system is in a basin or on a barrier, the energy of the next basin is $E_1(t^*)$, calculated via Eq. (18). The times $\tau_1(t^*)$, $\tau_2(t^*)$ and $\tau_3(t^*)$ are the trapping times of the three basins following t^* .

of configurations at the threshold level. As we discuss in the next section, the mapping to the TM only emerges on very large time-scales in a coarse-grained way.

We first present numerical results concerning the dynamics of the REM and EREM. For each series of simulation, we focus on three temperatures in the glassy phase, $T = 0.25, 0.50, 0.75$, and $T = 1.50$ in the disordered phase. We remind the reader that $T_c = \alpha$ (we choose $\alpha = 1$) in the EREM, and $T_c = \frac{1}{\sqrt{2 \log(2)}} \simeq 0.849$ in the REM.

The system sizes are $N = 8, 12, 16, 20, 24$. The length of the run is $t_{\text{run}} = 2^{1+A+N}$ MC steps. The value of A was decided according to the wallclock time availability. The amount of samples N_{sam} per parameter set, and other details of the simulations are described in Table 1.

3.1 Non-renewal behaviour

In the following we focus on several observables that in the TM show that the dynamics is a renewal process. This means roughly speaking that every time a basin is abandoned, the system should completely loose memory of its past. As we shall show, the behaviour we find is quite different.

T	N	A	t_{run} (MC steps)	N_{sam}
0.25, 0.50, 0.75	8	18	10^8	20000
0.25, 0.50, 0.75	12	18	10^9	12000
0.25, 0.50, 0.75	16	18	10^{10}	6000
0.25, 0.50, 0.75	20	18	10^{12}	1500
0.25, 0.75	24	18	10^{13}	200
1.50	16	0	10^5	100
1.50	20	0	10^6	100
1.50	24	0	10^7	100

Table 1: Parameters of the simulations. We performed the same simulations both for the Exponential and the Gaussian REM. The exact number of MC steps of each run is $t_{\text{run}} = 2^{1+A+N}$. In the column of t_{run} we display the order of magnitude of this number. N_{sam} is the number of independent samples simulated with the given set of parameters.

3.1.1 Energy of the next basin

If we fix a time t , the system will likely be in a basin. As shown in Sec. 4.1, the average energy $E(t)$ depends on t approximatively as described by the relation (6). So, indeed, the system is trapped in deeper minima at long times as in the TM. However, in the TM, once a basin is abandoned then the energy of the following one, $E_1(t)$ (see Fig. 2), should be completely uncorrelated, i.e. E_1 is independent of $E(t)$ (and hence of t). The situation is quite different in our simulations of the dynamics of REM and EREM as shown in figure 3, where $E_1(t)$ is instead clearly dependent on t and $E(t)$.

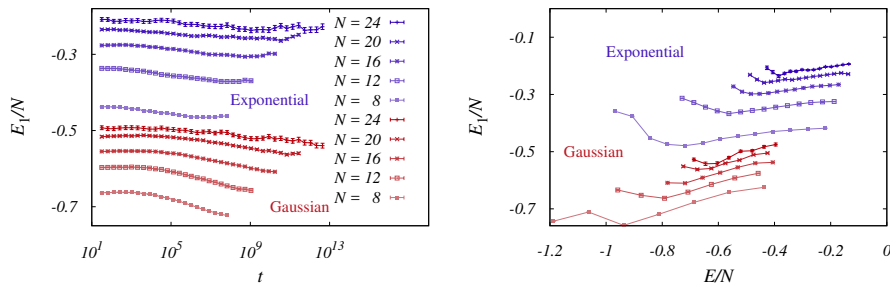


Figure 3: Energy of the next basin, E_1 , in the EREM (blue curves) and in the REM (red curves), at $T = 0.25$. On the left side we show it as a function of time, on the right side we show it as a function of the previous basin's energy. The data for the EREM levels out when the system has thermalised (thermalisation times can be deduced from figure 9). In the right plot, we removed the points at lowest energy because they were fluctuating. See also the caption of figure 4 for remarks on the measurement protocol.

3.1.2 Trapping time of the following basins

As well as the energy of the next basin, also the statistics of the trapping time should be time-independent if the dynamics is renewal. We call $\tau_1(t)$ the trapping time of the basin that immediately follows the one found at t , $\tau_2(t)$ the energy of the next basin, and τ_3 the one of the third basin after t . A schematic description of the definition of τ_1, τ_2 and τ_3 is provided in Fig. 2.

We see from figure 4 that τ_1, τ_2 and τ_3 are not time-independent in the aging regime, again in contradiction with the renewal property of a trap-like dynamics.

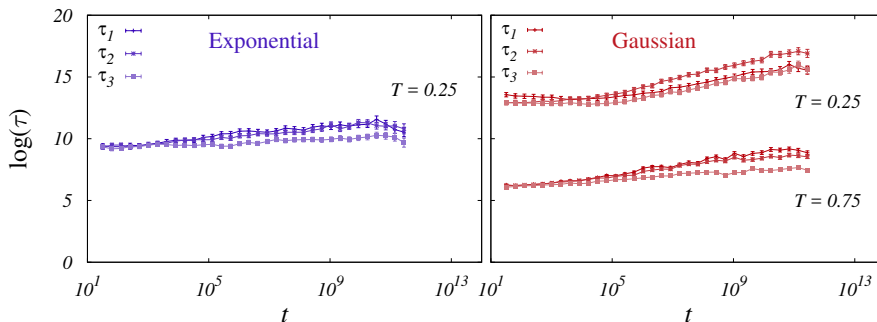


Figure 4: The trapping times τ_1, τ_2 and τ_3 in the EREM (left, $T = 0.25$) and REM (right, $T = 0.25, 0.75$), for $N = 20$. In the EREM, results for $T = 0.75$ are not reported, because no sizable growth of τ_1, τ_2 and τ_3 was found. The time dependence is fairly visible, especially in the Gaussian case. The growth of τ_1 is stronger than that of τ_2 and τ_3 , since it is the trapping time that preserves the highest memory of t . Our measuring procedure excludes basins that, at the end of the run, have not been abandoned. This gives a bias in the final points of the curves, since towards the end of a run large basins are rejected more often than small ones. The growth we observe is clearly visible despite this bias.

3.1.3 Rank of the deepest trap

After n_{traps} basins have been visited (we only count the last basin if the run finished on a barrier), one can ask when the longest trap has been visited. To be more precise, we assign a rank i to each basin, according to the order of occurrence. So, $i = 0$ indicates the first basin, $i = 1$ the second basin, and so on, until $i = n_{\text{traps}} - 1$ for the last basin. If the dynamics is renewal, the trapping time τ_i of basin i follows the same probability distribution for any i . Hence, any of the visited basins can be the one with the longest trapping time with equal probability. Calling i_{max} the rank of the basin with the longest τ , the distribution $P(i_{\text{max}}/n_{\text{traps}})$ is flat in TMs. In figure 5 we show that for the REM this is not the case.

3.2 Back and forth in the same basin

The non-renewal features of the previous section can be explained in a simple way in terms of *returns* to the same basin. While in the TM the phase space connectivity graph is fully connected, in the REM it is a hypercube, since a

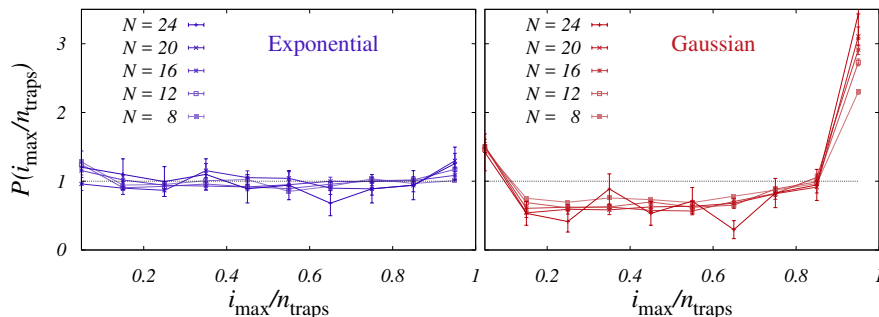


Figure 5: The distribution $P(i_{\max}/n_{\text{traps}})$ in the Exponential (left) and Gaussian (right) REM, at $T = 0.25$. In the Gaussian REM, $P(i_{\max}/n_{\text{traps}})$ is significantly non-flat, in contrast with the TM prediction.

movement is obtained by flipping one of the N spins. More explicitly, in a TM, from each configuration the system can reach any of the 2^N states. On the contrary, in the REM only N configurations are connected to each site, and of those only few (or one) of them are typically visited. The smaller amount of available paths in configuration space conveys that there is a finite probability that, when the system leaves a valley, the next visited basin will be again that same one. What happens is that when the system manages to jump out of a deep configuration it lands in a new configuration close to the threshold level. However, the dynamics and the configuration space at the threshold level is complicated (see later for a discussion) and before being able to wander far from the initial configuration, the system repeatedly visits it. If the system is in a very deep basin, it is likely that the next basin be very deep too, since it is likely that the same basin be visited again.

Let us now confirm these statements by simulation results. We call S the number of spins that are different between each basin and the following one. $S = 0$ means that the same basin was visited twice in a row. In figure 6 we show

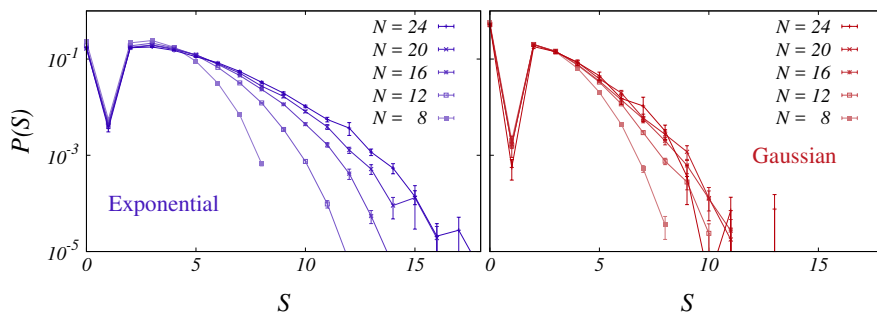


Figure 6: $P(S)$ in the Exponential (left) and Gaussian (right) REM, at $T = 0.25$. A large fraction of times the system comes back to a just visited basin ($S = 0$). This fraction does not show a visible dependence on N : already for $N = 20$ the size dependence seems to have saturated.

that $S = 0$ a finite fraction of the times. Note that $P(S)$ is an indicator of

the basin size and the distance between basins. When increasing N , the $P(S)$ seems to converge to a limiting function, independent of N . In other words, for large enough sizes $P(S)$ does not depend on N . Since the typical basin size is 1 by construction (in the REM two configurations differing by one spin-flip have uncorrelated energies), this indicates that the distance between basins is also $\mathcal{O}(1)$. A second indication of this is found in Sec.4.2, where the times spent on the barrier do not grow with N .

Finally, let us point out that one would expect that the distance between two different basins be at least 2, since there should be at least one barrier configuration between them. The fact that $P(S = 1)$ is positive, though small, descends from the fact that occasionally neighbouring configurations with $E < E_{th}$ may be counted as two separate basins at distance 1, because of our dynamic definition of basins. This happens when a transition to $E > E_{th}$ was accepted after visiting one, and before visiting the other.

We can quantify the phenomenon of immediate returns in the same basin by computing the probability that the system falls twice in a row in the same basin. We will focus on the simplest situation, that we call *immediate returns*, in which the system returns to a basin immediately after having left it. This particular case provides a lower bound to the frequency of returns, and can thus be used to state that returns occur when $N \rightarrow \infty$.

Until now, we have studied the structure of the configurations dynamically connected to a given C_0 , knowing that this structure does not depend on the chosen C_0 , due to the independence of the energies from the configurations in the EREM and REM. For the computation of the probability of immediate returns we will assume that the system just left a basin. In other words we will consider the case of a configuration C_0 on the barrier ($E_0 > E_{th}$) that is just one spin flip apart from a configuration C_1 that was deep down in the energy landscape ($E_1 < E_{th}$). Apart from this, the other $N - 1$ configurations C_2, \dots, C_N dynamically connected to C_0 follow the standard statistics described by (12) and (13). By construction, $E_1 \ll E_{th} \leq E_0$ (see figure 7).

We now estimate the probability of *immediate returns* by simple arguments. We call $P_i^{(acc)}$ the probability of accepting a move from C_0 to C_i . According to the Metropolis rule,

$$P_i^{(acc)} = \min[1, e^{\beta(E_0 - E_i)}]. \quad (19)$$

In a sequentially updated dynamics, the system returns straight back to C_1 when the other $N - 1$ attempted updates, leading to C_2, \dots, C_N , are rejected. Since each move is accepted independently, the probability of rejecting all the $N - 1$ moves is

$$P^{(rej)} = \prod_{i=2}^N (1 - P_i^{(acc)}). \quad (20)$$

Once any other move has been rejected we are sure that the system will go back to C_1 , since $E_1 < E_0$. Hence, to evaluate the probability of an immediate return, we need to estimate the probability of rejecting all the other moves, paying attention to the energy structure of the configurations C_2, \dots, C_N (there is no need to consider a sequential dynamics or that all moves are attempted before C_1 , since the only thing that matters is the scaling of $P^{(rej)}$ when $O(N)$ attempts are considered).

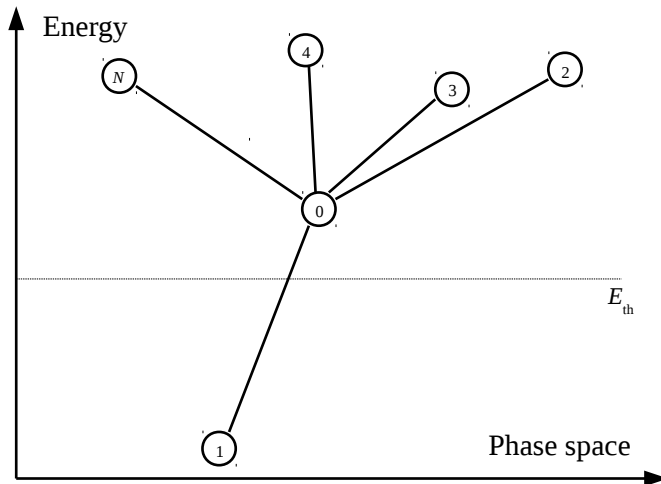


Figure 7: The diagram shows the situation we take in account for our computation. Each circle indicates a configuration in phase space, with its energy. The horizontal line marks the threshold energy E_{th} . We consider the case in which the system is found on the edge of a basin. It has just reached the configuration C_0 moving from C_1 which lies deep down in a basin. In order to go immediately back to C_1 the subsequent $\mathcal{O}(N)$ moves towards its other $N - 1$ neighbours need to be rejected.

Since the energy differences between E_0 and E_i scale in a different way in the REM and EREM, we consider the two cases separately.

- REM.** The energy E_0 is by definition typically of the same order (but slightly larger than) E_{th} . The neighbours of C_0 do not coincide with the configurations surrounding the initial configuration C_1 , the one deep down in the energy landscape (figure 7). Thus, in order to estimate the landscape around C_0 , one has to consider the minimal energies among the $N - 1$ other neighbours of C_0 . These follow eq. (12), since considering N or $(N - 1)$ neighbours leads to subleading corrections only. As a consequence, we obtain that the typical difference between E_0 and the energy of its lowest neighbouring configuration (excluding C_1) scales as $\delta^{(2)} \sim \sqrt{\frac{N}{\log N}}$ in the large N limit. The probability distribution of this difference can be obtained by using results from extreme value statistics, but for our purposes it suffices to say that, with finite probability $P^{up} < 1$, the lowest neighbour of C_0 (excluding C_1) has a higher energy, of order $E_0 + \mathcal{O}(\sqrt{\frac{N}{\log N}})$. In this case, since all other neighbours are even higher in the energy landscape, the probability of accepting a move to one of these neighbours is exponentially small in $\sqrt{\frac{N}{\log N}}$; therefore even after $O(N)$ attempts (the number of neighbours) it remains negligible, i.e. the probability of returning goes to one at large N . Note that with probability $1 - P^{up}$, instead, the lowest neighbour of C_0 (excluding C_1) has an energy

of order $E_0 - \mathcal{O}(\sqrt{\frac{N}{\log N}})$. In these cases there is no immediate return since the system directly goes to this neighbour if this move is attempted before the one towards C_1 ; depending on the neighbours of this neighbour, the system could then be forced to return to C_0 instead of exploring further the landscape.

- **EREM.** In the EREM the computation needs slightly more care, because the separation $\delta^{(2)}$ between the lowest energy levels is of order one. We will keep the reasoning simple and present just a rough (but essentially correct) argument. We call again P^{up} the finite probability less than one that the lowest neighbour of C_0 (excluding C_1) has a higher energy of order $\delta^{(2)} \sim \mathcal{O}(1)$. Let's say that in this case we divide the neighbours of C_0 (excluding C_1) in two groups: n which are the lowest in the landscape and have therefore positive energy differences of the order one and $N - n - 1$ which are typical configurations and have typical energies, hence a difference of the order $\alpha \log N$. Within this construction, n is finite. We can then estimate that

$$P^{(\text{rej})} > (1 - e^{-\beta\alpha})^n \left(1 - e^{-\beta\alpha \log(N)}\right)^{N-1-n}, \quad (21)$$

which in the large N limit goes like

$$P^{(\text{rej})} > (1 - e^{-\beta\alpha})^n e^{-N^{1-\beta\alpha}}. \quad (22)$$

In the high temperature regime, $\beta < 1/\alpha$ (where there is no aging) the lower bound goes to zero, while in the low temperature regime, $\beta > 1/\alpha$, (where there is aging) the lower bound remains finite and equal to $(1 - e^{-\beta\alpha})^n$. This implies that with finite probability the system returns back to the initial basin. A refined argument can be worked out and leads to the same conclusion, which is actually expected from the thermodynamics of the EREM: for $\beta < \beta_c$ the system is sucked up by configuration at zero intensive energies whereas for $\beta > \beta_c$ the system can return back to C_1 , but it does so with probability less than one since it can accept a move to a neighbour which has an energy difference of order one.

In cases where the lowest neighbour of C_0 (excluding C_1) has a lower energy of order $\delta^{(2)} \sim \mathcal{O}(1)$ an analogous scenario holds, except that now if the move to the neighbour with lower energy is attempted before the one to C_1 the system does not effectuate an immediate return with probability one (but it can return with finite probability in a finite number of steps).

Overall we have found that the dynamics after the jump out from C_1 is complicated and cannot be described as trap-like: it actually corresponds to a complex wandering on configurations at the threshold level and above. This explains the numerical results shown at the beginning of this section. We can further confirm that the returns are at the origin of the numerical findings by defining E_1, τ_1, τ_2 and τ_3 in a different way, that suppresses the effect from the returns. We define E_1^{nr} as the energy of the first basin encountered, after t , with $S > 0$, i.e. imposing that the basin is not the initial one. The definition of τ_1^{nr} is analogous. Regarding τ_2^{nr} and τ_3^{nr} , we prescribe that all the recorded basins since t be new ones.

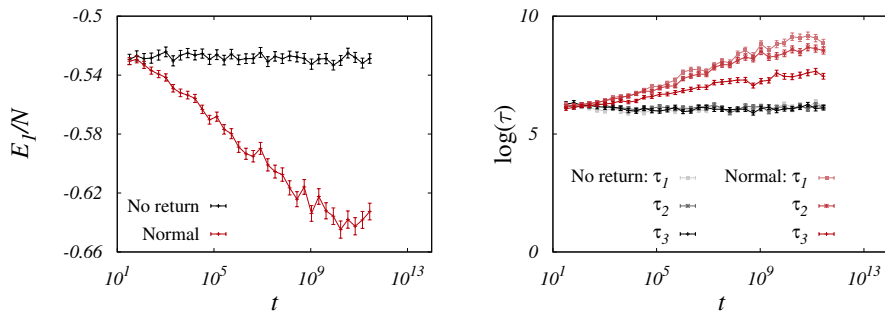


Figure 8: Left: comparison between E_1 in the REM, and its counterpart E_1^{nr} that does not take in account returns to the just-visited basin. While E_1 is logarithmic in time, E_1^{nr} is constant, recovering the TM prediction. Right: comparison between the trapping times τ_1, τ_2 and τ_3 , and their non-returning counterparts $\tau_1^{\text{nr}}, \tau_2^{\text{nr}}$ and τ_3^{nr} in the REM. Also in this case the curves become constant and collapse on each other once the returns are ignored. In both plots we have $N = 20$, $T = 0.25$.

In figure 8 we show that both E_1^{nr} and $\tau_1^{\text{nr}}, \tau_2^{\text{nr}}$ and τ_3^{nr} fulfill the trap prediction now that the returns are not being recorded. Note, however, that although this indicates trap-like dynamics, we do not expect to find renewal dynamics and hence all trap predictions to hold on the time-scale on which the system performs a complex wandering at the threshold level and above. This happens only on much larger time-scales as discussed and showed in the next section.

4 Trap-like dynamics on long time-scales

From the previous analysis we know that when the system is in a basin whose energy is extensively below zero, i.e. $\lim_{N \rightarrow \infty} E/N < 0$, it takes a time τ_J which diverges as $e^{-\beta E}$ to jump out of it. The dynamics that follows this first jump consists in a wandering on configurations at and above the threshold level and in returns to the initial basin. This exploration takes place on times that differ from τ_J by a multiplicative factor which vanishes at exponential leading order in N . For example, for the REM the barriers that the system has to overcome at the threshold level are of the order $\sqrt{N}/\log N$ therefore, even taking into account returns to the initial basin, the time-scale τ to wander at the threshold level is still $e^{-\beta E}$ at exponential leading order: $(\log \tau)/N = -\beta E/N \simeq -\beta E/N + \sqrt{1/N} \log N$ for N large.

As discussed previously, the dynamical processes following the jump out of the basins are not trap-like and, instead, correspond to a quite complex evolution in configuration space. However, if one observes motion on the time-scale $e^{-\beta E}$, which packs together all times τ that differ for a multiplicative factor vanishing at exponential leading order in N , then the wandering at the threshold level becomes a very fast process. On this time-scale, the system does many back and forth motion and explores new configurations. In particular, it visits a number of new configurations that diverges as a power law with N ; this should be enough to loose memory and, hence, effectively move to another uncorrelated basin.

Analogously, although there are correlations in the energies sampled during the dynamics at the threshold level, they do not extend on timescales and hence on a number of different visited configurations which diverge exponentially in N . All that suggests that once dynamics is coarse-grained on time-scales exponentially diverging with N an effective trap-like description should emerge, in particular we expect the fundamental pillars of trap-like dynamics to hold:

1. At time t the lowest intensive energy $e(t)$ sampled is given by the equation $t \sim e^{-N\beta e(t)}$.
2. The number of basins $\mathcal{N}(t)$ visited until time t is given at the exponential accuracy in N by the number of times one has to draw Gaussian random energies in order to get a minimum of the order of $e(t)$: $\mathcal{N}(t) \sim e^{Ne(t)^2}$.
3. Thermal equilibrium sets in for all configurations from energy zero to energy $e(t)$, i.e. at time t one gets an effective Boltzmann distribution cut at energies lower than $e(t)$.

These features imply the emergence of trap-like aging dynamics on exponentially diverging time-scales, as indeed very recently proven in [20].

In the following, we confirm this result through numerical simulations. As we shall show, important finite size-time effects correct the trap-like predictions. Low temperatures make the dynamics so slow that it is very difficult to probe the regime in which energies are extensively below E_{th} . At the same time, important finite size effects are observed for temperatures close to T_c . All that is particularly cumbersome in the REM, where the fact that trap-like aging emerges on an infinite spectrum of exponentially diverging time scales makes the analysis even more difficult.

Note that above we have stated the trap-like properties in terms of basins and not configurations. As discussed in the previous sections a basin typically contains just one configuration. For observables that probe typical behaviour as for example the aging function considering basins or configurations is equivalent; however for observables for which rare effects are enhanced this might be incorrect. For instance, the number of jumps done until time t behaves differently if one considers basins or configurations. In the former case the usual trap prediction holds, whereas in the latter the back and forth motions within rare basins formed by more than one configuration bias completely the statistics which, depending on the regimes, can be different from the trap prediction; see [20] for a detailed discussion and results. Moreover, also from the numerical point of view, considering basins or configurations is not equivalent. As we shall show, the former leads to smaller asymptotic corrections and is therefore preferable.

4.1 Logarithmic decrease of the energy

During aging, the trap prediction is that the intensive energy decreases logarithmically in time as $-T(\log t)/N$, see equation (6). Figure 9 shows that, at $T = 0.25$, for the largest system sizes this is fairly reproduced both for EREM and REM; the deviation from the logarithmic behaviour is mainly due to equilibration at long-times. The result is not as good when $T = 0.75$. We attribute this to the emergence of extremely strong finite-size effects as T approaches T_c . This explanation is confirmed by the fact that, despite being very big for

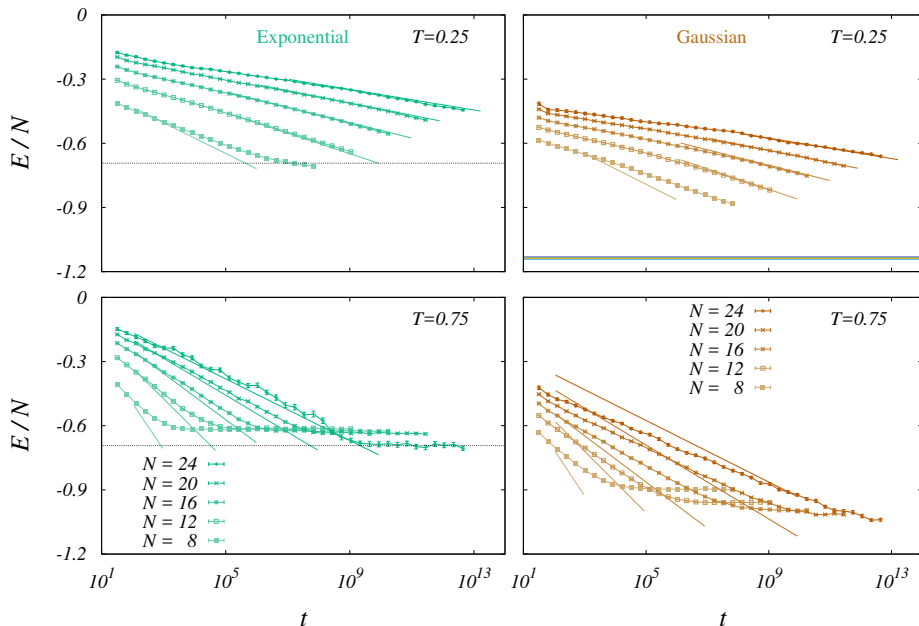


Figure 9: $E(t)$ in the EREM (left) and in the REM (right). The temperature is $T = 0.25$ on top, and $T = 0.75$ on the bottom. The solid lines represent equation (6). The dotted horizontal lines on the left side indicate the ground state energy. When the curve $E(t)$ reaches its equilibrium value, the system has thermalized, and it is no more in the aging regime. The solid horizontal lines in the top-right plot are the predictions of the equilibrium free energy according to reference [33], which takes into account logarithmic corrections. It was not possible to plot the same predictions for $T = 0.75$, because the logarithmic order was not enough to obtain reasonable predictions.

$T = 0.75$ and smaller for $T = 0.25$, deviations between TM predictions and observations decrease as N grows in both cases. As a matter of fact, the finite-size corrections to the free energy diverge as $T \rightarrow T_c$ [33], so the closer the system is to the critical temperature, the larger it needs to be.

4.2 Trapping times

The hallmark of a trap-like dynamics is the power-law distribution of the trapping times $\psi_{\text{exp}}(\tau)$. In the EREM case it follows the power-law behaviour of equation (5). At $T = 0.50, 0.75, 1.50$ the prediction is beautifully satisfied both for the basin and for the configuration trapping times, even for $T > T_c$ (figure 10). This is not the case for $T = 0.25$. At this temperature dynamics is so slow that the aging regime cannot be explored in extent, one needs longer times to enter the trap-like regime in which the typical configuration has $E < E_{\text{th}}$. Anyhow, the data are compatible with the trap prediction both in the exponent and in the fact that basin and configuration observables behave in the same way. As anticipated above, some tuning of the parameters (temperature, size of the system) is necessary in order to avoid big finite time-size corrections.

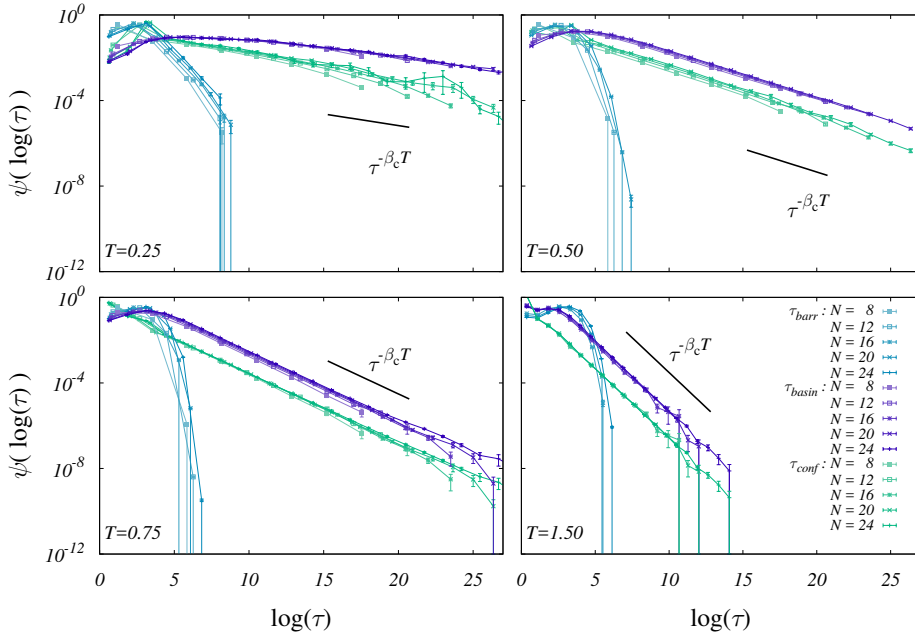


Figure 10: The distribution of the logarithm of the trapping times $\psi_{\text{exp}}(\log(\tau)) \propto \tau^{-T/\alpha}$ with $\beta_c = 1/\alpha$ in the EREM for $T = 0.25, 0.50, 0.75, 1.50$. The red data are the times τ_{conf} before each configuration is abandoned, the blue data represent the time τ_{basin} spent in each basin, and the green data are the times spent on the barriers. The black line is the trap prediction (5). The exponent is different because we are plotting ψ_{exp} as a function of $\log(\tau)$.

In the REM the trap prediction for $\psi_{\text{gauss}}(\tau)$ follows equation (9), which is not a power law.⁴ As found for the EREM, at $T = 0.25$ the simulations require too long times and it is not possible to compare ψ_{gauss} to the prediction. On the other side, at $T = 0.75$, the data fit very well both for the basins and for the configurations (figure 11).

A note on the consistency of the results at different temperatures

Since all temperatures under T_c are equivalent when one seeks the qualitative behaviour of the model, in principle showing results only for one temperature would be enough. There is a reason why one should focus on more than one temperature, and it is that there are two sources of quantitative (not qualitative) changes which depend on T , and we can only choose to rule out one or the other in a single simulation:

- When T_c is approached, the finite-size effects on the free and equilibrium energy are larger [33]. Thus, all other things being equal, close to T_c one has to consider larger system sizes.
- Many of the dynamical behaviors we are interested in only appear on exponentially long time scales, $t \gg \exp(a(T)N)$, with $a(T)$ growing as

⁴Only on a given exponentially diverging time-scale the usual power law is recovered as we recalled in Section 2.

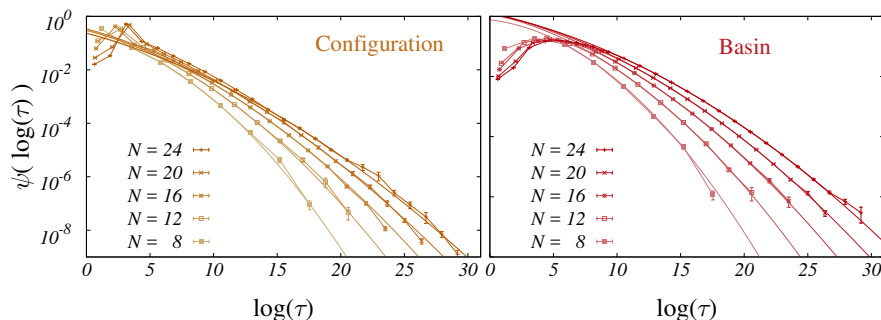


Figure 11: Distribution of trapping times $\psi_{\text{gauss}}(\tau)$ in the Gaussian REM, for $T = 0.75$. Configuration trapping times on the left, and basins on the right. The curves represent fits to the function $f(\log(\tau)) = A \exp(-\frac{T^2}{2N} \log(\frac{\tau}{\tau_0})^2)$, which is the long-time prediction of $\psi_{\text{gauss}}(\tau)$ [equation (9)]. Fits exclude the last point, because we expect that the finiteness of our simulations can bias on the sampling of the longest times.

T is lowered. As a consequence, the temperature has to be high enough otherwise interesting phenomena emerge on prohibitively large time-scales.

Given the competition between the two effects, one happening at low and the other at high temperature, we simulated several temperatures, in order to be able to be in the correct regime for each observable.

4.3 Aging functions

As discussed in section 2.1, all the relevant information of the correlation functions for trap dynamics is contained in the aging function $\Pi(t_w, t)$, which represents the probability of not changing trap. To study the trap behaviour of the aging dynamics in the REM and EREM, we observed the behaviour of this function by considering either basins and configurations for the sake of comparison (as done in [32]). We then defined the probability of not changing basin $\Pi_{\text{basin}}(t_w, t)$ or configuration $\Pi_{\text{conf}}(t_w, t)$ between a time t_w and $t = t_w(1 + w)$ ($w > 0$).

4.3.1 Aging functions in the EREM

In the large-time limit of aging dynamics of TMs, $\Pi(t_w, t_w(1 + w))$ converges to a value $H_x(w)$, with $x = T/\alpha$, that can be computed analytically [11, 16]. In figure 12 we show the aging functions $\Pi_{\text{basin}}(t_w, t)$ for different sizes N and different temperatures in the EREM (left) and REM (right).

We can distinguish two regimes, an initial plateau regime corresponding to the aging regime (associated to $t_w \rightarrow \infty$ after $N \rightarrow \infty$) and a long-time decay once the system thermalised (associated to $t_w \rightarrow \infty$ before $N \rightarrow \infty$). The plateau becomes more and more pronounced as the system size increases, as expected. In the plateau regime, the trap prediction given by $H_x(w)$ applies well to the basin aging functions and less well to the configuration aging function, indicating that activated dynamics in the EREM is effectively trap-like as long as we focus on basins. We interpret the discrepancy between the TM prediction and the

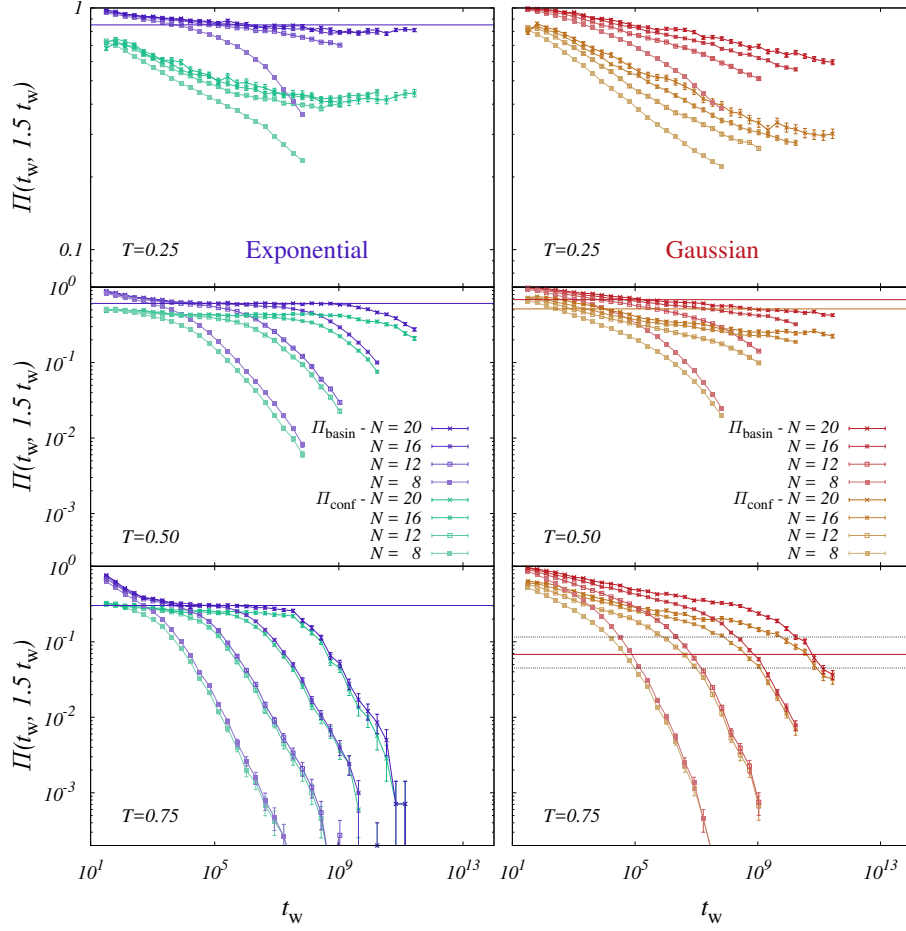


Figure 12: Aging functions REM for $w = 0.5$. From top to bottom: $T = 0.25, 0.50, 0.75 < T_c$. On the left we depict the Exponential, on the right the Gaussian REM. We show the basin and configuration aging functions, for different system sizes. Note the different scale for Π in the two top sets. In the left plots, the blue horizontal lines represent the infinite-time limit of the aging function for an infinitely large system [recall equation (7)]. The horizontal lines in the center-right plot show the effective value of the effective infinite-time prediction for the aging functions (see main text). The top one is calculated on the basin's $\psi_{\text{gauss}}(\tau)$, the lower one is on the configuration $\psi_{\text{gauss}}(\tau)$. The horizontal lines on the bottom-right plot represent a confidence interval for the same quantity, averaged between basins and configurations.

configuration aging function as a finite time effect: the lower the temperature, the larger the time needed to enter in the trap-like dynamics regime where the energy is intensively lower than E_{th} and a basin is really formed by just one configuration. Accordingly, we should observe for very large system sizes (for which the plateau of the aging function extends to very large time scales) a slow upwards drift of the plateau of the configuration aging function towards

the plateau of the basin aging function. A hint of this upward drift seems to appear in the long time limit of aging function obtained at the lowest observed temperature $T = 0.25$ and for the largest system size considered in figure 12.

4.3.2 Aging functions in the REM

In the case of the REM, the interpretation of the data is less straightforward, as it is seen in figure 12 (right), because the plateau regime is not well-defined. This is due to the fact that in the REM the trap-like dynamics is visible only if the exponentially diverging time-scales are well separated. This would only happen in the right scaling limit of infinitely large systems and times. We can try nevertheless to extract an effective aging function to compare with. Equation (9) can be seen as a power law with an exponent that depends on the simulation time: according to the length of the run, one can extract an effective exponent $x_{\text{eff}}(t) = \frac{T}{T_c^{\text{eff}}(t)}$.

Two equivalent ways can be used to extract it. The first method is to fit to a power law the final part of the curve $\psi(\log(\tau))$. By comparison with equation (5), $x_{\text{eff}}(t)$ is the exponent of the curve. A second way is to use the relation, valid in the asymptotic limit, $x_{\text{eff}}(t) = -\frac{TE(t)}{N}$ that only requires the knowledge of the average energy $E(t)$ reached in the simulations at time t .

The two methods of extracting the effective asymptotic values of the aging functions give compatible values, that are plotted in figure 12 (right), and show that numerical results are not incompatible with a trap behaviour. Further, as stated in section 4.2, at the lowest temperature, $T = 0.25$, the simulation time was too small to sample well the phase space. As a consequence, we could not extract an effective exponent x_{eff} . Clearly much larger size and times would be needed to enter the asymptotic regime. Remarkably, a recent numerical study of generalised trap models [28] has shown that the very same problem also arises in simpler models with a gaussian distribution of energies. The analysis developed in [28] could be applied to the REM case in order to find more stringent evidences of trap dynamics.

Finally in agreement with the results for the EREM, we find also in this case a more pronounced difference at low temperature between basin and configuration descriptions.

5 Conclusions

We have shown how the main features of TM-like dynamics emerge effectively on exponentially diverging time-scales in the case of the simplest mean-field glassy models (REM and EREM). Despite motion in configuration space is complex on intermediate time-scales, in particular at the threshold level and above, with frequent back and forth movements in configuration space, the very large separation of time-scales washes out correlations induced by the exploration of the landscape, and hence memory. As a result, by coarse-graining dynamical evolution in terms of basins, a full TM description of the dynamics holds, in agreement with very recent rigorous results [20]. Identifying traps with basins instead of configurations is, as we have seen, important both numerically, to reduce pre-asymptotic corrections, and theoretically (see [20]). Only in this way a complete TM description can be reached. This will be even more cru-

cial in studies of more complex and rich models, for which our results provide guidelines.

The natural next step is to study the p -spin model, that in the large p limit converges to the REM [18, 34]. In our study we found that in the activated dynamics regime the system has always to reach and wander at the threshold level in order to escape from a basin and loose memory of the past. Whether this holds in general is a very interesting question which would shed light on whether an effective TM description is valid more generally. An interesting possibility is that to escape from a deep minimum the system only needs to reach an energy level which is lower than the threshold, which is identified as the lowest energy reached during aging dynamics on time-scales of order one. This would be an important first piece of information to understand activated dynamics in realistic system such as super-cooled liquids whose energy landscape displays strong similarities with the one of mean-field glassy systems [3].

These questions can be addressed applying our approach to p -spin or more general models. Work is in progress in this direction [35].

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