A precise approximation for directed percolation in \( d = 1 + 1 \)

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Abstract

We introduce an approximation specific to a continuous model for directed percolation, which is strictly equivalent to \( 1 + 1 \) dimensional directed bond percolation. We find that the critical exponent associated to the order parameter (percolation probability) is \( \beta = \frac{1}{2} \left( 1 - \frac{1}{d} \right) = 0.276393202 \ldots \), in remarkable agreement with the best current numerical estimate \( \beta = 0.276486(8) \).

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Directed percolation (DP) is a useful paradigm for dynamical phase transitions between an active/spreading phase and an extinct/absorbing phase. Models in the DP class of universality are involved in the description of catalytic reactions, surface dynamics, porous systems, granular media, epidemics, Calcium dynamics in cells, developed turbulence and coupled maps. Recently, H. Hinrichsen summarized the large scope of possible physical applications of DP, which led P. Grassberger to conjecture that the DP universality class should describe any continuous phase transition from a fluctuating active phase into a single absorbing phase, in the absence of quenched disorder and special symmetries. In a sense, DP plays a similar role in the study of dynamical phase transitions as the Ising model for continuous equilibrium phase transitions.

Despite its ubiquity, DP is maybe the only major statistical physics model which has not yet been successfully solved in one spatial dimension (+ time), probably due to its lack of conformal invariance.

Let us recall the original model of directed bond percolation in \( d = 1 + 1 \), describing the propagation of a fluid in a 2d porous medium. On a square lattice tilted at \( 45^\circ \), a fraction of bonds \( p \) are chosen at random to be active, whereas the remaining bonds stay inactive or broken. The “fluid” starts from the top row, and propagates downward, only passing through the active bonds. One then defines the order parameter \( n(p,t) \), which measures the average density of occupied sites at row \( t \). \( n(p,t) \) happens to coincide with the probability that at least one site at row \( t \) is still active (percolation probability).

In the stationary limit \( t \to +\infty \), the order parameter tends to a constant value \( n(p) \), which is zero below \( p_c = 0.644700185(5) \), and behaves as \( n(p) \sim (p-p_c)^\beta \) near \( p_c \). This defines the universal critical exponent \( \beta = 0.276486(8) \). Defining \( n_i(t) = 1 \) (resp. \( n_i(t) = 0 \)), when the \( i \)th site at row \( t \) is active/occupied (resp. inactive/empty), \( n_i(t) \) satisfies the following recursion relation

\[
n_i(t+1) = a_i n_i(t) + b_i n_{i+1}(t) - a_i b_i n_i(t) n_{i+1}(t), \quad (1)
\]

where \( a_i(t) \) and \( b_i(t) \) are independent random variables taking the value 1 with probability \( p \) (if the corresponding ongoing bond is active), and 0 otherwise.

In relation to self-organized criticality (SOC), it has been recognized that directed bond percolation is strictly equivalent to a continuous dynamical model (SOCDP), involving no external parameter (like \( p \) in DP) [12–15].

On a 1d lattice, we define the continuous variables \( x_i(t) \) as satisfying the recursion relation

\[
x_i(t+1) = \min \left[ \max(x_i(t), z_i(t)), \max(x_{i+1}(t), z_i(t)) \right], \quad (2)
\]

where \( z_i(t) \) and \( z_i(t) \) are independent random variables uniformly distributed between 0 and 1. It can be easily shown that the \( n_i \)'s for directed bond percolation and the \( x_i \)'s are very simply related:

\[
n_i(t) = \theta(x_i(t) - p), \quad (3)
\]

where \( \theta(\cdot) \) is the usual Heaviside step function. In the large time limit, the \( x_i \)'s are distributed according to a stationary probability distribution \( \rho(x) \), and

\[
n(p) = \int_0^p \rho(x) \, dx. \quad (4)
\]

Now, using Eq. (1) (or equivalently Eq. (2)), we find in the stationary limit,

\[
\frac{2p - 1}{p^2} n(p) = \frac{2p - 1}{p^2} n_1 = \langle n_1 n_2 \rangle,
\]

\[
= \int_0^p \int_0^p \rho_2(x_1, x_2) \, dx_1 \, dx_2, \quad (5)
\]

where \( \rho_2(x_1, x_2) \) is the nearest neighbor correlation function of the \( x_i \)'s. In mean field (MF) theory, one makes the approximation \( \langle n_1 n_2 \rangle \approx \langle n_1 \rangle^2 \), leading to

\[
n_{MF}(p) = \frac{2p - 1}{p^2}, \quad \rho_{MF}(p) = \frac{2(1-p)}{p^2}. \quad (7)
\]

From now, we study the stationary state of DP in terms of the continuous model defined by Eq. (3). We first notice that

\[
x_1(t+1) = \min[x_1(t), x_2(t)], \quad \text{with probability} \quad p_{\text{min}} = \min[x_1(t), x_2(t)],
\]

\[
x_1(t+1) = \max[x_1(t), x_2(t)], \quad \text{with probability} \quad p_{\text{max}} = \max[x_1(t), x_2(t)][1 - \max[x_1(t), x_2(t)]],
\]
Hence, this prompts the introduction of the key approximate amplitude ratios between them, for all values of $p$ (at least for short range correlation functions like $\rho_1(x_1,x_2)$, noting that as $\rho(p)$ diverges near $p_c$ (since $\beta < 1$), $\rho(\min(x_1,x_2)) > \rho(\max(x_1,x_2))$, at least near $p_c$ (numerically $\rho(p)$ appears to be a strictly decreasing function, like in mean field theory). We expect that $f(x_1,x_2)$ is a smooth function of order unity. Indeed, contrary to the MF approach (where one assumes that $(n_1n_2) \sim [n(p)]^2$), correlation functions all behave as $n(p)$ near $p_c$. Indeed, as $p_c > 1/2$, Eq. (3) implies that $(n_1n_2) \sim n(p) \sim (p-p_c)^\beta$, such that $\rho_2(p,p) \sim \rho(p) \sim (p-p_c)^{-(1-\beta)}$ (instead of $\rho(p)^2 \sim (p-p_c)^{-2(1-\beta)}$, predicted by MF). A natural guess for $f(x_1,x_2)$ is provided by the general statement that although MF is inept at describing correlation functions near $p_c$, it still leads to reasonably accurate amplitude ratios between them, for all values of $p$ (at least for short range correlation functions like $(n_1n_2)$). Hence, this prompts the introduction of the key approximation

\[
f(x_1,x_2) \approx f_{MF}(x_1,x_2),
\]

\[
= \frac{\rho_{MF}(x_1)\rho_{MF}(x_2)}{\rho_{MF}(\min(x_1,x_2))},
\]

\[
= \rho_{MF}(\max(x_1,x_2)).
\]

In the following, we make the more general ansatz

\[
\hat{\rho}_2(x_1,x_2) = \rho(\min(x_1,x_2))f(\max(x_1,x_2)),
\]

where $f(p)$ is not necessarily equal to $\rho_{MF}(p)$, $\rho(p)$, $f(p)$ and $g(p)$ are not independent functions as they are related together by Eq. (3), and by the probability conservation constraint,

\[
\rho(p) = \int_0^1 \rho_2(x,p) \, dx.
\]

From Eq. (3) and Eq. (14), and after straightforward calculations, we obtain the two relations

\[
n(p) = \exp \left[ -\int_p^1 \frac{2f(x) - \rho_{MF}(x)}{n_{MF}(x) - g(x)} \, dx \right],
\]

\[
g(p) + \int_p^1 f(x) \, dx + \frac{n(p)}{\rho(p)} = 1.
\]

From Eq. (17) and Eq. (18), one can obtain a first order differential equation for $F(p) = f(p) - \rho_{MF}(p)$, involving only $p$ and $g(p)$. This equation can be shown to have only $F \equiv 0$ as a global solution satisfying the boundary conditions and the physical constraints. Hence, and in complete accordance with the physical argument given in Eq. (14), we find

\[
f(p) = \rho_{MF}(p).
\]

Quite remarkably, for this precise form for $f(p)$, Eq. (18) is now satisfied for any choice of $g(p)$, so that we are left with Eq. (17) as the only non trivial relation between $n(p)$ and $g(p)$.

Now, as $n(p)$ vanishes at $p_c$, we expect the function involved in the integral of Eq. (17) to develop a single pole at $p_c$, of residue $\beta$, so that $n(p) \sim (p-p_c)^\beta$, near $p_c$. This leads to

\[
g(p_c) = n_{MF}(p_c) = \frac{2p_c - 1}{p_c^\beta},
\]

\[
\beta = \left( 1 - \frac{g(p_c)}{\rho_{MF}(p_c)} \right)^{-1}.
\]

Note that Eq. (24) is in fact an exact identity, which does not rely on the present approximation on $\hat{\rho}_2(x,y)$. In order to achieve our goal of computing $n(p)$, we need a further relation for $g(p)$. This can be obtained by writing the exact stationary equation for $g(p)$. We first define (\bullet\bullet\bullet) as the probability of having three consecutive sites with $x_i = p$, divided by $\rho(p)$ (that is, conditional to having one site with $x_i = p$). In the same manner, we define (\bullet\bullet\bullet) (resp. (\bullet\bullet\bullet)) as the probability of having two consecutive sites with $x_1 = x_2 = p$, and $x_3 > p$ (resp. $x_3 < p$), divided by $\rho(p)$. Finally, in the stationary limit, we obtain the exact relation

\[
g(p) = (2p - p_c)^2(\bullet\bullet\bullet)
\]

\[
+ 2(2p - p_c^2) \left( p(\bullet\bullet\bullet) + p(1-p)(\bullet\bullet\bullet) \right)
\]

\[
+ p^2(\bullet\bullet\bullet) + p^2(1-p)(\bullet\bullet\bullet) + p^2(1-p^2)(\bullet\bullet\bullet)
\]

\[
\quad = (\bullet\bullet\bullet). \tag{22}
\]

For instance, the first term $(2p - p_c)^2(\bullet\bullet\bullet)$ represents the fact that a configuration \bullet\bullet\bullet $(x_1(t+1) = x_2(t+1) = p)$ at time $t+1$ can arise from a configuration \bullet\bullet\bullet $(x_1(t) = x_2(t) = x_3(t) = p)$ at time $t$, provided that $x_1$ and $x_2$ are preserved by the transformation of Eq. (3). This happens with probability

\[
(p_{\min} + p_{\max})^2 = (p + (1-p))^2 = (2p - p_c)^2, \tag{23}
\]

hence the coefficient in Eq. (22) ($p_{\min}$ and $p_{\max}$ have been defined in Eq. (3) and Eq. (4)).

Eq. (22) relates $g(p)$ to three-point correlation functions, and cannot be exploited unless an additional approximation is introduced. We will factor these three-point correlation functions into products of two-point correlation functions, according to the usual mean field.
scheme. Introducing \( p_+ \) as the probability that \( x_2 > p \), conditional to the fact that \( x_1 = p \), we obtain
\[
[1 - g(p)]p_+ = \int_p^1 \frac{\tilde{\rho}_2(x, p)}{\rho(p)} \, dx,
\]
(24)
\[
= \int_p^1 \frac{\rho pMF(x) \, dx}{(1-p)^2},
\]
(25)
where Eq. (24) is an exact identity. We give below a few examples of three-point correlation functions computed according to this MF factorization scheme:
\[
\langle \bullet \bullet \bullet \rangle = g(p)^2,
\]
(26)
\[
\langle \bullet \bullet \hat{p} \rangle = g(p)[1 - g(p)]p_+ = g(p) \frac{(1-p)^2}{p^2},
\]
(27)
\[
\langle p \bullet \hat{p} \rangle = [1 - g(p)]^2 p_+ (1 - p_+),
\]
(28)
\[
= \left[ \frac{2p-1}{p^2} - g(p) \right] \frac{(1-p)^2}{p^2}.
\]
(29)
Inserting the MF form for the three-point correlation functions into Eq. (22), we finally obtain a closed equation for \( g(p) \),
\[
g(p) = (1 - p + pg(p))^2 + (1 - p)^2 g(p)(1 - g(p)),
\]
(30)
which can be readily solved, leading to
\[
g(p) = \frac{(1-p)^2}{2p-1}.
\]
(31)
\[
\text{FIG. 1. We plot } \langle \bullet \bullet \bullet \rangle \text{ (two top lines) and } \langle \bullet \bullet \hat{p} \rangle \text{ as obtained from numerical simulations (full lines), and as given by Eq. (24) and Eq. (27) (dashed lines), where the numerical value of } g(p) \text{ has been inserted in these expressions. Note that } \langle \bullet \bullet \bullet \rangle \approx g(p)^2, \text{ especially near } p_c. \text{ These functions all vanish as } (1-p)^4 \text{ near } p = 1, \text{ as predicted by Eq. (24) and Eq. (27). Insert: comparison between the numerical } g(p) \text{ and the present theory. In all figures of this letter, we have simulated a system of } N = 300000 \text{ sites, averaged over 100 samples. Physical quantities in the stationary state have been estimated by averaging them between } t = 300000 \text{ and } t = 310000.\]

\[
\text{FIG. 2. We respectively plot } n(p) \text{ as given by mean field theory (dashed line), the present theory (dotted line), and numerical simulations (full line). Note that the three curves coincide near } p = 1, \text{ as mean field theory becomes exact in this limit. A fit of the numerical data to the functional form of Eq. (24), where } \tau \text{ becomes a fitting parameter, cannot be distinguished from the actual data.}\]

The fact that the relation \( \langle \bullet \bullet \bullet \rangle = g(p)^2 \) seems to be exactly satisfied numerically at \( p_c \) could explain this agreement, which also implies that the MF factorization of Eq. (23) is quantitatively correct near \( p_c \) (the three-point correlation functions appearing in Eq. (23) are not independent and are related to \( \langle \bullet \bullet \bullet \rangle \) by various sum rules). This is illustrated in Fig. 1, where the exact numerical \( g(p) = \langle \bullet \bullet \rangle, \langle \bullet \bullet \hat{p} \rangle, \text{ and } \langle \bullet \bullet \hat{\sigma} \rangle \) are plotted with their theoretical counterpart.

Now, \( f(p) \) and \( g(p) \) being known, the percolation probability can be easily computed by using Eq. (17):
\[
n(p) = p^{-2} \frac{(x - \tau)(2 + \tau - x)}{\tau}^{\beta}
\]
(34)
\[
\times \left[ \frac{(x - 1 + \tau)(x + 1 + \tau)}{1 + \tau} \right]^{1-\beta}.
\]
In Fig. 2, we compare this result with the numerically extrapolated stationary percolation probability, and to the MF result of Eq. (6).

Finally, in Fig. 3, in order to test the validity of our basic approximation Eq. (14), we plot $f(x_1, x_2)$ (defined in Eq. (11)) as a function of $\max(x_1, x_2)$. We find that this scatter plot is reasonably aligned around an effective curve, and that $f_{MF}(x_1, x_2) = \rho_{MF}(\max(x_1, x_2))$ appears to be a lower bound for the actual $f(x_1, x_2)$.

In conclusion, we have introduced a new approximation for a continuous model equivalent to directed bond percolation. In this language, this approximation amounts to properly modelizing the correlation function $\rho_2(p_1, p_2)$, relating the properties of directed bond percolation for two different percolation parameters $p_1$ and $p_2$. By assuming that amplitude ratios are correctly described by mean field theory, we end up with a precise description of the percolation probability. In particular, we find an exponent $\beta$ in remarkable agreement with the best available numerical simulations.

It would be interesting to exploit the present approach in order to describe the dynamical properties of DP. This study is currently in progress.

This approach could also prove useful in tackling the notably difficult problem of parity conserving branching annihilating walks [3]. This universality class is exemplified by the reaction-diffusion model of diffusing particles $A$, involving annihilation ($A + A \rightarrow \emptyset$) and branching ($A \rightarrow A + A + A$) processes. This problem has so far eluded all manner of theoretical approaches in $d = 1 + 1$.

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