

Theoretical and numerical studies of chemisorption on a line with precursor layer diffusion

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Abstract

We consider a model for random deposition of monomers on a line with extrinsic precursor states. As the adsorbate coverage increases, the system develops non-trivial correlations due to the diffusion mediated deposition mechanism. In a numeric simulation, we study various quantities describing the evolution of the island structure. We propose a simple, self-consistent theory which incorporates pair correlations. The results for the correlations, island density number, average island size and probabilities of island nucleation, growth and coagulation show good agreement with the simulation data.

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1 Introduction

Models of *random sequential adsorption* (RSA) have been used to describe the process by which particles are irreversibly deposited without overlap onto a surface. They are relevant to studying the adsorption of gas molecules [1] or colloid particles in solution [2] onto solid surfaces, or of large molecules on biological membranes [3]. One of the key assumptions, is that the particles bind strongly to the substrate so that desorption and surface mobility are negligible on the time-scale of the experiment. Although providing a very simplified picture, RSA models have the virtue of being exactly solvable in dimension $d = 1$, for monomer and for k -mer particles [4].

Many additional features have been added to this model in order to make it more physical, or to generalize its range of applications, and the literature in the field is vast [5]. The inclusion of diffusional relaxation on the substrate [6, 7], leading to equilibration, or of cooperative effects, such as multi-site exclusion [8], have been examined. The possibilities that the particles are reflected back to the fluid or desorbed [7, 9, 10], or that the surface comprises more than one chemical species [11] have also been considered.

Another important possibility, motivated by many experimental studies of gas-metal surfaces [12, 13], is that if the particles lose just enough of their kinetic energy they can become trapped (physisorbed) in a mobile, temporary *precursor state*, from which they can be adsorbed (chemisorbed) at a far site at a later time. The mechanism of precursor mediated chemisorption, first postulated by Taylor and Langmuir [14], was initially formulated as a statistical model by Kisliuk [15]. It was later adapted by King and coauthors [10, 13, 16, 1] to include other effects, such as temperature dependence, desorption, molecular dissociation and pair interactions between the adatoms. Variations on the Kisliuk model have also been studied by other authors, both numerically and analytically [9, 11, 17, 18, 19, 20, 22]. Most analytical treatments, however, are essentially mean-field like, in that they largely ignore the correlations between the state of occupation of different sites which arise through the precursor mediated deposition process, i.e. they assume that the rate of deposition at a given site only depends on the state of occupation of that site, and not on the neighbouring sites.

In this paper, we consider a lattice model which is both a simplified version of the Kisliuk model, and a slight complication of the RSA model. Contrary to most previous studies, we examine the case where the mobility of physisorbed atoms is only possible on the top of occupied chemisorption sites (*extrinsic* precursor states), and where there is neither reflection nor desorption back to the gas phase (later we shall comment on how scattering and desorption can in principle be included into

the equations of motion). Each deposition attempt will, therefore, result either in the direct occupation of an empty site, or in the diffusion on the top of an island of occupied sites until the particle finds an empty site at the edge of the island, where it is irreversibly deposited. For simplicity, we also consider that the deposited particles are *monomers*, or atoms, in which case the lattice will eventually become full. As a consequence of the (extrinsic) precursor diffusion, the edges of islands are preferential sites for chemisorption and the growth of the larger islands is favoured. This introduces non-trivial correlations between sites, especially at late-times when diffusion is the dominant deposition mechanism, and makes the model unsolvable even for the simplest case of monomer deposition on a *line*. A model similar to the one adopted here has been examined by Becker and Ben-Shaul in $2d$ [19]. Their analysis, however, only applies to the early kinetics, as they treat the islands as uncorrelated objects, completely neglecting the contribution of island coalescence to the growth process.

Regarding the motivation for our study, we recall a somewhat unrealistic feature of the RSA model, namely that once a deposition attempt fails, the particle's position is 'randomized' before another adsorption attempt is made. As the transport of particles to the surface is diffusive, one might expect that a failed deposition attempt would be followed by another nearby adsorption attempt [21]. Since the intrinsic precursor seems to play no role in this effect, the present model may account for it, at least partially. Furthermore, it has been previously suggested by Cassuto and King [16], that a kinetic deposition model without intrinsic precursor states and with negligible desorption, could well also explain the experimental data from some gas-solid systems, such as hydrogen on tungsten. Naively, one may also think that such a model could be able describe the slow deposition of liquid droplets on a plane surface, the spread of epidemics from immunized to non-immunized populations, or the growth of a forest where seeds have to be transported to an open field to find suitable conditions to develop.

Although island formation and structure kinetics have been studied (see e.g. [8, 22]), most work on RSA and precursor models, either analytical or simulation, has focused on determining the dependence of the coverage fraction θ and sticking probability on the exposure, i.e. on the time t , and its value at the *jamming limit* (which is trivial only for monomers). In the present model, we assume that the diffusion time scale is small enough compared to the time scale of deposition attempts, and so the coverage is proportional to time or, equivalently, the sticking probability is 1. We examine the case of a substrate of dimension $d = 1$, and concentrate on studying the quantities describing the evolving morphology of the occupied regions. Namely, the pair correlation functions, the total number of islands, and the prob-

ability distribution of the island sizes and its moments. We have measured these quantities in numerical simulations, and developed a minimal theory incorporating correlations, to calculate them approximately. The results show good agreement, while at the same time differing considerably from those of the RSA model for monomer deposition in $1d$ where there are no spatial correlations at all.

A brief summary of this paper is the following. In section 2 we describe the model and define the formalism. The results of the numerical simulation are discussed in section 3. The theory is presented in section 4, and the results are compared with the simulation data. Section 5 is dedicated to some conclusions.

2 Model

We consider a one dimensional lattice with N sites, with N large enough to neglect boundary effects, i.e. we take $N \rightarrow \infty$ in the calculations. At each site i we define a variable S_i , such that

$$\begin{aligned} S_i &= 0 && \text{if the site is empty} \\ &= 1 && \text{if the site is occupied .} \end{aligned}$$

Every time step $\delta t = 1$ a site is chosen randomly and a monomer deposition attempt is made. If the site is empty the particle is adsorbed and the site becomes occupied irreversibly. Otherwise, the particle diffuses on top of the occupied region until it reaches an empty site where it is adsorbed (this is appropriately modelled by a random walk with traps [23]). Only then is the next deposition attempted. The process repeats until the lattice is full.

We assume here that the particle diffusion is rapid enough to be over before the time of the next deposition attempt, independently of the size of the island where it takes place. This may sound as an unrealistic assumption, especially if the island is of the order of the system size. We note, however, that due to the randomness of the process it is unlikely that a diffusing particle would interact or compete with the next particle to be deposited. This argument fails, of course, in the limit when the lattice is almost full (intermediate coagulation may occur) or in the case when diffusion is very slow (a gas of net precursor particles develops).

Since each time step results in a deposition, either direct or mediated by diffusion, the time dependence of the adsorbate coverage, $\theta = \langle S_i \rangle$, is trivial in this case:

$$\theta(t) = t/N \quad , \quad 0 \leq t \leq N \quad . \quad (1)$$

One can easily incorporate in the model the possibility of *scattering* of a particle back into the gas phase at the instant of collision with the surface. If the scattering probability is $(1 - \alpha)$, whatever the state of occupation of the site, the rate of

adsorption per unit time must be multiplied by α . We may then set $\alpha = 1$ through a redefinition of the time scale. To account for the possibility that scattering may depend on the state of occupation of the site, we set the probability of no chemisorption to be $(1 - D\alpha)$ if a particle lands on an occupied site. In this case, the rate of variation of the number of occupied sites is (setting $\alpha = 1$)

$$Nd\theta/dt = 1(1 - \theta) + D\theta = 1 - (1 - D)\theta , \quad (2)$$

with initial condition $\theta(0) = 0$. The solution is

$$\theta(t) = \frac{1 - \exp[-(1 - D)t/N]}{1 - D} . \quad (3)$$

This reduces to (1) for $D = 1$, and to the RSA behaviour for $D = 0$ and for $t \ll N$. When $0 < D < 1$ the full coverage $\theta = 1$ is attained at $t = N \log(1/D)/(1 - D)$, larger than N , due to the larger scattering from the occupied regions. We may also account for *desorption*, simply by adding an extra negative term (linear in θ) to the rate equation (2), effectively decreasing the value of D . The coefficient of the term may incorporate the probability of desorption from the physisorbed or from the chemisorbed states, or both. Allowing for desorption from the chemisorbed states would not only affect the coverage, however, but will also imply extra terms in the rates of growth and coalescence (eq. (17)). For convenience we shall keep $D \neq 1$ in the future expressions, as it allows us to distinguish the diffusion from the direct deposition terms.

Next we define several quantities which will be useful to examine the morphology of the system. We shall call:

- $N(t)$ the total number of islands of occupied sites
- $N(L) = N(0L0)$ the number of islands with L sites, or number of rows of L consecutive occupied sites with at least one empty site on the left and right
- $N(L00) = N(0L00)$ the number of islands of L sites with at least two empty sites on the right
- $N(L0L') = N(0L0L'0)$ the number of islands of L sites separated by an empty site from an island of L' sites on the right
- $N(00)$ the number of pairs of empty sites

- $N(000)$ the number of trios of empty sites .

The generalization of the notation for more complicated configurations is obvious. The densities n associated with the above numbers, are defined by their ratio to the system size, e.g. $n(t) = N(t)/N$. Since the boundary effects can be neglected, the number of occupied regions (islands) is equal to the number of empty regions. On average the system has left and right symmetry, so $N(L00) = N(00L)$ and $N(L0L') = N(L'0L)$.

For a $1d$ substrate, the above quantities are easily defined in terms of the correlation functions of the local variables. Although the system is not in equilibrium we expect it to be translationally invariant, on average, due to the randomness of the deposition attempts. Using this property we have, for example:

$$n(t) = \sum_i \langle (1 - S_i) S_{i+1} \rangle / N = \theta - \langle S_i S_{i+1} \rangle \quad (4)$$

$$\begin{aligned} n(L) &= \sum_i \langle (1 - S_i) S_{i+1} \dots S_{i+L} (1 - S_{i+L+1}) \rangle / N \\ &= \langle (1 - S_0) S_1 \dots S_L (1 - S_{L+1}) \rangle \end{aligned} \quad (5)$$

$$\begin{aligned} n(L00) &= \sum_i \langle (1 - S_i) S_{i+1} \dots S_{i+L} (1 - S_{i+L+1}) (1 - S_{i+L+2}) \rangle / N \\ &= \langle (1 - S_0) S_1 \dots S_L (1 - S_{L+1}) (1 - S_{L+2}) \rangle \end{aligned} \quad (6)$$

$$n(00) = \sum_i \langle (1 - S_i) (1 - S_{i+1}) \rangle / N = 1 - 2\theta + \langle S_0 S_1 \rangle \quad (7)$$

$$n(000) + n(1) = \sum_i \langle (1 - S_i) (1 - S_{i+2}) \rangle / N = 1 - 2\theta + \langle S_0 S_2 \rangle \quad (8)$$

A number of normalization or hierarchical sum rules follow, some of which are:

$$N(t) = \sum_{L=1}^{\infty} N(L) \quad (9)$$

$$N(L) = \sum_{L'=0}^{\infty} N(L0L') = \sum_{L'=1}^{\infty} N(L0L') + N(L00) \quad (10)$$

$$N(00) = \sum_{L=0}^{\infty} N(L00) = \sum_{L=1}^{\infty} N(L00) + N(000) \quad (11)$$

$$N(0) = (1 - \theta)N = \sum_{L=0}^{\infty} N(0L0) = N(t) + N(00) \quad (12)$$

$$\theta N = \sum_{L=1}^{\infty} LN(L) . \quad (13)$$

It is easy to show that expressions (4) and (9) (with $n(L)$ replaced by (5)) are equivalent, using fixed or periodic boundary condition and the property $S_i^2 = S_i$. The probability of finding an island of size L and its moments, the average island size and its mean square deviation, are therefore:

$$P(L) = N(L)/N(t) \quad (14)$$

$$\langle L \rangle = \sum_{L=1}^{\infty} LP(L) = \theta/n(t) \quad (15)$$

$$\frac{\langle L^2 \rangle - \langle L \rangle^2}{\langle L^2 \rangle} = 1 - \theta^2 / \left(n(t) \sum_{L=1}^{\infty} L^2 n(L) \right) . \quad (16)$$

There are three basic mechanisms by which these numbers may change with the deposition process (where Δ denotes variation):

- *nucleation*: $\Delta N(000) = 1$
- *growth*: $\Delta N(L00) = -1$; $\Delta N(L+1) = 1$
- *coagulation*: $\Delta N(L0L') = -1$; $\Delta N(L+1+L') = 1$.

The rates of occurrence of each of these events per unit time are:

$$\begin{aligned} R_n &= \frac{N(000)}{N} = n(000) \\ R_g &= \sum_{L=1}^{\infty} \frac{1+DL/2}{N} [N(L00) + N(00L)] \\ &= 2(n(00) - n(000)) + D \sum_{L=1}^{\infty} Ln(L00) \\ R_c &= \sum_{L=1}^{\infty} \sum_{L'=1}^{\infty} \frac{1+D(L+L')/2}{N} N(L0L') \\ &= n(t) - n(00) + n(000) + D\theta - D \sum_{L=1}^{\infty} Ln(L00) \\ &= 1 - (1-D)\theta - 2n(00) + n(000) - D \sum_{L=1}^{\infty} Ln(L00) , \end{aligned} \quad (17)$$

where we have used (9)-(13). The factor $L/2$ accounts for a particle landing and diffusing on top of an island of size L towards the right (or left). For simplicity, we have assumed, and we will assume throughout, that in modeling the diffusion process the random walk with traps can be replaced by a random choice between right and left. We shall return to this point in section 3. As expected, the sum of the rates (17) yields the total rate of adsorption per unit time: $\sum_i R_i = 1 - (1-D)\theta = Nd\theta/dt$. The probabilities of nucleation, growth and coagulation per deposition event, P_n , P_g and P_c , are then defined by the ratios:

$$P_i = R_i / (Nd\theta/dt) . \quad (18)$$

It is then straightforward to write the exact equation for the rate of variation of the number of islands, given by the difference between the rates of nucleation and

coagulation:

$$\begin{aligned} \frac{dN(t)}{dt} &= R_n - R_c \\ &= 1 - (D + 1)\theta - 2n(t) + D \sum_{L=1}^{\infty} L n(L0) , \end{aligned} \quad (19)$$

where we have used (12). Setting $D = 0$ in (19) one obtains the RSA result $n(t) = \theta(1 - \theta)$. Using (15) and (2), it is possible to derive from (19) an exact equation for the average island size $\langle L \rangle$. It is also illustrative to look at the growth of a single, isolated island. Neglecting coagulation, the equation for the island size reads: $dL/dt = (2 + DL)/N$, which solution (with $L(0) = 0$) is $L(t) = (2/D)(\exp(Dt/N) - 1)$. This, however, only gives the expected growth at early times. When island coagulation becomes important the growth of L should be faster than exponential, and as $t \rightarrow N$ (for $D = 1$) L should become of order N .

3 Simulation

We performed numerical simulations of the model described in section 2. For convenience we used free boundary conditions, although the choice of boundary conditions should be irrelevant. The results were averaged over an ensemble comprising a great number (N_s) of system samples with different random deposition histories. For large coverages, the distribution $P(L)$ and its second moment proved particularly sensitive to finite sampling and size effects. This is easily understood, as the size of the larger island at late-times (which controls the dynamics at this stage) can fluctuate by as much as $N/2$. The smoothness of the curves for the probabilities P_n , P_g and P_c also depends on the number of samples used. We used $N = N_s = 50000$, and verified that the systematic finite sampling and size errors were almost eliminated as we increased N and N_s up to these values. For ease of analytical treatment (eq. 17) and for computational efficiency, we modelled the precursor layer diffusion with a random choice between right and left rather than with a random walk with traps. Although that mechanism does not take into account the starting point of the diffusion process, we expect this effect to be irrelevant because on average all sites on an island are equally likely to be chosen for a deposition attempt.

3.1 Results

The measurements of $n(t)$, $1/\langle L \rangle$, $(\langle L^2 \rangle - \langle L \rangle^2)/\langle L^2 \rangle$ and $P(L)$ (at six different coverages) are displayed in Figures 1 to 4 (the pair correlations are shown in section 4.1). We have divided the second moment by $\langle L^2 \rangle$ rather than $\langle L \rangle^2$ since the latter diverges.

It is interesting to observe the extent to which the RSA kinetics of deposition is modified by the additional diffusion mechanism. For comparison, we included in Figures 1 to 3 the plots (broken lines) of: $\theta(1 - \theta)$, $(1 - \theta)$ and $\theta/(1 + \theta)$, and in Figure 5 the plot of $\theta^{L-1}(1 - \theta)$, which correspond to the same quantities in the RSA model. We see from Figures 1 to 3, that the number of islands is smaller, and the average island size and island size fluctuations are larger with diffusion mediated deposition. This results from the increase in the rates of island growth and coagulation relative to the rate of island nucleation. Comparing Figures 4 and 5 for the island size distribution, we can see a close agreement at small coverages, when all islands are still small. At intermediate coverages, the distribution spreads out (larger islands) in the diffusion case. At large coverages the difference is even greater. The RSA curve becomes uniform (all island sizes are equally probable), while the diffusion curve shows that the majority of islands are still small in size, but there is a minority of very large islands. At the late stages of diffusion mediated growth, the number of large islands is small and their size can fluctuate enormously, therefore the smoothness of the $P(L)$ curve depends strongly on the sampling number.

We have also examined the mechanisms responsible for the structure kinetics. Figure 6 shows the evolution of the probabilities of nucleation, growth and coagulation per deposition event, P_n , P_g and P_c (eq. (18)). For comparison we have included the corresponding RSA curves (broken lines): $P_{n,RSA} = (1 - \theta)^2$, $P_{g,RSA} = 2\theta(1 - \theta)$ and $P_{c,RSA} = \theta^2$ (cf. (17)-(18) with $D = 0$). The most obvious difference occurs as $\theta \rightarrow 1$. In the RSA case the probability of growth vanishes. In the diffusion case, however, the probability of growth tends to a finite value $1 - P_c$, showing that a considerable number of gaps with two or more empty sites still exist at large coverages.

4 Theory

Due to the diffusion on top of the occupied regions, the probability of adsorption at an empty site depends on the state of occupation of its neighbouring sites and even of far located sites, if the site is at the edge of a large island. These correlations develop in the system as the coverage increases with time and diffusion mediated deposition becomes more likely. As a result, the density numbers defined in section 2 obey an infinite set of hierarchical coupled equations which cannot be solved exactly (an example of an equation in the top of the hierarchy is given by (19)). One must, therefore, look for approximate solutions by truncating the hierarchy with some closure scheme. The simplest approximation consists in incorporating only the *pair correlations* in the system, which are then determined self-consistently. We shall

see that, by carefully choosing the multi-site functions decoupling scheme, such a simple approach, despite neglecting multiple correlations, is capable of capturing many qualitative and quantitative features of the system's behaviour.

Let us denote the pair correlations between sites with different separations, as:

$$\begin{aligned} p &= q^{(1)} = \langle S_i S_{i+1} \rangle \\ q &= q^{(2)} = \langle S_i S_{i+2} \rangle \\ q^{(n)} &= \langle S_i S_{i+n} \rangle, \quad n \geq 1. \end{aligned} \quad (20)$$

We expect the correlations to decay with distance, i.e.:

$$p \geq q \geq q^{(3)} \geq \dots \geq \theta^2. \quad (21)$$

The quantities depending on the local variables of *two sites* only, can be expressed exactly in terms of the pair correlations. From (4)-(8) we have:

$$n(t) = \theta - p \quad (22)$$

$$n(00) = 1 - 2\theta + p \quad (23)$$

$$n(000) + n(1) = 1 - 2\theta + q, \quad (24)$$

and the average island size (15) is given by

$$\langle L \rangle = \frac{\theta}{\theta - p}, \quad (25)$$

To write the *multi-site* functions approximately, in terms of the pair correlations, we decouple the higher order correlators into a product chain of pair correlators, each associated with an adjacent bond, as follows:

$$\begin{aligned} \langle S_1 S_2 \dots S_n S_{n+m} \rangle &\simeq \frac{\langle S_1 S_2 \rangle \langle S_2 S_3 \rangle \dots \langle S_{n-1} S_n \rangle \langle S_n S_{n+m} \rangle}{\langle S_2 \rangle \dots \langle S_{n-1} \rangle \langle S_n \rangle} \\ &= \frac{p^{n-1} q^{(m)}}{\theta^{n-1}}, \quad (n \geq 2, m \geq 1). \end{aligned} \quad (26)$$

The normalization factors in the denominator assure that the RSA result is recovered in the decoupling limit. Then, from (5)-(8) we have:

$$n(L) \simeq \frac{p^{L-1}}{\theta^L} (\theta - p)^2, \quad (L \geq 1) \quad (27)$$

$$n(000) \simeq 1 - 3\theta + 2p + q - p^2/\theta. \quad (28)$$

$$n(L00) \simeq \frac{p^{L-1}}{\theta^{L+1}} (\theta - p) [\theta^2 - \theta(p + q) + p^2], \quad (L \geq 1) \quad (29)$$

Note that the limits of $(n(00) - n(L))$ and $(n(000) - n(L00))$ when $L \rightarrow 0$, which are $(p - \theta^2)/p$ and $(p - \theta^2 + \theta(q - p))/p$, respectively, although non-zero have a small

value. Using these expressions and (22)-(23), we can then write the deposition rates (17) as:

$$\begin{aligned}
R_n &= 1 - 3\theta + 2p + q - p^2/\theta \\
R_g &= (D + 2)(\theta - p) + 2(p^2/\theta - q) + D\theta\frac{p - q}{\theta - p} \\
R_c &= Dp + q - p^2/\theta - D\theta\frac{p - q}{\theta - p},
\end{aligned} \tag{30}$$

and the mean square deviation of the island sizes (16) as:

$$\frac{\langle L^2 \rangle - \langle L \rangle^2}{\langle L^2 \rangle} = \frac{p}{\theta + p}. \tag{31}$$

For the theory to be self-consistent, we must check that the expressions for the densities, although approximated are properly normalized. In fact, summing (27), with L from 1 to ∞ , gives $\theta - p$ (cf. (9) and (22)). Also, the sum $\sum_{L=1}^{\infty} Ln(L)$ gives θ (cf. (13)). Hence, the probability distribution for the island sizes (14), obtained by dividing (27) by $n(t)$,

$$P(L) = (p/\theta)^{L-1} (1 - p/\theta), \tag{32}$$

is normalized to 1. This is a geometric distribution, as in the RSA case, but with θ replaced by p/θ . Adding (28) and (29), with L from 1 to ∞ , gives $1 - 2\theta + p$ (cf. (23) and (11)). Other more complicated density numbers turn out to be consistently normalized too, as their sums yield the correct density number within the pair approximation. This is the case for $n(L0L')$, which satisfies the sum rule (10).

The next step of our approach is to determine the pair correlations self-consistently. There is an infinite hierarchy of coupled equations for the pair correlations, even within the pair approximation. The first two are the equations for p and q . We will now derive the exact form of these two equations.

The p equation follows immediately from equation (19) for $n(t)$, using (22) and (2). It is more instructive, however, to write it down by inspection. pN is the average number of pairs 11, of two neighbouring occupied sites, which increases by one with island growth and by two with coalescence. Hence, using (17), we have:

$$\begin{aligned}
N\frac{dp}{dt} &= 2R_c + R_g \\
&= 2(D + 1)\theta - 2p - D\sum_{L=1}^{\infty} Ln(L00).
\end{aligned} \tag{33}$$

Within the pair approximation, using (30) and (1), and setting $D = 1$, we obtain:

$$\frac{dp}{d\theta} = 3\theta - p - \theta\frac{p - q}{\theta - p}. \tag{34}$$

The exact equation for q , although more complicated can also be written down by inspection. qN is the average number of pairs 1 – 1, of two occupied sites separated by a site in any state. It can increase by 1 or 2 with island nucleation, growth or coalescence. A careful consideration of all possibilities leads to the following equation:

$$\begin{aligned}
N \frac{dq}{dt} &= 2 \sum_{LL'=1}^{\infty} \frac{1}{N} N(L00L') + 2 \sum_{LL'=2}^{\infty} \frac{1 + D(L + L')/2}{N} N(L0L') \\
&+ 2 \sum_{L=1, L'=2}^{\infty} \frac{1 + DL'/2}{N} [N(L00L') + N(L'00L)] \\
&+ \sum_{L=1}^{\infty} \left\{ \frac{1}{N} [N(L0000) + N(0000L)] + \frac{1 + D/2}{N} [N(L001) + N(100L)] \right\} \\
&+ \sum_{L=2}^{\infty} \left\{ \frac{1 + DL/2}{N} [N(L000 + N(000L))] + \frac{1 + D(1 + L)/2}{N} [N(L01) + N(10L)] \right\}. \tag{35}
\end{aligned}$$

Employing the hierarchical relations (9)-(12) and some obvious generalizations, this equation then simplifies to

$$N \frac{dq}{dt} = 2 - 2(1 - D)\theta - (2 + D)n(1) - 2n(000) - D \sum_{L=1}^{\infty} L [n(L000) + n(L010)]. \tag{36}$$

Evaluating the sum of $L[n(L000) + n(L010)]$ within the pair approximation, using (27) and (28) and putting $D = 1$, yields

$$\frac{dq}{d\theta} = 2(\theta + p - q) - p^2/\theta + \frac{\theta q^{(3)} - pq}{\theta - p}. \tag{37}$$

One can check that $n(L000)$ and $n(L010)$ are properly normalized within the pair approximation (their sums give $n(000)$ and $n(1)$, respectively). The approach also yields $\sum_{L=0}^{\infty} [n(L000) + n(L010)] = 1 - 2\theta + q$, which is the exact result (cf. (24)).

It is straightforward to derive the behaviour of p and q for small and for large coverage. As $\theta \rightarrow 0$ the system is RSA like, so p , q and $q^{(3)}$ should behave as θ^2 . Equations (34) and (37) reduce to $p' = 3\theta + O(\theta^2)$ and $q' = 2\theta + O(\theta^2)$, where primes indicate derivatives with respect to θ . Hence

$$p = 3/2 \theta^2 + O(\theta^3) \tag{38}$$

$$q = \theta^2 + O(\theta^3) \quad , \quad (\theta \rightarrow 0). \tag{39}$$

The factor 3/2 (confirmed by the simulations, Figure 7) shows that even in this regime there is an increase in the correlations relative to the RSA case. It results from the diffusion mediated growth of single site islands: with RSA there 2 possibilities for growth, and with diffusion there is a third one; therefore, there are 3/2 as many double site islands as in RSA. Let us now consider the limit when $\theta \rightarrow 1$. At

$\theta = 1$, (34) and (37) yield the equations: $(2 - p'(1))(1 - p'(1)) = p'(1) - q'(1)$ and $(2 - q'(1))(1 - p'(1)) = q'(1) - q^{(3)'}(1)$. The first equation and inequality (21) imply that $1 \leq p'(1) \leq q'(1) \leq q^{(3)'}(1) \leq \dots \leq 2$. Hence, the solution is

$$\begin{aligned} p'(1) &= 2 - (2 - q'(1))^{1/2} = 2 - (2 - q^{(3)'}(1))^{1/3} \\ q'(1) &= 2 - (2 - q^{(3)'}(1))^{2/3} . \end{aligned} \quad (40)$$

The actual value of the derivatives depends on the higher derivative $q^{(3)'}$, and thus on the truncation scheme adopted. The simulation results (Figure 7) suggest that $p'(1) = q'(1) = 2$, which implies (via (40) that $q^{(3)'}(1) = 2$, and that

$$p = q = q^{(3)} = 2\theta - 1 + O(1 - \theta)^2 = \theta^2 + O(1 - \theta)^2 \quad , \quad (\theta \rightarrow 1) . \quad (41)$$

This is consistent with the truncation schemes: $q = \theta^2$ or $q^{(3)} = \theta^2$.

4.1 Results

The p equation (34) involves q ; the q equation (37) involves $q^{(3)}$, etc. The nature of the approximation depends on how we close the hierarchy. In the simplest, *first approximation* we neglect the correlations beyond the nearest neighbours, i.e. we set $q = q_1 = \theta^2$, solve numerically equation (34) for $p = p_1$, with initial condition $p(0) = 0$, and substitute p_1 and q_1 in the quantities of interest. In the *second approximation* we neglect the correlations beyond the second neighbours, i.e. we set $q_2^{(3)} = q^{(3)} = \theta^2$, and solve the system of equations (34)-(37) for $p = p_2$ and $q = q_2$, with initial conditions $p(0) = q(0) = 0$.

Figure 7 shows the simulation data, p_s and q_s , and the predictions from the first theory, p_1 and $q_1 = \theta^2$. The correlations from the simulation decay with distance as in (21), and the differences $p_s - \theta^2$, $q_s - \theta^2$ are small but non-zero, as they should be since they establish the difference of behaviour relative to the RSA model. The agreement between p_1 and p_s is quite good. As expected, it gets slightly worse as θ approaches 1 and the correlations between sites further apart become more relevant, but the correct asymptotic behaviour is obtained. Consequently, there is also good agreement in the results for the density of islands $n(t)$ (Figure 8) and the average island size $\langle L \rangle$ (Figure 9), which are (exact) functions of p only. The agreement between q_1 and q_s (Figure 7) is, of course, less satisfactory. Figure 10 compares the theoretical and simulation plots for the probabilities of nucleation, growth and coagulation, P_n , P_g and P_c (eq. (30) with $D = 1$; cf. Figure 6). There is good quantitative agreement up to $\theta = 0.5$, and there is still some qualitative agreement for larger coverages. The theory fails, however, to give the correct asymptotic behaviour as $\theta \rightarrow 1$: although there is a region where $P_g \simeq 1 - P_c$, the theory gives $P_g \rightarrow 0$ as

$\theta \rightarrow 1$. From (30) we can see that $P_g \simeq (1 - q')/(1 - p') - 1$ close to $\theta = 1$. Hence the limit $P_g = 0$ is a consequence of having $p'(1) = q'(1)$, which follows (via (40) from the truncation $q = \theta^2$.

Nearest-neighbour correlations are sufficient to probe the presence of island boundaries. Since the above quantities depend essentially on island counting, their predictions are fairly accurate. Except in the early stages of deposition, however, long-range correlations need to be accounted for to correctly describe the spectrum of island sizes. Hence, as a result of the pair decoupling approximation (26), the theory predicts a geometric distribution for the island size probability (eq. (32)), the same as in the RSA case but with θ replaced by p/θ . The plots of $P(L)$ are analogous to the RSA ones (Figure 5), though since $p/\theta > \theta$ each curve appears to correspond to a slightly larger coverage. The second moment of $P(L)$ (eq. (31)) is, of course, also incorrect: its limit as $\theta \rightarrow 1$ is $1/2$ as in RSA, rather than 1 as in the simulation. From the plots of $P(L)$ and its second moment, we find that the theory breaks down for these quantities for coverages over 0.2 .

Figure 11 compares the simulation data with the predictions from the second theory, p_2 , q_2 and $q_2^{(3)}$. As before, p_2 fits p_s quite well, and q_2 fits q_s even better. Consequently, the agreement in the results for $n(t)$ (Figure 8) and $1/\langle L \rangle$ (Figure 9) is also good: there is no major difference between the two theories, apart from the fact that now the curves lay below the simulation plots. A similar difference between theories is found (Figure 10) for the probability of nucleation $P_n = n(000)$, an approximate function of both p and q (eq. (28)). A worse agreement with the simulation than in the first theory is obtained, however, for the probabilities of growth and coagulation. We have also tested other plausible choices for the closure scheme, as the ones employed above are not unique, but found the results were either largely unchanged or incorrect.

Finally, we note that p_2 lies over p_s , while p_1 lies below p_s . q_2 , on the other hand, lies between q_s and q_1 . Hence, the p correlations are underestimated in the first theory and overestimated in the second theory. This seems to indicate that the approach cannot be systematically improved by higher order truncations in the pair correlation hierarchy. This fact comes as no surprise given the uncontrolled nature of a self-consistent approach. Moreover, the results are more likely to be affected by the pair decoupling approximation (26) than by the order of truncation in the pair correlation hierarchy.

5 Conclusions

We have studied numerically and analytically a simple, but non-trivial model for the deposition of monomers on a line. The particles can diffuse on the extrinsic precursor layer until they reach the edge of the island, where they are irreversibly deposited. As time progresses, island nucleation becomes less frequent, while the larger islands grow rapidly and merge with other islands. During this process, the precursor particles migrate for larger and larger distances with ever increasing probability, establishing correlations between sites further and further apart. As a result, the system develops a structure characterized by strong correlations whose range grows to the system size at full coverage.

In the simulation, we looked at the evolving structure pattern by measuring the island density number, the island size probability distribution and its first and second moments, and some of the pair correlations. We also looked at the interplay between the direct and the diffusion mediated mechanisms of deposition, by measuring the nucleation, growth and coalescence probabilities per unit time. As expected, the results (Figs. 1-7) differ considerably from the RSA model, especially at large coverages, due to the increasing correlations.

To explain and interpret these measurements, we proposed a simple, self-consistent theory which incorporates correlations to a minimum extent, i.e. local pair correlations. We considered two levels of approximation, depending on the closure of the hierarchical equations for the pair correlations. Altogether, the lowest level of approximation, accounting for nearest-neighbour correlations only, gave the best fit to the simulation data. The predictions of the theory (Figs. 7-9) proved very accurate for the nearest-neighbour correlator p , island density number $n(t)$ and average island size $\langle L \rangle$. A fairly good prediction was also obtained for the probabilities of nucleation, growth and coagulation (Fig. 10). While the nearest-neighbour correlations are sufficient to distinguish occupied from non-occupied regions, the full range of correlations is required to distinguish the sizes of those regions. An accurate determination of the island size distribution $P(L)$ and its second moment is, therefore, beyond the scope this theory, and would, in principle, require the use of more complicated methods.

We end with a comment on the $2d$ systems, which are of more interest to experimentalists. In this case, however, the same quantities are not easily expressed in terms of the local lattice variables, and one is faced with basic difficulties in the development of a useful formalism (most of section 2 would not be applicable) and in the analytical treatment. To illustrate the problem, we note that instead of the nearest-neighbour correlations, some non-local operator would be required to probe

the domain boundaries. It is desirable and possible, nonetheless, to perform numerical simulations of the $2d$ model, which would also enable the study of richer phenomena, such as percolating clusters.

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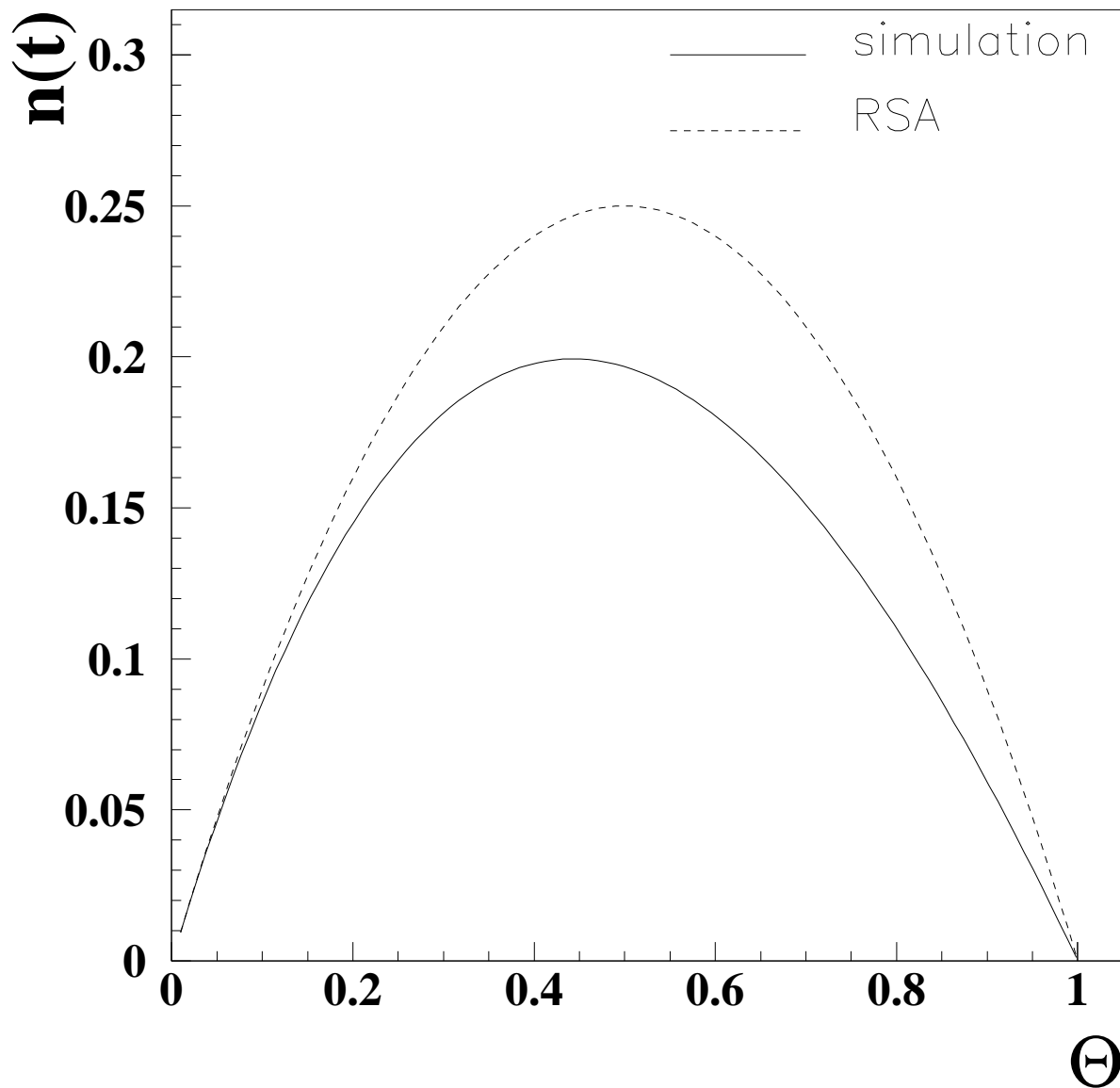


Figure 1: Density number of islands, $n(t)$, as a function of the coverage, θ . Simulation data and RSA model.

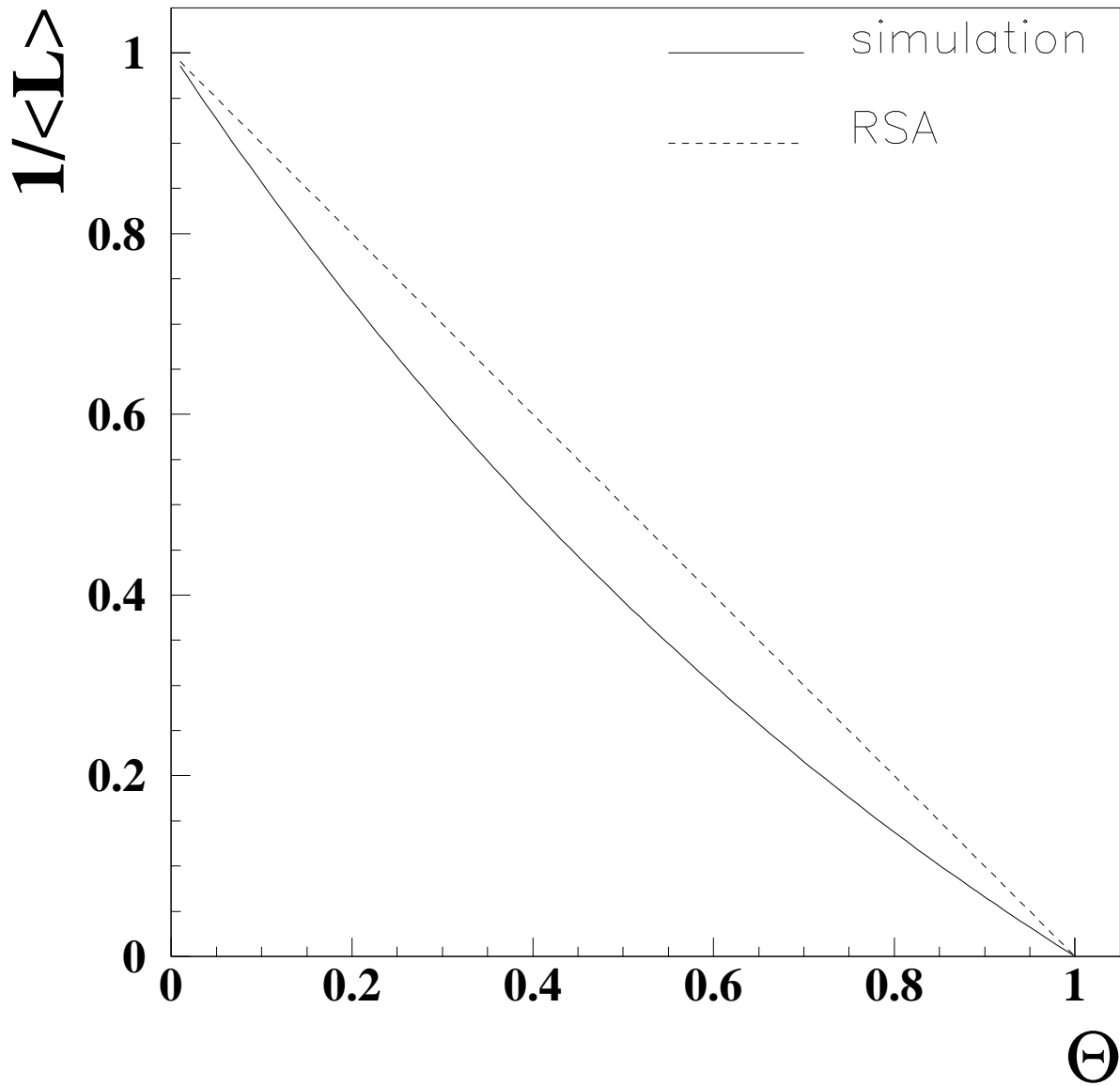


Figure 2: Inverse of the average island size, $1/\langle L \rangle$, as a function of the coverage, θ . Simulation data and RSA model.

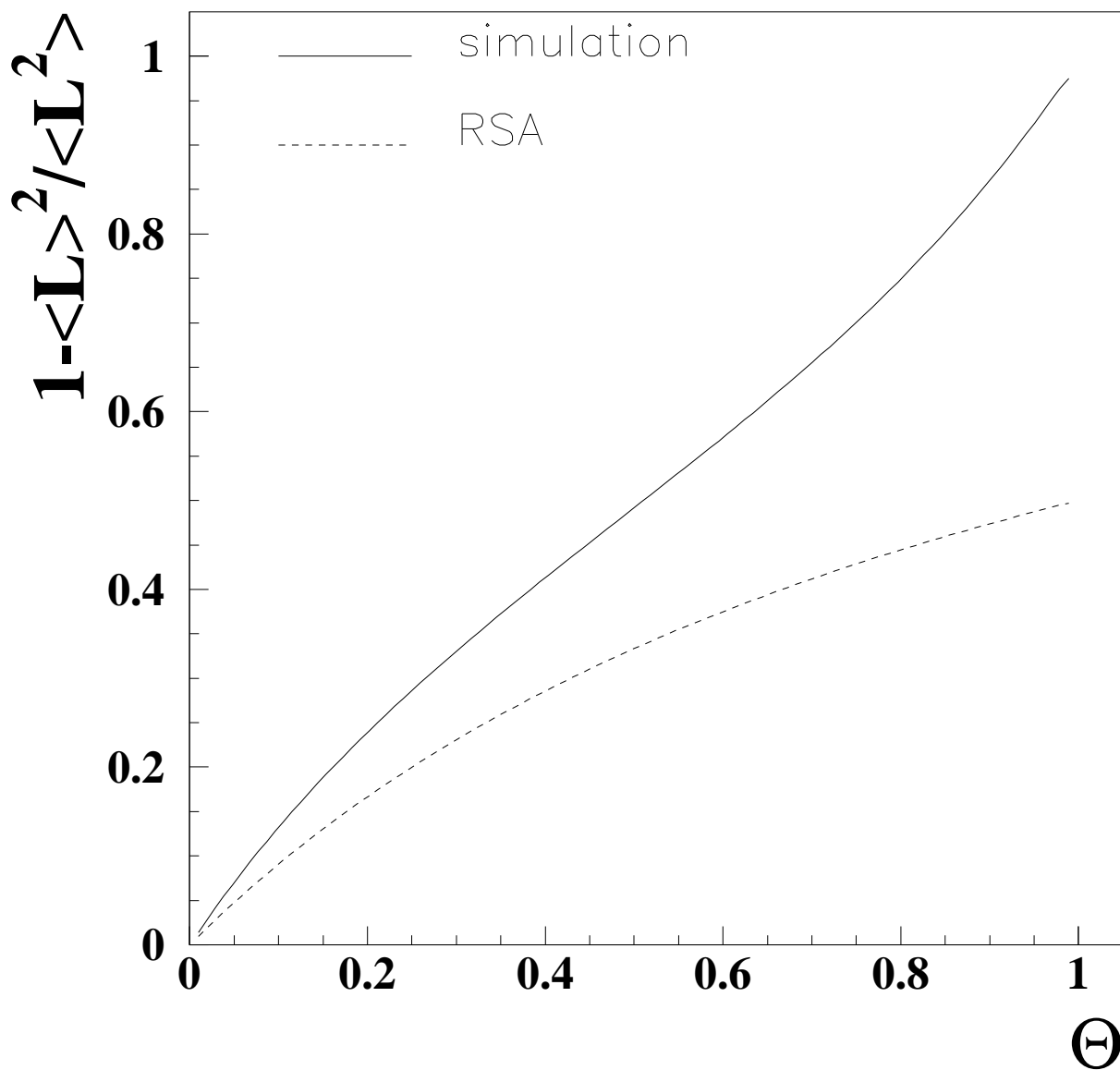


Figure 3: Mean square deviation of the average island size, $1 - \langle L \rangle^2 / \langle L^2 \rangle$, as a function of the coverage θ . Simulation data and RSA model.

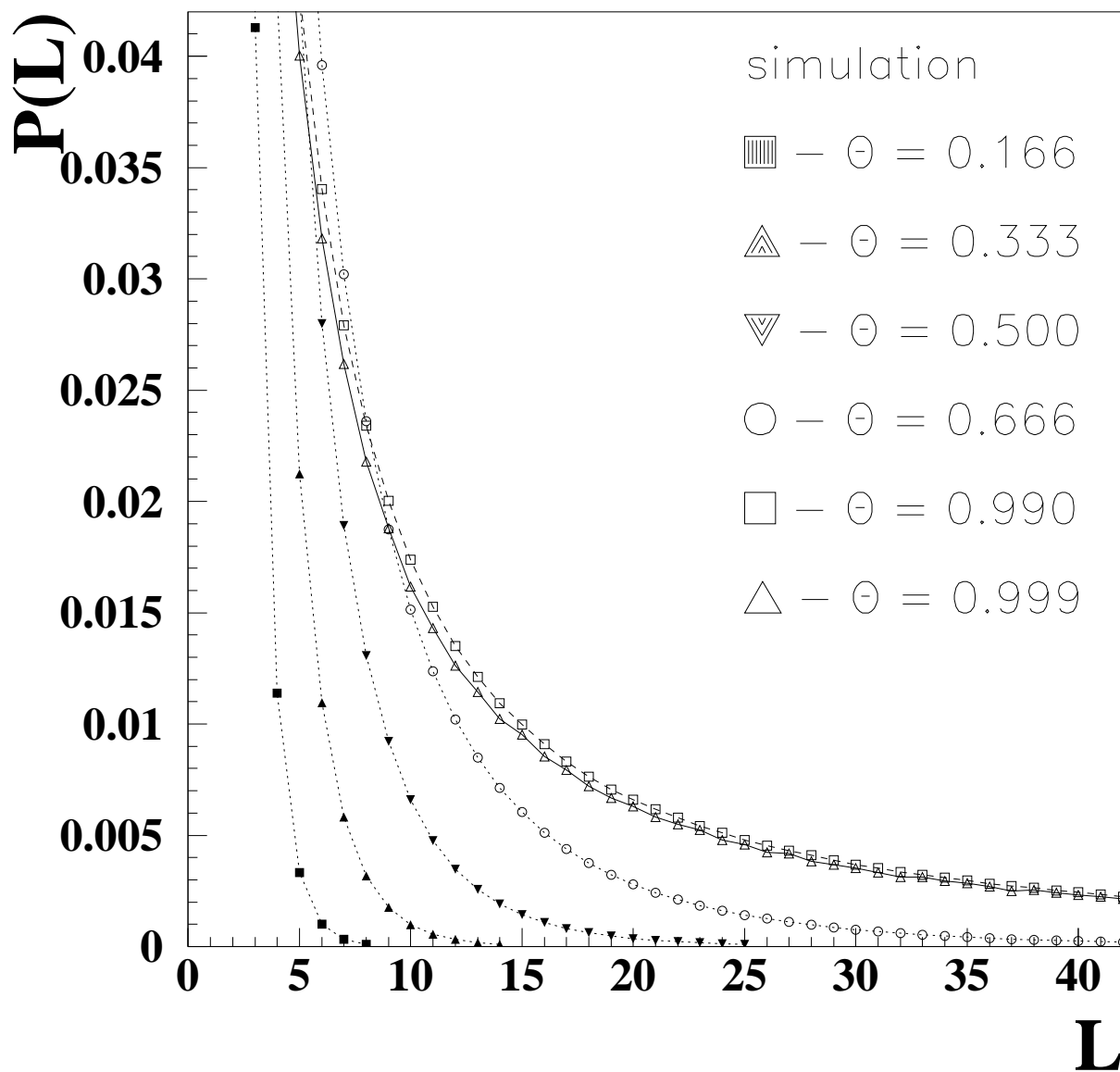


Figure 4: Probability distribution for the island sizes, $P(L)$, at six different coverages, $\theta = 0, 0.166, 0.333, 0.500, 0.666, 0.990, 0.999$. Simulation data.

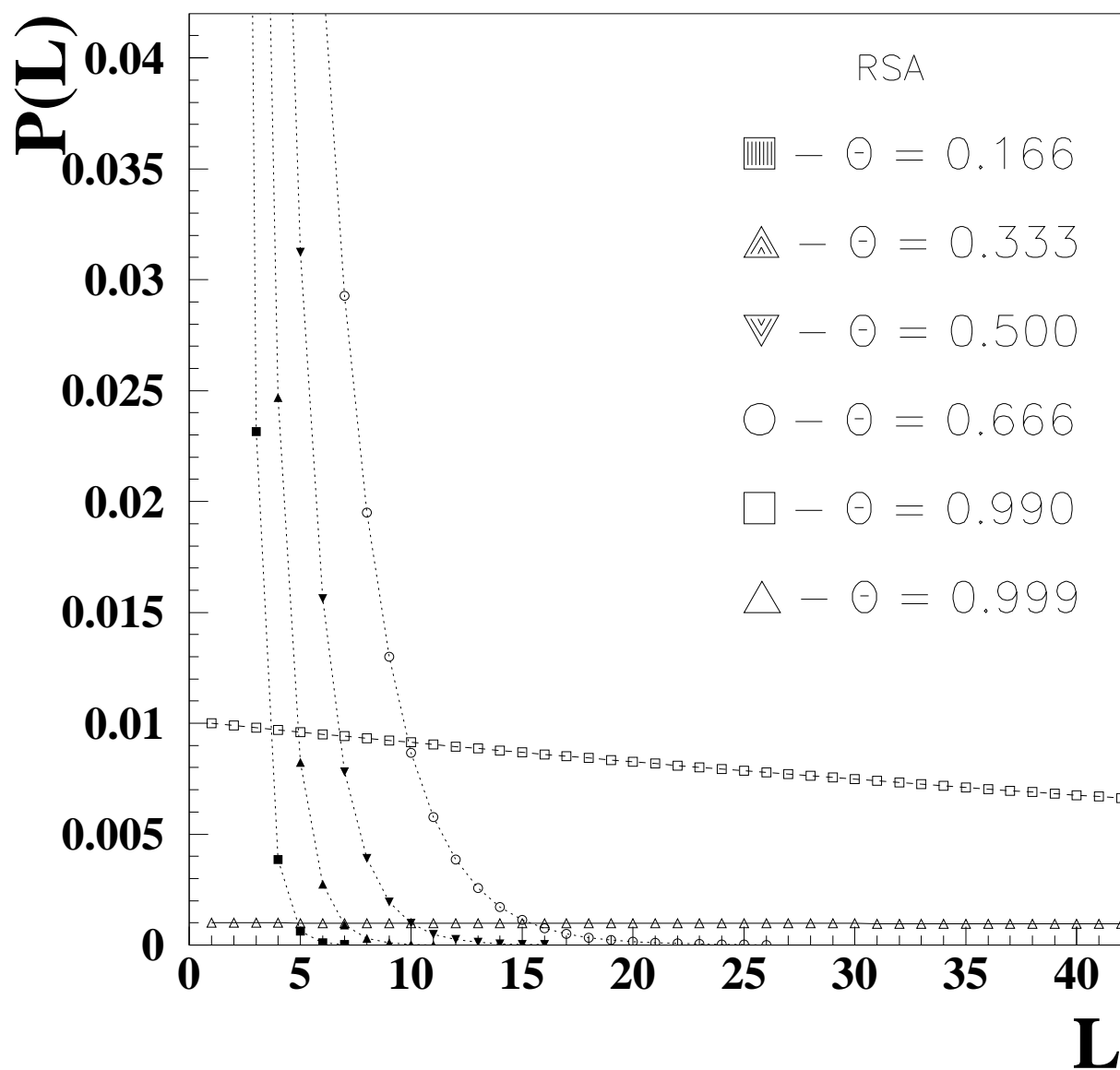


Figure 5: Probability distribution for the island sizes, $P(L)$, at six different coverages, $\theta = 0, 0.166, 0.333, 0.500, 0.666, 0.990, 0.999$. RSA model.

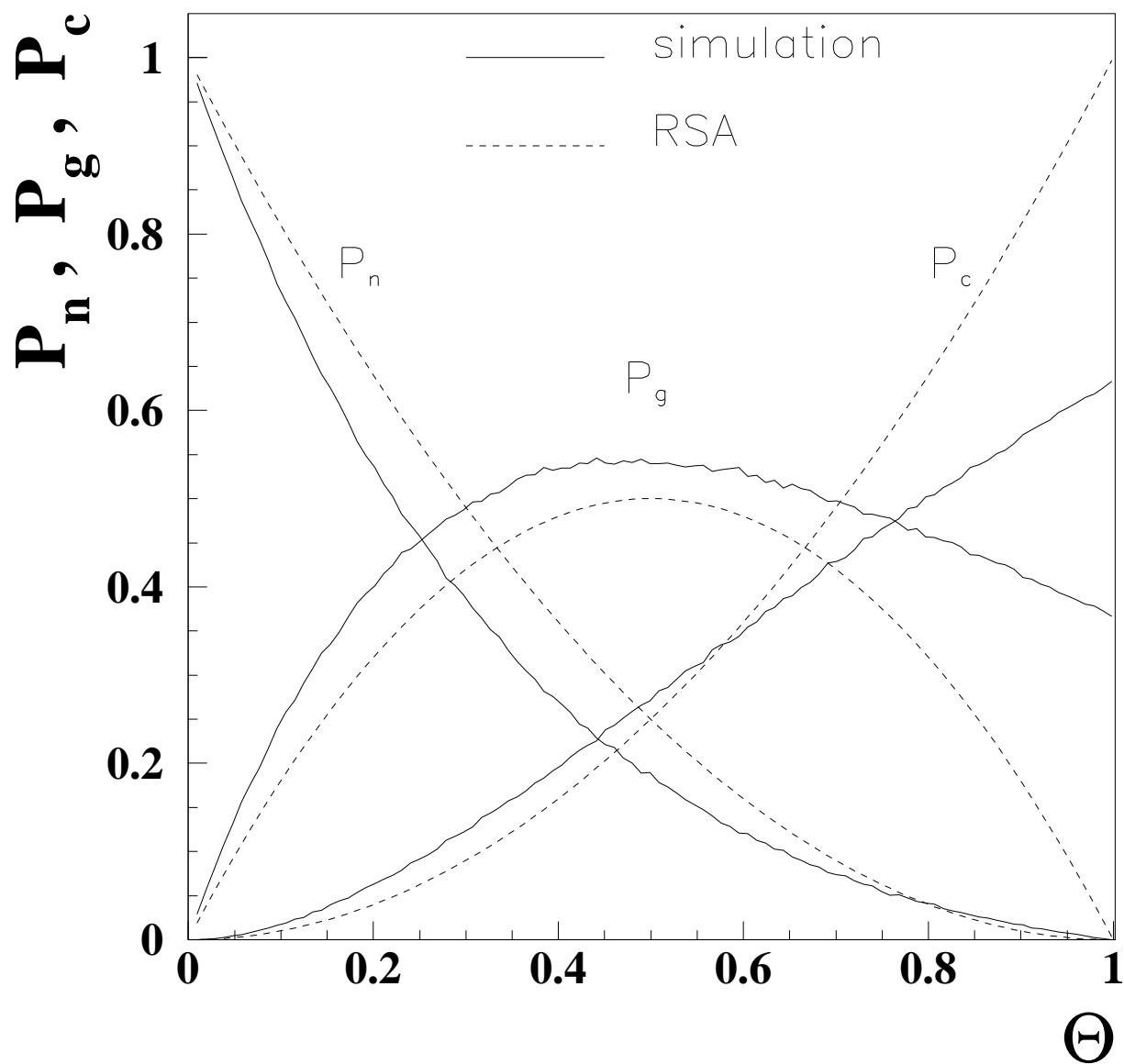


Figure 6: Probabilities of nucleation, growth and coagulation, P_n , P_g and P_c , as a function of the coverage θ . Simulation data and RSA model.

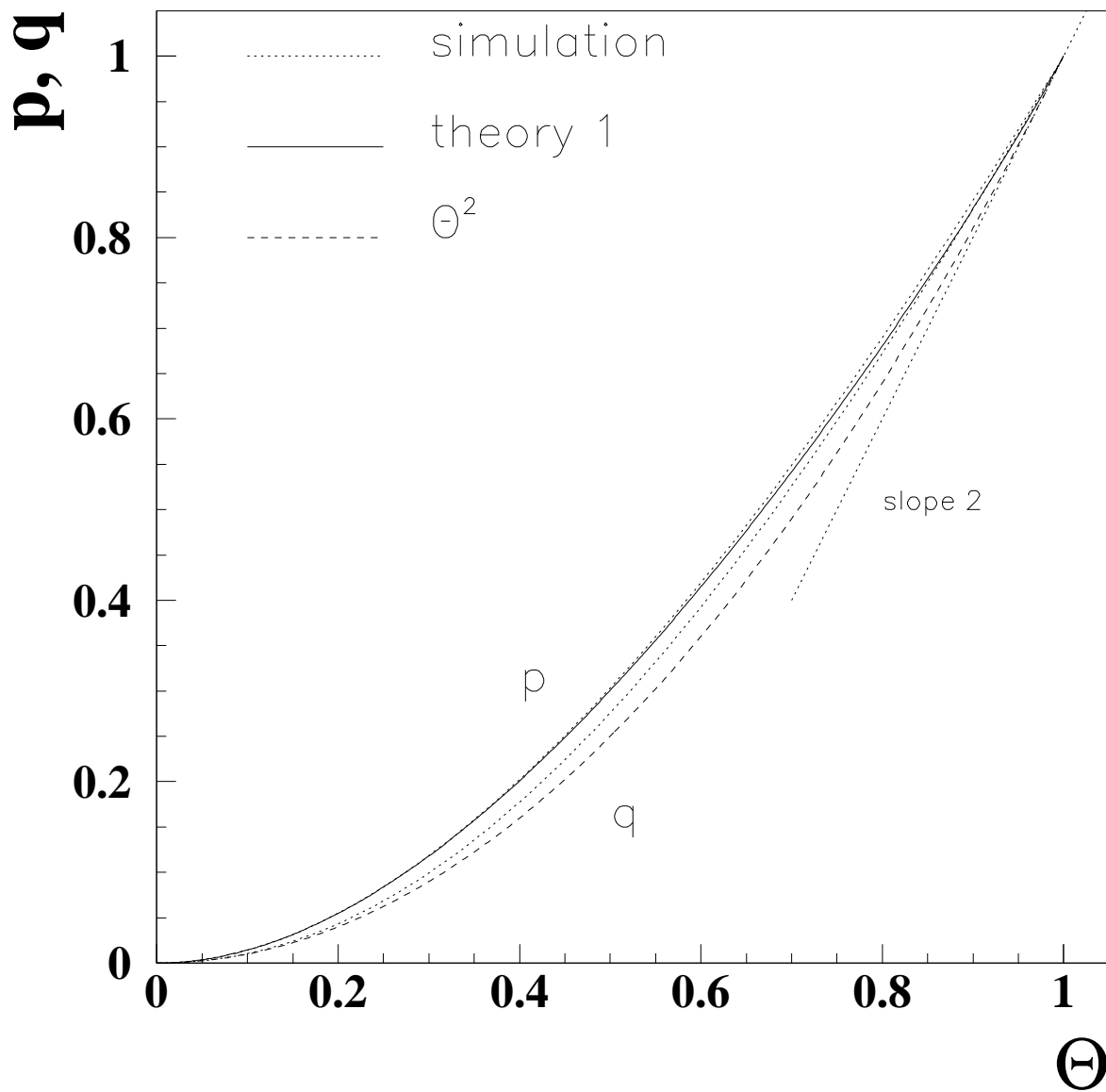


Figure 7: Nearest-neighbour and second nearest neighbour pair correlations, p and q , as a function of the coverage θ . First theory ($q = \theta^2$) and simulation data.

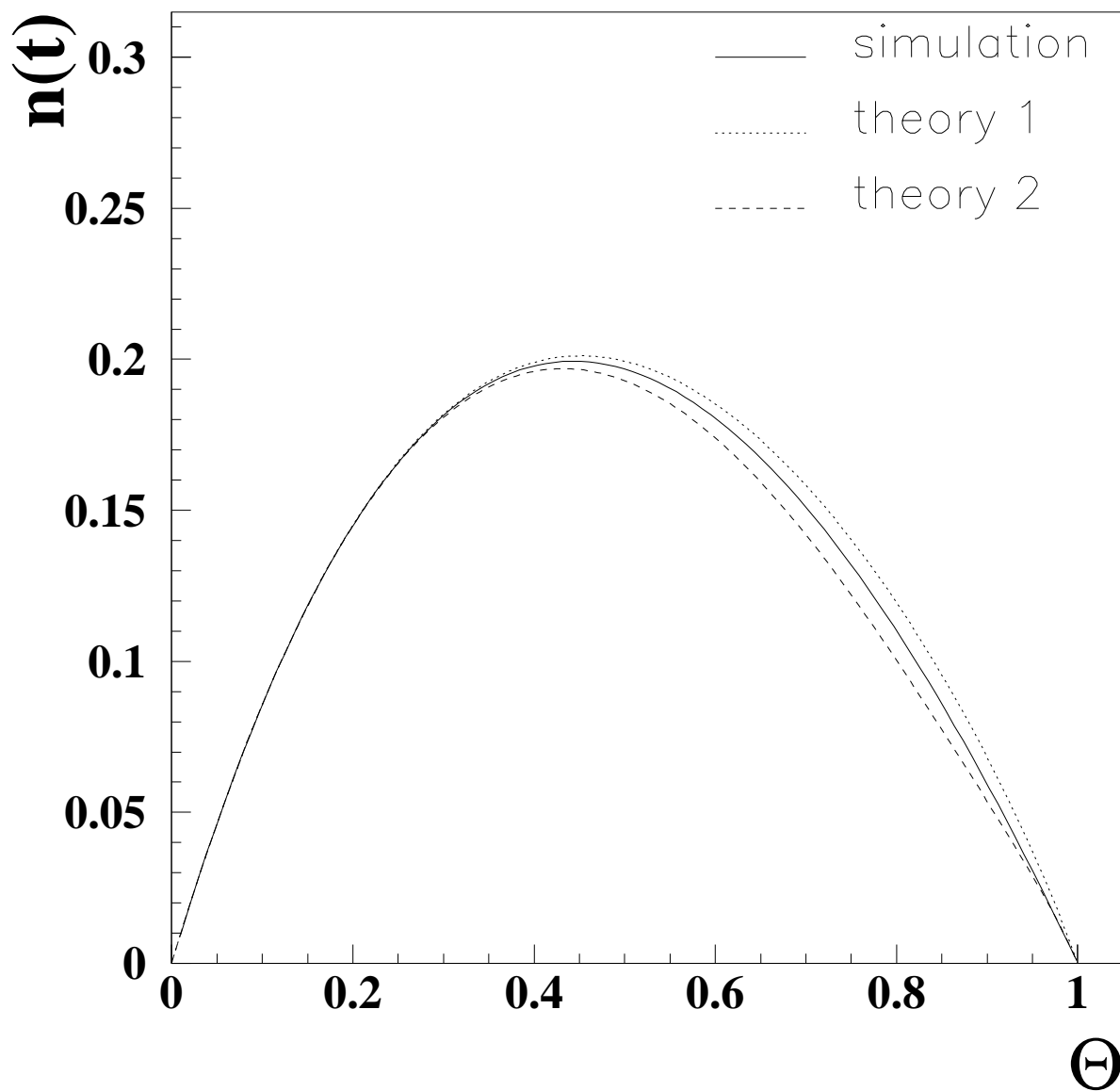


Figure 8: Density number of islands, $n(t)$, as a function of the coverage, θ . First theory, second theory and simulation data.

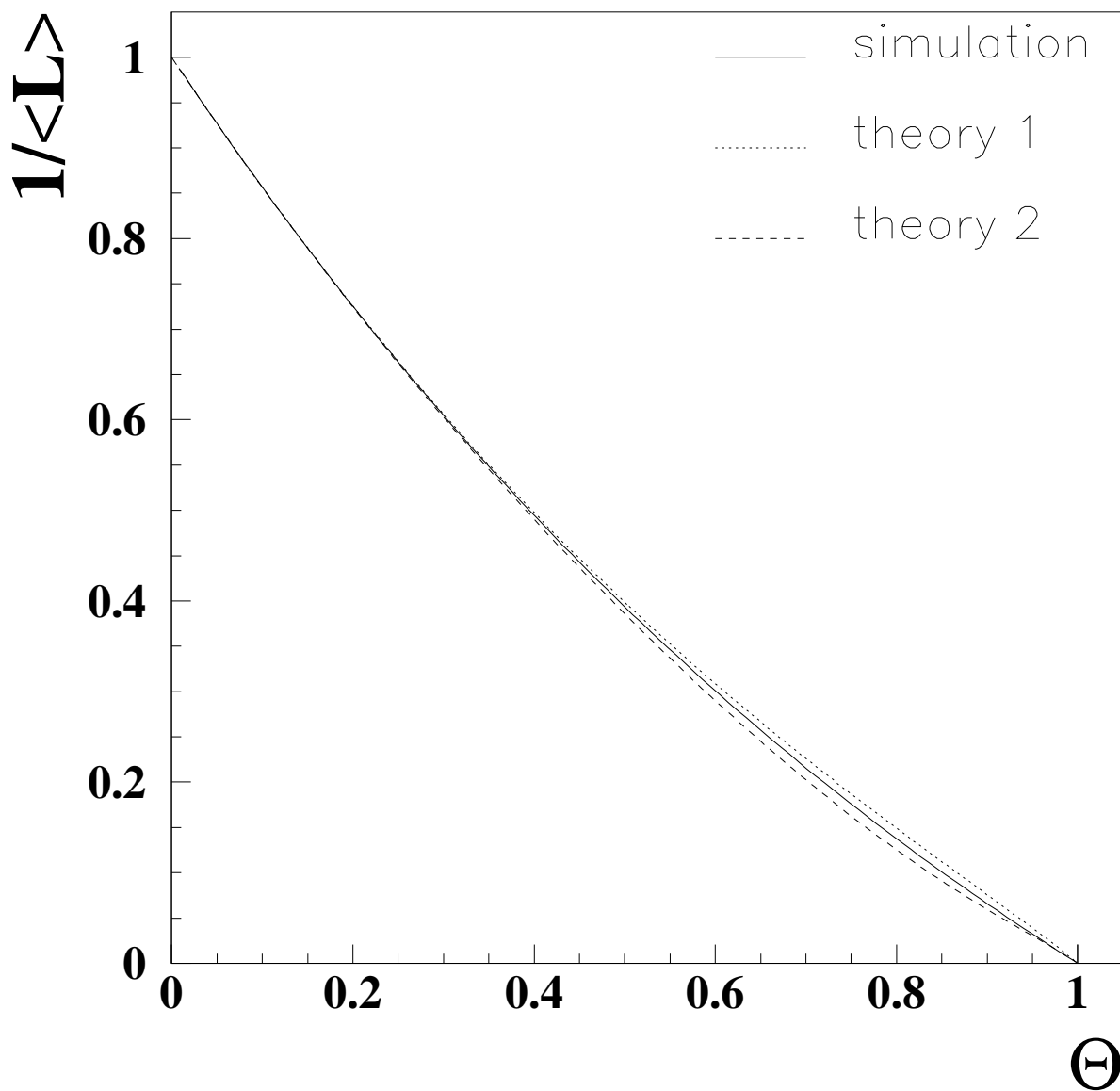


Figure 9: Inverse of the average island size, $1/\langle L \rangle$, as a function of the coverage, θ . First theory, second theory and simulation data.

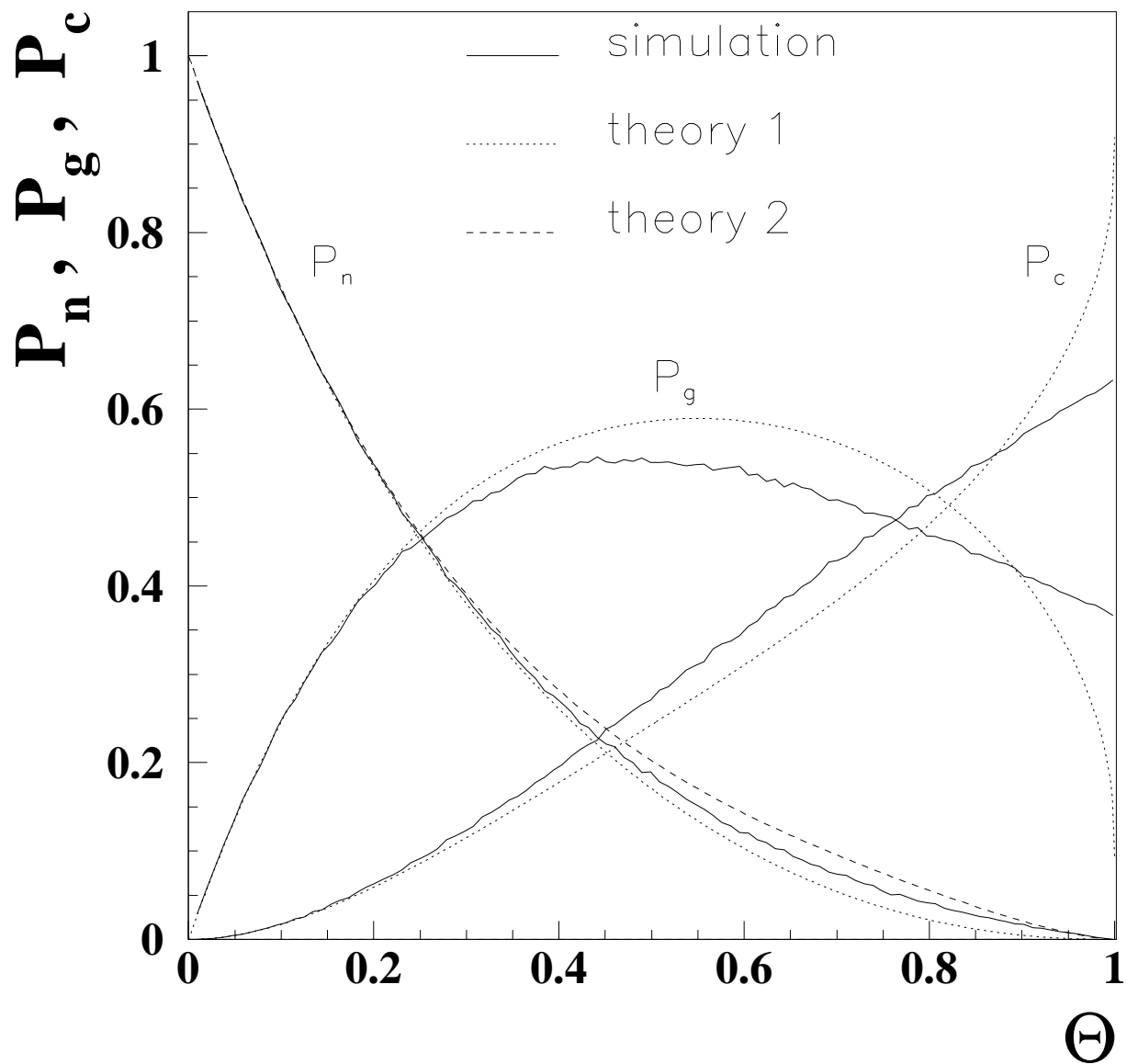


Figure 10: Probabilities of nucleation, growth and coagulation, P_n , P_g and P_c , as a function of the coverage θ . Simulation data and RSA model. P_n : first theory, second theory and simulation data. P_g and P_c : first theory and simulation data.

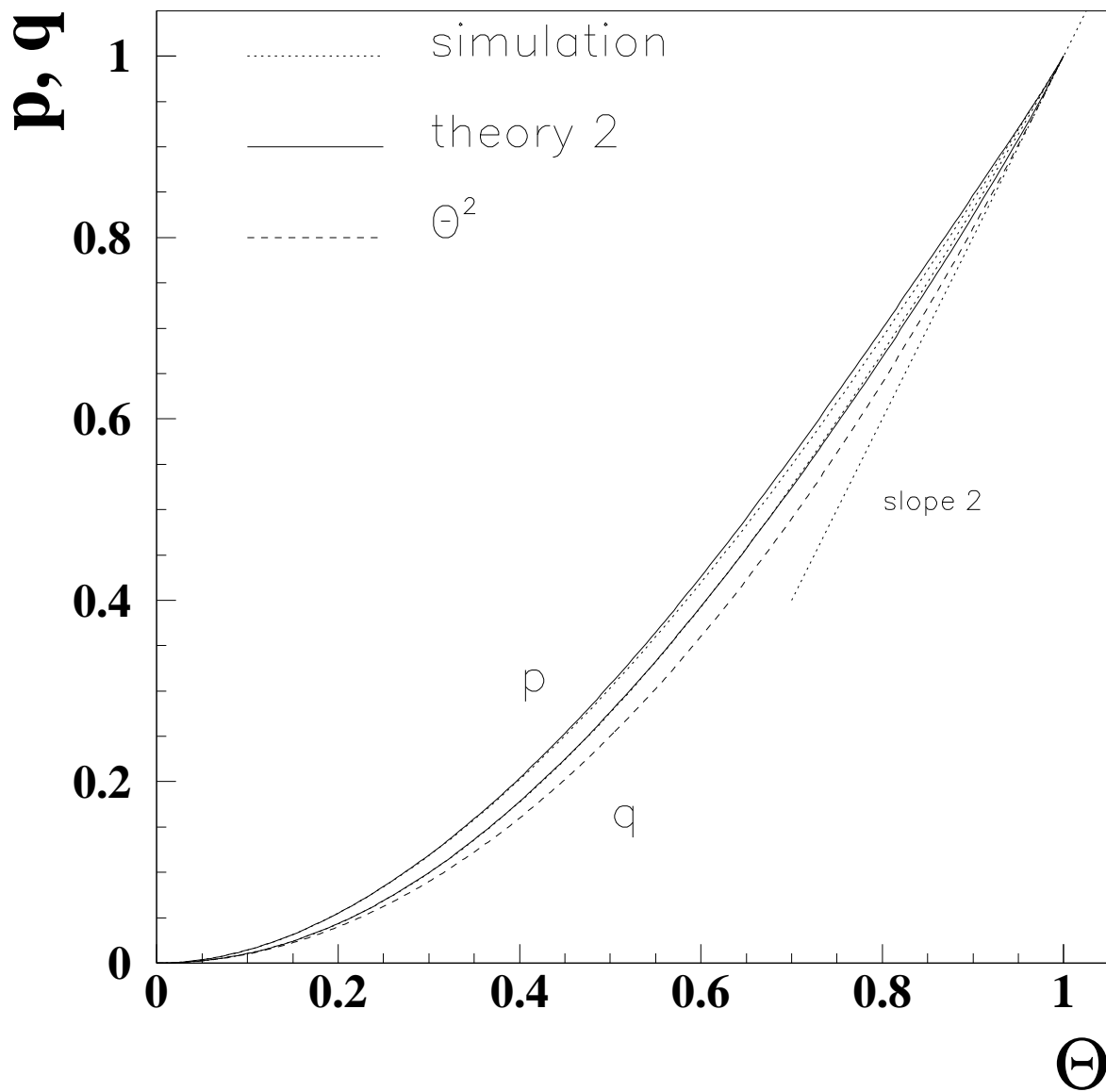


Figure 11: Nearest-neighbour and second nearest neighbour pair correlations, p and q , as a function of the coverage θ . Second theory ($q^{(3)} = \theta^2$) and simulation data.