



Elucidating esterification reaction during deposition of cutin monomers from classical molecular dynamics simulations

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Abstract

The structural behavior of some cutin monomers, when deposited on mica support, was extensively investigated by our research group. However, other events, such as esterification reaction (ER), are still a way to explore. In this paper, we explore possible ER that could occur when these monomers adsorb on support. Although classical molecular dynamics simulations are not able to capture reactive effects, here, we show that they become valuable strategies to analyze the initial structural configurations to predict the most favorable reaction routes. Thus, when depositing aleuritic acid (ALE), it is observed that the loss of capacity to form self-assembled (SA) systems favors different routes to occur ER. In pure ALE bilayers systems, an ER is given exclusively through the –COOH and primary –OH groups. In pure ALE monolayers systems, the ER does not happen when the system is self-assembled. However, for disorganized systems, it is able to occur by two possible routes: –COOH and primary –OH (route 1) and –COOH and secondary –OH (route 2). When palmitic acid (PAL) is added in small quantities, ALE SAMs can now form an ER. In this case, ER occurs mostly through the –COOH and secondary –OH groups. However, when the presence of PAL is dominant, ER can occur with either of both possibilities, that is, routes 1 and 2.

Keywords Molecular dynamics · Esterification reaction · Self-assembly · Aleuritic acid · Palmitic acid

Introduction

Fatty polyhydroxyacids (C_{16} , C_{18}) are the building monomers of ubiquitous biopolyesters such as cutin and suberin in higher plant tissues [1]. Both constitute the fourth most abundant plant biopolymer in nature after cellulose, hemicellulose, and lignin. In addition to the natural abundance, their extraordinary barrier capacity has attracted the interest for the

obtaining of synthetic mimetic polymers to replace hazardous and non-biodegradable petroleum-derived materials [2, 3]. Fatty polyhydroxyacids have been found to have the capability to self-assemble and self-esterify at an ambient condition to form nanometer-size particles named as cutinsomes [4]. Such nanoparticles have been detected *in planta* [5] and they can further aggregate and spontaneously polymerize at standard environmental conditions to form a continuous polyester layer, thus providing a plausible mechanism for *in vivo* cutin biosynthesis [6, 7]. Such hypothesis is gaining recognition since analogous nanometer-size structures have also been recently detected in the elucidation of the molecular architecture of suberin [8].

The structure of nanoparticles constituted by fatty hydroxyacid is primarily conditioned by intermolecular interactions. Those interactions have been studied by our group both theoretically and experimentally by the formulation of confined and geometrically well-defined systems such a self-assembled (SA) layers on flat inert surfaces.

Several analytical techniques such as atomic force microscopy (AFM) and scanning tunneling microscopy (STM) combined with molecular dynamics simulations were essential to understanding the behavior of these monomers [9]. With these

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techniques, it was possible to show that the deposition of polyhydroxy acids leads to the formation of self-assembled bilayers (SABs) and monolayers (SAMs), depending on the presence and position of the functional groups ($-\text{COOH}$, and $-\text{OH}$) in the aliphatic chain.

The importance of these studies lies in the transference of this seminal knowledge to the design and manufacturing of layered biodegradable biomimetic materials with potential applications such as protective films for food and cosmetics packaging [10–13].

Therefore, to investigate other properties, in addition to the structural ones, the feasibility of the esterification reaction (ER) is of major importance for this purpose. Our first ER studies by attenuated total reflection Fourier transform infrared (ATR-FT-IR) spectroscopy, X-ray photoelectron spectroscopy (XPS), and molecular dynamics (MD) techniques in self-assembled (SABs) systems of aleuritic acid (ALE) bilayers show that the esterification reaction occurs as a consequence of the formation of SA [14]. However, in this preliminary study, it was not satisfactorily clarified experimentally whether the water detected originates from esterification, or from the environment. Our results from MD simulations suggested that it comes from esterification.

This reaction could be classified as a rare event since it would present a high activation energy barrier and requires catalysts to happen [15]. In this work, this possibility will be more explored by using MD simulations of different nanostructures observed in AFM images in pure ALE systems. Additionally, we also study mixed systems of ALE and palmitic acid (PAL) with different compositions (25:75 and 75:25% of ALE:PAL). PAL molecule has only a terminal $-\text{COOH}$, and when mixing with ALE modifies the capability of self-assembling, resulting in systems with different degrees of disordering, which constitutes an essential aspect in the understanding of the ER mechanisms.

Experimental section

AFM operation

AFM images were obtained with a Nanotec Cervantes microscope (Nanotec, Spain) using either a long-range ($70 \times 70 \text{ m } \mu\text{m}^2$) scanner to check for reproducible sample preparation or a shorter one ($10 \times 10 \text{ m } \mu\text{m}^2$) to collect more accurate topographic data. Every sample was studied at four distant points using both scanners, and images were acquired, processed, and analyzed using the WSxM software. Scanners were calibrated using NT-MDT TGT01 and Nanosensors H8 gratings for x - y - and z -directions, respectively.

The AFM microscope was operated in the air in dynamic, contact, and jumping modes. Dynamic is selected for height measurement because it exerts a minimal

perturbation of the sample by the scanning probe. In this mode, silicon cantilevers (PPP-FMR Nanosensors) with nominal 75-kHz resonance frequency and 2.8 N/m force constant (k) were used. The lever was oscillated ($A_0 \sim 90 \text{ nm}$ peak to peak) at its free resonance frequency. The feedback signal was set to a 20% reduction of the free oscillation amplitude and both topographic and phase-shift images were recorded simultaneously. For additional friction and adhesion mapping, the AFM was operated in contact and jumping modes using Si_3N_4 levers (TR400PSA Olympus, $k = 0.08 \text{ N/m}$, and RC800PSA Olympus, $k = 0.1 \text{ N/m}$, respectively) at low set points.

Sample preparation

Self-assembled layers of aleuritic (9,10,16 trihydroxyhexadecanoic, TCI, >98.0%) acid were prepared from a 0.5 mM solution in chloroform (Merck, 99.8%) by spin coating. A 20- μl drop was placed on top of a piece of freshly cleaved mica (muscovite) and spun at 40 rps for 20 s. Samples were stabilized for 3–4 h at RT inside a covered glass Petri dish before being transferred to the AFM microscope. For reproducibility, each preparation was repeated at least four times.

Computational methods

Molecular dynamics (MD) simulations were carried out using the Classic DL_POLY [16] code in the canonical ensemble NVT [17]. The temperature was kept constant at 300 K by applying a Nosé-Hoover thermostat [18]. The equations of motion were integrated by using the velocity Verlet algorithm with a time step of 1.0 fs.

The $-\text{CH}_3$ and $-\text{CH}_2-$ groups were treated as pseudo atomic units using the CHARMM19 force field [19] (Fig. 1 and Eq. 1). Lennard-Jones (LJ) and Coulomb interactions were truncated at 15 Å in our simulations. Periodic boundary conditions were applied along the x - and y -directions. The Ewald method with periodic boundary conditions in two dimensions was used to treat the electrostatic interactions. Computational boxes with lengths of $L_x = 60.0 \text{ Å}$ and $L_y = 34.6 \text{ Å}$ were considered to simulate all systems. The substrate was represented as a flat surface using two models: (1) a 12–3 Lennard-Jones type external potential (Eq. 2) and (2) a short-range homogeneous flat potential.

$$E_1 = \sum_b K_b [b - b_0]^2 + \sum_\theta K_\theta [\theta - \theta_0]^2 + \sum_\varphi K_\varphi [1 + d \cos \varphi_0] + \sum_{i,j} 4\epsilon_{ij} \left[\left(\frac{\sigma_{ij}}{r_{ij}} \right)^{12} - \left(\frac{\sigma_{ij}}{r_{ij}} \right)^6 \right] + \sum_{i,j} \frac{q_i q_j}{\epsilon r_{ij}} \quad (1)$$

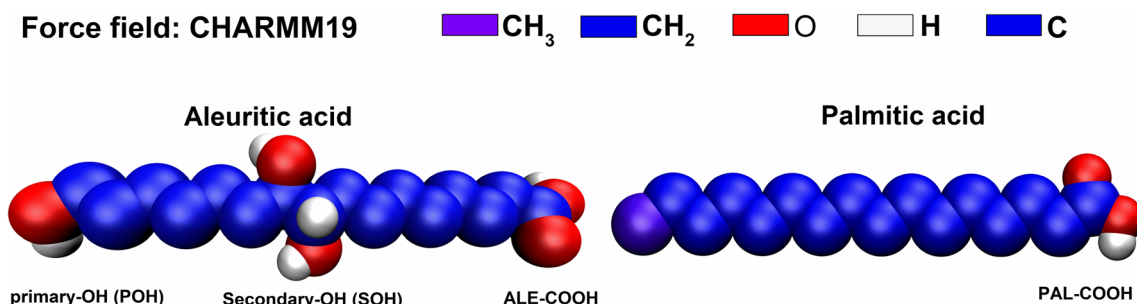


Fig. 1 Representation of ALE and PAL molecules using the CHARMM19 force field

$$E_2 = \sum_{i,j} \frac{C_{12}}{[z-z_0]^{12}} - \frac{C_3}{[z-z_0]^3} \quad (2)$$

where C_{12} and C_3 parameters for $-\text{CH}_2-$, $-\text{CH}_3$, C, and O were taken from the literature [20–22], z is the distance to the surface, and z_0 is a limit approach distance for each center. Furthermore, b_0 , θ_0 , and φ_0/φ_i are equilibrium bond length, bond angle, and dihedral angle, respectively. K_b , K_θ , and K_φ are the corresponding constants, and b and θ are the actual bond length and bond angle, respectively. ε_{ij} and σ_{ij} decide the minimum and zero values of the van der Waals term, which is described by 12–6 Lennard-Jones (L-J) potential, and r_{ij} is the distance between two particles with charges q_i and q_j . L-J potential parameters between different crossed atoms were calculated using the Lorentz-Berthelot combining rules. In the short-range homogeneous flat potential model, the parameters were taken from the literature [23–25] and the atoms were frozen for stability as recommended.

A total of 10 systems were defined to model mixtures of ALE compositions in weigh of 25, 75, and 100%, named as 25:75, 75:25, and 100:00 (ALE:PAL), respectively. Based on experimental results, these systems were chosen as representative cases for (1) segregated, (2) dispersed, and (3) pure systems, respectively. Thus, initial configurations were constructed by placing a total of 96 molecules of ALE and/or PAL oriented vertically with the carboxylic group near the support.

For each of these compositions, in a hexagonal network, the dispersed (D), segregated (S), and pure (P) systems were built. For a 75:25 composition, two dispersed configurations, named H75D1 and H75D2, and two segregated configurations, denoted as H75S1 and H75S2, were established. Similarly, for a 25:75 composition, there are two dispersed (H25D1 and H25D2) and two segregated (H25S1 and H25S2) systems. For pure ALE systems, monolayer and bilayer systems were defined and named as H100M and H100B. The nomenclature and details for each type of system are compiled in Fig. 2.

The initial configurations, created with high structural tension, were submitted to a relaxation process using the “zero” technique implemented in the DL_POLY code for 1 ns. Next, the equilibration process was carried out

gradually at 100, 200, and 300 K, during 1 ns for each temperature and verifying that equilibrium was reached at each stage. Finally, all systems were simulated for 30 ns. The structural properties were analyzed along the trajectory in the following intervals: (1) 9.5–10.0 ns, (2) 19.5–20.0 ns, and (3) 29.5–30.0 ns. Configurations were saved every 50 timesteps collecting 10,000 configurations at each interval for statistical analysis. In all of those intervals, the behavior of the structural properties remained unaltered. For the analysis of hydrogen bonding (H-bond), it was considered the 10–30-ns trajectory.

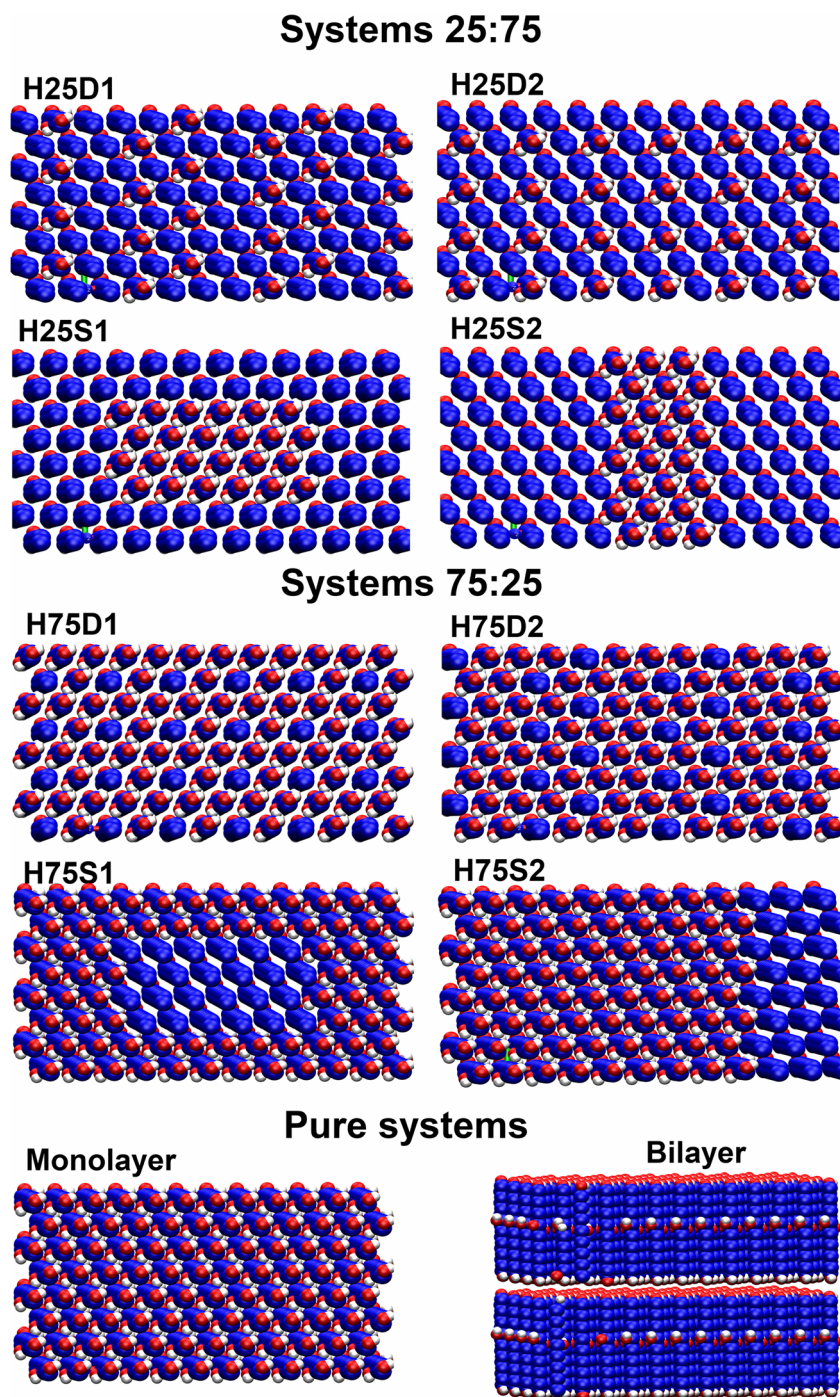
Results and discussion

Pure ALE systems

AFM images of pure self-assembled ALE (Fig. 3a, right) show mostly isolated compact island ≈ 2.4 nm high (L2), as well as scattered detached tinny rounded agglomerates. Occasionally, bilayers (L3) can also be observed. Higher resolution scanning (Fig. 3a, left) reveals another flat and compact phase characterized by a lower height (L1). Additional AFM measurements (not shown) indicate that L1, if compared with L2, L3, and mica background, is a low friction and low adhesion moieties. Previous work has demonstrated that L1 corresponds to a disordered and esterified arrangement of ALE molecules [26]. This study also justifies the predominance of vertically packed monolayer islands (L2) vs. bilayers (L3) due to the enhancement of the 2D vs. the 3D growing mechanism conditioned by the presence of the two secondary hydroxyls in the molecular skeleton.

Simulations on these systems (L1, L2, and L3, Fig. 3b) reveal two different ways for the $-\text{COOH}$ and $-\text{OH}$ groups to interact and to conduct to esterification reactions, namely route 1 (R1): when involving interactions between $-\text{COOH}$ and primary $-\text{OH}$ groups (POH), and route 2 (R2): for $-\text{COOH}$ and secondary $-\text{OH}$ groups (SOH) interactions (Fig. S1).

Fig. 2 Initial configurations of all simulated systems. Systems with ALE:PAL compositions of 25:75, 75:25, and pure systems



ALE bilayer systems

In the bilayers (denoted as L3 in the AFM images, Fig. 3), the only way to favor an ER is through route R1, that is, when $-\text{COOH}$ and POH groups interact (Fig. S2). Figure 4 shows the radial distribution functions (RDF) for these groups. The plot for carbon and oxygen distances ($g_{\text{C-O}}$) presents two peaks at 3.2 and 3.7 Å, and three peaks at 1.8, 3.0, and 3.8 Å for hydrogen and

oxygen distances ($g_{\text{H-O}}$). We analyze these distances (named $\delta 1$ and $\delta 2$) to understand which of those peaks are related to an ER. Taking advantage of the fact that $\delta 1$ and $\delta 2$ would be correlated if an ER occurs, combined distribution functions (CDF) are used to poll zones where ER can happen. We use the TRAVIS [27] code to perform all analyses, and they are illustrated in Fig. 5. There are three predominant zones: zone 1 ($1.6 < \delta 1 < 2.5$ Å and $3.5 < \delta 2 < 4.0$ Å); zone 2 ($2.5 < \delta 1 < 3.3$ Å and $3.0 < \delta 2 <$

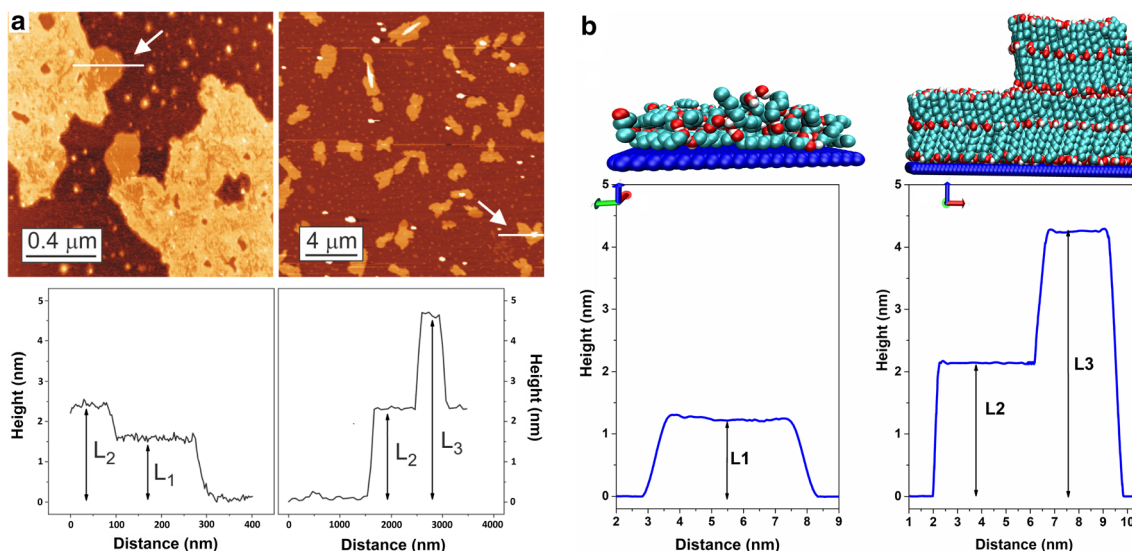


Fig. 3 Experimental and theoretical height profile of self-assembled island. **a** Topographic AFM images of self-assembled island of pure ALE on mica. **b** Molecular dynamics simulations of the self-assembled island

height profile of disorganized ALE monolayer (L1), self-assembled ALE monolayer (L2), and ALE bilayer (L3)

3.8); and zone 3 ($3.5 < \delta_1 < 4.5 \text{ \AA}$ and $3.4 < \delta_2 < 4.1 \text{ \AA}$). It is also clear when comparing the relative intensity of all zones that zone 1 is rarely visited and the system remains the longest in zones 2 and 3.

Analyzing the configurational arrangements existing in those three zones, they can be schematically summarized as represented in Fig. 6. In zone 1, there is a configuration with $\delta_1 = 3.7 \text{ \AA}$ and $\delta_2 = 1.8 \text{ \AA}$, where $-\text{COOH}$ and POH groups get an adequate configuration to favor an ER. For this reason, the presence of these peaks is only monitored in the next systems. It is likely that this configuration is stabilized by H-bond interactions (H-bond 1) formed between the POH group and the protonated oxygen of the $-\text{COOH}$ group.

In zone 2, we find two different configurations that are also possibly stabilized by H-bond interactions. In this zone, the POH group interacts with the unprocessed oxygen of $-\text{COOH}$ (H-bond 2). This type of H-bond prevents an ER from occurring since it would not be possible to get a fitting initial configuration.

Similarly, the main configuration in zone 3 may also be stabilized by H-bond interactions. The $-\text{OH}$ of the $-\text{COOH}$ group interacts with the oxygen of the POH (H-bond 3). This configuration also prevents an ER from occurring.

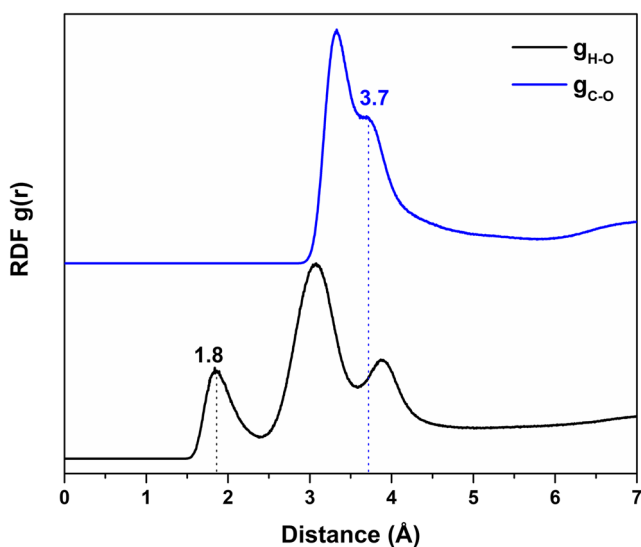


Fig. 4 Radial distribution function (RDF). Analysis of the R1 route for an ER. L3-ALE bilayer system

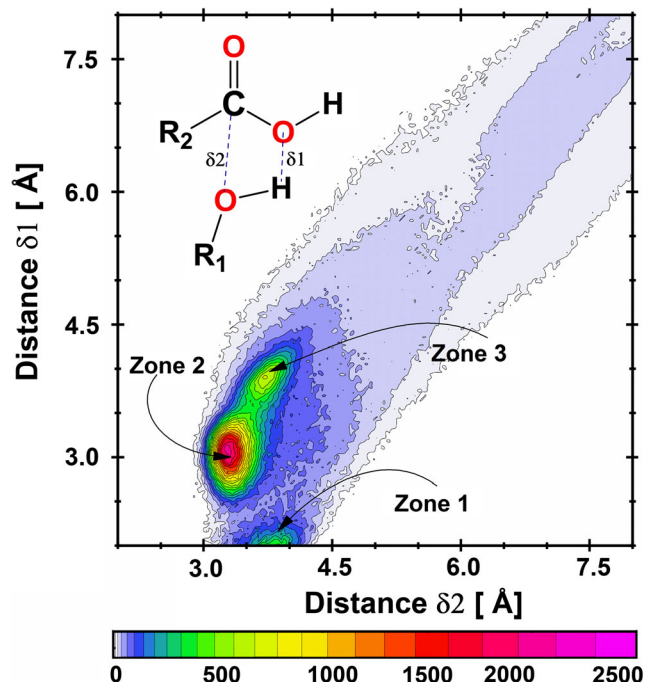
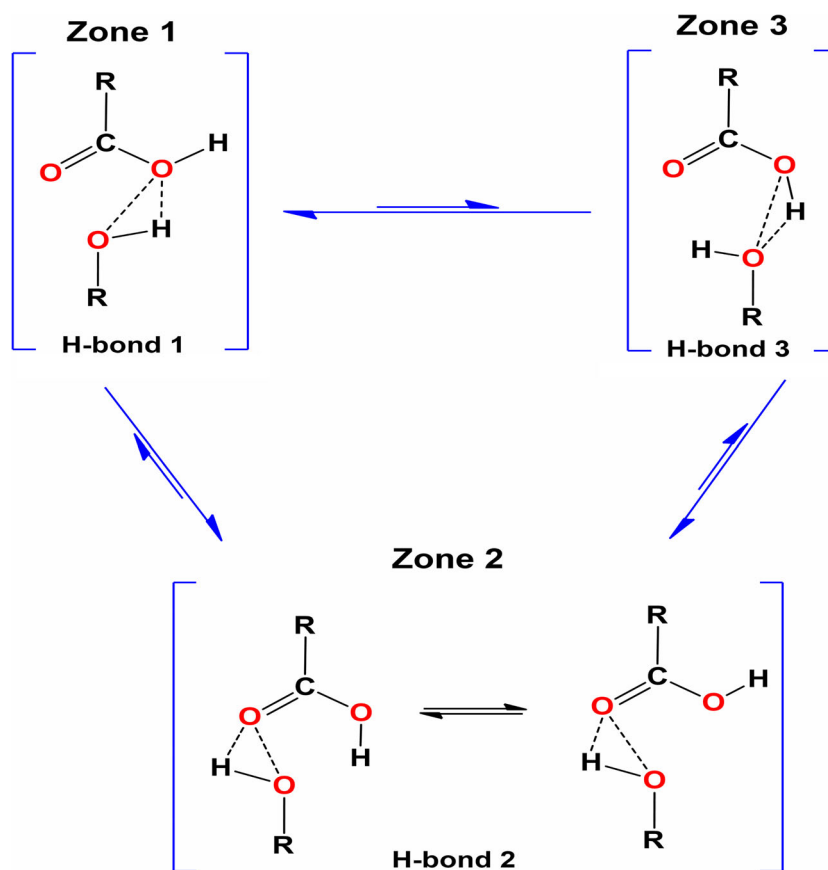


Fig. 5 Combined distribution functions (CDF) for bilayers L3 with two channels: δ_1 and δ_2 distances

Fig. 6 Schematic arrangements found when the $-\text{COOH}$ and $-\text{OH}$ groups interact with each other in ALE self-assembled bilayers (L3 structures in Fig. 3)



All these results show that to favor an ER it is necessary to meet two requirements: the hydrogen of the $-\text{COOH}$ group has to be oriented upwards, and also, H-bond interactions of type 1 (H-bond 1) must occur.

The presence of the three types of H-bonds (H-bond 1, H-bond 2, and H-bond 3) was confirmed by combined distribution functions (CDF). Specifically, radial and angular distribution functions (RDF and ADF) were used to monitor the presence of H-bond ($r_{\text{O-O}}$ and α) and the results are presented in Fig. 7.

Analyzing all possible H-bonds, it is clear that, effectively, all zones are stabilized by H-bond interactions ($r_{\text{O-O}} \leq 3.0 \text{ \AA}$ and $\alpha \leq 20^\circ$). Also, it was confirmed that the local minimum in zone 1 is the least stable by having a lower number of H-bonds (H-bond number ratios: H-bond2/H-bond1 ~ 2.5 and H-bond3/H-bond1 ~ 4.3). This result confirms that the local minimum that favors an ER (zone 1) is the least visited and is not the most stable region, making clear that an ER could be poorly favored.

ALE monolayer systems

MD simulations of ALE monolayers starting from initial configurations where molecules were vertically aligned or disorganized allowed us to assign the L1 and L2 structures observed in the AFM images from Fig. 3. Thus, L2 structures with higher height correspond to ALE self-assembled monolayers. Analyzing the density distribution along the z -direction

of the $-\text{COOH}$, POH, and SOH groups, it is observed that the way they distribute prevents any possibility of favoring an ER (Figs. 3b and S3). Therefore, it is expected that an ER does not occur in monolayers of vertically packed ALE molecules as deduced from AFM results [26].

On the other hand, the lower height structures (L1) can be explained from simulations of disorganized monolayers. In these systems (Figs. 3b and S4), the RDF results for both ER routes show similar behaviors (Fig. 8). Thus, $g_{\text{(C-O)}}$ has two peaks at 3.2 and 3.7 \AA and $g_{\text{(H-O)}}$ plot shows three peaks at 1.8, 3.0, and 3.8 \AA (Fig. 8a, b). All these peaks form three local minima that are most visited throughout the simulation trajectory (Fig. 4). Zone 1 (about $d(\text{C-O}) = 3.7 \text{ \AA}$ and $d(\text{H-O}) = 1.8 \text{ \AA}$) corresponds to the region that favors an ER, where the presence of the peak at $d(\text{C-O}) = 3.7 \text{ \AA}$ ensures a required approximation distance favorable to a nucleophilic attack. Also, the peak at $d(\text{O-H}) = 1.8 \text{ \AA}$ indicates a propitious distance for the formation of the water molecule.

Mixed systems

75:25 (ALE:PAL) composition

In a previous study [28], we analyzed the self-assembling process of mixtures of 75:25 composition, and we showed that the ALE molecules are still well organized when compared with the pure

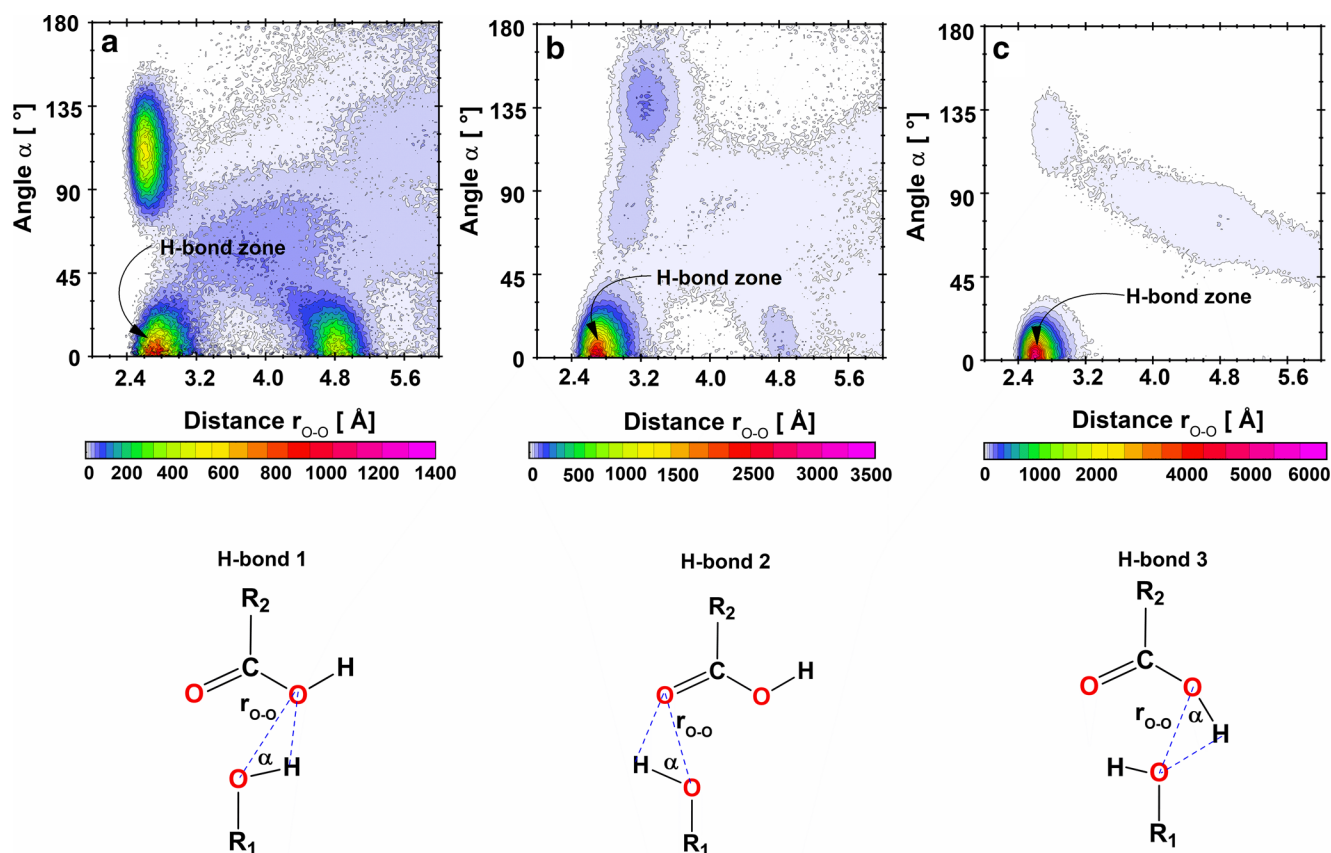


Fig. 7 Analysis of possible H-bond interactions between the -COOH and -OH groups using Combined Distribution Functions (CDF) with channels distance and α angle. (a) For H-bond 1. (b) For H-bond 2. (c) For H-bond 3. L3-ALE bilayer system

ALE monolayers. That work shows that the density profile along the z -direction, $\rho(z)$ of the active groups, shows a slight dislocation of the molecules that could favor interaction between the -COOH and SOH groups, both for segregated and dispersed systems. Also, $\rho(z)$ shows that there is not a convenient approach between the other reacting groups, and therefore, the interactions between -COOH and POH groups are hampered. Under these conditions, an ER could occur exclusively between the -COOH and SOH groups. These suggestions were also analyzed using the radial distribution functions (RDF), which are depicted in Fig. 9 for the H75D1 system. The other systems have the same behavior, and they are shown in the supplementary material (Fig. S5).

It can be seen that all studied systems (dispersed and segregated) for R2 route present characteristic peaks for $g_{(C-O)}$ and $g_{(O-H)}$ at 3.7 and 1.8 Å, respectively (Fig. 9b), whereas the absence of these peaks for R1 route (Fig. 9a) indicates that both groups are not arranged conveniently to favor an ER. These results imply that under high ALE concentrations, ER can only occur through the R2 route.

It was also observed that the -COOH groups of the ALE and PAL molecules exhibit different behaviors, indicating a different role in the ER through the R2 route. They were analyzed by defining the angle β formed between the z -direction and the normal vector to the plane formed by the

C-O-O atoms of the -COOH group. The β distributions are shown in Fig. S6. It is observed that even though the molecules remained confined in a compacted and orderly configuration, the -COOH groups, both in ALE and PAL molecules, can reorient in any possible orientation, mainly PAL -COOH groups.

The ALE -COOH groups have different orientations when compared between segregated and dispersed systems. The angles at 30° and 140° are more pronounced in the most stable (segregated) systems and slightly less in the dispersed ones. On the other hand, PAL -COOH groups are randomly distributed in all systems (Fig. S6). This fact suggests that PAL -COOH groups could be interacting to a lesser extent than ALE -COOH groups with other functional groups such as SOH. In ALE rich systems, the lower tendency of PAL -COOH to interact with SOHs of ALE molecules is compatible with the ideal behavior observed by AFM, i.e., a low degree of mixing and the occurrence of segregation between ALE and PAL phases [28].

25:75 (ALE:PAL) composition

In these systems, due to a large number of PAL molecules, the self-assembling of ALE molecules is almost entirely lost. The

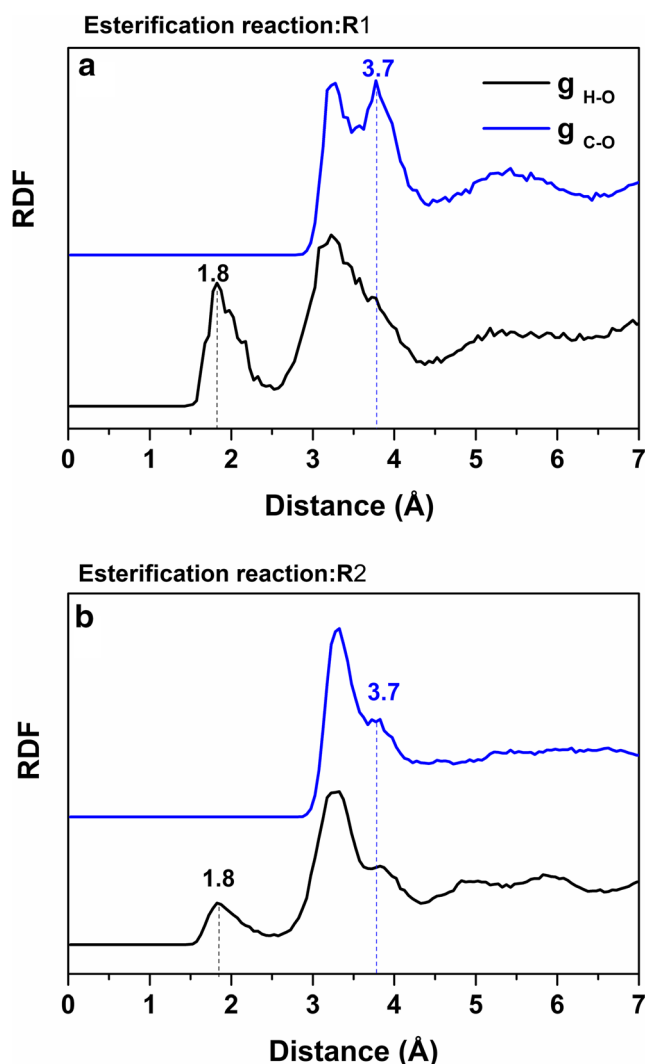


Fig. 8 Radial distribution functions (RDF) for interactions in the ALE monolayer L1 system between the following: **a** –COOH and POH groups (R1 route) and **b** –COOH and SOH groups (R2 route)

density profile $\rho(z)$ analysis shows that there is a dislocation of ALE molecules along the z -direction, facilitating interactions between all groups, that is, (–COOH)–(POH) and (–COOH)–(SOH) (Fig. S7). Again, RDF was used to check whether these interactions could effectively favor possible ER. The corresponding plots for the H25D2 system are reported in Fig. 10, whereas the other systems have the same behavior and are shown in the supplementary material (Fig. S8).

As expected, due to the increased disorder in the systems, an ER via R2 route is even more favored, as indicated from the intensity of the peak at 3.7 Å for $g_{\text{C-O}}$ (Fig. 10b). On analyzing the R1 route, unlike the previous case for 75:25 systems, it is now observed that ER also occurs via R1 route (Fig. 10a). This is evidenced by the peaks in $g_{\text{C-O}}$ and $g_{\text{O-H}}$ at 3.7 and 1.8 Å, respectively. Also, the higher height of the first peak and lesser overlapping with the close peak at 3.3 Å suggests that this route (R1) would be slightly more favored compared with the R2

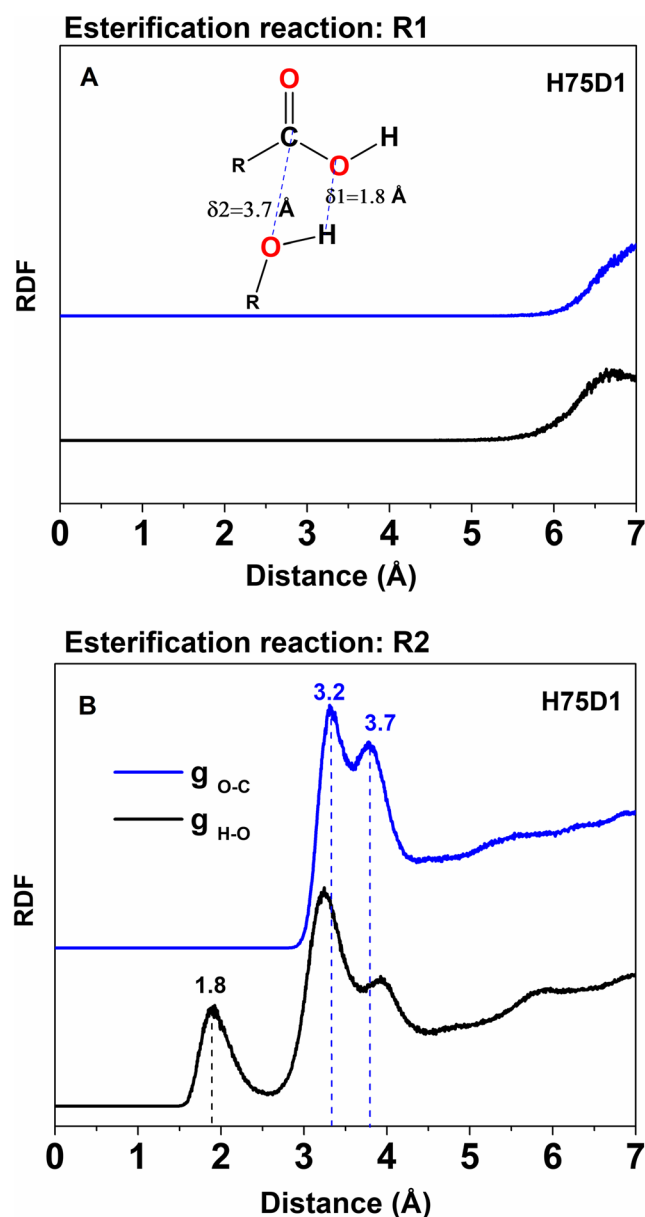


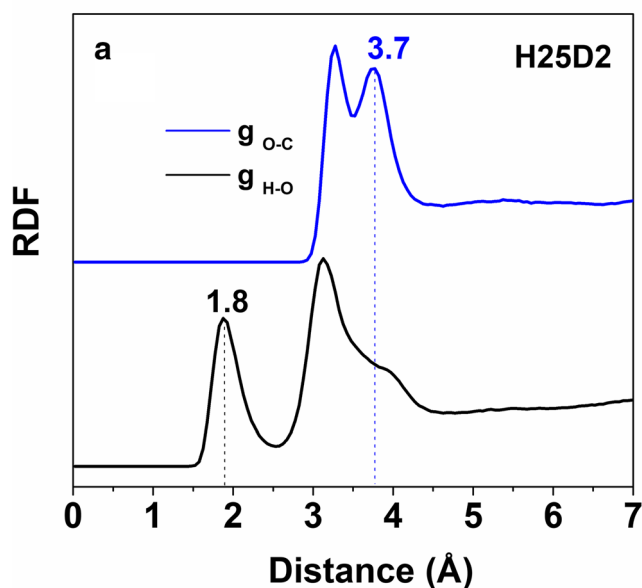
Fig. 9 Radial distribution functions (RDF) for interactions in the H75D1 system between the following: **a** –COOH and POH groups (R1 route) and **b** –COOH and SOH groups (R2 route)

route, only in disordered systems. The potentiation of both R1 and R2 mechanisms in PAL-rich mixtures would support the reported non-ideal behavior and the enhanced development of the mixed configuration in 25:75 (ALE:PAL) compositions, as revealed by AFM data. The nature of such a mixed phase (phase II) would be constituted by both pure assembled PAL and PAL-ALE interacting molecules [28].

Conclusions

Despite the limitations of classical molecular dynamics simulations to study chemical reactions, we present valuable

Esterification reaction: R1 route



Esterification reaction: R2 route

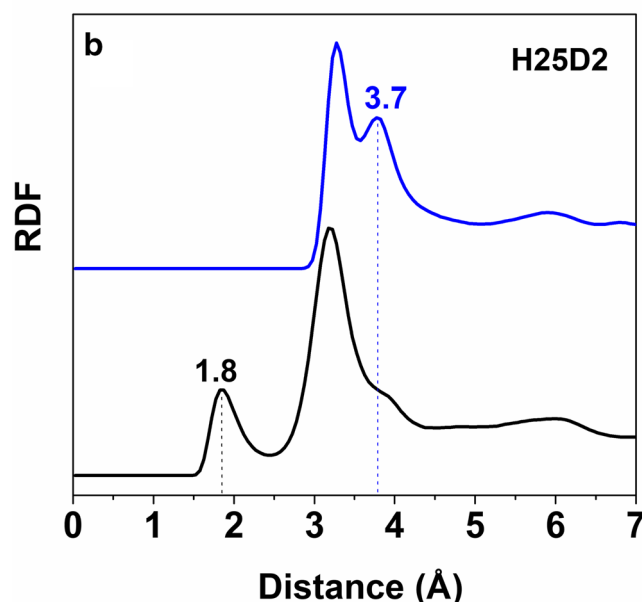


Fig. 10 Radial distribution functions (RDF) for interactions in the H25D2 system between the following: **a** –COOH and POH groups (R1 route) and **b** –COOH and SOH groups (R2 route)

parameters allowing us to infer possible routes for ER occurring when pure ALE and ALE/PAL mixtures are deposited on a support.

Systems formed from some cutin monomers have the ability to form self-assembled structures. The loss of ability for self-assembling of these systems leads to favoring esterification reactions, and as the disorder increases, the number of possible esterification routes increases.

For example, the 75:25 system keeps significantly a well-organized structure and the ER is only favored through R2

route (–COOH and SOH interactions). For the 25:75 ALE system, having lost all its self-assembly ability, it is observed that both R1 and R2 routes are favored. In pure ALE systems, it becomes evident that self-assembled monolayers do not have the ability to form ER. However, when monolayers are disorganized, the R1 and R2 routes are favored. Finally, for bilayer pure ALE systems, where the system is well organized, only the R1 route is favored due to interactions between –COOH from the upper layer with the POH in the lower layer.

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Authors’ contributions All authors contributed equally to this work.

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Data availability The data that supports the findings of this study are available within the article [and its [supplementary material](#)].

Compliance with ethical standards

Conflict of interest The authors declare that they have no conflict of interest.

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