Theoretical Insight into the Mechanism of an Efficient L-Proline-catalyzed Transamidation of Acetamide with Benzylamine

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The detailed mechanisms of the efficient L-proline and pyrrolidine catalyzed transamidation of acetamide with benzylamine have been investigated using density functional theory (DFT) calculations. Our calculated results show: (1) the mechanisms of two catalytic cycle reactions are similar. However, the rate-determining steps of their reactions are different for the whole catalytic process. One is the intramolecular nucleophilic addition reaction of 1-COM, the other is hydrolysis reaction of 2-C. (2) COOH group of L-proline is essential for efficient transamidation. The computational results are in good agreement withthe experiment finding and mechanism resported by Rao *et al.* for L-proline-catalyzed synthesis of amidesin good to excellent yields.

Key Words: Transamidation, L-Proline-catalydst, Pyrrolidine-catalydst, Acetamide, Benzylamine

Introduction

The amide bond is arguably one of the most important gifts in nature, as it is essential to sustain living system and exits in all natural peptides and protein structures. Amides are also often used as synthetic intermediates for the preparation of natural products, pharmaceuticals molecules, agrochemicals, polymers, etc.^{2,3} Amides are surprisingly robust compared to their related derivatives due to the structural characteristics of amide bonds. Therefore, there has been much effort aimed at the development of efficient syntheticmethodologies for improving the preparation of amides. In general, amides are prepared by the coupling of amines with carboxamides, alcohols, aldehydes, carboxylic acid derivatives⁶ etc., or by the reaction of primary amides with aryl and alkyl halides.7 Alternatively, the transamidation process is an attractive tool in the field of synthetic organic chemistry and represents an excellent unusual route for the functionalization of a given carboxamide. 8 So far, elegant methods for transamidation reactions (i.e., transamidation processes catalyzed by copper,9 cerium,10 hydroxylamine hydrochloride,¹¹ ammonium bromide,¹² and boric acid,¹³ hypervalent iodine¹⁴) have been reported. Unfortunately, these methods to accomplish the desired amide exchange process suffer from certain demerits² (i.e. the excessive use of an expensive or waste-generating activation reagents, reversible reactions giving mixtures of amides). In view of environmental friendliness, it is desirable to develop benign and metal-free transamidation procedures with high yield and selectivity.15

Recently, Rao and his co-workers¹⁶ reported a novel solvent-free L-proline-catalyzed transamidation of carbox-amides and amines. In their experiment, acetamide and benzylamine were selected as a model system. In the presence of L-proline (1-L) (10 mol %), the reaction was carried

out in a sealed tube at 100 °C for 36 h and proceeded smoothly to lead to the formation of the desired transamidation product P in 99% yield. When the reaction was performed in the pyrrolidine (2-L) (10 mol %), an 10% conversion was observed (see Scheme 1). They concluded the L-proline is inexpensive, readily available and conveniently. What's more, it could be efficiently applied to transamidation of a wide range of amines containing benzylic, aromatic, aliphatic, propargylic, and heteroaromatic. The proposed mechanism by Rao is shown in Scheme 2.

To the best of our knowledge, there are no theoretical studies for the mechanism of the L-proline catalyzed transamidation process reported by Rao *et al.* Hence, we report a thorough density functional theory (DFT) computational investigation on the mechanisms of the L-proline and pyrrolidine-catalyzed transamidation of acetamide (R1) with benzylamine (R2) based on the experimental evidence reported by Rao *et al.* In the present study, our aim is to understand the mechanism details of the reactions at the molecular level by providing the geometric and energetic information of the intermediates and transition states involved.

Scheme 1. Rao's L-proline and pyrrolidine-catalyzed transamidation of acetamide with benzylamine.

Scheme 2. Sketch of the catalytic cycle for the transamidation of acetamide with benzylamine.

Computation Methods

All the calculations were performed using the Gaussian 03 program package.¹⁷ The geometries of the reactants, complex, intermediates, products and transition states were fully optimized by hybrid density functional theory (DFT)¹⁸⁻²⁰ using the 6-31g(d, p)²¹ basis set. This hybrid DFT method has generally been shown to perform quite well in the mechanistic studies of transition metal-or non-transition metalcatalyzed reactions.²² The 6-31G(d, p) basis set has been popularly used in the mechanistic studies of Au-catalyzed reactions and is shown to be quite reliable for optimizing both the geometries and energies.²³ Vibrational frequencies calculated were used to further identify all of the stationary points as minima (intermediates (zero imaginary frequencies) or transition states (one imaginary frequencies)), zero-point energy (ZPE). The calculations of intrinsic reaction coordinate (IRC)²⁴ were performed to further confirm that the transition states connect between designated intermediates. The relative energies were thus corrected for the vibrational zero-point energies (ZPE, not scaled). In order to obtain more reliable energies of various structures, the B3LYP/6-31G (d, p) optimized geometries were used for single-point calculations at MP2²⁵/6-31g(d, p) level. Therefore, we expect that the MP2//B3LYP calculations reported in this work are reliable.

Results and Discussion

In this sections, we present the calculated results for transamidation reaction of acetamide (R1) with benzylamine (R2) catalyzed by the L-proline and pyrrolidine according to the proposed catalytic cycle shown in Scheme 2, respectively.

The Transamidation Reaction Catalyzed by L-Proline. We first discuss the mechanism of L-proline-catalyzed transamidation reaction of R1 with R2. The optimized geometries with selected key geometry parameters for the reactants, complex, intermediates, transition states and product(P) are given in Figure 1. The potential energy surface profiles for the transamidation reaction are depicted in Figure 2. It should be noted that the energy of R1 + L-proline (1-L) is set at zero as a reference for other species.

From the energy profile we can find the first step of this mechanism starts from the L-proline (1-L) catalyst approaching to one of the reactants (R1), forming a weakly-bound complex 1-COM (-5.71 kcal/mol, ΔE), which is lower than the reactants (R1+1-L). Subsequently, there is nucleophilic attack of N¹- atom of the L-proline (1-L) on the positively charged C9- atom of R1 to lead to the formation of the intermediate 1-IM (-0.16 kacl/mol) via surmounting a 38.24 kcal/mol energy barrier, transition states 1-TS1 (32.53 kcal/ mol). During the process of generating 1-IM, the distance N¹-C⁹ is gradually shorten. As shown in Figure 1, the distances N¹-C⁹ in 1-COM, 1-TS1 and 1-IM is 3.664 Å, 1.695 Å and 1.458 Å, respectively. Furthermore, the C⁹–O¹ bond gradually strenches and losts its double bond character (1-COM: 1.230 Å, 1-TS: 1.343 Å, 1-IM: 1.434 Å). The fact that is L-proline (1-L) catalyst shows some activity and is sufficiently efficient. And at the same time, the H¹ atom is migrating from the N¹ to O¹. From the intermediate 1-IM, there are two different reaction pathway a and b, both of which involve in different intramolecular H-migration reac-

In the pathway a, the 1-IM (-0.16 kcal/mol) firstly isomerizes to another intermediate 1-IM* (4.44 kcal/mol) for dehydration. And then, the 1-IM* transforms to intermediate 1-A (3.09 kcal/mol) *via* 1-TS2 (16.16 kcal/mol) by the elimination of H₂O molecular with a barrier of 11.72 kcal/mol, followed by a conjugated Π-bond (N-C-N) forming, which makes 1-A lie only 3.25 kcal/mol above the 1-IM. It should be noted that the intermediate 1-IM lies lower than that of 1-IM*. In 1-IM*, it is evident that the N¹-C³ is significantly stronger than that of 1-IM as demonstrated by their bond lengths of 1.425 and 1.458 Å, respectively. This can give a good explanation of the intermediation 1-IM* lies 4.6 kcal/mol above 1-IM.

On the other hand, as shown in the pathway b of Figure 2. 1-IM converts to 1-A though another transition state 1-TS2* (50.08 kcal/mol). In 1-TS2*, The H^3 atom is migrating from N^2 to O^1 . The N^2 - H^3 and H^3 - O^1 bones are 1.139 Å and 1.449 Å, respectively. The energy barrier of this process is 50.24 kcal/mol. By comparing the H- atom migrations of 1-TS2 with 1-TS2* in Figure 1, one can find the energy barrier of pathway a is obviously different from that of pathway b. This may be due to different ability of the H atom donors, COOH and NH_2 group. According to the difference of the two energy barrier heights, we can rule out the pathway b is not the best path.

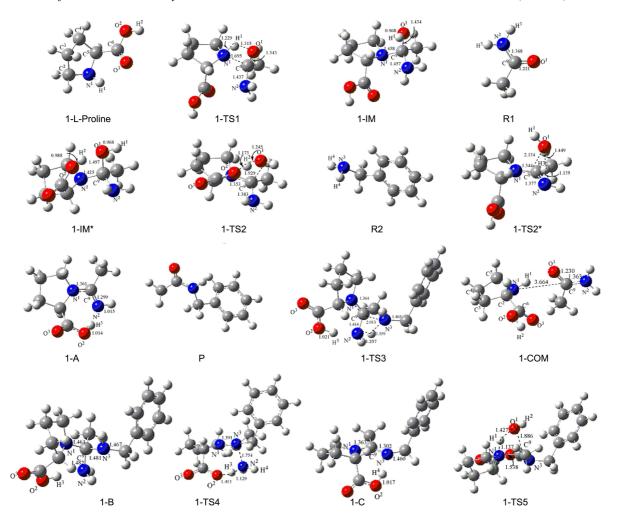


Figure 1. Optimized structures for the species involved in the L-proline-catalyzed (1-L) transamidation of acetamide with benzylamine. Bond distance are given in Å.

The nucleophilic attack of N³- atom of another reactants (R2) on the positively charged C⁹ of 1-A leads to the formation of a new intermediate 1-B through 1-TS3. The formation of 1-B is an exothermic process (The energy of reaction for the 1-B was -5.63 kcal/mol with respect to 1-A, as shown in Figure 2). The activation energy of this step is 35.0 kcal/mol. Subsequently, 1-B can evolve into 1-C via 1-TS4, a transition state of the NH₃ molecular elimination. From the optimized geometrical parameters of 1-TS4, it is apparent that H³ atom is moving from O² to N² atom. As shown in Figure 2, the energy barriers of the process is 12.91 kcal/mol and the formation of 1-C is an endothermic process (The energy of reaction for the 1-C was 2.03 kcal/mol with respect to 1-B, as shown in Figure 2). In 1-C, the N¹-C⁹ and N³-C⁹ bonds are shorten to 1.363 Å and 1.302 Å, respectively. Clearly, the N3-C9 bond becomes double bond again and the length of the N¹-C9 bond between single and double bonds. There is subsequent hydrolysis reaction of 1-C to result in the formation of the desired product P (-5.30 kcal/ mol, lower than reactants (R1+1-L)) and regeneration of the catalyst (1-L) through 1-TS5. Figure 2 shows that the energy barrier of this step $(1-C\rightarrow 1-TS5\rightarrow P)$ is calculated to be

35.55 kcal/mol. This process is exothermic by 4.79 kcal/mol. In total, the whole catalytic process is exothermic by -5.30 kcal/mol, lower than reactants (R1+1-L). It is clear that the intramolecular nucleophilic addition reaction of 1-COM is unfavorable kinetically. Notably, this step is rate-determining for the whole catalytic process.

According to the Figure 2, it can be directly seen that there are two key energy-demanding steps in the catalytic cycle reaction. One is the intramolecular nucleophilic addition reaction of 1-COM *via* 1-TS1 with a barrier of 38.24 kcal/mol. The other is the hydrolysis reaction of 1-C with a barrier of 35.55 kcal/mol. It is the reason why the experimental was carried out at a higher temperature (100 °C) for 36 h. Futhermore, the energy barriers of the two steps labeled with the blue line (11.72 kcal/mol and 12.91 kcal/mol, respectively) are very lower than other reaction steps in the Figure 2, indicating that COOH group of L-proline is essential for efficient transamidation, which is consistent with the experiment finding proposed by Rao *et al.*

The Transamidation Reaction Catalyzed by Pyrrolidine. Then, we extend our study to the transamidation reaction catalyzed by pyrrolidine (2-L) (see Scheme 1). The calculated

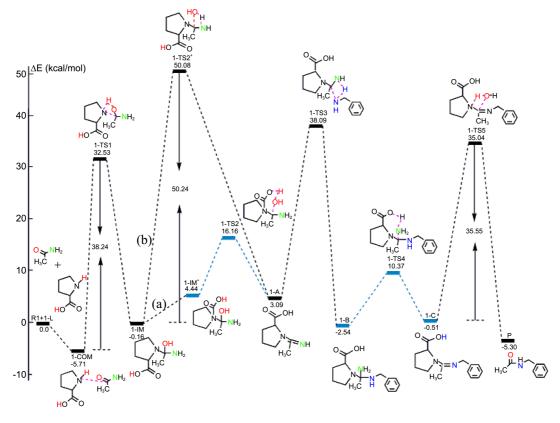


Figure 2. The potential energy surface (PES) for L-proline-catalyzed transamidation reaction of acetamide and benzylamine with MP2/6-31G (d, p).

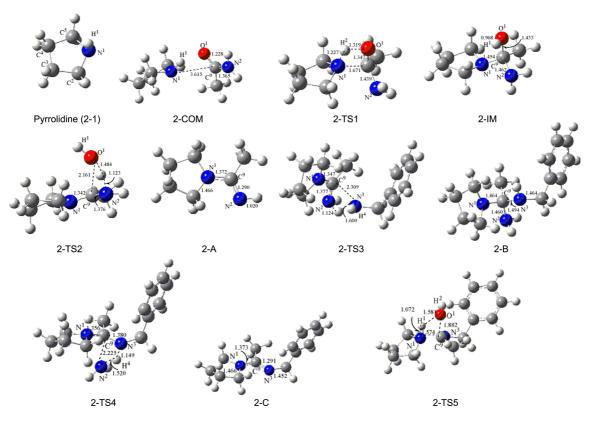


Figure 3. Optimized structures for the species involved in the transamidation reaction catalyzed by pyrrolidine (2-L) of acetamide with benzylamine. Bond distance are given in Å.

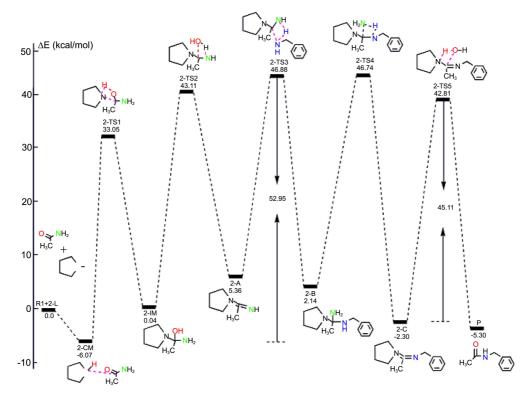


Figure 4. Potential energy surface of pyrrolidine-catalyzed (2-L) transamidation reaction of acetamide and benzylamine with MP2/6-31G (d, p).

mechanism detail is very similar to the those discussed above, so we only describe the catalytic cycle and the relative energy surface profile briefly. The optimized geometries with selected key geometry parameters are depicted in Figure 3. The calculated relative energy surface profiles are given in Figure 4. Note that the energy of R1 + pyrrolidine (2-L) is set at zero as a reference for other species.

At the entrance of the reaction, the reactants (R1+2-L) firstly form a weakly-bound complex 2-COM though barrierfree exothermic reaction of $\Delta E = 6.07$ kcal/mol. Then, COM undergoes an intramolecular nucleophilic addition to form 2-IM via 2-TS1 with a energy barrier of 39.12 kcal/mol, which only higher 0.88 kcal/mol than the corresponding value to form 1-IM. After that, 2-IM will proceed with an intramolecular H-migration through 2-TS2 to 2-A. According to Figure 4, the barrier for the step of the forming 2-A is 43.07 kcal/mol, which is much higher than the corresponding value (16.32 kcal/mol) for the step of forming 1-A. In the following step, 2-A undertakes an intermolecular nucleophilic addition with R2 to form 2-B via 2-TS3. The energy of activation for this step is calculated to be 41.52 kcal/mol for 2-TS3. Moreover, the barriers to form 2-C through 2-TS4 with a second intramolecular H-migration is 44.60 kcal/mol, which is much larger than the corresponding value (12.91 kcal/mol) for the step of forming 1-C. The barriers of the final step, hydrolysis reaction from 2-C to final product P, is 45.11 kcal/mol. Clearly, the higher barrier calculated for 2-TS5 indicates that this step is rate-determining for the whole catalytic process, and every step of the catalytic cycle reaction is high energy-demanding. But above all, the overall energy barrier of the catalytic reaction is calculated to be 52.95 kcal/mol. That can explain the low conversion, the high temperature and rate of the pyrrolidine-catalyzed transamidation reaction of acetamide and benzylamine.

Conclusion

In summary, the mechanisms of the L-proline and pyrrolidine-catalyzed transmidation process of acetamide and benzylamine have been investigated by using DFT calculation. According to our calculations, the rate-determining steps of the two catalytic cycle reactions are different for the whole catalytic process although the mechanisms of both of the reactions is similar. The former is the intramolecular nucleophilic addition reaction of 1-COM *via* 1-TS1 with a barrier of 38.24 kcal/mol. The latter is hydrolysis reaction of 2-C with a whole barrier of 45.11 kcal/mol and every step of its is high energy-demanding. This may be the reason why the conversion of the former is obviously higher than that of the latter. In addition, COOH group of L-proline is favored for efficient transamidation, which are consistent with the experimental observations of Rao *et al.*

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