DFT studies of the conversion of four mesylate esters during reaction with ammonia

Andrzej Nowacki · Karol Sikora · Barbara Dmochowska · Andrzej Wiśniewski

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Abstract The energetics of the Menshutkin-like reaction between four mesylate derivatives and ammonia have been computed using B3LYP functional with the 6-31+G** basis set. Additionally, MPW1K/6-31+G** level calculations were carried out to estimate activation barrier heights in the gas phase. Solvent effect corrections were computed using PCM/B3LYP/6-31+G** level. The conversion of the reactant complexes into ion pairs is accompanied by a strong energy decrease in the gas phase and in all solvents. The ion pairs are stabilized with two strong hydrogen bonds in the gas phase. The bifurcation at C2 causes a significant activation barrier increase. Also, bifurcation at C5 leads to noticeable barrier height differentiation. Both B3LYP/6-31+G** and MPW1K/6-31+G** activation barriers suggest the reaction 2 (2a+NH₃) to be the fastest in the gas phase. The reaction 4 is the slowest one in all environments.

Keywords Ammonium salts · DFT calculations · Menshutkin-like reaction · Nucleophilic substitution · THF conformation

Introduction

Menshutkin was the first scientist to describe the reaction leading to the formation of quaternary ammonium salts (QASs) [1]. In Menshutkin’s classical method (the Menshutkin reaction, MR) an alkyl halide (an electrophile) is treated with a tertiary, heterocyclic amine (acting as a nucleophile). The MR has been used to obtain QASs from different classes of organic compounds. Twenty years after Menshutkin’s original work, Fisher and Raske proved its usefulness for the synthesis of N-glycopyranosyl quaternary salts [2].

Detailed studies of the MR have answered the questions regarding both the mechanistic aspects of this reaction and the experimental conditions required for it to occur. Now we have a profound understanding of the factors influencing this reaction: the solvent, the nucleophile and the leaving group. It has been established that polar solvents stabilize both the transition state and the ionic products, thereby speeding up the reaction. In contrast to polar solvents, the MR is dramatically retarded in less polar media.

The alkyl halide can be replaced by a sulfonate ester, which leads to a reaction analogous to the classical MR. This kind of reaction has been used to synthesize N-glycoammonium and N-glycopyridinium tosylates [3–7].

The classical MR has been the subject of extensive theoretical study [8–20]; at first, the reaction between ammonia and methyl halide was mostly explored in this way. Such a simple model enables calculations to be performed in both the gas phase and in solution. Recently, however, thanks to the increase in computational efficiency, a significantly extended model has been used for the calculations [19].

Our group has long been interested in the synthesis of quaternary aminium salts [3–6] via the halide and sulfonate ester derivatives of monosugars and alditols. In recent years we have been concentrating on the theoretical aspects of the reaction between sulfonate esters and tertiary amines, both aliphatic and heterocyclic [21–23]. In continuation of our theoretical studies, we now present the results concerning the ammonia-assisted conversion of four mesylate derivatives. Surprisingly, no theoretical studies of the reaction between ammonia and sulfonate esters have yet been carried out, in contrast to the classical MR. The structures studied and the IUPAC names of the three mesylates

A. Nowacki (✉) · K. Sikora · B. Dmochowska · A. Wiśniewski
Faculty of Chemistry, University of Gdańsk, Sobieskiego 18, 80-952 Gdańsk, Poland
e-mail: anowacki@chem.univ.gda.pl

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are shown in Fig. 1. The salts under discussion are analogues of (+)-muscarnine, the principal alkaloid in some poisonous fungi. There has been a resurgence of interest in muscarine in recent years following the discovery of a relationship between a cholinergic deficit and the pathology of Alzheimer’s disease [24]. We also did the calculations for the conversion of methyl mesylate. Our aim was to produce a kinetic and thermodynamic description of this type of modified MR. As before, we also wanted to assess the influence of the branching three bonds distant from the reaction center (the methyl or methoxy group bound to C5 of the THF ring is cis-oriented in relation to C1). With this work we have completed the review of different nucleophilic agents typically used in the classic variant of the MR.

Computational details

All the calculated structures were prepared in the MOLDEN program [25]. The geometries were then optimized using density functional theory (DFT) based on Becke’s three-parameter hybrid exchange functional [26] involving the gradient-corrected correlation functional of Lee, Yang and Parr [27], with the split-valence basis set including polarized and diffuse functions [28, 29] — (B3LYP/6-31+G** method). Optimizations for reactant complexes and transition states were additionally done using the MPW1K/6-31+G** method [30]. This functional (Pedrew-Wang 1-parameter model for kinetics, MPW1K) gives remarkably accurate activation state. The energy difference between optimization barrier heights. The optimization was considered satisfactory if the energy difference between optimization cycles was less than 1×10−6 Hartree and a gradient of <1×10−4 a.u. was achieved. The convergence of all the systems studied was checked by harmonic vibration analysis. No imaginary frequencies were observed for the ground state and there was only one for the transition state.

Solvent effects were included in the calculations employing the self-consistent reaction field SCRF-PCM solvation model [31] for the reactions studied in chloroform (ε=4.9) and water (ε=78.39) at the B3LYP/6-31+G** level. Implicit solvent calculations imply the generation of a vacuum cavity inside a continuous and homogeneous dielectric field. In PCM model the cavity is built up by a series of interlocking atomic spheres. We used UA0 with scale factor alpha=1.2 for water and 1.4 for chloroform [18]. Individual spheres were centered on acidic hydrogen atoms.

All DFT calculations were done with the aid of the Gaussian 03 program [32].

Results and discussion

General characteristic of the reaction pathway

The studied reactions together with atom numbering order are shown in Scheme 1. The atom numbering order presented in Scheme 1 is not compatible with the names shown in Fig. 1 and IUPAC recommendations, but we use it in order to make the presentation of our results clearer.

Reaction 1 is closely related to the classical MR, differing only in the leaving group. We carried out calculations for this reaction to compare the influence of leaving group exchange on the reaction.

The next two reactions (2 and 3) involve anhydroalditols, whereas the last one (4) relates to the glycoside. The spatial arrangement of the aglycone (here, the OCH3 group) in relation to the THF ring in glycosides is governed by the steric factor and by the so-called exo-anomeric effect. This effect forces the aglycone to be located in a position where an acute torsion angle is formed in relation to the endocyclic oxygen atom. This requirement is satisfied in two orientations: −sc and +sc (Fig. 2), but the steric handicap present in the +sc conformation disqualifies it, so the preferred one is −sc.

The H2C–O5–C5–O2 torsion angle ranges from −65° to −82° all along the reaction pathway (Table 1). However, this is not a golden rule, since we showed previously that in the reaction of mesylate derivative 4a with a bulky trimethylamine the −sc orientation of the methoxyl group appeared to be unfavorable because of steric hindrance and thus changed to ap [21].

The steps of the reactions under discussion are analogous to those reported previously [21–23]. The energy (E0) and pseudochemical potential (U0) [33] profiles for the conversion of the mesylates into the corresponding ammonium salts (Fig. 3) in the gas phase and in solvents consists of an asymmetric double-well potential with five stationary points, corresponding to the separated reactants (R), reactant complex (RC), transition state (TS), product complex (PC) and separated ionic products (P). Two reactants (electrophile — mesylate derivative, and nucleophile — ammonia) approach one another, forming a van der Waals reactant complex. This complex converts into an ionic pair, which requires an activation barrier to be overcome. Finally, the constituents of the ionic pair are separated to an infinitely great distance. The main difference between the energy diagrams shown in Fig.3 and those presented elsewhere [21–23] is on the product side. Previously, the last stage of the reaction was endergonic in the gas phase and in chloroform, whereas it was exergonic in polar solvents. In the reactions being studied here, separation of the ion pair constituents requires the application energy in the gas phase and in all solvents.
Gas-phase calculations

Table 1 lists the activation and reaction energies for both the gas state and solutions, together with important geometrical parameters of all the stationary points on PES. A scaling factor of 0.9877 was used for the zero-point vibrational energy (ZPVE) correction of the calculated total energies [34].

To discuss the conformational details of the THF ring, the Altona–Sundaralingam (AS) pseudorotational phase angle ($P$) and the AS puckering amplitude ($\phi_m$) were considered [35, 36]. The THF ring conformations, $P$ and $\phi_m$ values are given in Table 2 together with the set of the endocyclic torsion angles $\phi_0$–$\phi_4$, whereas the definition of these angles is shown in Fig. 4. The conformational descriptors that we adopted differ from the classical ones because of the different atom numbering scheme. Table 2 also lists two torsion angles ($\chi$) describing the spatial disposition of the exocyclic groups.

The calculated geometries together with selected bond distances, valence angles and relative energies corresponding to all the stationary points along the reaction pathway are presented in Figs. 5 and 6. The relative energies refer to the sum of the separate reactant energies.

The first point on the energy curves (Fig. 3) corresponds to the separate reactants, i.e., mesylate derivative and ammonia. In the case of reaction 2 (Table 2) the THF ring takes the $E_4$ conformation, ($P=349^\circ$, $\phi_m=36^\circ$), in which C1 atom is in the pseudo-equatorial position ($\chi_1=-144.5^\circ$). The same ring conformation is found for the individual mesylate in reaction 4, ($P=336^\circ$, $\phi_m=36^\circ$), whereas the $4E$ conformation is observed for the separate
mesylate in reaction 3 (\(\phi_m = -36^\circ\)). The \(E_4\) conformation is free of 1,3-diaxial-like steric interactions and the eclipsed orientation of the substituents, as we described previously [23]. The mesylate C1 atom in reaction 4 is in the pseudo-equatorial orientation (\(\chi_1 = -138.5^\circ\)) whereas the OCH$_3$ group is in the pseudo-axial position (\(\chi_2 = 84.6^\circ\)). Thus the THF ring avoids the unfavorable 1,3-diaxial-like steric interactions between these two groups. The THF ring in reaction 3 bears two bulky substituents on the same side of the ring; hence, two reverse conformations should be considered, namely, those in which the C1 or C6 atom is in the pseudo-equatorial orientation (the \(3^E\) or \(4^E\) conformation respectively). Both conformations have features thought to stabilize five-membered rings because they hold two bulky substituents on the same side of the ring; hence, two reverse conformations should be considered, those in which the C1 or C6 atom is in the pseudo-equatorial orientation (the \(3^E\) or \(4^E\) conformation respectively). Both conformations have features thought to stabilize five-membered rings because they hold two bulky substituents on the same side of the ring; hence, two reverse conformations should be considered, namely, those in which the C1 or C6 atom is in the pseudo-equatorial orientation (the \(3^E\) or \(4^E\) conformation respectively).

The next point on the energy diagrams represents the reactant complex. In all the cases studied, the approach of the individual reactants is accompanied by a slight energy decrease (Table 1, Figs. 5 and 6) in the gas phase. Gibbs free energies, however, predict that reactant complex formation will be unfavorable (\(\Delta G = 5.0 - 6.2 \text{ kcal mol}^{-1}\)).

The small value of the complexation energy, \((-2.9 \text{ kcal mol}^{-1}, -2.2 \text{ kcal mol}^{-1}, -2.1 \text{ kcal mol}^{-1}\) and \(-2.4 \text{ kcal mol}^{-1}\) for 1a, 2a, 3a and 4a respectively) indicates that the interaction (hydrogen bond) between these two molecules in this complex is relatively weak. The intermolecular interaction between mesylate and ammonia in reactant complexes is slightly stronger than the one for trimethylamine [21], and comparable with that for pyridine [22, 23]. This may be related to the type of intermolecular interaction. A typical S–O⋯H–N hydrogen bond stabilizes the reactant complex in these reactions, whereas an atypical S–O⋯H–C hydrogen bond interaction was observed for the reactions with trimethylamine [21]. After decades of the controversy on whether O⋯H–C hydrogen bonds really exist, it is accepted now that they do exist, although O⋯H–C hydrogen bonds are mostly weak in comparison with classical ones [37].
The shape of the THF ring does not change while reactants are approaching one another. The $E_4$ conformation is observed for THF ring in reactant complexes 2a and 4a, whereas the $^4E$ conformation is noted in 3a (Table 2). A noteworthy fact is that in reaction 4, two different conformations ($E_4$ for the separate mesylate and $^2E$ for the reactant complex) were preferred for the THF ring when the conversion with pyridine was being studied and described [23].

The next stationary point on the energy curve, shown in Fig. 3, corresponds to the transition state. The relative energy values matching the transition states with respect to the separated reactants are given in Table 1 and in Figs. 5 and 6, whereas the activation barriers relating to the reactant complexes are listed in Table 3.

The geometry of the relevant transition state can be characterized by the C···O and C···N distances, OCN valence angles and two torsion angles (A, B, Table 1). Whereas the C···O and C···N distances are roughly the same in all TSs (about 2.06 Å and 2.08 Å respectively), the C···N distance in reaction 1 is much shorter (1.93 Å).

Table 1 and Fig. 5 indicate that the transition state for reaction 1 is linear, the valence angle O···C···N being 178.8°. The other three TSs are slightly bent, the cited valence angle ranging from ~156° to ~164°, which is the result of the steric hindrance.

The approach of the ammonia molecule to C1 induces a change in the O–C1–C2–C3 torsion angle (Table 1). In the reactant complex this angle is about ~65°, whereas in the transition state it takes values of ~13.0°, 0.4° and ~15.9° for reactions 2, 3 and 4 respectively. The observed differences in values of the torsion angle come from the variation of the THF ring conformation. The preferred conformation of the THF ring in the transition state corresponding to conversions 2
and 4 is $^3T_2 (P = 2^\circ, \phi_m = 39^\circ$ and $P = 358^\circ, \phi_m = 40^\circ$ for 2 and 4 respectively, Table 2). This conformation is free of the eclipsed orientation of the substituents and 1,3-diaxial-like steric interactions (in reaction 4) as C1 is moved away from the THF ring ($\chi_1 = -154^\circ$, Table 2). In consequence, the MSO leaving group is also located beside the THF ring. In reaction 3 the THF ring adopts the $^4T_3$ conformation ($P = 174^\circ$, $\phi_m = -37^\circ$), in which the C6 atom is moved away from the ring ($\chi_2 = 153^\circ$, Table 2). In this conformation the MSO leaving group is located above the THF ring.

The C2–C1–H1s–H1t deformation torsion angle (B, Table 1) reflects the planar placing of substituents at the C1 atom in the transition state. This indicates that the transition state geometry is exactly halfway between the reactant and product complex for reactions 2, 3 and 4, whereas the late transition state is observed for reaction 1.

The calculated barriers are higher than those for the reactions with trimethylamine [21] and pyridine [22, 23], which corresponds to the lower basicity of ammonia in a vacuum (Fig. 7). The proton affinity of ammonia is 204.0 kcal mol$^{-1}$, but is 225.1 kcal mol$^{-1}$ and 220.8 kcal mol$^{-1}$ for trimethylamine and pyridine respectively [38]. The energy barrier is the lowest for reaction 2 but the highest for reaction 4. Interestingly, the energy barrier for reaction 1 is higher than for reactions 2 and 3 according to both B3LYP and MPW1K methods. This stands in contrast to the reactions of the same mesylate derivatives with other nucleophiles studied earlier [21–23], where the barrier was the lowest for the reaction of methyl mesylate with the corresponding nucleophile (Fig. 7). Presumably, hydrogen bond formation between the endocyclic oxygen atom and the hydrogen atom attached to the nitrogen atom stabilizes the transition state geometry. Such an interaction cannot occur in reaction 1.

The second minimum on the reaction pathway corresponds to ion pairs. Both $\Delta E$ and $\Delta G$ predict that the conversion of the reactant complexes to the respective ion pairs is accompanied by an energy decrease in the gas phase. This effect is smallest in reaction 1 and the energy of the ion pair 1b is ca 10 kcal mol$^{-1}$ lower than that of the reactant complex. Interestingly, $\Delta E$ and $\Delta G$ at this stage were expected to be unfavorable for the reaction of methyl mesylate and pyridine [22]. On the other hand, ion pair formation was slightly favorable ($\Delta E = -1.5$ kcal mol$^{-1}$) when trimethylamine was the nucleophile [21]. The conversion of complexes into ion pairs is rather more favorable for the remaining three reactions (about 12 kcal mol$^{-1}$) than for reaction 1. The constituents of ion pair 1b are oriented in such a way that the whole geometry has the Cs symmetry (Fig. 8), with two carbons, oxygen, sulfur and nitrogen being planar. The anion and the cation are held together due to two strong hydrogen bonds, which may be

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Table 2: Selected torsion angles and calculated values of the pseudorotational phase angle ($P$) and of the puckering amplitude ($\phi_m$) of the THF ring for all stationary points for conversions 2–4

| Reaction 2 | $P$ | $\phi_m$ | $\phi_0$ | $\phi_1$ | $\phi_2$ | $\phi_3$ | $\phi_4$ | $\chi_1$ | $\chi_2$ |
|-----------|-----|----------|----------|----------|----------|----------|----------|----------|----------|
| R         | $E_d$ | 349     | 36       | 35.6     | -26.0    | 5.2      | 18.2     | -33.6    | -144.5   |
| RC        | $E_d$ | 345     | 36       | 36.2     | -24.1    | 2.3      | 20.8     | -34.9    | -142.6   |
| TS        | $^3T_4$ | 2      | 39       | 39.1     | -33.4    | 14.3     | 11.0     | -31.6    | -154.5   |
| IP        | $^3T_4$ | 4      | 36       | 36.2     | -31.9    | 14.6     | 9.1      | -28.7    | -150.6   |
| P         | $^3T_4$ | 2      | 36       | 36.5     | -31.1    | 13.4     | 10.4     | -29.5    | -148.1   |

*Definition of the torsion angles: $\phi_0$ = C5–C4–C3–C2; $\phi_1$ = C4–C3–C2–O2; $\phi_2$ = C3–C2–O2–C5; $\phi_3$ = C2–O2–C5–C4; $\chi_1$ = C1–C2–C3–C4; $\chi_2$ = R–C5–C4–C3, where R represents the substituent attached to C5.

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Fig. 4: Definition of the endocyclic torsion angles $\phi_0$–$\phi_4$.
responsible for the stronger stability of the ion pair. Such interactions are also found in the ion pairs formed in the other reactions studied.

In ion pair 2b the THF ring has the $^3T_d$ conformation, $(P=4^\circ, \phi_m=36^\circ)$, as in the corresponding transition state, whereas in ion pair 3b the almost ideal $E_5$ conformation of the THF ring $(P=117^\circ, \phi_m=-39^\circ)$ is preferred (Table 2). This means that a conformational change occurs on going downhill from the energy maximum to the valley. In the case of reaction 4 the THF ring adopts the same conformation as in the reactant complex, $E_4$, $(P=342^\circ, \phi_m=36^\circ)$.

The last stage of the reactions studied consists in the separation of the ion pair constituents. Again, this process is extremely unfavorable in the gas phase. More than 90 kcal mol$^{-1}$ in relation to the sum of the energies of the individual reactants must be supplied to move the ions to an infinite distance from one another (Table 1). Ion pair dissociation is more endoenergetic here than it was in the reactions with trimethylamine and pyridine [21–23].

Calculations in solution (SCRF-PCM)

Tomasi’s polarizable continuum model (PCM) was used to investigate the influence of the liquid phase on the course of the reactions under scrutiny [31]. The PCM model permits the self-consistent computation of free energies of solvation, including polarized solute/solvent interactions and non-electrostatic terms in the Hamiltonian. On the other hand it should be emphasized that reaction field models are incapable of modeling specific (short range) solute/solvent interactions, that is, those occurring in the first solvation sphere. Thus, the conclusions drawn based on the calculations where such interactions occur should be interpreted with care. Although PCM operates better in aprotic solvents it has also been used to predict the solvation effect in protic solvents [15, 39].

Keeping in mind the limitations of the implicit solvent models PCM approach was applied to both minimum energy structures and saddle point configurations. In our previous papers [23] we showed that almost the entire solvent effect is achieved after single point PCM calculations, and no significant energy changes were observed during the optimization in water. Moreover, we showed that the TS geometry changes were not so profound as those experienced with the classical Menshutkin reaction [17, 40]. Although previously we studied the solvent effect in three solvents this time we decided to carry out full optimization only in two solvents, that is, chloroform and water. We resigned from doing the calculation in ethanol because energy changes in ethanol and water were roughly the same.
The results of calculations in chloroform and in water are listed in Table 4 whereas Fig. 9 illustrates the geometries of the transition states optimized in water.

Significant changes in the energy diagrams are observed along the reaction path in both solvents, as compared with the gas phase (Fig. 3), especially on the product side. In chloroform, a solvent of low polarity, the energy sum of the separated ions decreases dramatically, but the overall process is still endothermic, relative energy ranges from 23 kcal mol\(^{-1}\) (reactions 2 and 3) to almost 29 kcal mol\(^{-1}\) (reaction 1), with respect to individual reactants (Table 4). The Gibbs free energy also predicts the process to be slightly unfavorable \(\Delta G \approx 4\) kcal mol\(^{-1}\). In very polar solvent, in turn, the reactions are exothermic and favorable. The energies of the separated ions are less than those of the individual reactants by about \(-2\) kcal mol\(^{-1}\) (reaction 1) and about 5 kcal mol\(^{-1}\) for other reactions.

According to the \(\Delta U_0\) values, complexation is slightly exothermic in chloroform (Table 4). On the other hand, \(\Delta G s\) indicate that this step of the reaction is strongly unfavorable, about 6 kcal mol\(^{-1}\), on average, must be supplied to form the reactant complex. Also, in water medium the reactant complex formation is unfavorable. It means that interactions between the constituents of the reactant complex are rather weak. The same conclusions were drawn previously [21–23].

Castejon et al. [17] studied the relationship between the stability of the reactant complexes (and ion pairs as well) and solvent polarity for the Menshutkin reaction. They noticed that neither the reactant complex nor the ion pair were energy minima in very polar solvents, like dimethyl sulfoxide. However, in our case they appeared to be real energy minima in water, although certain geometry differences were observed. The geometries of reactant complexes and ion pairs are roughly the same within the mesylate derivatives in the gas phase and in water. However, the reactant complexes differ in the relative positions of their constituents. The C−−N distance is 3.488 Å on average in the gas phase, whereas in water it is much longer, about 3.7 Å. The reactant complexes and ion pairs formed in the reactions under discussion owe their stability to the hydrogen bonds formed between H and O atoms from the nucleophile and mesylate moiety respectively.

Both \(\Delta U_0\) and \(\Delta G\) showed that ion pair formation was an exoenergetic process in solvents with respect to the reactant complex, as in a vacuum. In reaction 1 taking place in chloroform the decrease in energy is \(-12.9\) kcal mol\(^{-1}\) (\(\Delta G\)), whereas in water the conversion of the reactant complex into the ion pair is more exoenergetic \((-16.8\) kcal mol\(^{-1}\)) than in chloroform. The free energy changes accompanying ion pair formation in the other three reactions are roughly the same.
earlier stage of the reaction.

developed geometrical changes shifts transition states back to an classical Menshutkin reaction \cite{17,40}. Like in chloroform, described geometrical changes are not as profound as in the chloroform, and the C···N distance decreases from 2.060 Å in the gas phase to 1.967 Å in chloroform, and the C···N distance increases from 1.932 Å to 2.040 Å (Table 4). This means that the transition state is shifted toward an earlier stage of the reaction. For the other three reactions the transition state geometry changes are even less.

In turn, more significant changes in transition state geometry, in relation to the gas phase, take place during optimization in water, especially for reactions 1 and 4. In reaction 1 the C···O distance decreases to 1.909 Å whereas the C···N distance increases to 2.113 Å with respect to the values found in the gas phase. Analogous geometry changes in transition state geometries occur for reactions 2 and 3, however, here the C···O distance decreases by about 0.05 Å, whereas the C···N distance increases by about 0.1 Å. The described geometrical changes are not as profound as in the classical Menshutkin reaction \cite{17,40}. Like in chloroform, these geometrical changes shifts transition states back to an earlier stage of the reaction.

Optimization of transition states in chloroform leads to rather insignificant geometry changes. These changes are the greatest for the reaction 1, where the C···O distance decreases from 2.060 Å in the gas phase to 1.967 Å in chloroform, and the C···N distance increases from 1.932 Å to 2.040 Å (Table 4). This means that the transition state is shifted toward an earlier stage of the reaction. For the other three reactions the transition state geometry changes are even less.

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Apart from the geometry changes at the reaction center, there is an additional variation of the transition states in reactions 2–4, particularly in water. In reaction 2 the O–C1–C2–C3 torsion angle (A, Table 4) increases from −13.0° in the gas phase to 53.8° in water. The value is similar for reaction 3 (43.2°), whereas for conversion 4 it is 6.0°. This indicates that in two TSs (reactions 2 and 3) the leaving group is shifted above the THF ring. Rotation about the C1–C2 bond induces the ring conformation switch (T3 → E3) in reaction 3. In reactions 2 and 4 the THF ring conformation changes are not significant (T4 → E4).

It is well recognize that the transfer from the gas phase to solvent leads to a significant energy barrier drop. Indeed, the B3LYP level barriers calculated in both solvents are much lower than those calculated for the gas phase. In the case of the reaction 1 occurring in chloroform, calculated barrier is lower by about 8 kcal mol\(^{-1}\) whereas in water it is ca 13 kcal mol\(^{-1}\) less. For the other three reactions the differences between barriers in the gas phase and in solvents are significantly lower. For the reaction 4 barriers are lower by about 5 kcal mol\(^{-1}\) and 8 kcal mol\(^{-1}\) in chloroform and water, respectively.

A different barriers height order is found in the gas phase and in solvents. In the gas phase the lowest barrier was found for the reaction 2. In turn, for the reaction 1 taking place in chloroform the barrier is by about 3 kcal mol\(^{-1}\) lower than for the reaction 2. The difference between barrier height are even greater in water. This may suggest that hydrogen bond stabilizing TS in the gas phase is weaker in solvents.

Like in the gas phase also in solvents the barrier height depends on the type of the substituent at C5. The presence of a spatial group attached to this carbon atom slightly increases the barrier, although the differences are rather small in chloroform. It is difficult to judge which reaction, 2 or 3 should be faster, based on the free energy barriers calculated in water.

The transfer from the gas phase to the solvent leads to a significant energy drop in the final step of these reactions. While separating the ion pair constituents requires about 100 kcal mol\(^{-1}\) (\(\Delta G\)) in a vacuum, in chloroform over 30 kcal mol\(^{-1}\) need to be supplied to accomplish this step of the reaction. In water, in turn, only about 5 kcal mol\(^{-1}\) are required.
Table 4  Geometry parameters, relative energies and relative Gibbs free energies of relevant stationary points on the FES at B3LYP/6-31+G** calculated for reactions 1-4 in chloroform and water

|                | Reaction 1          | Reaction 2          | Reaction 3          | Reaction 4          |
|----------------|---------------------|---------------------|---------------------|---------------------|
|                | (R) (RC) (TS) (IP) (P) | (R) (RC) (TS) (IP) (P) | (R) (RC) (TS) (IP) (P) | (R) (RC) (TS) (IP) (P) |
| Chloroform     |                     |                     |                     |                     |
| R(C–O)         | 1.451               | 1.469               | 1.459               | 1.469               |
| R(C–N)         | ∞                   | 3.506               | 2.040               | 2.040               |
| ΔR             | ∞                   | -2.045              | -0.073              | 2.158               |
| θ OCN          | -                   | 100.1               | 177.1               | 39.5               |
| ΔU₀            | 0.0                 | -1.5                | 22.7                | -9.5               |
| ΔG             | 0.0                 | 6.9                 | 32.0                | 6.9                |
| ΔU₀²           | 24.3                | 27.0                | 27.7                | 28.9               |
| ΔG²            | 25.1                | 29.4                | 29.6                | 30.7               |
| Water          |                     |                     |                     |                     |
| R(C–O)         | 1.454               | 1.462               | 1.461               | 1.467               |
| R(C–N)         | ∞                   | 3.598               | 2.113               | 1.495               |
| ΔR             | ∞                   | -2.136              | -0.204              | 1.965               |
| θ OCN          | -                   | 100.9               | 176.7               | 45.6               |
| ΔU₀            | 0.0                 | -0.2                | 18.4                | -9.2               |
| ΔG             | 0.0                 | 7.6                 | 28.5                | 23.9               |
| ΔU₀²           | 18.6                | 23.9                | 23.9                | 24.8               |
| ΔG²            | 20.9                | 27.2                | 27.2                | 27.7               |

All energy values in kcal mol⁻¹, R in Å and angles in deg. Reaction coordinate; ΔR = R(C–O) - R(C–N). The smallest value of the distance between C and O was taken to define the reaction coordinate. (R) separate reactants, (RC) reactant complex, (TS) transition state, (IP) – ion pair; (P) separate ions

A torsion angle: (O–C1–C2–C3) for R, RC and TS; (N–C1–C2–C3) for IP and P. In R, this torsion angle corresponds to mesylate, whereas in P this angle corresponds to the cation.

B deformation angle C2–C1–H1a–H1b (for the reaction 1: H–C–H2–H3) describing the planarity of the transition state geometry

C torsion angle defining the position of the aglycone in relation to THF ring (H2C–O–C5–O2)
a the O atom closest to the reaction center carbon atom was used to obtain this value
Conclusions

In this work we continued our DFT study into ammonium salt formation in a Menshutkin-like reaction between ammonia and four mesylate derivatives: methyl mesylate (1a), \((S)\)-1,4-andydro-2,3-dideoxy-5-\(O\)-mesylpentitol (2a), (2S,5S)-2,5-andydro-1,3,4-trideoxy-6-\(O\)-mesylhexitol (3a) and methyl 2,3-dideoxy-5-\(O\)-mesyl-\(\beta\)-D-pentofuranoside (4a). The reactions were investigated using the B3LYP functional with the 6-31+G** basis set in the gas phase and in solvents. Additionally, MPW1K/6-31+G** level calculations were carried out to estimate activation barrier heights. Apart from that, the reactions were studied in two solvents (chloroform and water) using the PCM model.

The energy diagrams presented in Fig. 3 exhibit two minima corresponding to the reactant complex and the ion pair in the gas phase. Clearly, ion pairs are more stable than the respective reactant complexes, but this is not a golden rule. For example, in the reaction between methyl mesylate and pyridine the reactant complex is more stable. The two strong hydrogen bonds formed between the mesylate anion and the ammonium cation are responsible for the extraordinary stability of the ion pair with respect to the individual reactants. The ion pair stability is even greater in solvents. This unusual stabilization of ion pairs causes the final step of the reaction to be endergonic in solvents.

Energy and Gibbs free energy values indicate that the overall process is highly unfavorable in the gas phase. In chloroform it is still unfavorable, but in more polar solvents the sum of the energies of the individual ions is less than that of the separate reactants.

According to B3LYP and MPW1K functionals, the energy barrier is the lowest for reaction 2 and not reaction 1, in contrast to the reactions with trimethylamine or pyridine. However, reaction 1 should be the fastest one in solvents used in the calculations. In turn, reaction 4 seems to be the slowest, both in the gas phase and in chloroform. In water, however, the difference between calculated barriers are insignificantly small thus do not indicate univocally the reaction 4 to be the slowest one. It should be emphasized once again at this place that reaction field models do not describe hydrogen bonding interaction which could affect the energetics of reactions studied thus results of the calculations in water should be taken with care. Possibly the solution for this problem would be achieved based on the calculation of the model with discrete water molecule.

We have also discussed the conformational behavior of the THF ring along the reaction pathway, noting that in reactions 2 and 4 the THF ring adopts a conformation from the narrow region of the northern half of the pseudorotational circle. In turn, the THF ring is solely in conformations located in the southern half in reaction 3. The THF ring conformation switch (\(T_3 \rightarrow E_5\)) was observed for the reaction 3 transition state geometry during the optimization in water, which is the result of rotation about the C1–C2 bond.

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