Conformational analysis of flephedrone using quantum mechanical models

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Abstract Flephedrone is an analogue of cathinone - chemically similar to ephedrine, cathine and other amphetamines. Conformations of all isomers of flephedrone have been studied at the quantum-chemical level. Calculations have been performed using DFT and MP2 methods with two basis sets - 6-31G and 6-31G(d,p). Results show that there are low energy conformers for the ortho, meta, and para isomers that are connected by way of low-barrier transition states. Boltzmann distribution of population predicts the highest population for the 1-meta conformer with a 10 % increase in solution. The molecular electrostatic potential surface data for each molecule has been calculated revealing likely reaction sites.

Keywords Conformations · Flephedrone · Molecular electrostatic potential

Introduction

In recent years, there has been a rapid development of “legal highs” - various psychotropic substances being sold and distributed around the world via the internet. Initially, these substances were freely available to buy legally and without restriction as fertilizers and incense and could be purchased using the internet or from “legal high” shops throughout the UK and Ireland [1, 2]. In 2010, the UK government responded to concerns about the safety of the drugs by banning their use, supply, or possession in the UK. The governments of other countries immediately followed with similar action, but there are still countries where one can buy them legally. Although there is no extensive scientific data on the structure of some of these psychoactive compounds, it is easy to find some basic information. The method of synthesis is easily accessed through web based sites and forums that are specifically designed to provide users means of acquiring “legal high” drugs. Because this information is so readily available to the general public, there is a real danger of overdose or poisoning from consumption. Since only a few of these compounds have been scientifically investigated [1–4] there exists a need to identify structural properties, determine the mechanism of action, and elucidate toxicological effects.

One such compound that produces psychoactive effects is flephedrone (4-fluoromethcathinone) (Fig. 1) [1, 2] which is an analogue of cathinone [4, 5]. Cathinone is an alkaloid found in the shrub Catha edulis (khat). It is chemically similar to ephedrine and other amphetamines. Flephedrone is easily synthesized from ephedrine [6], and it is believed that cathinone induces the release of dopamine from striatal preparations that are pre-labeled either with dopamine or its precursors [5].

The primary aim of this study is to identify all possible conformers of flephedrone so that these structures may be compared to the cathinone structures. There are two
additional isomers of fluoromethcathinone which are 2-fluoromethcathinone and 3-fluoromethcathinone (Fig. 1). These isomers with 4-fluoromethcathinone are considered and calculated using ab initio and density functional treatments. Additionally, the molecular electrostatic potential isosurfaces (MEPS) are calculated.

**Computational methods**

Potential energy surfaces were scanned, and all minima were optimized to find all possible conformers of cathinone, norephedrine and norpseudoephedrine. All structures have been obtained using a relaxed dihedral angle scan to generate a potential energy surface. Points were then optimized using density functional theory’s [7] B3LYP [8, 9] hybrid functional with 6-31G and 6-31G(d,p) basis sets. Second-order Moller Plesset (MP2) [10] calculations were then performed on the generated (DFT/B3LYP) surfaces. The thermodynamic parameters of all compounds were calculated using statistical mechanics expressions.

Solvation free energies ($\Delta G_{solv}$) were determined at the DFT/B3LYP level with COSMO [11]. The $\Delta G_{solv}$ values were calculated in aqueous continuum ($\varepsilon=78.39$), and the molecular cavities were built using the united atom model for Hartree–Fock (UAHF). In principle, the free energy surfaces are determined by a rigorous combination of free energy perturbation/umbrella sampling approaches [12, 13]. These surfaces reflect nonequilibrium solvation. However, such calculations are very challenging when performed within the ab initio framework. Thus, a more practical, simplified approach has been made. The relative free energies ($\Delta G$) are determined as [14]:

$$\Delta G = \Delta H_{gas}(298 K) - T \Delta S - RT \ln(\omega) + \Delta G_{solv}$$

$$\approx \Delta G_{solv}(298 K) + \Delta G_{solv}.$$  \hspace{1cm} (1)

In Eq. (1), $\Delta H_{gas}(298 K)$ is enthalpy at 298 K, $\Delta S$ is the gas-phase entropy, $\Delta G_{gas}(298 K)$ is the Gibb’s free energy at 298 K, and $\Delta G_{solv}$ is the relative free energy of solvation. The contribution of the $RT \ln(\omega)$ term is zero, as the electronic degeneracy term, $\omega$, for the singlet state is unity. The $\Delta G$ values are used to calculate the relative population of the various conformers of the molecules under study in aqueous medium.

Since the molecules are much less rigid in solution than in the gas phase, we have used another definition of free energy, ($\Delta g_{flex}$), to compare the intrinsic flexibility of different conformers, when embedded in aqueous medium. This is the practical implementation of the more general expression used by Warshel and coworkers [14, 15]. The $\Delta g_{flex}$ could be expressed as:

$$\Delta g_{flex} = \Delta E_{solute} + \Delta ZPE + \Delta G_{solv} - \alpha T \Delta S,$$ \hspace{1cm} (2)

where $\Delta E_{solute}$ and $\Delta ZPE$ in Eq. (2) are the relative gas-phase energy separations and the zero-point energies of the various conformers. The scale factor $\alpha$ is usually taken as zero [15], and expression (2) can be simplified as:

**Table 1** Relative energy and structural properties of low energy isomers of 2-FMC calculated at DFT and MP2 levels with 6-31G and 6-31G(d,p) basis sets. All energies are given in kcal mol$^{-1}$.

| 2-FMC | 6-31G | 6-31G(d,p) |
|-------|-------|------------|
|       | $\Delta H_0$ | $\Delta G$ | $\Delta H_0$ | $\Delta G$ | $\tau_1$ | $\tau_2$ | $\tau_3$ | $\mu$ |
| DFT 1o | 0.00 | 0.00 | 0.00 | 0.00 | 39.5 | 22.3 | 179.5 | 3.62 | 6-31G |
|       | 4.86 | 4.42 | 3.94 | 3.31 | 79.8 | 71.0 | 177.8 | 3.62 | 6-31G(d,p) |
| MP2 1o | 0.00 | 0.00 | 0.00 | 0.00 | 27.7 | 37.3 | 179.2 | 3.67 | 6-31G |
|       | 3.16 | 3.32 | 2.64 | 2.72 | 79.8 | 71.0 | 177.8 | 3.99 | 6-31G |
|       | 70.6 | 89.4 | 178.4 | 3.22 | 6-31G(d,p) |
\[ \psi_{\text{gflex}} = \Delta H_0^{\text{gas}} + \Delta G_{\text{solv}}. \]  

(3)

All terms in Eqs. (1) and (3) are available from the thermochemical analyses based on statistical mechanics expressions using the ideal gas, rigid rotator, and harmonic oscillator approximations [16]. All calculations were carried out using the GAUSSIAN 03 structure calculation software [17], and molecular graphics were generated using the GAUSS VIEW visualization program [18].

**Results and discussion**

2-FMC

Results show that 2-FMC, 3-FMC and 4-FMC have two, four, and two low energy conformers, respectively. Thermodynamic and structural properties of 2-FMC have been collected in Table 1 and include relative enthalpy and free energy changes as well as angles of rotation. The two

| 3-FMC | 6-31G | 6-31G(d,p) | 6-31G | 6-31G | \( \tau_1 \) | \( \tau_2 \) | \( \tau_3 \) | \( \mu \) |
|-------|-------|-----------|-------|-------|--------|--------|--------|-------|
| DFT   |       |           |       |       |        |        |        |       |
| 1 m   | 0.00  | 0.01      | 0.00  | 0.00  | -157.1 | -176.3 | -179.3 | 3.38  |
|       |       |           |       |       | -156.9 | -175.9 | -179.7 | 2.87  |
| 2 m   | 0.02  | 0.00      | 0.11  | 0.08  | -157.2 | 3.6    | 179.9  | 1.36  |
|       |       |           |       |       | -156.9 | 4.1    | 179.9  | 1.60  |
| 3 m   | 3.25  | 1.35      | 4.32  | 4.32  | 52.2   | -1.1   | 178.9  | 1.58  |
|       |       |           |       |       | 49.9   | 16.9   | 179.3  | 1.83  |
| 4 m   | 3.53  | 2.89      | 4.28  | 4.26  | 52.6   | -172.6 | -179.3 | 4.29  |
|       |       |           |       |       | 50.4   | -164.3 | -179.8 | 3.31  |
| MP2   |       |           |       |       |        |        |        |       |
| 1 m   | 0.00  | 0.00      | 0.00  | 0.00  | -161.7 | -169.2 | -179.6 | 3.20  |
|       |       |           |       |       | -160.2 | -172.2 | -179.7 | 2.75  |
| 2 m   | 0.03  | 0.00      | 0.11  | 0.07  | -161.7 | 12.8   | 179.5  | 1.31  |
|       |       |           |       |       | -160.2 | 9.6    | 179.7  | 1.54  |
| 3 m   | 3.22  | 3.31      | 3.79  | 3.93  | 41.2   | 29.4   | 178.9  | 1.69  |
|       |       |           |       |       | 41.0   | 29.5   | 179.1  | 1.78  |
| 4 m   | 3.29  | 3.37      | 3.75  | 3.87  | 41.5   | -152.1 | -179.4 | 3.75  |
|       |       |           |       |       | 41.4   | -152.1 | -179.8 | 3.09  |
conformers have a 2.64 to 4.86 kcal mol\(^{-1}\) difference in \(\Delta H_0\) depending on the basis set and method. Analysis of the structures shows that the structural properties of the conformers do not vary much with the changing of basis sets or method. Figure 2 is the energetic profile for the 2-FMC isomer with two conformers connected through a rotational transition state. The barrier for transition is equal to 5.66 kcal mol\(^{-1}\) for gas phase and 7.54 kcal mol\(^{-1}\) in aqueous solution.

3-FMC

Thermodynamic and structural parameters of 3-FMC isomers have been collected in Table 2 with four low energy conformers connected through five transition states (Fig. 3). Calculations using larger basis sets show that relative energy differences for each conformer increase. MP2 calculations demonstrate lower differences between conformers than the DFT results. Changing basis set or method does not have a significant effect on structural properties. Calculations in solution do not change relative differences between conformers, but it lowers the energy barrier for the transition of 1 m <-> 2 m (Fig. 3). The Boltzmann distribution of population in Table 3 shows that the conformation with the highest population is the 1 m. In solution its population is increased by about 10 %. The higher dipole moment can contribute to the stabilization of the 1 m structure and influence the reactivity of the molecule in solution. Changing methods does not change the population percentages.

4-FMC

The 4-FMC isomer has two conformers connected via one rotational transition state. Calculations using MP2 and DFT methods with 6-31G and 6-31G(d,p) basis sets show the same energetic trend for each conformer, although MP2 calculations predict lower energies. Table 4 contains all calculated structural and energetic properties of 4-FMC conformers with a graphical representation of each conformer shown in the Supporting information. Structural properties do not change much with higher basis set or different method of calculation. The rotational barrier for gas phase is equal to 8.78 kcal mol\(^{-1}\) for the lowest energy conformer (Fig. 4). In solution, this difference decreases to 6.66 kcal mol\(^{-1}\). The energy trend for the gas phase shown in Fig. 4 is the same in solution. The energy difference for conformers is between 3.5 kcal mol\(^{-1}\) and 4 kcal mol\(^{-1}\) depending on the basis set and method for the gas phase. Solution decreases the energy by about 0.20 kcal mol\(^{-1}\). Of the three isomers, cathinone has an energetic profile most similar to 4-FMC.
Free energy differences between conformers of cathinone and 4-FMC are from 0.98 to 1.71 kcal mol\(^{-1}\) while other isomer conformations have energy differences 1.5 to 3.5 kcal mol\(^{-1}\) higher (Table 5).

**Table 4** Relative energy and structural properties of 4-FMC calculated using DFT and MP2 with 6-31G and 6-31G(d,p) basis sets. All energies are given in kcal mol\(^{-1}\).

| 4-FMC | 6-31G | 6-31G(d,p) |
|-------|-------|------------|
|       | \(\Delta H_0\) | \(\Delta G\) | \(\Delta H_0\) | \(\Delta G\) | \(\tau_1\) | \(\tau_2\) | \(\tau_3\) | \(\mu\) |
| DFT   |       |       |           |       |       |   |       |       |       |
| 1p    | 0.00  | 0.00  | 0.00      | 0.00  | -157.0 | 3.3 | 180.0 | 2.48  | 6-31G |
|       |       |       | 160.0     | 3.8   | 180.0  | 2.12 | 6-31G(d,p) |
| 2p    | 3.48  | 3.76  | 3.30      | 4.15  | 51.37  | 8.6 | 179.1 | 1.52  | 6-31G |
|       |       |       | 51.5      | 14.4  | 179.3  | 1.69 | 6-31G(d,p) |
| MP2   |       |       |           |       | 157.0  | 3.3 | 180.0 | 2.18  | 6-31G |
| 1p    | 0.00  | 0.00  | 0.00      | 0.00  | 160.2  | 9.6 | 179.7 | 1.79  | 6-31G |
|       |       |       | 41.6      | 14.4  | 179.1  | 1.20 | 6-31G |
| 2p    | 3.30  | 3.41  | 3.81      | 3.95  | 51.4   | 8.6 | 179.1 | 1.42  | 6-31G |

Molecular electrostatic potential surface analysis

Molecular electrostatic potential maps (MEP) use local electronic charge density to systematically determine a molecule’s ability to interact. Widely used in drug design, MEP maps provide a method for identification of ligand-protein attractive forces. Generation of MEP maps for this study is ideal for predicting the nature of interactions and provides justification for molecular phenomena. The molecular electrostatic potential isosurface (MEP) for 2-FMC shows that the most active site of the molecule for reaction is on the oxygen atom (Fig. 6). The second negative region is localized around the nitrogen atom of the amine group. The negative potential zone around the oxygen atom is much stronger than for fluorine. Oxygen’s high potential is expected due to lone pair electrons, but nevertheless provides a primary site for strong hydrogen bonding. The active conformers have positioning of the nitrogen as a secondary amine group. This restricts the ability of nitrogen to give a strong negative potential.

Figure 7 is the MEP isosurface of all low-energy conformers of 3-FMC. The negative potential region is maximized around the oxygen atom. Because of the shape of the molecule there appears to be one more active region of negative potential on the fluorine atom, but the negative potential is much less negative than for the oxygen atom. For structure 1 m, the nitrogen atom also has negative potential. Figure 8 gives the MEP for the 4-FMC conformers. As with the conformers of 2-FMC and 3-FMC, areas of potential exist near the oxygen, nitrogen, and fluorine with the higher electron density around oxygen. As compared to FMC

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Fig. 4 Energy profile for 4-FMC in gas phase. Energies in solution (gflex) are given in parenthesis.
isomers, cathinone will possibly form a stronger N–H bond, but FMC’s higher number of potential binding sites will make FMC a competitive receptor ligand.

**Conclusions**

All isomers have at least two low energy conformers connected through low-barrier transition states. In this work we carried out conformational analysis of isomers of fluoromethcathinone at the DFT-B3LYP/6-31G, B3LYP/6-31G (d,p), MP2/6-31G and MP2/6-31G(d,p) level. The dependence of the low-energy conformers of fluoromethcathinone on aqueous solvation was studied using the DFT/MP2/COSMO model and the MEP surfaces were generated for these low-energy surfaces to interpret the nature of practical reaction regions. Our calculations indicate that 2-FMC, 3-FMC, and 4-FMC have several low-energy conformers in the gas phase as well as in aqueous medium. These low-energy conformers are connected through low-barrier transition states, indicating that they can be converted from one form to the other. The energy trend is conserved for MP2/DFT calculations. In one case, (3-FMC) in aqueous medium, one transition barrier has been lowered in comparison with the gas phase. The MEP surfaces revealed three negative potential regions that can take place in reaction. Because of the geometry of these compounds there are few structures where the negative potential around nitrogen is weak. In closing, this study reveals the importance of low-energy conformers of flephedrone in exhibiting its specific biological properties. Obvious sites of reactivity exist which can participate in hydrogen bonding or charge transfer interactions. This implies that flephedrone, as with other psychostimulants, can act as an exogenous receptor agonist by receptor binding and

| Cathinone | 6-31G | 6-31G(d,p) |
|-----------|-------|------------|
|           | ΔH₀   | ΔG        | ΔH₀   | ΔG  | τ₁      | τ₂      | τ₃      | μ    |
| DFT       |       |           |       |     |         |         |         |      |
| 1 k       | 0.00  | 0.00      | 0.00  | 0.00| -179.8  | 175.8   | 158.2   | 3.09 |
|           |       |           |       |     | -179.9  | 175.1   | 157.3   | 3.10 |
| 2 k       | 1.18  | 0.98      | 1.31  | 1.12| -179.6  | 7.1     | 78.4    | 3.11 |
|           |       |           |       |     | -179.7  | 6.9     | 81.0    | 2.99 |
| 4-FMC     |       |           |       |     |         |         |         |      |
| MP2       |       |           |       |     |         |         |         |      |
| 1 k       | 0.00  | 0.00      | 0.00  | 0.00| 179.2   | 166.3   | 165.4   | 3.67 |
|           |       |           |       |     | 179.5   | 156.3   | 166.4   | 3.36 |
| 2 k       | 1.68  | 1.71      | 1.34  | 1.40| 179.3   | 19.4    | 66.2    | 3.66 |
|           |       |           |       |     | 179.1   | 25.1    | 66.9    | 3.22 |

![Fig. 5 Energy profile for cathinone in gas phase.](image)
Fig. 6 Molecular electrostatic potential surface of 2-FMC conformers. Red regions of the map are the most electron-rich regions of the molecule and blue regions are electron-poor. Order of increasing electron density is blue < green < yellow < orange < red.

Fig. 7 Molecular electrostatic potential surface of 3-FMC conformers. Red regions of the map are the most electron-rich regions of the molecule and blue regions are electron-poor. Order of increasing electron density is blue < green < yellow < orange < red.

Fig. 8 Molecular electrostatic potential surface of conformers of the 4-FMC isomer. Red regions of the map are the most electron-rich regions of the molecule and blue regions are electron-poor. Order of increasing electron density is blue < green < yellow < orange < red.
indirect or direct activation. Understanding the structure offers a starting point for determining the mechanism of action and provides for the development of antagonist drugs as treatment of overdose, poisoning, or addiction.

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