

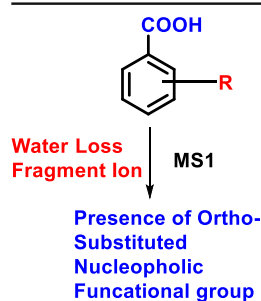
RESEARCH ARTICLE

Identification of *ortho*-Substituted Benzoic Acid/Ester Derivatives via the Gas-Phase Neighboring Group Participation Effect in (+)-ESI High Resolution Mass Spectrometry

William D. Blincoe,¹ Agustina Rodriguez-Granillo,² Josep Saurí,¹ Nicholas A. Pierson,¹ Leo A. Joyce,¹ Ian Mangion,¹ Huaming Sheng¹

¹Analytical Research and Development, Merck and Co., Inc., Rahway, NJ 07065, USA

²Chemistry Modeling and Informatics, Merck and Co., Inc., Rahway, NJ 07065, USA



Abstract. Benzoic acid/ester/amide derivatives are common moieties in pharmaceutical compounds and present a challenge in positional isomer identification by traditional tandem mass spectrometric analysis. A method is presented for exploiting the gas-phase neighboring group participation (NGP) effect to differentiate *ortho*-substituted benzoic acid/ester derivatives with high resolution mass spectrometry (HRMS¹). Significant water/alcohol loss (>30% abundance in MS¹ spectra) was observed for *ortho*-substituted nucleophilic groups; these fragment peaks are not observable for the corresponding *para* and *meta*-substituted analogs. Experiments were also extended to the analysis of two intermediates in the synthesis of suvorexant (Belsomra) with additional analysis conducted with nuclear magnetic

resonance (NMR), density functional theory (DFT), and ion mobility spectrometry-mass spectrometry (IMS-MS) studies. Significant water/alcohol loss was also observed for 1-substituted 1, 2, 3-triazoles but not for the isomeric 2-substituted 1, 2, 3-triazole analogs. IMS-MS, NMR, and DFT studies were conducted to show that the preferred orientation of the 2-substituted triazole rotamer was away from the electrophilic center of the reaction, whereas the 1-substituted triazole was oriented in close proximity to the center. Abundance of NGP product was determined to be a product of three factors: (1) proton affinity of the nucleophilic group; (2) steric impact of the nucleophile; and (3) proximity of the nucleophile to carboxylic acid/ester functional groups.

Keywords: Benzoic acid/ester derivatives, Gas phase neighboring group participation effect, HRMS, (+)-ESI, DFT, IMS, NMR, Structural elucidation

Received: 26 October 2017/Revised: 21 December 2017/Accepted: 23 December 2017/Published Online: 27 February 2018

Introduction

Substituted benzoic acid/ester derivatives are important building blocks used in many pharmaceutical compounds and in organic synthesis. They have notably been used as preservatives for food, cosmetics, and pharmaceutical products because of their antimicrobial and antifungal properties [1–3]. Compounds that contain the benzoic acid/benzamide moiety

have been fundamental to the pharmaceutical industry in products that cover a bevy of therapeutic benefits, including inflammation (aspirin, salicylic acid), schizophrenia (amisulpride), insomnia (suvorexant), and cancer therapy (pemetrexed, imatinib).

Different positional isomers substituted at *ortho*, *meta*, and *para* positions as well as conformational isomers such as *E/Z* isomers can be generated during the synthesis of the aromatic substituted benzoic acid/ester derivatives. Traditionally, analytical methods such as nuclear magnetic resonance (NMR) and high performance liquid chromatography-high resolution tandem mass spectrometry (HPLC-HRMS/MS) are able to differentiate these sets of isomers. However, each method has limitations. NMR studies can often be time-consuming, and

Electronic supplementary material The online version of this article (<https://doi.org/10.1007/s13361-017-1884-8>) contains supplementary material, which is available to authorized users.

Correspondence to: Huaming Sheng; e-mail: huaming.sheng@merck.com

depend on the purity of samples making complex mixture analysis difficult [4–9]. HPLC-HRMS/MS has the advantages of chromatographic separation of mixtures, but identification of each chromatographically resolved peak often requires the retention time match with an authentic sample because most positional isomers of aromatic substituted benzoic acid/ester derivatives have indistinguishable fragmentation patterns under tandem mass spectrometric analysis [10–16]. The neighboring group participation (NGP) effect in organic chemistry has been defined by IUPAC as the intramolecular interaction of a reaction center with a lone pair of electrons in an atom or the electrons present in a σ -bond or π -bond [17]. While NGP influences many reactions in solution, the gas-phase NGP effect has also been extensively studied to explain many fragmentation patterns during collision induced dissociation (CID) [18–24]. Cooks et al. showed δ -cleavage from gas-phase aryl participation in several azulene compounds after electron ionization [23]. Bigler and Hesse identified cinnamic acid derivatives of diamines based on electrospray ionization (ESI)-MS fragmentation and noted the difference in reactivity correlated with ring strain of cyclized products [19]. While these previous studies evaluated the NGP effect under CID conditions, this study mainly focuses on the NGP effect that can take place during MS¹ analysis of protonated *ortho*-substituted benzoic acid/ester derivatives with nucleophilic functional groups containing lone pair or π -bond electrons. The high energy barrier for intramolecular cyclization as seen in CID could be overcome if the nucleophile and electrophile is in small spatial proximity as is the case for *ortho*-substituted benzoic acid/ester derivatives. These nucleophilic groups can attack the protonated carboxylic acid/ester moiety to facilitate the water/alcohol neutral molecule loss. On the other hand, no significant water/alcohol neutral loss would be expected for the *meta*- and *para*-substituted benzoic acid/ester derivatives due to the larger spatial distance and steric penalty for nucleophilic attack. This observation can be used to differentiate some *ortho*-substituted benzoic acid/ester derivatives from their *meta*- and *para*-substituted isomers.

In the present study, 22 *ortho*-, *meta*-, and *para*-substituted benzoic acid scaffolds (**1–22**) (Figure 1) were chosen to study this effect. These compounds were incubated in 0.05% H₂SO₄ in methanol (MeOH), ethanol (EtOH), and isopropanol (*i*-PrOH) solutions in order to form the corresponding ester products. The resulting 65 acid and ester products were separated by LC followed by (+)-ESI/MS analysis. Three factors were determined to impact the differentiation of isomers based on MS¹ analysis: (1) the effects of the competing protonation of the carboxyl group versus the other nucleophilic group during ESI; (2) the steric effects of cyclization; and (3) the nucleophilic group proximity to the electrophilic center. Each compound was tested to see if neutral loss of either H₂O for the acids or MeOH, EtOH or *i*-PrOH for the esters occurred during MS¹ analysis. These losses will be referred to as neutral group losses. Furthermore, efforts were made toward understanding the gas-phase

NGP effect in two triazole benzoic acid isomers (**23–24**) as shown in Figure 1 generated during the synthesis of suvorexant (trade name Belsomra) and analyzed here by HRMS, MS/MS, Ion mobility spectrometry-mass spectrometry (IMS-MS), NMR spectroscopy, and density functional theory (DFT) calculations.

Experimental

Chemicals and Reagents

Methanol, ethanol, isopropanol, acetonitrile, water with 0.1% formic acid solvents, and sulfuric acid were obtained from Fischer Scientific (Optima LC-MS grade, Waltham, MA, USA). Compounds were sourced as follows: **1a**, **4**, **7a**, **12a**, **13a**, **15a**, **17a**, and **21a** were obtained from Sigma-Aldrich (St. Louis, MO, USA); **2a**, **6a**, **8a**, **19a**, and **20a** were obtained from Enamine LLC (Monmouth, NJ, USA); **3a**, **11a**, and **14a** were obtained from Maybridge (Loughborough, UK); **5a** was obtained from Rieke Metals (Lincoln, NE, USA); **9** was obtained from ChemBridge (San Diego, CA, USA); **10a** was obtained from Matrix Scientific (Elgrin, SC, USA); **16a** was obtained from Chem-Impex International (Wood Dale, IL, USA); **18a** was obtained from Apollo Scientific (Manchester, UK); **22a** was obtained from ArkPharm (Libertyville, IL, USA); **23a** and **24a** were synthesized according to literature procedure [25–27].

Sample Preparation

Samples were prepared in 2 mL HPLC vials using an Andrew Alliance pipetting robot. The 24 analyte compounds were prepared in 0.05% H₂SO₄ in three different solvents (methanol, ethanol, and isopropanol) for esterification to the **b**, **c**, and **d** compounds by adding a few crystals of each analyte compound to the vials and mixing. Dissolved analytes were stored at room temperature for 2 d to allow for esterification of the carboxylic acid to occur.

Instrumentation

HPLC/UV LC separation was performed on a Waters Acquity UPLC system, consisting of a binary pump system, a sample manager, and a PDA detector (Waters Corporation, Milford, MA, USA). The output signal was monitored and processed with MassLynx software designed by Waters Corporation (Milford, MA, USA).

Separation was carried out on a Waters BEH C18 column (2.1 × 50 mm, 1.7 μ m particle sizes). The mobile phase consisted of water with 0.1% formic acid (mobile phase A) and acetonitrile (mobile phase B). The injection volume was 3 μ L. Analytes were eluted using a gradient method consisting of an initial hold at 1% mobile phase B for 0.5 min, followed by a linear gradient to 90% B over 3.5 min, then a linear gradient to 0% B over 0.1 min, a hold at 0% B for 0.4 min, and finally a

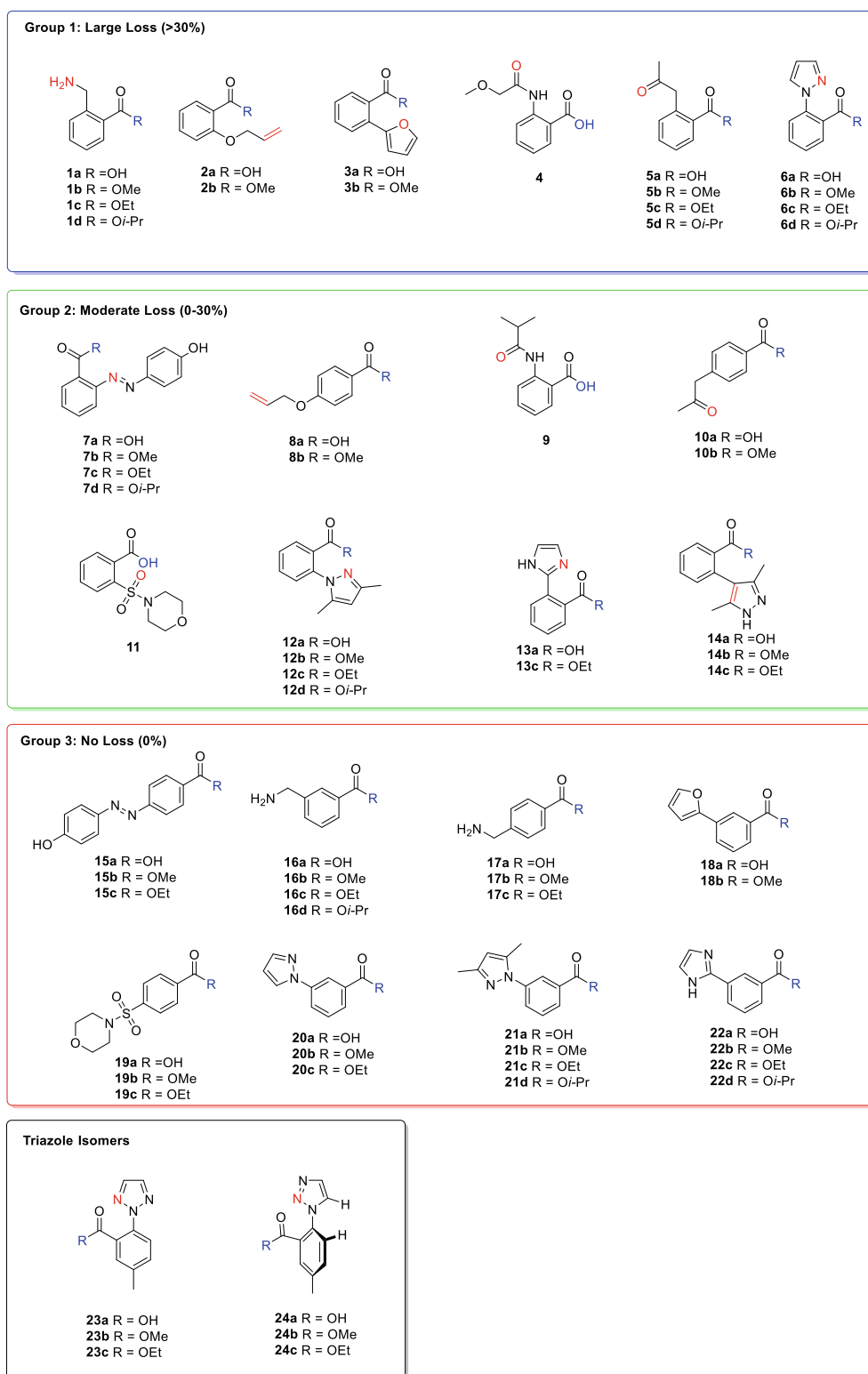


Figure 1. Model compounds grouped by the extent of water/alcohol loss. Nucleophilic groups are marked in red and the R group is marked in blue. Two triazole compounds were also tested with **23** having large loss (>30%) and **24** having no loss (0%)

hold at 1% B for 0.5 min affording a total run time of 5 min. The flow rate was 0.5 mL/min. The column temperature was maintained at 40 °C. The PDA detector was scanned from 200 to 400 nm.

Mass Spectrometry The eluent was introduced directly into the mass spectrometer via an electrospray ionization source. MS analysis was performed on a Waters Premier quadrupole time-of-flight (Q-ToF) mass spectrometer operating in positive

ion mode. Source temperature and desolvation temperature were set to 120 °C and 400 °C, respectively. Nitrogen was used as both the cone gas (50 L/h) and desolvation gas (800 liters/h). The capillary voltage was set to 3 kV. The cone voltage applied was 10 V. Leucine enkephalin was used as the lock mass (m/z of 556.2772) for accurate mass calibration and was introduced using the lock spray interface at 20 $\mu\text{L}/\text{min}$ at a concentration of 0.5 mg/mL in 50% aqueous acetonitrile containing 0.1% formic acid. During MS scanning, data were acquired in centroid mode from m/z 50 to 1000.

Ion Mobility Spectrometry (IMS) Instrumentation IMS-MS experiments were conducted on Agilent 6560 uniform-field ion mobility-Q-ToF instrument (Agilent Technologies, Santa Clara, CA, USA). Samples were infused by syringe pump (5 $\mu\text{L}/\text{min}$) and ionized by ESI (3.5 kV capillary voltage). The instrument was tuned for fragile ions, an automated procedure that lowers ion optics voltages to preserve fragile ion structures as they are transported through the instrument from the source to detector. Sheath and desolvation gas temperatures were set to 150 °C and 100 °C, respectively. Ion mobility measurements were performed in both helium and nitrogen drift gases (4.0 Torr) with a drift tube entrance voltage of 1200 V and a drift tube exit voltage of 250 V. Experimental nitrogen and helium collision cross-sections (CCS) were measured by single-field calibration with the Agilent Tune Mix ions m/z 118 and 322. Theoretical CCS values were calculated by the trajectory method in MOBCAL [28] from energy-minimized structures obtained from the DFT calculations described below.

NMR NMR data were acquired in DMSO- d_6 at room temperature using a 3 mm tube at 500 MHz from a Bruker instrument equipped with a triple-resonance (TCI) Prodigy Cryo-Probe. For the 1D selROE experiment acquired for isomer **24a**, a 20 ms Gaussian selective pulse was used to excite proton H_b. A mixing time of 300 ms was used for the ROESY step.

Density Functional Theory (DFT) Calculations An ensemble of molecular mechanics (MM) conformations for each intermediate were generated with Openeye's Omega [29, 30], followed by DFT optimization using Jaguar 9.1 [31, 32]. All calculations were performed at the B3LYP-D3/6-31++G**level in the gas phase [33–37].

Results and Discussion

MS¹ Spectra of Benzoic Acids and Esters

Table 1 summarizes the MS¹ data obtained for all 65 model compounds. The stability of the ester compounds was not uniform and some compounds were unable to form all three ester products, as shown in Figure 1. The compounds were divided into four groups also shown in Figure 1, with Group 1

including compounds that exhibited a neutral group loss greater than 30%, Group 2 including compounds with a neutral loss between 0% and 30%, and Group 3 with no neutral group loss observed. Examples of the MS¹ spectra observed for **6a-b** and **20a-b** are shown in Figure 2. All other MS¹ spectra can be found in the [Supplementary material](#).

The compounds in Group 1 (**1-6**) showed significant neutral group loss greater than 30%. These compounds all included nucleophilic groups *ortho* to the acid/ester functional group. These nucleophiles ranged from amine groups in **1**, to oxygen nucleophiles in **3** and **5**, to π -bonds in **2**. Group 2 compounds (**7-14**) showed moderate neutral group loss with *ortho* nucleophilic groups and two *para* nucleophilic groups in **8** and **10**. Interestingly, two *para* compounds include longer chain nucleophiles that might allow nucleophilic attack to bridge the aromatic ring. Another explanation for the neutral group loss observed in MS¹ analysis of **8** and **10** is that the carboxylic acid/ester is the main protonation site during ionization due to the relative low proton affinity of the *para* substitution group. Group 3 compounds (**15-22**) included only compounds that had nucleophilic groups *meta* and *para* to the acid/ester, which did not show neutral loss during ionization, as expected.

A proposed mechanism to explain the neutral group loss from the model compounds can be found in Scheme 1. The reaction proceeds via two pathways, a 5- or 6-member ring formation depending on the placement of the nucleophilic group. In the case of carboxylic acid, the reaction is initiated by protonation on the carbonyl group of the carboxylic acid during ionization. The NGP effect then occurs as the nucleophilic substituent attacks the electrophilic carbonyl with subsequent elimination of the neutral leaving group to form a cyclic product ion. The three controlling factors governing this cyclization are proposed as follows: (1) the proton affinity of the carboxylic acid/ester competing with other basic sites on the compound; (2) the steric penalty of forming the 5- or 6-member ring; and (3) the proximity of the nucleophilic group to the carbonyl.

Proton Affinity (PA) of Nucleophilic Group and Carboxylic Acid/Ester

The gas-phase PA of each functional group within a molecule is closely related to the probability of protonation during (+)-ESI ionization [38]. As shown in Table 1, it was observed that the proton affinity (PA) of the neighboring nucleophilic groups and the carboxylic acid/ester closely correlate with the extent of neutral loss observed experimentally. On one hand, the higher PA of the leaving group (carboxylic acid/ester) led to more neutral loss. For example, methyl ester group (203.3 kcal/mol) in **1b** had higher PA than carboxylic acid group (196.2 kcal/mol) in **1a** [38]. Hence, more protonation was expected on the methyl ester, resulting in a larger extent of neutral loss (100% for **1b** vs 22% for **1a**). On the other hand, the higher PA of the neighboring substituent led to less neutral loss. For example, a larger extent of neutral loss was observed for **3a** (100%) compared with **1a** (22%) because the benzyl amino

Table 1. MS¹ Data for benzoic acid and ester model compounds. Nominal *m/z* provided with relative abundance in parenthesis. Compounds in blue are >30% abundance. Compounds in green are between 0% and 30% abundance relative to the base peak. Compounds in red are 0% abundance (not detected)

#	R	[M+H] ⁺ (<i>m/z</i>)	[M+H-ROH] ⁺ (<i>m/z</i>)	[M+H-NH ₃ /OR] ⁺ (<i>m/z</i>)
1	a (H-)	152 (100%)	134 (22.1%)	135 (34.0%)
	b (Me-)	166 (0.6%)	134 (100%)	135 (7.0%)
	c (Et-)	180 (0.2%)	134 (100%)	135 (8.6%)
	d (<i>i</i> -Pr-)	194 (0.4%)	134 (100%)	135 (6.6%)
2	a (H-)	179 (100%)	161 (57.7%)	162 (4.9%)
	b (Me-)	193 (85.5%)	161 (100%)	162 (11.2%)
3	a (H-)	189 (62.0%)	171 (100%)	172 (11.0%)
	b (Me-)	203 (15.0%)	171 (100%)	---
4	(H-)	210 (100%)	192 (59.0%)	193 (20.2%)
5	a (H-)	179 (27.3%)	193 (32.9%)	162 (5.0%)
	b (Me-)	161 (100%)	161 (100%)	162 (10.7%)
	c (Et-)	207 (27.5%)	161 (100%)	162 (11.4%)
6	a (H-)	189 (100%)	171 (60.5%)	172 (7.0%)
	b (Me-)	203 (100%)	171 (69.5%)	172 (8.5%)
	c (Et-)	217 (100%)	171 (51.2%)	172 (6.7%)
	d (<i>i</i> -Pr-)	231 (6.4%)	171 (100%)	172 (12.1%)
7	a (H-)	243 (100%)	225 (15.9%)	---
	b (Me-)	257 (100%)	(225) 15.2%	---
	c (Et-)	271 (100%)	225 (7.3%)	---
	d (<i>i</i> -Pr-)	285 (100%)	225 (4.5%)	---
8	a (H-)	179 (100%)	161 (10.7%)	---
	b (Me-)	193 (100%)	161 (9.4%)	---
9	(H-)	208 (55.3%)	190 (4.0%)	191 (0.8%)
10	a (H-)	179 (100%)	161 (0%)	---
	b (Me-)	193 (100%)	161 (1.4%)	---
11	(H-)	272 (100%)	254 (29.1%)	255 (4.1%)
	a (H-)	217 (100%)	199 (13.3%)	200 (1.9%)
	b (Me-)	231 (100%)	199 (21.2%)	200 (2.6%)
	c (Et-)	245 (100%)	199 (3.2%)	200 (0.5%)
12	d (<i>i</i> -Pr-)	259 (100%)	199 (9.3%)	200 (9.6%)
	a (H-)	189 (100%)	171 (11.9%)	172 (1.5%)
	c (Et-)	217 (100%)	171 (0%)	---
	a (H-)	217 (100%)	199 (2.6%)	---
14	b (Me-)	231 (100%)	199 (5.2%)	---
	c (Et-)	245 (100%)	199 (0%)	---
	a (H-)	243 (100%)	225 (0%)	---
15	b (Me-)	257 (100%)	225 (0%)	---
	c (Et-)	271 (100%)	225 (0%)	---
	a (H-)	152 (53.5%)	134 (0%)	135 (100%)
16	b (Me-)	166 (52.6%)	134 (0%)	149 (100%)
	c (Et-)	180 (76.5%)	134 (0%)	163 (100%)
	d (<i>i</i> -Pr-)	194 (39.3%)	134 (0%)	177 (100%)
	a (H-)	152 (54.5%)	134 (0%)	135 (100%)
17	b (Me-)	166 (39.6%)	134 (0%)	149 (100%)
	c (Et-)	180 (55.9%)	134 (0%)	163 (100%)
	a (H-)	189 (100%)	171 (0%)	---
	b (Me-)	203 (100%)	171 (0%)	---
19	a (H-)	272 (100%)	254 (0%)	---
	b (Me-)	286 (100%)	254 (0%)	---
	c (Et-)	300 (100%)	254 (0%)	---
	a (H-)	189 (100%)	171 (0%)	---
20	b (Me-)	203 (100%)	171 (0%)	---
	c (Et-)	217 (100%)	171 (0%)	---
	a (H-)	217 (100%)	199 (0%)	---
	b (Me-)	231 (100%)	199 (0%)	---
21	c (Et-)	245 (100%)	199 (0%)	---
	d (<i>i</i> -Pr-)	259 (100%)	199 (0%)	---
	a (H-)	189 (100%)	171 (0%)	---
	b (Me-)	203 (100%)	171 (0%)	---
22	c (Et-)	217 (100%)	171 (0%)	---
	d (<i>i</i> -Pr-)	231 (33.5%)	171 (0%)	---
	a (H-)	204 (3.6%)	186 (100%)	187 (12.8%)
	b (Me-)	218 (1.5%)	186 (100%)	187 (12.1%)
23	c (Et-)	232 (29.4%)	186 (100%)	187 (12.7%)
	a (H-)	204 (100%)	186 (2.5%)	---
	b (Me-)	218 (100%)	186 (4.4%)	---
24	c (Et-)	232 (100%)	186 (0%)	---

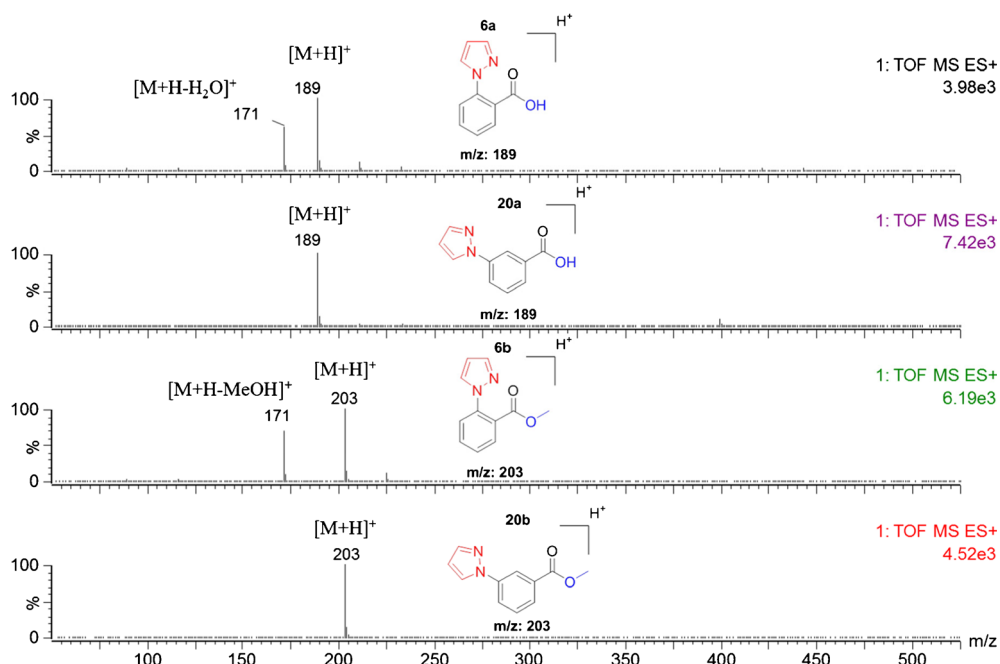


Figure 2. The MS¹ spectra of **6a** and **20b** under (+)-ESI. The pyrazole substitution group is highlighted in red. The acid/ester functional group is highlighted in blue

group (218.3 kcal/mol) has higher PA than furan (192.0 kcal/mol) [38]. Interestingly, compounds **1a**, **16a-d**, and **17a-c** with *ortho*, *meta*, and *para*-amino substitutions showed more NH₃ loss rather than water/alcohol loss, which is another indication that high PA of the neighboring substituent is not favored in cases where the NGP effect occurs.

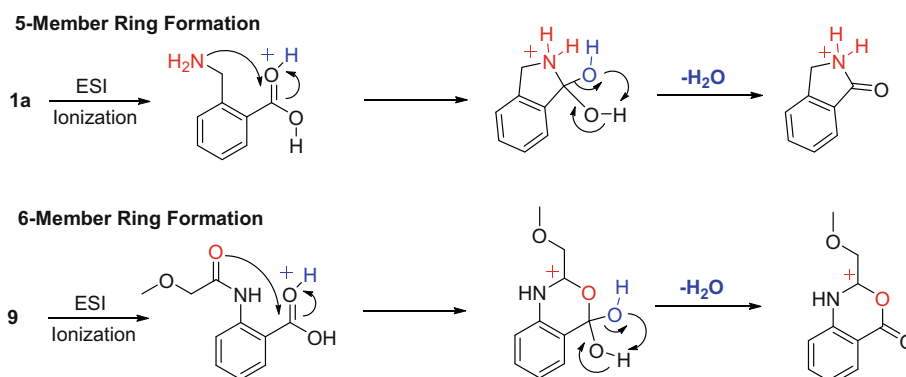
Steric Effect of Ring Formation

The formation of the 5- or 6-member rings following nucleophilic attack of the carbonyl was also seen to be impacted by the bulkiness of the groups attached to the nucleophile. As seen in **4** and **9**, the water loss was much greater in **4** (59.0%) than in **9** (4.0%). This difference in reactivity can be attributed to the steric penalty of the bulky isopropyl group attached to **9** compared with the smaller methoxyl group attached to **4**. It was also observed that **12a-d** had reduced neutral group losses (3.2%–21.2%) compared with **6a-d** (51.2%–100%). The differences in

these structures are the dimethyl substitution of the pyrazole rings in **12a-d** where as **6a-d** do not have these substituents. These methyl groups also create steric interference that reduces the stability of the ring formation.

Proximity of the Nucleophilic Group to the Carbonyl

The *ortho*, *meta*, and *para* relationships of the nucleophilic groups were also investigated in this study. As shown for **1a-d**, **16a-d**, and **17a-c**, the *ortho*-substituted primary amine exhibited significant neutral group loss that was not seen in the *meta*- and *para*-amines. Cyclization resulting from nucleophilic attack appears to be constrained to *ortho*-positioned nucleophilic groups. This effect was visible in numerous other positional isomeric compound pairs in this study, including **2a-b** and **8a-c**, **3a-b** and **18a-b**, **5a-d** and **10a-c**, **7a-d** and **15a-d**, and **12a-d** and **21a-d**. This differentiation based on nucleophilic group



Scheme 1. Proposed mechanism for neutral group loss based on the NGP effect. The protonation is on the carbonyl group to facilitate the subsequent intramolecular nucleophilic addition

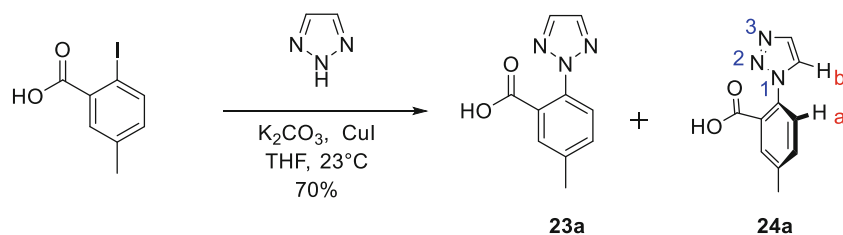


Figure 3. Intermediate step in synthesis of suvorexant [25]

placement proves to be an effective means for differentiating *ortho*-placed nucleophilic groups from *meta* and *para* groups by neutral group loss.

Studies of Two Triazole Acid **23a** and **24a** (Synthetic Intermediates for Suvorexant)

This research was also extended to an analysis of a benzoic acid building block of suvorexant (Belsomra), a commercially available drug produced by MSD approved by the FDA in 2014 for the treatment of insomnia. The synthesis of suvorexant includes two benzoic acid isomer intermediates as shown in Figure 3 after the triazole acid addition step [25–27]. These intermediates (shown in Figure 1) were also converted to esters and tested by UPLC-HRMS in order to investigate if MS spectrum differences can be found based on NGP effects. The MS¹ spectra and IMS drift time distributions for **23a** and **24a** can be found in Figure 4. The products were also studied with IMS, NMR, and density functional theory (DFT) calculations.

MSI Data

Table 1 summarizes the MS¹ data obtained for **23a-c** and **24a-c**. Among all acids and esters, it can be seen that significant neutral group loss of 100% occurs for **23a-c** while a minimal loss of 0%–4.4% occurs for **24a-c**. Model compound analysis suggests that significant neutral group loss would occur in both compounds due to the *ortho*-triazole ring to the carboxylic acid in both cases. Two factors were proposed to explain the low

reactivity of **24a-c**: (1) The main protonation site is on position 3 nitrogen instead of position 2 due to the PA difference, which would inhibit the following proton transfer from triazole to acid/ester group; (2) the triazole and phenyl ring were not favored to be synperiplanar due to the steric hindrance of the two hydrogens (Figure 5). A rotational barrier must be overcome to form cyclized neutral loss product for **24a** but not for **23a**. This hypothesis was further investigated using NMR spectroscopy, IMS-MS, and DFT analysis.

NMR Data

To investigate the orientation of **24a** towards the acid group, we acquired a 1D selective ROE applying selective inversion at the aromatic proton Hb. As shown in Figure 5 and Figure S49 (in Supporting Information), a strong ROE correlation was observed between protons Hb and Ha, indicating that they are close in the space (~2–3 Å distance), which would be consistent with the nitrogen at the 2-position oriented towards the carboxylic acid moiety. However, it must be noted that NMR could not determine the exact rotamer of **24a** in solution because the C–N bond was expected to rotate in solution. Evaluation of potential hydrogen bonding was carried out by acquiring 1D proton NMR for both **23a** and **24a**. Assuming that there is no hydrogen bonding in **23a** and due to the fact that proton spectra revealed that the OH proton had the same chemical shift in both isomers (Figures S49–S51), it was concluded that there was no hydrogen bonding in isomer **24a**

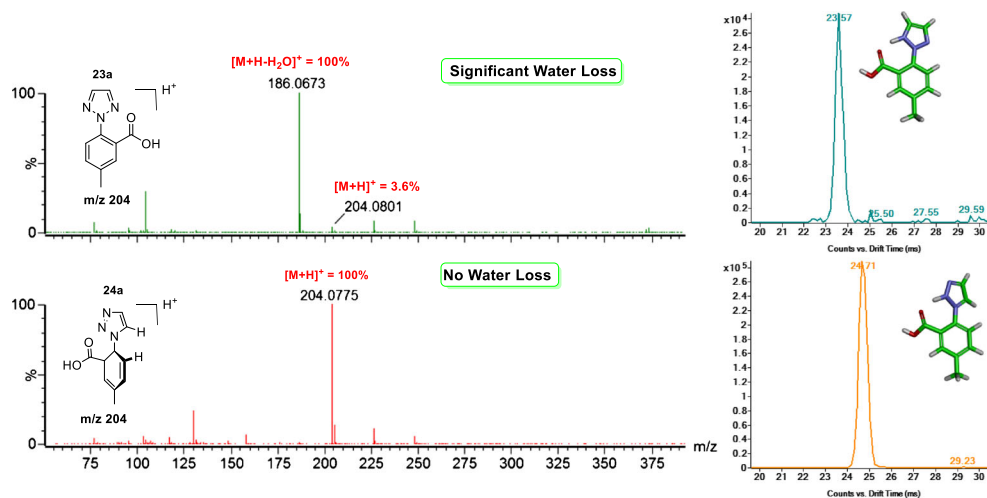


Figure 4. (Left) MS¹ spectra of **23a** and **24a** under (+)-ESI. (Right) IMS difference of **23a** and **24a** in drift time

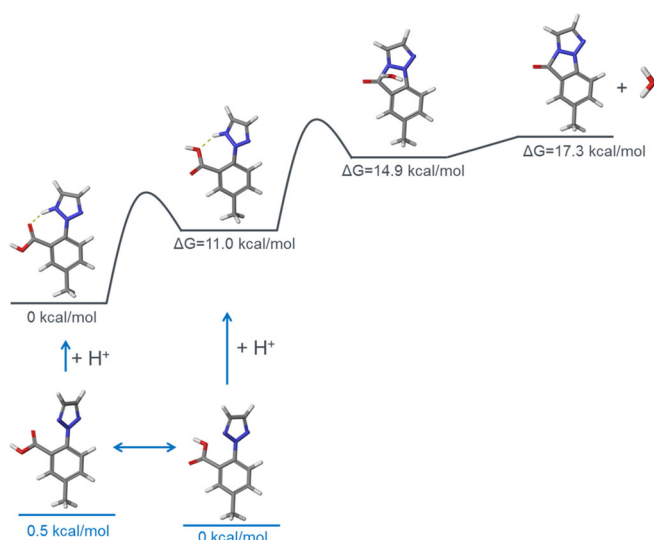


Figure 5. DFT optimization (B3LYP-D3-631++G**, gas phase) of relevant species for **23a**

either. Note that in the case of hydrogen bonding, the OH proton would have resonated at a different, more downfield chemical shift. In summary, the 1D selROE data were consistent with the nitrogen at the 2-position in **24a** pointing toward the carboxylic acid group. However, this could not be further confirmed by hydrogen bonding formation as it was not observed in proton NMR spectra.

DFT Calculations

The protonation and cyclization processes of **23a** and **24a** were also investigated using DFT calculations with results shown in Figures 5 and 6. The most stable species of **24a** in gas phase corresponds to an anti-planar conformation, where the triazole group is orthogonal to the phenyl ring. Therefore, the lack of neutral group loss in **24a** can be partially attributed to the position of the nucleophilic triazole group, which is not aligned

with the carbonyl reactive site. Thus, no nucleophilic attack can occur without a conformational rearrangement. Additionally, the 3-position is the most basic site in **24a**. As shown in Figure 6, in order for the proton transfer to occur, the nitrogen at the 2-position needs to be protonated, which is 6.5 kcal/mol higher in energy than the protonated isomer at position 3. This pathway was compared with the corresponding one for **23a** in which significant water loss is observed experimentally. DFT calculations show that the neutral species of **23a** has the triazole pointing towards the carbonyl group. Initial protonation of the triazole is likely leading to a rapid proton transfer to the hydroxyl portion of the carboxylic acid with subsequent addition and elimination of the water.

Ion Mobility Spectrometry-Mass Spectrometry Analysis

Ion mobility spectrometry (IMS) provides an additional dimension of analysis through gas-phase shape and charge separation of ionized analytes prior to MS detection [39]. In this study, IMS-MS was utilized to compliment NMR and computational efforts to distinguish isomer structures **23a** and **24a**. Ion mobility drift time distributions extracted from precursor m/z 204 are shown in Figure 4 for both triazole benzoic acid isomers analyzed in nitrogen drift gas. The faster drift time for isomer **23a** indicates a more compact structure compared to **24a**. The proposed mechanism for intramolecular cyclization and subsequent neutral loss prevalent in MS¹ of compound **23a** suggests the amine-containing neighboring group is positioned in close proximity to the acidic functional group, which is supported by the observed relative ordering of a higher mobility isomer **23a**. Despite the close structural similarity in these isomers, IMS allows for differentiation between the two protonated precursor ions. When combined with computational modeling, experimentally and theoretically derived collision cross-section (CCS) values can be compared to proposed all-atom structures

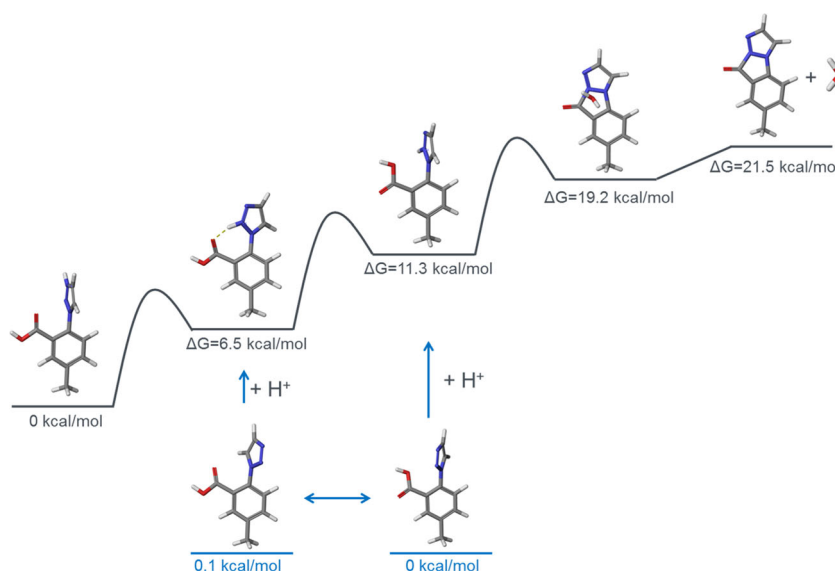


Figure 6. DFT optimization (B3LYP-D3-631++G**, gas phase) of relevant species for **24a**

[40]. Energy minimization of the DFT-generated protonated **23a** and **24a** structures, followed by calculation of CCS values by the trajectory method in MOBCAL yields theoretical CCS values in agreement with the experimentally measured CCS values; **23a** (77.7 \AA^2) has a smaller calculated trajectory method CCS than **24a** (80.1 \AA^2). MOBCAL calculates CCS with helium as the neutral collision gas, and thus experimentally derived CCS values in helium are reported for direct comparison with the calculations with values of 78.3 \AA^2 for **23a** and 79.7 \AA^2 for **24a**. Experimental and theoretical CCS (trajectory method) values are typically considered to be in agreement when values are within 2% relative difference [41]; this is the case for these two isomers, with **23a** (0.8% difference) and **24a** (0.5% difference). Characterization of subtle structural differences between these two triazole benzoic acid isomers through a combination of NMR, IMS-MS, and computational modeling helps reinforce the conclusions drawn from each of these complementary structure elucidation approaches, and supports the hypothesis of water loss due to a gas-phase NGP effect for the scaffolds investigated in this study.

Conclusion

Ortho-substituted benzoic acid/ester derivatives were identified among other positional isomers using HRMS¹ by the neighboring group participation (NGP) effect. A total of 65 model compounds were tested, and it was determined that the NGP effect depends on three factors: the proton affinity of the nucleophile, the bulkiness of the nucleophile, and the positional isomer (*ortho*- compared to *meta*- and *para*-nucleophile). The trends found for the model compounds were then extended to investigate two triazole isomers that form during the synthesis of the active pharmaceutical ingredient (API) suvorexant (Belsomra®). MS¹ spectra differentiated the isomers based on the NGP effect, as **23** showed large neutral group loss whereas **24** did not. Further IMS-MS, NMR and DFT studies were also conducted to show the non-reactivity of **24** (in terms of neutral loss during MS¹) can be attributed to two main factors: 1) higher PA of the nitrogen at the 3-position of the triazole; and 2) the protonated triazole in **24** is likely oriented 90 degrees, orthogonal to the carboxylic acid in the gas phase, disfavoring nucleophilic attack by the nitrogen at the 2-position. The results from this work aid structural elucidation of substituted benzoic acid/ester isomers through facile interrogation of the gas-phase NGP effect in MS¹ analysis.

Acknowledgments

The authors thank the MSD Summer Internship Program for funding, Dr. Gary Martin for helpful discussions, and Dr. R.

Thomas Williamson and Dr. Caroline McGregor for their managerial support.

References

1. Chipley, J.R.: 2 Sodium benzoate and benzoic acid. In: Davidson, P.M., Sofos, J.N., Branen, A.L. (eds.) Antimicrobials in food, 3rd ed., p. 11. CRC Press Taylor and Francis Group, Boca Raton (2005)
2. Shrewsbury, R.P.: University of North Carolina Eshelman School of Pharmacy. The Pharmaceutics and Compounding Laboratory. Available: http://pharmlabs.unc.edu/appendix_resources.htm. Accessed 11 Oct 2017
3. United States Food and Drug Administration (FDA). Code of Federal Regulations Title 21. Volume3. Available: <http://www.accessdata.fda.gov/scripts/cdrh/cfdocs/cfcfr/CFRSearch.cfm?fr=184.1021>. Accessed 11 Oct 2017
4. Malz, F., Jancke, H.: Purity assessment problem in quantitative NMR-impurity resonance overlaps with monitor signal multiplets from stereoisomers. *Anal. Bioanal. Chem.* **385**, 760–765 (2006)
5. Szafran, Z.: Structural Isomer Identification via NMR. *J. Chem. Educ.* **62**, 260–261 (1985)
6. Queiroz, S.L., de Araujo, M.P., Batista, A.A., Macfarlane, K.S., James, B.R.: Synthesis of [RuCl₂(dppb)(PPh₃)] and identification of the cis- and trans-[RuCl₂(dppb)(phen)] geometrical isomers via 31P{1H} NMR spectroscopy. *J. Chem. Educ.* **78**, 87–88 (2001)
7. Kikuchi, K., Nakahara, N., Wakabayashi, T., Suzuki, S., Shiromaru, H., Miyake, Y., Saito, K., Ikemoto, I., Kainosho, M., Achiba, Y.: NMR characterization of isomers of C₇₈, C₈₂, and C₈₄ fullerenes. *Nature*. **357**, 142–145 (1992)
8. Forseth, R.R., Schroeder, F.C.: NMR-spectroscopic analysis of mixtures: from structure to function. *Curr. Opin. Chem. Biol.* **15**, 38–47 (2011)
9. Caldearelli, S.: Chromatographic NMR: a tool for the analysis of small molecules. *Magn. Reson. Chem.* **45**, 548–555 (2007)
10. Slabizki, P., Legrum, C., Meusinger, R., Schmar, H.G.: Characterization and analysis of structural isomers of dimethyl methoxy-pyrazines in cork stoppers and ladybugs (*Harmonia axyridis* and *Coccinella septempunctata*). *Anal. Bioanal. Chem.* **406**, 6429–6439 (2014)
11. Babushok, V.I.: Chromatographic retention indices in identification of chemical compounds. *Trends Anal. Chem.* **69**, 98–104 (2015)
12. Pacakova, V., Feltl, V.: Chromatographic retention indices, an aid to identification of organic compounds. E. Horwood, New York (1992)
13. Wenig, P., Odermatt, J.: Efficient analysis of Py-GC/MS data by a large scale automatic database approach: an illustration of white pitch identification in pulp and paper industry. *J. Anal. Appl. Pyrolysis.* **87**, 85–92 (2010)
14. Zhang, L., Tang, C., Cao, D., Zeng, Y., Tan, B., Zeng, M., Fen, W., Xiao, H., Liang, Y.: Strategies of structure elucidation of small molecules using gas chromatography-mass spectrometric data. *Trends Anal. Chem.* **47**, 37–46 (2013)
15. Alam, M.S., Stark, C., Harrison, R.M.: Using variable ionization energy time-of-flight mass spectrometry with comprehensive GCxGC to identify isomeric species. *Anal. Chem.* **88**, 4211–4220 (2016)
16. Zhang, L., Cheng, X.L., Liu, Y., Liang, M., Dong, H., Lv, B., Yang, W., Luo, Z., Tang, M.: Identification of the related substances in ampicillin capsule by rapid resolution liquid chromatography coupled with electrospray ionization tandem mass spectrometry. *J. Anal. Methods Chem.* **2014**, 1–15 (2014)
17. International Union of Pure and Applied Chemistry (IUPAC). Compendium of Chemical Terminology, 2nd ed. (the “Gold Book”). Compiled by A. D. McNaught and A. Wilkinson. Blackwell Scientific Publications: Oxford (1997) XML on-line corrected version: <http://goldbook.iupac.org> (2006-) created by M. Nic, J. Jirat, B. Kosata; updates compiled by A. Jenkins. Accessed 11 July 2016
18. Capon, B.: Neighboring group participation. *Q. Rev. Chem. Soc.* **18**, 45–111 (1964)
19. Bigler, L., Hesse, M.: Neighboring group participation in the electrospray ionization tandem mass spectra of polyamine toxins of spiders. Part 1: α,ω -Diaminoalkane compounds. *J. Am. Soc. Mass Spectrom.* **6**, 634–637 (1995)
20. Darshan, D.V., Chandar, B.G.N., Srujan, M., Chaudhuri, A., Prabhakar, S.: Electrospray ionization tandem mass spectrometry study of six

- isomeric cationic amphiphiles with ester/amide linker. *Rapid Commun. Mass Spectrom.* **28**, 1209–1214 (2014)
21. Tu, Y.-P., Harrison, A.G.: Fragmentation of protonated amides through intermediate ion-neutral complexes: neighboring group participation. *J. Am. Soc. Mass Spectrom.* **9**, 454–x (1998)
 22. Hu, W., Reder, E., Hesse, M.: Neighboring-group participation in the mass-spectral decomposition of 4-hydroxycinnamoyl-spermidines. *Helv. Chir. Acta.* **79**, 2137–2151 (1996)
 23. Cooks, R.G., Wolfe, N.L., Curtis, J.R., Petty, H.E., McDonald, R.N.: Neighboring-group participation reactions in the mass spectral fragmentations of some azulenes. Comparisons with solvolytic processes. *J. Organomet. Chem.* **35**, 4048–4054 (1970)
 24. Reid, G.E., Simpson, R.J., O’Hair, R.A.J.: Leaving group and gas phase neighboring group effects in the side chain losses from protonated serine and its derivatives. *J. Am. Soc. Mass Spectrom.* **11**, 1047–1060 (2000)
 25. Mangion, I.K., Sherry, B.D., Yin, J., Fleitz, F.J.: Enantioselective synthesis of a dual orexin receptor antagonist. *Org. Lett.* **14**, 3458–3461 (2012)
 26. Strotman, N.A., Baxter, C.A., Brands, K.M.J., Cleator, E., Krska, S.W., Reamer, R.A., Wallace, D.J., Wright, T.J.: Reaction development and mechanistic study of a ruthenium catalyzed intramolecular asymmetric reductive amination en route to the dual orexin inhibitor Suvorexant (MK-4305). *J. Am. Chem. Soc.* **133**, 8362–8371 (2011)
 27. Baxter, C.A., Cleator, E., Brands, K.M.J., Edwards, J.S., Reamer, R.A., Sheen, F.J., Stewart, G.W., Strotman, N.A., Wallace, D.J.: The first large-scale synthesis of MK-4305: a dual orexin receptor antagonist for the treatment of sleep disorder. *Org. Process. Res. Dev.* **15**, 367–375 (2011)
 28. Mesleh, M.F., Hunter, J.M., Shvartsburg, A.A., Schatz, G.C., Jarrold, M.F.: Structural information from ion mobility measurements: effects of the long-range potential. *J. Phys. Chem.* **100**, 16082–16086 (1996)
 29. OMEGA 2.5.1.4: OpenEye Scientific software, Santa Fe, NM. Available: <http://www.eyesopen.com>. Accessed 3 Mar 2017
 30. Hawkins, P.C.D., Skillman, A.G., Warren, G.L., Ellingson, B.A., Stahl, M.T.: Conformer generation with OMEGA: algorithm and validation using high quality structures from the Protein Databank and Cambridge Structural Database. *J. Chem. Inf. Model.* **50**, 572–584 (2010)
 31. Jaguar, version 9.1, Schrödinger, Inc.: New York, NY (2016)
 32. Bochevarov, A.D., Harder, E., Hughes, T.F., Greenwood, J.R., Braden, D.A., Philipp, D.M., Rinaldo, D., Halls, M.D., Zhang, J., Friesner, R.A.: Jaguar: a high-performance quantum chemistry software program with strengths in life and materials sciences. *Int. J. Quantum Chem.* **113**, 2110–2142 (2013)
 33. Lee, C.T., Yang, W.T., Parr, R.G.: Development of the Colle-Salvetti correlation-energy formula into a functional of the electron-density. *Phys. Rev. B.* **37**, 785–789 (1988)
 34. Becke, A.D.: A new mixing of Hartree-Fock and local density-functional theories. *J. Chem. Phys.* **98**, 1372–1377 (1993)
 35. Johnson, B.G., Gill, P.M.W., Pople, J.A.: The performance of a family of density functional methods. *J. Chem. Phys.* **98**, 5612–5626 (1993)
 36. Grimme, S., Antony, J., Ehrlich, S., Krieg, H.: A consistent and accurate ab initio parametrization of density functional dispersion correction (DFT-D) for the 94 elements H-Pu. *J. Chem. Phys.* **132**, 154104 (2010)
 37. Grimme, S., Ehrlich, S., Goerigk, L.: Effect of the damping function in dispersion corrected density functional theory. *J. Comput. Chem.* **32**, 1456–1465 (2011)
 38. National Institute of Standards and Technology (NIST). Proton affinity search. Available: <http://webbook.nist.gov/chemistry/pa-ser.html>. Accessed 27 July 2016
 39. Wyttenbach, T., Pierson, N.A., Clemmer, D.E., Bowers, M.T.: Ion mobility analysis of molecular dynamics. *Annu. Rev. Phys. Chem.* **65**, 175–196 (2014)
 40. Kim, H., Kim, H.I., Johnson, P.V., Beegle, L.W., Beauchamp, J.L., Goddard, W.A., Kanik, I.: Experimental and theoretical investigation into the correlation between mass and ion mobility for choline and other ammonium cations in N₂. *Anal. Chem.* **80**, 1928–1936 (2008)
 41. Ho, K.-M., Shvartsburg, A.A., Pan, B., Lu, Z.-Y., Wang, C.-Z., Wacker, J.G., Fye, J.L., Jarrold, M.F.: Structures of medium-sized silicon clusters. *Nature.* **392**, 582–585 (1998)