

Ether Cleavage Re-Investigated: Elucidating the Mechanism of BBr₃-Facilitated Demethylation of Aryl Methyl Ethers

Talon M. Kosak,^[a] Heidi A. Conrad,^[a] Andrew L. Korich,^{*[a]} and Richard L. Lord^{*[a]}

Keywords: Reaction mechanisms / Ether cleavage / Density functional calculations

One of the most well-known, highly utilized reagents for ether cleavage is boron tribromide (BBr₃), and this reagent is frequently employed in a 1:1 stoichiometric ratio with ethers. Density functional theory calculations predict a new mechanistic pathway involving charged intermediates for ether cleavage in aryl methyl ethers. Moreover, these calculations predict that one equivalent of BBr₃ can cleave up to three

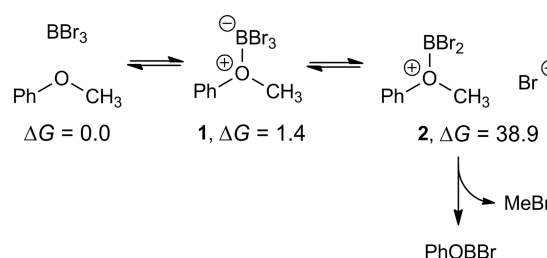
equivalents of anisole, producing triphenoxyborane [B(OPh)₃] prior to hydrolysis. These predictions were validated by gas chromatography analysis of reactions where the BBr₃:anisole ratio was varied. Not only do we confirm that sub-stoichiometric equivalents may be used for ether demethylation, but the findings also support our newly proposed three cycle mechanism for cleavage of aryl methyl ethers.

Introduction

Boron tribromide is a versatile reagent utilized in diverse areas ranging from polymer chemistry to natural product synthesis.^[1] Owing its high reactivity to the Lewis acidic boron center, BBr₃ reactions include haloborylation,^[2] boron–silicon exchange,^[3] and rearrangement of 7,7-diphenylhydromorphone derivatives.^[4] While there is no shortage in the diversity of BBr₃-mediated reactions, many of the mechanisms for these transformations have not been fully elucidated. In this report we investigate the mechanism of ether cleavage by BBr₃^[5–10] in anisole.

Conceptually, demethylation of anisole is initiated by the formation of an ether adduct **1** followed by the loss of bromide. Free bromide nucleophilically attacks the methyl group of the cationic intermediate (**2**) cleaving the C–O bond and producing PhOBBr₂, which undergoes hydrolysis upon aqueous work-up. While this pathway (Scheme 1) at first appears to be viable, we calculated that the formation of **2** and bromide in dichloromethane is thermodynamically inaccessible ($\Delta G = +38.9$ kcal/mol).

Recently, alternative mechanisms for ether cleavage were proposed by Sousa and Silva that involve unimolecular or bimolecular rate-determining steps that circumvent formation of bromide in solution (Scheme 2).^[11] While a unimo-



Scheme 1. Conceptual reaction mechanism for demethylation of anisole is thermodynamically inaccessible due to formation of **2** and bromide. Gibbs energies are in kcal/mol.

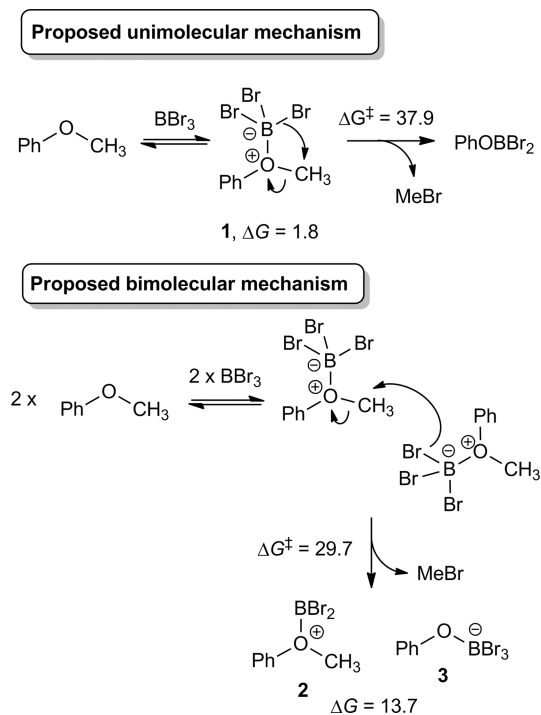
lecular process is kinetically favored for ethers containing one or more substituents (e.g. branched alkyl) that stabilize carbocation character in an S_N1-like transition state, this barrier for demethylation of primary C atoms, like in the methyl group of anisole, lies too high on the potential energy surface to be accessible under reported reaction conditions. They found that a bimolecular process (Scheme 2, bottom) decreases the kinetic barrier for anisole demethylation significantly. During this reaction pathway, one of the bromides of the first ether adduct nucleophilically attacks the methyl group of the second ether adduct. This is analogous to an S_N2 reaction with 180° attack of the methyl group by a bromide in the nucleophilic ether adduct. However, this bimolecular pathway produces two highly charged intermediates **2** and **3** that Sousa and Silva did not investigate. Their computational investigation stopped with the calculation of the initial kinetic barrier.^[11] We speculate that these charged intermediates may undergo a similar bimolecular reaction to yield two equivalents of PhOBBr₂ and MeBr. Moreover, if charged intermediates are formed then we believe an important set of mechanistic pathways may have been overlooked, namely, those where Lewis acidic BBr₃ abstracts bromide from the ether complex to form

[a] Grand Valley State University, Department of Chemistry
1 Campus Drive, Allendale, MI 49401, USA
E-mail: koricha@gvsu
lordri@gvsu.edu
http://www.gvsu.edu/chem

Supporting information for this article is available on the WWW under <http://dx.doi.org/10.1002/ejoc.201501042>.

© 2015 The Authors. Published by Wiley-VCH Verlag GmbH & Co. KGaA. This is an open access article under the terms of the Attribution Non-Commercial License, which permits use, distribution and reproduction in any medium, provided the original work is properly cited and is not used for commercial purposes.

BBr_4^- in a mechanism related to the pathway introduced in Scheme 1.



Scheme 2. Previously proposed unimolecular and bimolecular pathways proposed by Sousa and Silva.^[11] Gibbs energies are in kcal/mol.

In this manuscript, we re-investigate the mechanisms of anisole ether cleavage proposed by Sousa and Silva and compare them to new mechanisms involving charged intermediates using density functional theory coupled to a continuum solvation model. The kinetically favored pathway explains mechanistic intricacies regarding the reactivity of all three bromides on BBr_3 , a finding that we confirm with ^1H NMR and GC experiments run with different stoichiometries. Our proposed mechanism rationalizes the key observation of 0.33:1 BBr_3 :ether stoichiometry reported in the original methodology paper^[5] and is supported by additional experimental findings.

Results and Discussion

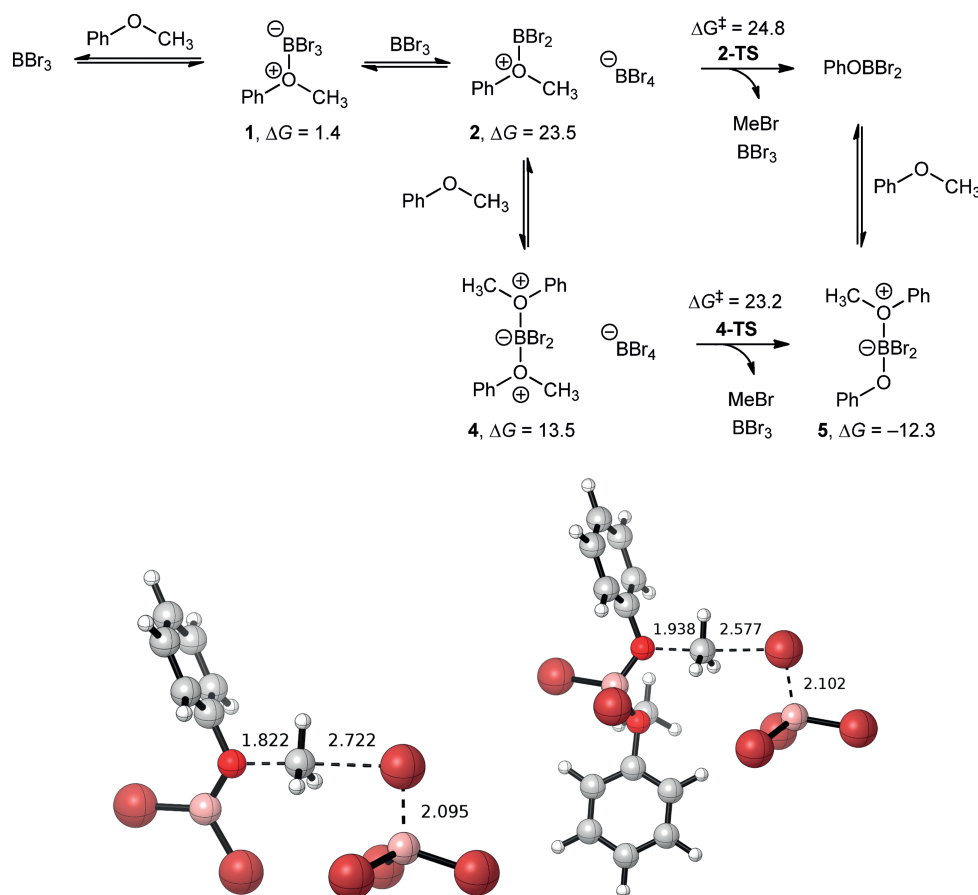
The computational investigation began by replicating the uni- and bimolecular pathways proposed by Sousa and Silva for the demethylation of anisole. Our methodology differs slightly from their method because we did not use a pseudopotential for Br, did not scale computed frequencies when evaluating zero-point and thermal corrections, and we assumed a temperature of 298 K vs. 250 K (though this difference has a negligible impact). We also avoided mixed basis sets in our triple-zeta energy refinements by employing 6-311+G(d,p) for all elements. Our method reproduces well their geometry (Figure S1 in the Supporting Information) and kinetic barrier ($\Delta G^\ddagger = 37.9$ kcal/mol vs. 38.8 kcal/mol) for the unimolecular pathway. We were also able to replicate

a similar geometry for the bimolecular pathway (Figure S2 in the Supporting Information), however, we were unable to replicate their low barrier of 13.3 kcal/mol. Formation of the ether adduct is only slightly uphill in Gibbs energy ($\Delta G = 1.4$ kcal/mol) and therefore anisole, BBr_3 , and the ether adduct are predicted to be in equilibrium in solution. The loss of translational entropy due to two of the ether adducts coming together at the bimolecular transition state suggests that ca. 10 kcal/mol of this barrier is entropic.^[12] Consequently, nucleophilic attack of the methyl group must only invoke a ca. 3 kcal/mol enthalpic barrier based on their reported value, a finding that we struggled to rationalize. We calculated an enthalpic barrier (relative to two ether adducts) of 14.9 kcal/mol and an entropic barrier of 12.0 kcal/mol for an overall Gibbs energy barrier of 26.9 kcal/mol (or 29.7 kcal/mol vs. free BBr_3 and anisole). This transition state leads to MeBr and two charged intermediates, **2** and **3**, that are uphill by 13.7 kcal/mol. Thus, while this bimolecular pathway is a significant improvement over the unimolecular pathway, its kinetic barrier is high enough to warrant the investigation of alternative mechanisms based on the experimental conditions. Three important concepts from the Sousa and Silva bimolecular mechanism are: (i) ether adduct formation makes bromide a nucleophile without generation of free bromide, (ii) ether adduct formation makes the methyl group more susceptible to nucleophilic attack, and (iii) the local geometry of the methyl group at the transition state is $\text{S}_{\text{N}}2$ -like.

A New Demethylation Mechanism

Using the key insights from Sousa and Silva's mechanism, we envisioned that BBr_3 could abstract bromide from an ether adduct to afford **2** and BBr_4^- (Scheme 3, top). These charged intermediates are computed to be thermodynamically unfavorable compared to formation of two ether complexes (23.5 kcal/mol vs. 2.8 kcal/mol) but still accessible under the experimental conditions. Importantly, this energy is below the previously proposed bimolecular kinetic barrier by ca. 6 kcal/mol. Because boron tetrabromide is less stable, this masked bromide should be a more potent nucleophile than the bromide in **1**. The barrier for nucleophilic attack of the methyl group in **2** by BBr_4^- is found to be only 1.3 kcal/mol higher in energy than these intermediates, or 24.8 kcal/mol (**2-TS**, Scheme 3 bottom left). This reaction produces MeBr , BBr_3 , and PhOBBR_2 with an overall reaction energy of -21.8 kcal/mol. The critical feature of our mechanism is the dynamic equilibrium at the boron center that stabilizes bromide until it is needed for nucleophilic attack.

This idea of dynamic equilibrium at boron made us consider further possibilities for stabilization of the charged intermediates. Boron is coordinatively-saturated in BBr_4^- , however, three-coordinate boron in **2** could be stabilized by an additional Lewis base. Thus, we evaluated the thermodynamics of Lewis base stabilization of **2** by anisole to afford **4**, which is favorable by 10.0 kcal/mol (+13.5 kcal/mol vs.



Scheme 3. Top: calculated mechanism for demethylation of anisole. Bottom left: transition state (**2-TS**) for demethylation of mono-ether adduct **2**. Bottom right: transition state (**4-TS**) for demethylation of di-ether adduct **4**. Gibbs energies are in kcal/mol and bond lengths are listed in Å.

reactants). Demethylation of **4** by BBr_4^- is predicted to have a slightly lower kinetic barrier (**4-TS**, Scheme 3 bottom right) of 23.2 vs. 24.8 kcal/mol for **2** and generates MeBr, BBr_3 , and **5** with an overall reaction energy of -12.3 kcal/mol. PhOBBr_2 may be formed via this pathway upon dissociation of anisole from **5** (-9.5 kcal/mol). Within the intrinsic errors of our computational methodology one cannot predict whether the barrier involving **2** or **4** is preferred and both may be operative. However, both pathways are favored over the previously proposed bimolecular mechanism. The transition state structures for the mono- and di-ether adduct pathways are shown at the bottom of Scheme 3. Both feature $\text{S}_{\text{N}}2$ -like geometries with a trigonal bipyramidal carbon center. **2-TS** is earlier than **4-TS**, as evidenced by the shorter C–O bond length of 1.82 vs. 1.94 Å and longer C–Br bond length of 2.72 vs. 2.58 Å. This finding is consistent with the stability of the cation adducts **2** and **4**; the more reactive intermediate **2** has an earlier transition state. **2-TS** and **4-TS** are much earlier than the bimolecular transition state reported by Sousa and Silva that had C–O and C–Br bond lengths of 2.09 and 2.42 Å (Figure S2 in Supporting Information), respectively.^[11]

Bromide loss from the ether adduct should become more accessible with more stabilizing substituents on boron, like phenoxide. Thus, while we computed **5** to be slightly uphill

in energy relative to PhOBBr_2 , this adduct is still thermodynamically accessible and features a tetracoordinate boron center containing two bromides. The loss of bromide from **5** to BBr_3 yields BBr_4^- and **6**. **6** may either undergo demethylation facilitated by BBr_4^- to produce $(\text{PhO})_2\text{BBr}$ or may coordinate an additional anisole to generate intermediate **7** prior to demethylation to form **8** (Scheme 4, top). One could envision similar reactivity occurring until all bromides are lost from the boron center. This concept of multiple demethylation cycles per BBr_3 equiv. is consistent with the original methodology paper that reported a 3:1 ether: BBr_3 stoichiometry.^[5] Because the reverse barrier from PhOBBr_2 to **2** is disfavored by 44.9 kcal/mol, we now reference our thermodynamics to PhOBBr_2 , anisole, and BBr_3 . Bromide abstraction from **5** to produce **6** and BBr_4^- is computed to be uphill by 22.5 kcal/mol. Similar to the first reaction cycle that we considered (Scheme 3, top), **6** can be stabilized by anisole to generate **7**. This reaction is only downhill by 3.9 kcal/mol ($+18.6$ kcal/mol vs. PhOBBr_2) as compared to the stabilization of **2** to form **4** by 10.0 kcal/mol. Already we see a marked difference from the first cycle; **6** is much more stable, as compared to **2**, due to the π -donor ability of the phenoxide substituent. The transition state involving **6** is 25.3 kcal/mol (**6-TS**) while the transition state for **7** is 32.5 kcal/mol (**7-TS**). Thus, the mono-ether adduct path-

way is kinetically favored in this second reaction cycle. **6-TS** produces MeBr, BBr₃, and (PhO)₂BBr with a reaction energy of -17.0 kcal/mol. A summary of this reaction mechanism is shown in Scheme 4 (top) and the rate-limiting transition state is shown in Figure 1. The C–O and C–Br bond lengths in **6-TS** of 1.87 and 2.64 Å, respectively, are intermediate between those of the transition states in the first cycle and further highlight the stability provided by the phenoxide substituent.

One can further envision a third cycle that would react the remaining bromide in (PhO)₂BBr (Scheme 4, bottom). Because the reverse reaction from the product of cycle 2 to form **6** is uphill by 36.4 kcal/mol, all energies are referenced to (PhO)₂BBr, anisole, and BBr₃ in cycle 3. Formation of **8** from (PhO)₂BBr and anisole is uphill by 11.1 kcal/mol. Abstraction of bromide from **8** by BBr₃ generates **9** and BBr₄[−] and is further uphill by 4.9 kcal/mol [16.0 kcal/mol vs. (PhO)₂BBr]. Intermediate **9**, which features a three-coordinate boron center, may bind anisole to form **10** and become coordinatively saturated. This adduct formation only slightly stabilizes **9** by 1.1 kcal/mol [14.9 kcal/mol vs. (PhO)₂BBr]. Similar to the second cycle, the kinetic barrier

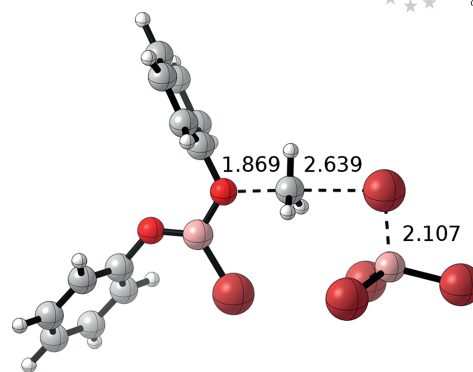
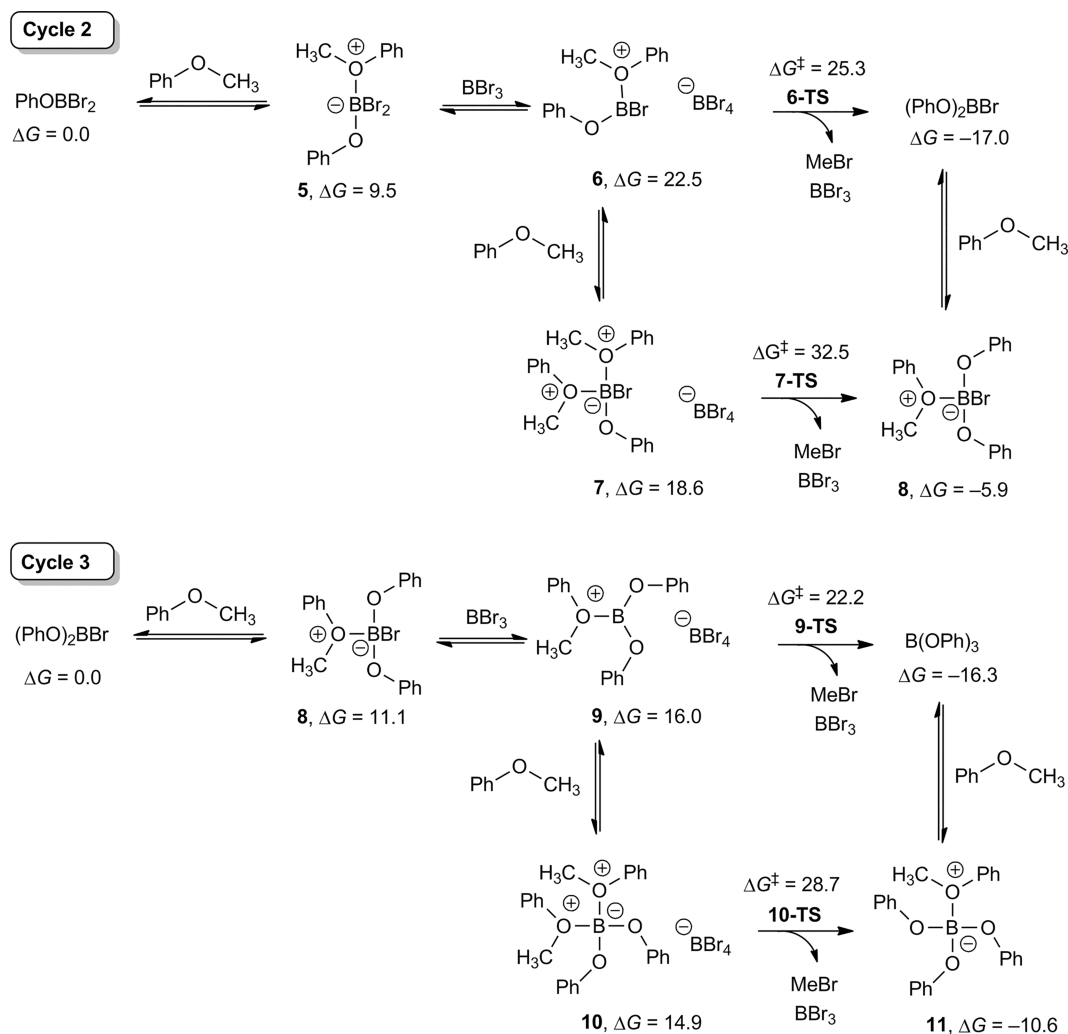


Figure 1. Optimized structure for **6-TS**. Bond lengths are listed in Å.

for the mono-ether adduct intermediate **9** is much lower than the di-ether adduct intermediate **10** (22.2 vs. 28.7 kcal/mol for **9-TS** and **10-TS**). The favored kinetic pathway **9-TS** generates MeBr, BBr₃, and B(OPh)₃, and is calculated to be exergonic by 17.6 kcal/mol. The C–O and C–Br bond lengths in **9-TS** of 1.92 and 2.60 Å, respectively, indicate a



Scheme 4. Cycles 2 and 3 for proposed mechanism for BBr₃-facilitated ether cleavage. Gibbs energies are in kcal/mol.

later transition state than cycle 2 with bond lengths nearly equal to that of **4-TS** in cycle 1. Two questions come to mind from these calculations: (i) why does the third cycle have the lowest barrier and (ii) why does the di-ether adduct pathway (**4-TS**, **7-TS**, and **10-TS**) become kinetically disfavored after cycle 1? (Figure 2).

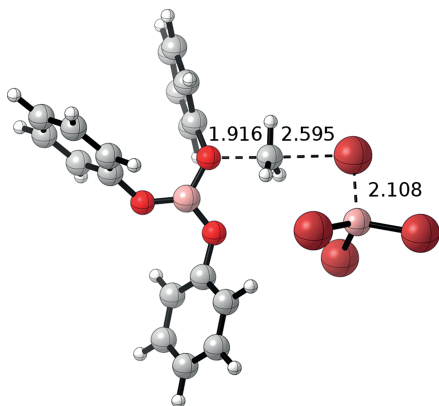


Figure 2. Optimized structure for **9-TS**. Bond lengths are listed in Å.

Analysis of Reaction Barrier Heights

To answer the first question, we decomposed the rate-limiting barrier for the mono-ether adduct pathways for all three cycles in terms of the individual reaction steps. The first step towards demethylation is formation of the Lewis acid/base adduct in each cycle (Table 1, column 1). Our calculations predict that this reaction becomes increasingly unfavorable as bromides are substituted with phenoxides, as evidenced by the increasing reaction energy of $1.4 < 9.5 < 11.1$ kcal/mol. This trend can be rationalized by the π -donor capability of phenoxide that reduces the Lewis acidity of the boron center. Next, each of these mono-ether adducts (**1**, **5**, and **8**) loses bromide to BBr_3 . Additional phenoxide substituents bound to the boron center that loses bromide should stabilize the cationic boron species formed in this reaction (**2**, **6**, and **9**). This prediction is observed in the reaction free energies of 22.1, 13.0, and 4.9 kcal/mol for cycles 1, 2, and 3, respectively (Table 1, column 2). Finally, these cationic intermediates react with boron tetrabromide in the rate-limiting steps **2-TS**, **6-TS**, and **9-TS** to form MeBr , BBr_3 , and $(\text{PhO})_x\text{BBr}_{3-x}$. This rate-limiting barrier increases with the number of phenoxide substituents because a less Lewis acidic boron center will make the oxygen of the coordinated anisole less positive, thereby disfavoring

demethylation (Table 1, column 3). While each of these individual steps are easily rationalized by the ratio of bromide:phenoxide substituents at the boron center, the overall barrier trend of 24.8, 25.3, and 22.2 kcal/mol for cycles 1, 2, and 3, does not dominantly arise from one of the three fundamental reaction steps (i.e. none of these individual reaction steps govern the rate for each cycle).

To answer the second question, we decomposed the binding energy of the second barrier starting from the cationic intermediates **2**, **6**, and **9**, whose thermodynamics are already summarized in the first two columns of Table 1. The subsequent steps are summarized in Table 2. Adduct formation between the cationic intermediates and a second equivalent of anisole steadily decreases in exergonicity from -10.0 to -1.1 kcal/mol as the number of phenoxide substituents at the boron center increases (Table 2, column 1). The kinetic barrier, as measured from these di-ether adduct intermediates **4**, **7**, and **10**, increases slightly for cycles 2 and 3. This finding suggests that additional phenoxide substituents are counterproductive to demethylation, as evidenced by the demethylation energies in Table 2 ($+9.7$ to $+13.8$ kcal/mol) vs. Table 1 ($+1.3$ to $+6.2$ kcal/mol). One can rationalize this based on the reduced Lewis acidity with increasing numbers of phenoxides and with increasing of the coordination number at boron from three to four. The trend in binding of the first anisole and formation of the charged species (Table 1, sum of columns 1 and 2) results in energies of 23.5, 22.5, and 16.0 kcal/mol for cycles 1–3, respectively. When combined with the second equivalent of anisole (Table 2, column 1), one observes overall energies of 13.5, 18.6, and 14.9 kcal/mol to reach **4-TS**, **7-TS**, and **10-TS**. The more costly energetic penalties for **7-TS** and **10-TS** (13.8 kcal/mol) increases the gap for the di-ether adduct barrier in cycles 2 and 3 as compared to cycle 1. One might ask why **10-TS** is not higher in energy? While **10** maintains a weakly coordinated anisole, both **10-TS** and **11** dissociate the additional anisole to form a weak van der Waal complex. Thus, while cycle 1 has competitive kinetics for the mono- and di-ether adduct pathways, the boron center with

Table 2. Breakdown of rate-limiting barrier free energies for the di-ether adduct pathways of cycles 1, 2, and 3 in terms of the fundamental reaction free energies leading to that barrier. All energies are in kcal/mol.

Second anisole binding $\text{C} + \text{PhOMe} \leftrightarrow \text{E}$	Demethylation $\text{E} + \text{BBr}_4^- \rightarrow \text{E-TS}$	Total ΔG^\ddagger
C = 2 , E = 4 -10.0	4-TS $+9.7$	23.2
C = 6 , E = 7 -3.9	7-TS $+13.8$	32.5
C = 9 , E = 10 -1.1	10-TS $+13.8$	28.7

Table 1. Breakdown of rate-limiting barrier free energies for the mono-ether adduct pathways of cycles 1, 2, and 3 in terms of the fundamental reaction free energies leading to that barrier. All energies are in kcal/mol.

Anisole binding $\text{A} + \text{PhOMe} \leftrightarrow \text{B}$	Bromide loss $\text{B} + \text{BBr}_3 \leftrightarrow \text{C} + \text{BBr}_4^-$	Demethylation $\text{C} + \text{BBr}_4^- \rightarrow \text{C-TS}$	Total ΔG^\ddagger
A = BBr₃ , B = 1 $+1.4$	C = 2 $+22.1$	2-TS $+1.3$	24.8
A = PhOBBr₂ , B = 5 $+9.5$	C = 6 $+13.0$	6-TS $+2.8$	25.3
A = (PhO)₂BBr , B = 8 $+11.1$	C = 9 $+4.9$	9-TS $+6.2$	22.2

phenoxide substituents in cycles 2 and 3 is not sufficiently Lewis acidic for the di-ether adduct pathway to be viable because decreased Lewis acidity (i) disfavors binding of the second ether equivalent and (ii) raises the barrier height for the demethylation step.

Experimental Validation

In order to validate our computational predictions, we aimed to reproduce and expand the findings disclosed in the original methodology report by varying the BBr_3 :anisole ratio. We began our investigation using standard conditions^[10] for this reaction: CH_2Cl_2 as the reaction solvent, near ambient temperature, and a reaction time of ca. 18 h. However, the reaction solvent was varied from CH_2Cl_2 to CDCl_3 for NMR analysis and to *o*-dichlorobenzene for the high temperature reactions.

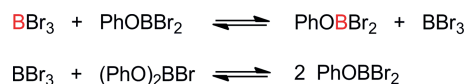
Table 3 shows the quantitative GC results of varied the molar equivalence of BBr_3 :anisole. Not surprisingly a 1:1 ratio results in near quantitative conversion to phenol after hydrolysis. Decreasing the ratio to 0.66:1 only decreases the efficiency of the reaction from 99.8 to 95.8%, which is within our experimental error and demonstrates more than one cycle must be operative. Monitoring the reaction by ^1H NMR shows disappearance of the methoxy resonance at 3.6 ppm and formation of the MeBr resonance at 2.6 ppm after 18 h at room temperature (Figure S3 in Supporting Information). However, when the ratio is decreased further to 0.33:1 a marked decrease in the conversion to 56.9% is observed. Our GC data demonstrates that this inefficiency is due to unreacted anisole rather than formation of by-products. Our computational mechanism predicts that we should observe full conversion at 0.33:1 because the second cycle had the highest barrier of 25.3 kcal/mol, and if there is sufficient energy to overcome the second barrier then the third cycle, with a lower barrier of 22.2 kcal/mol, should be easier to traverse. Unfortunately, conversion of anisole to phenol does not surpass 67% even when run at slightly elevated temperatures (50 °C in CH_2Cl_2). This finding suggests that only cycles 1 and 2 are accessible, which contradicts our computational prediction.

Table 3. Gas chromatography analysis of product mixture run in CH_2Cl_2 after hydrolysis.

BBr_3 [equiv.]	Temp. [°C]	Anisole [%]	Phenol [%]
1.0	30	0.20	99.8
0.66	30	4.23	95.8
0.33	30	43.1	56.9
0.33	50	33.2	67.8

Is this simply a failure of density functional theory or is there additional chemistry missing from our computational model? In recent years there has been an increasing interest in the dynamic covalent chemistry of boron–oxygen bonds.^[13] One relevant pathway to consider is the output of

each individual cycle [PhOBBr_2 , $(\text{PhO})_2\text{BBr}$, and $\text{B}(\text{OPh})_3$] equilibrating with BBr_3 . Roy reported that MeOBBr_2 and $(\text{MeO})_2\text{BBr}$ can be synthesized by equilibrating different ratios of $\text{BBr}_3/\text{B}(\text{OMe})_3$.^[14] With the well-established dynamic nature of boron–oxygen bonding, we envisioned similar equilibria may impact our proposed reaction mechanism. The first possible equilibrium would be between BBr_3 and the product of cycle 1, PhOBBr_2 (Scheme 5, top). Even if this reaction occurs it does not complicate our mechanism because: (i) the reaction is thermoneutral and (ii) it does not generate new species. The second equilibrium between BBr_3 and the product of cycle 2, $(\text{PhO})_2\text{BBr}$ (Scheme 5, bottom), potentially complicates our mechanism by generating two equivalents of PhOBBr_2 . Our computational model predicts this reaction to be exergonic by 4.8 kcal/mol, thus any $(\text{PhO})_2\text{BBr}$ generated in the presence of BBr_3 will disproportionate to form PhOBBr_2 . With this in mind, an additional energetic penalty should be factored into the barrier height of cycle 3 because two equivalents of PhOBBr_2 must disproportionate to form BBr_3 and $(\text{PhO})_2\text{BBr}$. In turn, the barrier rises from 22.2 to 27.0 kcal/mol and makes cycle 3 the least kinetically accessible. Therefore, additional thermal energy may make full conversion through cycle 3 possible. We tested this hypothesis by switching solvents from CH_2Cl_2 to *o*-dichlorobenzene, which allowed for this reaction to be run at 100 °C. Under this set of reaction conditions using 0.33:1 we observed full conversion of anisole to phenol as measured by GC. This result not only supports our mechanism, but also confirms the results disclosed in the original report of full conversion with a 0.33:1 ratio at elevated temperatures.^[5]



Scheme 5. Possible dynamic covalent equilibria between BBr_3 and products of cycles 1 and 2.

One may ask: why can anisole not be demethylated through cycle 2 once PhOBBr_2 is generated (Scheme 5, bottom)? A critical feature of our mechanism is that catalytic BBr_3 is required for each of the cycles. In the 1:1 and 0.66:1 experiments the possibility of demethylating through cycle 2 multiple times is viable because excess BBr_3 remains after the disproportionation equilibrium has consumed the $(\text{PhO})_2\text{BBr}$ generated from cycle 2. However, the 0.33:1 reaction would consume all of the BBr_3 by the time the disproportionation reaches equilibrium, thereby disabling cycle 2.

Conclusions

While the cleavage of highly branched aliphatic ethers most likely proceeds through the unimolecular process proposed by Sousa and Silva, we believe our alternative bimolecular mechanism for demethylation in aryl methyl ethers, which differs from the classical mechanism of bromide attack in terms of the bromide source, is operative. Our

computational predictions indicate that demethylation of anisole proceeds through a three-cycle mechanism that is underpinned by previously reported and new experimental findings. The results of this study show that sub-stoichiometric amounts of BBr_3 can be used in place of one full equivalent and may enable the use of BBr_3 in total synthesis when multiple moieties are susceptible to attack. However, there are a number of lingering questions that require further investigation. (i) Does the new mechanism for ether cleavage extend to alkyl methyl ethers and, if so, are all three cycles operable? (ii) Is the multi-cycle mechanism proposed for aryl methyl ether cleavage general for BX_3 reagents ($\text{X} = \text{Br}, \text{Cl}$)? The reaction of BCl_3 with ethers and alcohols suggests that two- and three-cycle mechanisms are accessible under the reported conditions.^[15,16] (iii) Can a multi-cycle extension of Sousa and Silva's unimolecular mechanism be viable for branched ethers? Work is ongoing in our laboratories to answer these and related questions about BX_3 reactivity.

Computational Methods

Geometry optimizations were performed in the Gaussian09 program (G09.D01)^[17] at the B3LYP/6-31G(d) level of theory.^[18–22] The effect of implicit solvation was included during geometry optimization using the SMD model for dichloromethane.^[23,24] Stationary points on the potential energy surface were characterized as minima or first-order saddle points (transition states) by evaluating harmonic frequencies at the optimized geometries.^[25] From these frequency calculations performed with a double-zeta basis set, electronic energies [$E(\text{SCF})_{\text{DZ}}$] and Gibbs free energies [$G(\text{sol})_{\text{DZ}}$] based on standard thermodynamic approximations were tabulated.^[26] Single-point energy refinements with the 6-311+G(d,p) basis set, implicit solvation, and Grimme's empirical dispersion corrections (with Becke–Johnson damping)^[27,28] allowed for improved triple-zeta electronic energies [$E(\text{SCF})_{\text{TZ}}$]. For select species, single point energy refinements with aug-cc-pVTZ produced similar results (see Supporting Information) and the more efficient 6-311+G(d,p) basis was therefore employed. Approximate triple-zeta free energies were obtained by $G(\text{sol})_{\text{TZ}} \approx G(\text{sol})_{\text{DZ}} - E(\text{SCF})_{\text{DZ}} + E(\text{SCF})_{\text{TZ}}$. Visualizations were made with GaussView 5.0.9^[29] and CylView.^[30]

Experimental Methods

General: Anisole was purchased from Acros Organics and used without additional purification. BBr_3 was purchased from Aldrich as a 1 M solution in CH_2Cl_2 and used only for two weeks after opening to ensure purity. Dry CH_2Cl_2 was obtained from a PureSolv solvent purification system that passes solvent over two columns of neutral alumina. ^1H NMR were taken on a 300 MHz JEOL OXFORD spectrometer in CDCl_3 that was stored over 4 Å molecular sieves. Spectra were referenced to the deuterated solvent. Gas chromatography was performed on Thermo Scientific GC-Focus Series instrument which was fitted with a Supelco MDN-5 column. The injector (Thermo Scientific AL3000) temperature was set

at 185 °C. An initial column temperature of 80 °C was held for four minutes and ramped at a rate of 10 °C per minute until a final temperature of 280 °C was reached. The detector temperature was set to 280 °C.

Representative Experimental Procedure: To a dry 5.00 mL thick walled vial equipped with a stir bar and septum was added anisole followed by the addition of dry dichloromethane (1 mL per 1 mL of BBr_3 solution). The vial was allowed to purge under nitrogen for approximately 5 min, after which BBr_3 was added slowly through the septum with stirring. The reaction was left to stir overnight before the contents were poured into ca. 1 mL of deionized H_2O . The organic layer was separated and an aliquot was analyzed by GC using dichloromethane as the solvent.

Acknowledgments

T. M. K. acknowledges a Ott-Stiner fellowship and both T. M. K. and H. A. C. thank the Weldon fund for financial support. A. L. K. and R. L. L. recognize Grand Valley State University start-up funds (GVSU-OURS, GVSU-CSCE, GVSU-CLAS) and the National Science Foundation (NSF) (computational support through CHE-1039925 to the Midwest Undergraduate Computational Chemistry Consortium) for financial support. Special thanks go to Jacob Dillon and Donovan Anderson for help translating references 15 and 16.

- [1] J. R. Lawson, E. R. Clark, I. A. Cade, S. A. Solomon, M. J. Ingleson, *Angew. Chem. Int. Ed.* **2013**, *52*, 7518.
- [2] V. Bagutski, A. Del Grosso, J. A. Carrillo, I. A. Cade, M. D. Helm, J. R. Lawson, P. J. Singleton, S. A. Solomon, T. Marcelli, M. J. Ingleson, *J. Am. Chem. Soc.* **2013**, *135*, 474.
- [3] Y. Qin, G. Cheng, A. Sundararaman, F. Jäkle, *J. Am. Chem. Soc.* **2002**, *124*, 12672.
- [4] P. Gao, P. S. Portoghese, *J. Org. Chem.* **1996**, *61*, 2466.
- [5] E. Dillon, O. F. N. Dame, *J. Am. Chem. Soc.* **1942**, *64*, 1128.
- [6] S. Punna, S. Meunier, M. G. Finn, *Org. Lett.* **2004**, *6*, 2777.
- [7] J. F. W. McOmie, M. L. Watts, D. E. West, *Tetrahedron* **1968**, *24*, 2289.
- [8] E. H. Vickery, L. F. Pahler, E. J. Eisenbraun, *J. Org. Chem.* **1979**, *44*, 4444.
- [9] E. Paliakov, L. Strekowski, *Tetrahedron Lett.* **2004**, *45*, 4093.
- [10] C. Pasquini, A. Coniglio, M. Bassetti, *Tetrahedron Lett.* **2012**, *53*, 6191.
- [11] C. Sousa, P. J. Silva, *Eur. J. Org. Chem.* **2013**, 5195.
- [12] L. Watson, O. Eisenstein, *J. Chem. Educ.* **2002**, *79*, 1269.
- [13] A. L. Korich, P. M. Iovine, *Dalton Trans.* **2010**, *39*, 1423.
- [14] C. D. Roy, *Aust. J. Chem.* **2006**, *59*, 657.
- [15] E. Wiberg, W. Sütterlin, *Z. Anorg. Allg. Chem.* **1931**, *202*, 22.
- [16] E. Wiberg, W. Sütterlin, *Z. Anorg. Allg. Chem.* **1931**, *202*, 31.
- [17] M. J. Frisch, G. W. Trucks, H. B. Schlegel, G. E. Scuseria, M. A. Robb, J. R. Cheeseman, G. Scalmani, V. Barone, B. Mennucci, G. A. Petersson, H. Nakatsuji, M. Caricato, X. Li, H. P. Hratchian, A. F. Izmaylov, J. Bloino, G. Zheng, J. L. Sonnenberg, M. Hada, M. Ehara, K. Toyota, R. Fukuda, J. Hasegawa, M. Ishida, T. Nakajima, Y. Honda, O. Kitao, H. Nakai, T. Vreven, J. A. Montgomery Jr., J. E. Peralta, F. Ogliaro, M. Bearpark, J. J. Heyd, E. Brothers, K. N. Kudin, V. N. Staroverov, R. Kobayashi, J. Normand, K. Raghavachari, A. Rendell, J. C. Burant, S. S. Iyengar, J. Tomasi, M. Cossi, N. Rega, J. M. Millam, M. Klene, J. E. Knox, J. B. Cross, V. Bakken, C. Adamo, J. Jaramillo, R. Gomperts, R. E. Stratmann, O. Yazyev, A. J. Austin, R. Cammi, C. Pomelli, J. W. Ochterski, R. L. Martin, K. Morokuma, V. G. Zakrzewski, G. A. Voth, P. Salvador, J. J. Dannenberg, S. Dapprich, A. D. Daniels, Ö. Farkas, J. B. Foresman, J. V. Ortiz, J. Cioslowski, D. J. Fox, *Gaussian 09*, revision D.01, Gaussian, Inc., Wallingford, CT, USA, **2009**.

- [18] S. H. Vosko, L. Wilk, M. Nusair, *Can. J. Phys.* **1980**, *58*, 1200.
- [19] A. D. Becke, *Phys. Rev. A* **1988**, *38*, 3098.
- [20] C. Lee, W. Yang, R. G. Parr, *Phys. Rev. B* **1988**, *37*, 785.
- [21] A. D. Becke, *J. Chem. Phys.* **1993**, *98*, 5648.
- [22] P. Stephens, F. J. Devlin, C. F. Chabalowski, M. J. Frisch, *J. Phys. Chem.* **1994**, *98*, 11623.
- [23] V. Marenich, C. J. Cramer, D. G. Truhlar, *J. Phys. Chem. B* **2009**, *113*, 6378.
- [24] R. F. Ribeiro, A. V. Marenich, C. J. Cramer, D. G. Truhlar, *J. Phys. Chem. B* **2011**, *115*, 14556.
- [25] H. B. Schlegel, *WIREs Comput. Mol. Sci.* **2011**, *1*, 790.
- [26] C. J. Cramer, *Essentials of Computational Chemistry: Theories and Models*, 2nd ed., Wiley, **2004**.
- [27] Ehrlich, S. J. Moellmann, S. Grimme, *Acc. Chem. Res.* **2013**, *46*, 916.
- [28] S. Gimme, S. Ehrlich, L. Goerigk, *J. Comput. Chem.* **2011**, *32*, 1456.
- [29] R. Dennington, T. Keith, J. Millam, C. Y. *GaussView*, **2009**.
- [30] C. Y. Legault, *CylView*, **2009**.

Received: August 11, 2015
Published Online: October 23, 2015