Synthesis and structure of a magnesium–amidoborane complex and its role in catalytic formation of a new bis-aminoborane ligand[†]

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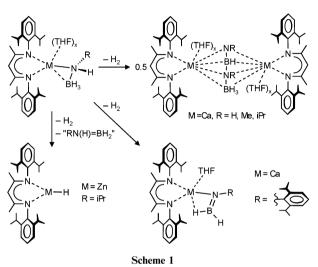
A synthetic route to a magnesium-amidoborane complex and its role in the catalytic conversion of a substituted ammonia-borane RNH₂BH₃ into HB(NHR)₂ is discussed.

The recently introduced early main group metal amidoborane compounds $LiNH_2BH_3$, $NaNH_2BH_3$ and $Ca(NH_2BH_3)_2$ show high potential as future hydrogen storage materials.^{1,2} The elimination of molecular hydrogen in these compounds displays several advantages over that in NH_3BH_3 . Among these are a significantly lower hydrogen release temperature and near thermoneutrality which might facilitate development of a reversible hydrogen up-take process.

We recently initiated studies on heteroleptic calcium amidoborane complexes containing a large DIPP-nacnac ligand (Scheme 1, DIPP-nacnac = $CH\{(CMe)(2,6-iPr_2C_6H_3N)\}_2$).³ Solubilization of the calcium amidoborane complex allowed investigation of the hydrogen release process under homogeneous conditions and crystallization of the decomposition products. In addition, we investigated the effect of substituents on nitrogen.⁴ A large substituent inhibits dimerization during hydrogen release and we could isolate a complex containing a borylamide anion $RN=BH_2^-$ (Scheme 1).

Attempted isolation of a zinc amidoborane complex, however, gave quantitative conversion into a zinc hydride species (Scheme 1).⁵ In order to understand the effect of the metal on the stability and decomposition of metal amidoborane complexes in general, we here report our preliminary investigations on the synthesis of a magnesium amidoborane complex with a large substituent on nitrogen.

Addition of one equivalent of (DIPP)NH₂BH₃ to a solution of (DIPP-nacnac)MgN(SiMe₃)₂ in benzene at 20 °C immediately gave vigorous evolution of H₂ gas. As all (DIPP)NH₂BH₃ had reacted but the starting material (DIPP-nacnac)MgN(SiMe₃)₂ was hardly consumed, the formation of H₂ gas was not due to the instability of the expected magnesium amidoborane complex but rather to a catalytic decomposition of (DIPP)NH₂BH₃. This decomposition reaction is clean and the only product was fully characterized as HB[NH(DIPP)]₂ by NMR and X-ray



diffraction (Fig. 1).[‡] The N and B atoms display close to planar conformations and could be regarded as sp^2 -hybridized. Their coplanar arrangement allows for delocalization of the *p*-lone pairs on both N atoms into the empty p-orbital on B and explains the short B–N bond distances. This species represents a hitherto unknown bis(amino)borane, a class of compounds which has received considerable attention: double deprotonation gives access to dianionic bora-amidinate (bam) ligands HB(NR)₂²⁻ which are isolobal to the mono-anionic amidinate HC(NR)₂⁻⁶

Scheme 2 shows the proposed mechanism of formation. In the first step the expected magnesium amidoborane complex is formed. This product could react with the acidic ammonia– borane (DIPP)NH₂BH₃ and eliminate H₂. A 1,3-H shift from one B to the other affords HB[NH(DIPP)]₂ and (DIPP-nacnac)MgBH₄. The latter borate complex could react with (DIPP)NH₂BH₃ to give (DIPP-nacnac)MgNH(DIPP)BH₃ and BH₃. Alternatively, β -hydride elimination could afford (DIPP-nacnac)MgH, HB[NH(DIPP)]₂ and BH₃. The highly reactive magnesium hydride complex could either deprotonate (DIPP)NH₂BH₃ or react with BH₃ to give (DIPP-nacnac)-MgBH₄.

Although speculative, the proposed mechanism is underscored by several observations.

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⁽i) Cooling of the reaction mixture gave a small crop of colourless crystals which could be identified as (DIPP-nacnac)-MgBH₄ by NMR and X-ray diffraction (*vide infra*).

⁽ii) Addition of catalytic amounts of (DIPP-nacnac)MgBH₄ to a solution of (DIPP)NH₂BH₃ likewise resulted in quantitative formation of HB[NH(DIPP)]₂. This makes it a likely intermediate in the catalytic cycle.

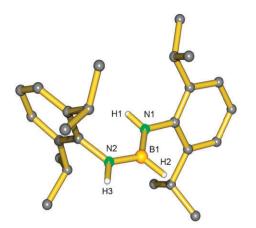
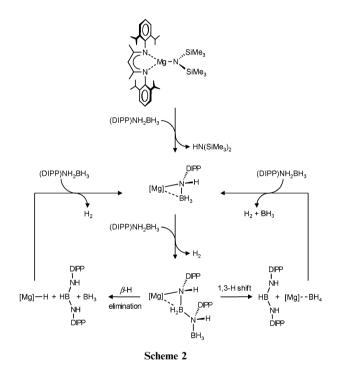
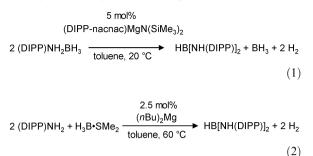


Fig. 1 The molecular structure of HB[NH(DIPP)]₂; most hydrogen atoms have been omitted for clarity. Selected bond distances (Å) and angles (°): B1–N1 1.415(2), B1–N2 1.414(2), N1–B1–N2 121.9(1).



(iii) The by-product "BH₃" could be identified in the ¹H NMR spectrum as B_2H_6 (the product HB[NH(DIPP)]₂ does not react with diborane).

Although the Mg-mediated conversion of (DIPP)NH₂BH₃ into HB[NH(DIPP)]₂, BH₃ and H₂ (eqn (1)) is a quantitative and clean reaction, we developed a more atom-efficient route. It was found that (DIPP)NH₂ and H₃B·SMe₂ in a 2/1 ratio can be converted quantitatively to HB[NH(DIPP)]₂ and H₂ (eqn (2)). Commercially available *n*-Bu₂Mg works equally well as a catalyst, however, a higher temperature (60 °C) is essential to prevent precipitation of the salt Mg(BH₄)₂. This simple route allows an economically large scale preparation of this bis(amino)borane. Its twofold deprotonated form, the bora-amidinate HB[N(DIPP)]₂²⁻, would be the first bam ligand with a H-substituent in the backbone and is isolobal to a versatile sterically congested amidinate: HC[N(DIPP)]₂⁻. The scope of this dianionic bam ligand in organometallic chemistry is currently under investigation.



Crystals of (DIPP-nacnac)MgBH₄, isolated from Mgmediated catalytic formation of HB[NH(DIPP)]₂, were analyzed by X-ray diffraction.[‡] A solvent-free dinuclear complex with terminal chelating DIPP-nacnac ligands and bridging BH₄⁻ ions is observed (Fig. 2; hydrogen atoms in BH4⁻ have been located and refined). The BH4⁻ ions coordinate to both Mg²⁺ centers in a bidentate fashion in which one of the hydride atoms bridges both metals. This bridging bonding mode has never been observed in magnesium tetrahydridoborate complexes but is known in lithium chemistry (*e.g.* in $[(TMEDA)LiBH_4]_2)$.⁷ It differs from $Mg \cdots D_2BD_2 \cdots Mg$ bridging in $[Mg(BD_4)_2]_{\infty}$.⁸ Heavier alkaline– earth metal borate complexes typically show (η^3) HBH₃···M²⁺ contacts (M = Ca, Sr, Ba).^{9,10} The bridging nature of the BH_4^- ions in the current structure results in Mg···B distances (average 2.538(2) Å) that are much longer than that of 2.241(2) Å in (DIPP-nacnac)MgBH₄·OEt₂.¹¹ In the latter structure BH_4^- is bound to Mg^{2+} in η^3 -fashion.

The attempted synthesis of the amidoborane complex (DIPP-nacnac)MgNH(DIPP)BH₃ resulted in catalytic conversion of (DIPP)NH₂BH₃ into HB[NH(DIPP)]₂. This is likely due to the low reactivity of the Mg-N(SiMe₃)₂ functionality: slow formation of the magnesium amidoborane complex allows for its subsequent reaction with the substrate (DIPP)NH₂BH₃. The recently published magnesium hydride complex [(DIPP-nacnac)MgH]₂,¹² however, is much more reactive and deprotonates (DIPP)NH₂BH₃ in toluene immediately already at 0 °C. Under these conditions no side reactions are observed and (DIPP-nacnac)MgNH(DIPP)BH₃ could be isolated in 59% crystalline yield. Although it crystallizes in the form of small cubes, a structure could be determined (Fig. 3).[†] The monomeric solvent-free complex shows side-on coordination of the amidoborane anion to Mg^{2+} . This magnesium amidoborane complex is quite stable towards elimination of hydrogen. Decomposition in either benzene or THF solution starts at temperatures well over 100 °C. The stability of this magnesium amidoborane complex underscores the mechanism proposed for catalytic formation of HB[NH(DIPP)]₂ in Scheme 2 and excludes a mechanism with an intermediate containing the boryl amide $(DIPP)N = BH_2^{-}$. Further details on the decomposition of (DIPP-nacnac)MgN(DIPP)HBH₃ as well as that of other magnesium amidoborane complexes will be described in a forthcoming paper.¹³

In summary, reaction of the heteroleptic magnesium complex $(DIPP-nacnac)MgN(SiMe_3)_2$ with $(DIPP)NH_2BH_3$

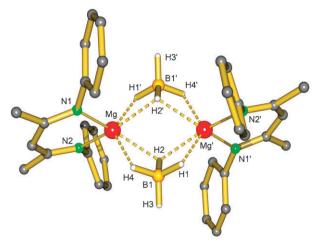


Fig. 2 The molecular structure of $[(DIPP-nacnac)MgBH_{4}]_{2}$ (the *i*Pr substituents and most hydrogen atoms have been omitted for clarity). Selected bond distances (Å): Mg–N1 2.035(1), Mg–N2 2.041(1), Mg–H1' 1.95(2), Mg–H2' 2.20(2), Mg–H2 2.34(2), Mg–H4 1.96(2), Mg···B1 2.541(2), Mg···B1' 2.535(2).

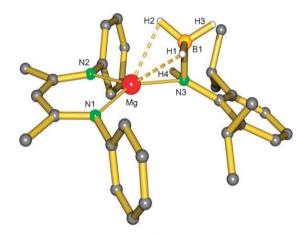


Fig. 3 The molecular structure of (DIPP-nacnac)MgNH(DIPP)BH₃ (the *i*Pr substituents on the DIPP-nacnac ligand and most hydrogen atoms have been omitted for clarity). Selected bond distances (Å): Mg–N1 2.004(5), Mg–N2 2.024(5), Mg–N3 2.083(4), Mg···B1 2.291(7).

resulted in a catalytic decomposition to the bis(amino)borane HB[NH(DIPP)]₂. As this compound can be doubly deprotonated to a potentially useful bam ligand, a convenient atom-efficient route was developed. Magnesium amidoborane complexes are accessible by use of a much more reactive magnesium hydride reagent.

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Notes and references

‡ Crystallographic data for HB[NH(DIPP)]₂: C₂₄H₃₇BN₂, M = 364.37, monoclinic, space group $P2_1$, a = 10.2052(9), b = 10.3128(9), c = 11.5048(11) Å, $\beta = 108.727(5)^\circ$, U = 1146.71(18) Å³, Z = 2, $D_c = 1.055$ g cm⁻³, F(000) = 400, μ (Mo-K $\alpha) = 0.060$ mm⁻¹, $1.9 \le \theta \le 30.1^\circ$, 14398 reflections measured, 6317 unique ($R_{int} = 0.054$) were used in all calculations, $R_1 = 0.0474$ [$I > 2\sigma(I)$, 5329 reflections] and $wR_2 = 0.1133$ (all data); max/min. residual electron density: 0.17/-0.21.

Crystallographic data for [(DIPP)nacnacMgBH₄]₂: $C_{58}H_{90}B_2Mg_2N_4$, M = 913.58, monoclinic, space group $P2_1/n$, a = 14.3266(11), b = 13.9025(8), c = 15.1584(13) Å, $\beta = 106.287(6)^\circ$, U = 2898.0(4) Å³, Z = 2, $D_c = 1.047$ g cm⁻³, F(000) = 1000, μ (Mo-K α) = 0.079 mm⁻¹, $3.5 \le \theta \le 25.6^\circ$, 18.615 reflections measured, 5409 unique ($R_{int} = 0.049$) were used in all calculations, $R_1 = 0.0460$ [$I > 2\sigma(I)$, 43.69 reflections] and $wR_2 = 0.1298$ (all data); max/min. residual electron density: 0.46/-0.20.

Crystallographic data for (DIPP-nacnac)MgNH(DIPP)BH₃: C₄₁H₆₂BMgN₃, M = 632.06, monoclinic, space group $P_{21/c}$, a = 18.295(4), b = 11.8119(14), c = 18.846(3) Å, $\beta = 103.004(15)^\circ$, U = 3968.2(12) Å³, Z = 4, $D_c = 1.058$ g cm⁻³, F(000) = 1384, μ (Mo-K α) = 0.075 mm⁻¹, $3.5 \le \theta \le 25.0^\circ$, 17455 reflections measured, 6903 unique ($R_{int} = 0.109$) were used in all calculations, $R_1 = 0.0924$ [$I > 2\sigma(I)$, 2340 reflections] and w $R_2 = 0.2375$ (all data); max/min. residual electron density: 0.34/-0.30.

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