

## X-RAY CRYSTAL STRUCTURE OF A CHIRAL OXAZABOROLIDINE CATALYST FOR ENANTIOSELECTIVE CARBONYL REDUCTION

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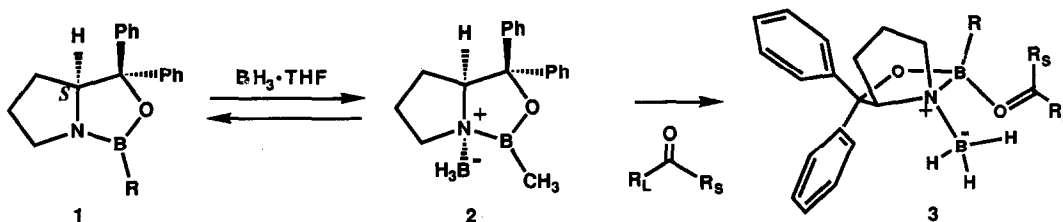
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**Summary:** The three dimensional structure of the chiral complex 2, which is an effective reagent for the enantioselective reduction of ketones, has been determined by X-ray crystallography to be as shown in Figure 1.

Chiral oxazaborolidines 1, with R = H, CH<sub>3</sub> or alkyl, (or enantiomers) catalyze the reduction of ketones by borane or catecholborane as stoichiometric reductant to produce secondary alcohols with high enantiofacial selectivity.<sup>1</sup> This process is extremely useful in synthesis because only 5-10 mole % of 1 generally suffices to ensure >20/1 enantioselectivity and because the catalyst precursor (*S*)- or (*R*)-2-[diphenylhydroxymethyl]-pyrrolidine can be recovered efficiently for reuse. The reductions using borane and 1, R = CH<sub>3</sub> for example, proceed by way of complexation of BH<sub>3</sub> and 1, R = CH<sub>3</sub>, to give adduct 2 which has been characterized previously by <sup>1</sup>H, <sup>13</sup>C and <sup>11</sup>B NMR spectroscopy.<sup>1</sup> Reaction of 2 with the ketonic substrate is considered to occur by way of transition state assembly 3, a model which correctly predicts the absolute configuration of the product for a very large number of examples.<sup>1</sup> This oxazaborolidine-catalyzed reduction of ketones to

epoxides,<sup>1c</sup> PAF antagonists,<sup>2</sup> ginkgolide B,<sup>3</sup> bilobalide,<sup>4</sup> forskolin,<sup>5</sup> fluoxetine,<sup>6</sup> 1-deuterated primary alcohols,<sup>7</sup> isoproteranol,<sup>8</sup>  $\alpha$ -hydroxy acids,<sup>1e</sup> denopamine,<sup>9</sup> 2-hydroxymethylpyrrolidines (precursors for their own catalytic enantioselective synthesis),<sup>10</sup> and  $\alpha$ -amino acids.<sup>11</sup> The structure of the reactive complex 2 has now been verified by a single-crystal X-ray diffraction study which revealed interesting three-dimensional detail and which is described herein.

Reaction of (*S*)-2-[diphenylhydroxymethyl]-pyrrolidine with methylboronic acid was carried out as previously described<sup>1b,1c</sup> to give 1, R = CH<sub>3</sub>, which was converted to the borane adduct 2<sup>12</sup> by treatment in toluene with gaseous B<sub>2</sub>H<sub>6</sub><sup>13</sup> and filtration of the resulting crystalline precipitate. Slow recrystallization of these colorless crystals from a CH<sub>2</sub>Cl<sub>2</sub>-hexane bilayer at -20 °C gave clear, colorless cubes of 2.<sup>14</sup> One such crystal (0.36 x 0.32 x 0.4 mm), orthorhombic, space group P<sub>2</sub><sub>1</sub>2<sub>1</sub>2<sub>1</sub>, 4 molecules per unit cell, was subjected to X-ray diffraction analysis and yielded the structure shown in Figure 1,<sup>15</sup> which is in complete agreement with previous



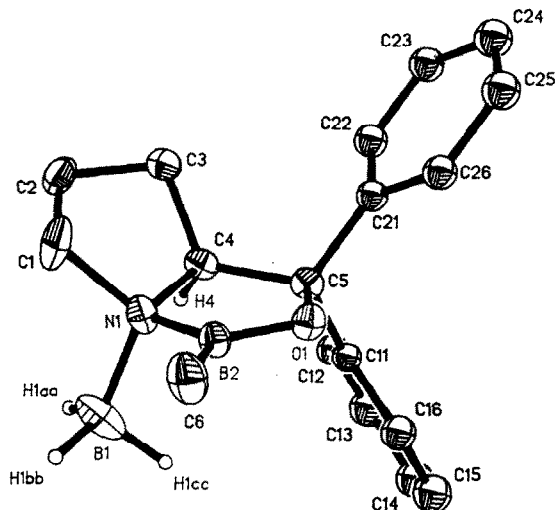


Figure 1. X-Ray structure of borane complex 2.

proposals.<sup>1</sup> The N<sub>1</sub>B<sub>1</sub> bond distance (1.62 Å) is close to that reported for ammonia borane (1.604 Å) and N<sub>1</sub>B<sub>2</sub> is considerably shorter (1.486 Å). The O<sub>1</sub>B<sub>2</sub> distance (1.335) is definitely shorter than the OB distance in the phenylboronate ester of catechol (1.394) indicating substantial O<sub>1</sub>B<sub>2</sub> π-bonding in 2. The results of the present study provide additional evidence for transition state assembly 3 for the catalytic CBS reduction.<sup>1,16</sup>

#### References and Notes

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- All operations were carried out in an argon atmosphere because of sensitivity to H<sub>2</sub>O and O<sub>2</sub>.
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- <sup>1</sup>H NMR (400 MHz, CD<sub>2</sub>Cl<sub>2</sub>) δ 7.56 (m, 2H), 7.3 - 7.1 (m, 8H), 4.58 (m, 1H), 3.14 (m, 1H), 1.89 (m, 2H), 1.58 (m, 1H), 1.6 - 1.8 (br.), 1.25 (m, 1H), 0.69 (s, 3H); <sup>11</sup>B NMR (96 MHz, CD<sub>2</sub>Cl<sub>2</sub>) δ 40.47, -12.87 (quartet, J<sub>BH</sub> = 98.2 Hz).
- A total of 3364 reflections were collected at -70 °C; unit cell dimensions *a* = 9.010(4) Å, *b* = 12.987(3) Å, *c* = 14.386(3) Å, *Z* = 4. Least squares refinement of the data using 1987 reflections converged on the structure shown in Fig. 1 with R = 0.0846 and a goodness of fit 0.91.
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