



Self-assembly of a library of polyborate chiral anions for asymmetric catalytic quinoline reduction



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ABSTRACT

The 'template' polyborate BOROX catalysts are shown to mediate the asymmetric transfer hydrogenation of 2-quinolines. The rapid and simple generation of a large family of BOROX catalysts with significantly altered asymmetric pockets is described. A transition state model that explains the enantioselectivity is proposed.

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We have identified, by crystallographic and NMR studies, the active catalyst in our catalytic asymmetric aziridination reaction (AZ reaction).¹ The catalyst is a polyborate chiral anion that contains a boroxine ring in which one of the borons is four coordinate producing a boroxinate core (Fig. 1).² These boroxinate or BOROX catalysts, are *self-assembled* by a substrate of the catalyst, which is an imine or amine in the case of the aziridination reaction (Scheme 1).³ The base induced assembly of the BOROX catalyst can be achieved from: (1) the ligand, water, and triphenylborate,^{3–5} (2) the ligand and triphenoxyboroxine^{1b} and (3) the ligand, borane dimethylsulfide complex, phenol, and water.^{4,6}

Theoretical studies of the AZ reaction have shed light on the mode of catalysis of this chiral polyborate Brønsted acid catalyst.^{1c,d} In addition to rendering proton catalysis enantioselective, it was proposed that the VAPOL/VANOL–BOROX counterion also plays the role of orchestrating the orientation of the substrates at the key transition state via multiple H-bonding interactions with oxygens O1, O2, and O3 in the boroxinate core.^{1c,1d} In addition to aziridinations, the BOROX catalyst is also involved in an asymmetric catalytic aza-Cope rearrangement in the aminoallylation of aldehydes,⁷ in a catalytic asymmetric 3-component Ugi reaction⁸

and presumably in a heteroatom Diels–Alder reaction as well.⁹ Since the BOROX catalyst is self-assembled in situ when the

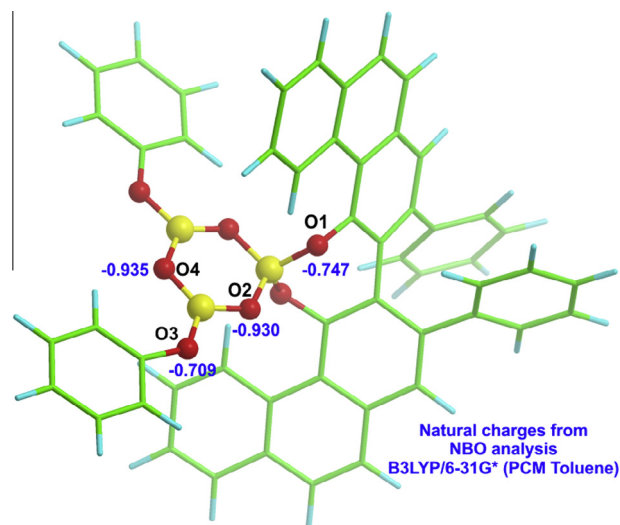
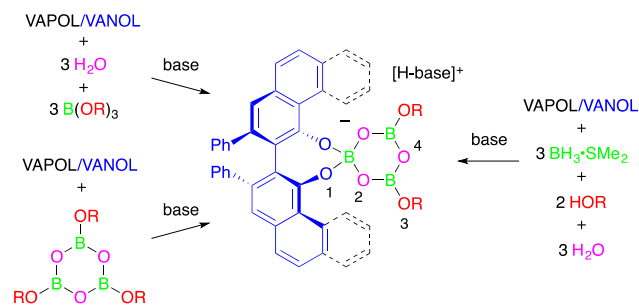


Figure 1. (S)-VAPOL–BOROX anion.

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Scheme 1. Self-assembly of boroxinate (BOROX) catalyst.

substrate is added, it thus becomes possible to generate a vast library of the BOROX catalysts on the spot by the incorporation of different alkoxy groups in the boroxinate core.¹⁰ Although a few borate esters are commercially available, a far greater avenue for maximizing diversity is to directly assemble the catalyst in the presence of a commercially available alcohol or phenol.⁸ Herein we report our efforts toward utilizing the diversity of this ‘template’ BOROX catalyst in the asymmetric transfer hydrogenation (ATH) of quinolines with Hantzsch’s ester.^{11,12} In addition, we propose a transition state model that accounts for the experimental observations.

We expected the quinolines **1a** and **1b** chosen for the present study to smoothly self-assemble the polyborate BOROX catalysts, similar to the imines and other bases in our previous studies.^{1,3} The ¹¹B and ¹H NMR spectra of the VAPOL–BOROX complex with **1a** were distinctive (Fig. 2). The two 3-coordinate borons appear as a broad peak at 15.97 ppm, and the 4-coordinate boron as a sharp peak at 5.76 ppm, with an integration of 2:1, respectively

(not shown). The most distinctive peak in the ¹H NMR spectrum of the VAPOL–BOROX-**1a** complex is the bay region doublet at 10.49 ppm (H_b in VAPOL). These observations are in accord with those made previously for the BOROX catalysts in our aziridination studies.^{1,3}

Initial attempts toward developing the catalytic asymmetric transfer hydrogenation reaction utilized the 2-pentylquinoline substrate **1a** and the Hantzsch ester **2a** as the hydride transfer reagent (Table 1). The 2-pentylquinoline was prepared by a titanium mediated multi-component coupling of an aniline, an alkyne, and a isonitrile.¹³ While the VANOL–BOROX catalyst **4** afforded the tetra-hydroquinoline product **3a** with a 40:60 er (entry 1), the corresponding BOROX catalyst **5** prepared from the VAPOL ligand provided **3a** in quantitative yield and with a promising er of 86:14 (entry 2). To optimize this protocol, different Hantzsch ester derivatives **2a–d** were subsequently screened, albeit without any big improvements in the er (entries 3–5). Additives were found to be deleterious in this reaction, and an extensive screen of 13 solvents established benzene as the solvent of choice.¹⁴ In addition, a number of substituted VANOL and VAPOL ligands were evaluated but none were superior to VAPOL.¹⁴ High yields and similar trends for the enantiomeric ratios were also observed for the ATH of 2-phenylquinoline **1b** (entries 6 and 7). These initial results were promising and we decided to pursue them for our catalyst optimization studies.

The BOROX catalyst in most of our aziridination studies was generated from B(OPh)₃ as the boron source,^{1,3,4} which was also the method in the preliminary screens in Table 1. It was found in the present study that replacing B(OPh)₃ with the combination of BH₃·SMe₂ and phenol provided identical results (compare result with phenol in Scheme 2 with that in entry 2 of Table 1). That the BOROX polyborate catalyst could be successfully generated

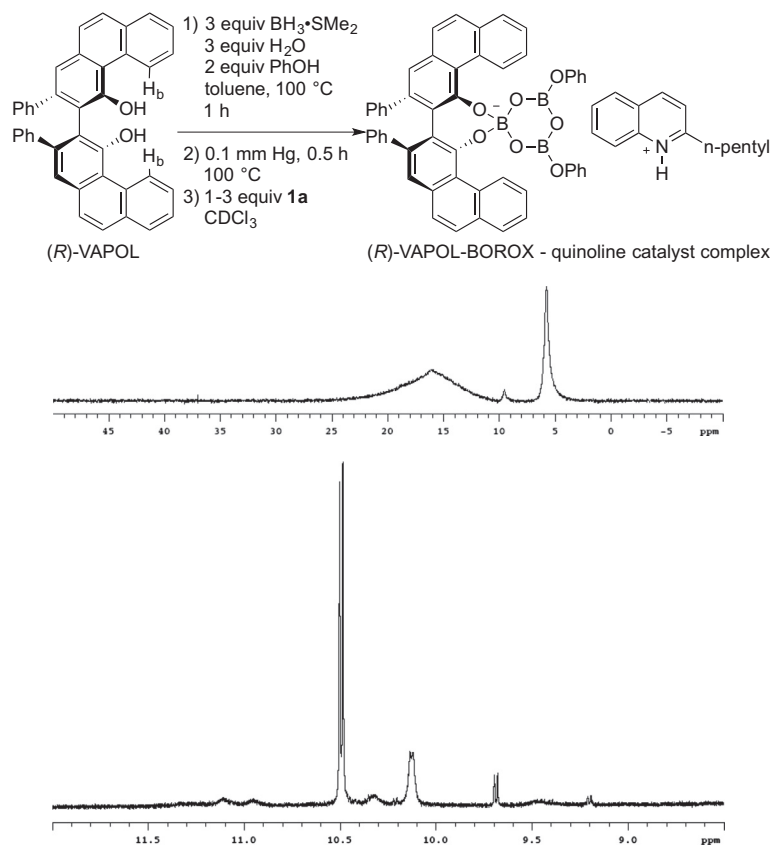
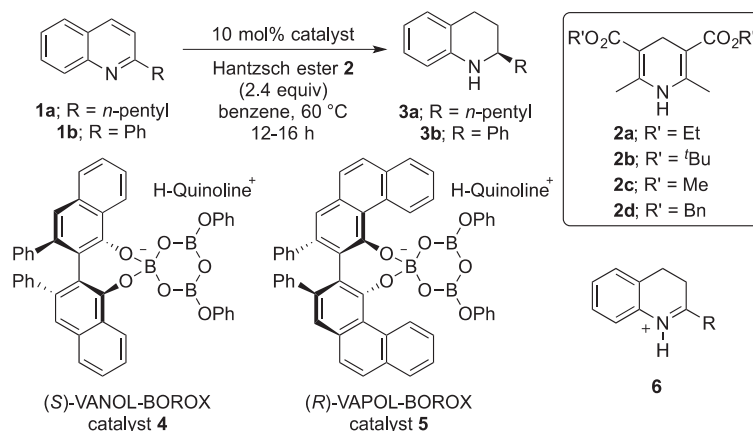


Figure 2. Preparation of (R)-VAPOL-BOROX-**1a** complex and its ¹¹B NMR (CDCl₃, 160 MHz) and ¹H NMR (CDCl₃, 500 MHz).

Table 1
Quinoline reduction with the VAPOL/VANOL-BOROX catalysts^a



Entry	1	2	Catalyst	3	Yield-3 ^b (%)	er-3 ^c (%)
1	1a	2a	4a	3a	88	40:60
2 ^d	1a	2a	5a	3a	>99	86:14
3 ^e	1a	2b	5a	3a	ND ^f	72:28
4	1a	2c	5a	3a	ND ^f	88:12
5	1a	2d	5a	3a	ND ^f	88:12
6	1b	2a	4b	3b	>99	32:68
7 ^g	1b	2a	5b	3b	>99	84:16

^a Unless otherwise specified, all reactions were run with 0.05 mmol of **1a** and 0.1 mmol of **1b**. The catalyst was prepared by heating 1 equiv of ligand, 4 equiv of B(OPh)₃, and 1 equiv of water in toluene at 80 °C for 1 h, followed by removing all volatiles under high vacuum (0.1 mm Hg) at 80 °C for 0.5 h. Conversion = 100% (TLC). Reaction with (S)-VANOL-BOROX catalyst **4** gives ent-**3**.

^b Isolated yield after column chromatography on regular silica gel.

^c Determined by HPLC on isolated **3**.

^d Average of 3 runs.

^e Average of 4 runs.

^f Product **3** isolated by pipette column chromatography on regular silica gel, usually in >99% isolated yield.

^g Reaction time was 58 h.

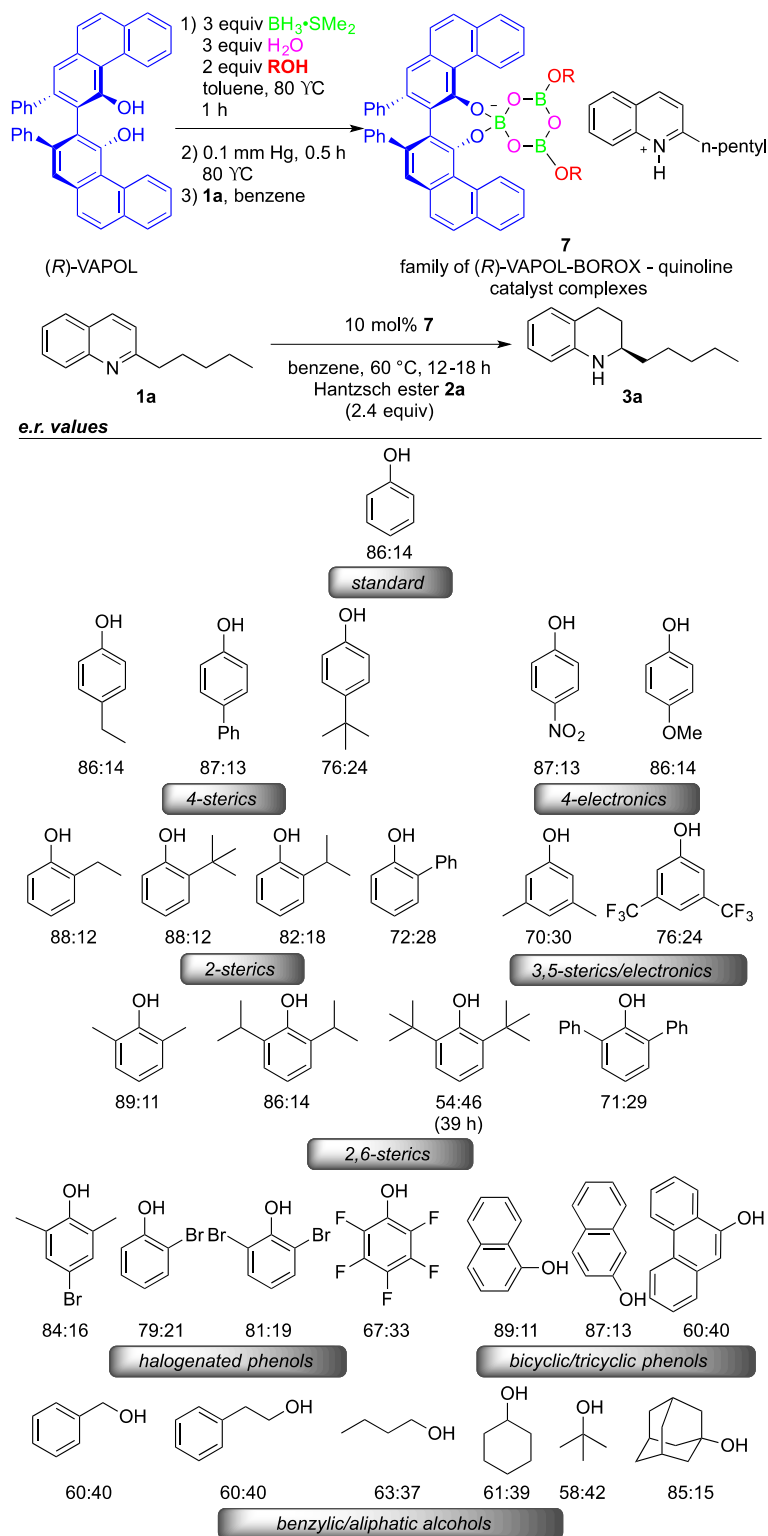
with a quinoline via the BH₃·SMe₂ route then lent us the unique opportunity to generate, in a rapid manner, a large library of these BOROX catalysts **7** (Scheme 1) by simply incorporating different phenols and alcohols during the catalyst self-assembly process. Such a family of BOROX catalysts would offer significantly altered active site polyborate cores; and such a self-assembly process would automatically forgo the tedious synthesis normally associated with preparing catalyst derivatives.

A proof of principle for this concept has been established for the ATH reaction and is shown in Scheme 2;¹⁵ a broad range of commercially available and sterically/electronically different phenols and alcohols were utilized to self-assemble a family of BOROX catalysts (**7**), which were subsequently evaluated for the ATH reaction of 2-pentylquinoline **1a**. The results were gratifying in that the gamut of asymmetric inductions was observed, from an er of 54:46 for the BOROX catalyst prepared from 2,6-di-*tert*-butylphenol to an er of 89:11 for that prepared from 2,6-dimethylphenol and 1-naphthol, thus establishing the importance of this region in the active site polyborate core of the BOROX catalyst.

The mechanism of the Hantzsch ester reduction of quinolines, mediated by a Brønsted acid catalyst, involves a two-step process where the enantioselective step is the reduction of the dihydroquinolinium ion **6** (Table 1).^{12a,16} The hydride transfer to the dihydroquinoline **6**, in the reaction of **1a** and **2a** catalyzed by the (R)-VAPOL-BOROX catalyst, was modeled using ONIOM(B3LYP/6-31G*:AM1) calculations.¹⁷ The main goal of this theoretical

study was to understand the catalyst-substrate interactions that stabilize this transition state. The reported relative energies are from free-energy estimates based on the unscaled harmonic frequencies obtained from the ONIOM calculations.

The lowest energy transition structures leading to the *R* enantiomer (TS1) and the *S* enantiomer (TS2) of **3a** are shown in Figure 3. Transition structures TS1 and TS2 correspond to hydride transfer from **2a** to the iminium carbon of the O3-bound dihydroquinoline **6** (see Fig. 1 for numbering of boroxinate oxygen atoms). The orientation of **2a** as it transfers the hydride is determined by a strong (~1.9 Å) H-bond interaction between the N-H of **2a** and O1 of the catalyst core.¹⁸ The key difference between the two transition structures is that hydride transfer occurs to the pro-*R* face of **6** in TS1 and to the pro-*S* face in TS2. Transition structure TS1 is 0.4 kcal/mol lower in energy than TS2. This prediction, though in the correct direction, is an underestimation of the experimental enantioselectivity (experimental difference is ~1.1 kcal/mol for 86:14 er at 60 °C). This discrepancy can possibly be attributed to inaccuracies arising as a consequence of (a) the size of the computational system, (b) the level of theory used and, (c) not including a solvent model in our calculations. A close examination of TS1 and TS2 also leads to a more interesting explanation for this discrepancy—the hydride is ‘more transferred’ in TS1 and this transition structure is characterized by a larger imaginary frequency (–985) as compared to a smaller value (–707) for the ‘less transferred’ hydride in TS2. The magnitude of the imaginary frequency is



Scheme 2. Self-assembly of a family of BOROXY catalysts (**7**). All reactions were run at 0.05 mmol scale of **1a** and the conversion was 100% (TLC), the product **3a** was isolated (usually in >99% yield) by column chromatography on silica gel.

inversely proportional to the width of the reaction barrier. This suggests that the 0.4 kcal/mol preference for TS1 over TS2 is likely an underestimation of the relative rates of formation of the two enantiomers—hydrogen tunneling would make a larger contribution to the rate of the reaction proceeding via TS1 (narrow barrier) as compared to TS2 (wider barrier).

A pair of transition structures was also located for the reversed H-bonding scenario, with **6** bound to O1 and **2a** bound to O3. These structures were found to be 4.5 (*R ent.*) and 4.8 (*S ent.*) kcal/mol higher in energy than TS1 (see Supporting information). Finally, the importance of both H-bonding interactions in stabilizing the transition state was illustrated when a pair of transition structures

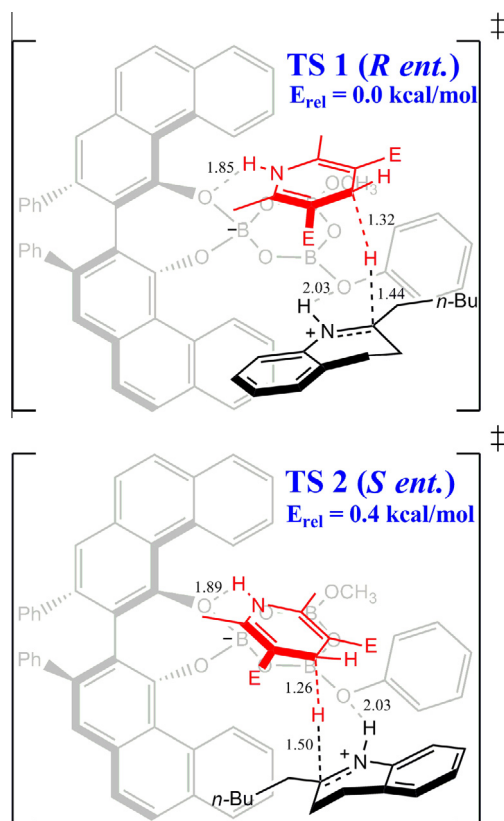


Figure 3. Key transition structures for hydride transfer to O3-bound **6**. All distances are in angstroms. (E = CO₂Et).

corresponding to hydride transfer to the pro-*R* and pro-*S* faces of the O2-bound **6** was located. These structures, lacking the second H-bonding interaction between the catalyst and **2a**, were found to be 16.4 (*R ent.*) and 9.0 (*S ent.*) kcal/mol higher in energy than TS1 (see [Supporting information](#)).

In conclusion, we have shown that our ‘template’ polyborate BOROXY catalyst catalyzes the asymmetric transfer hydrogenation of 2-quinolines with reasonable levels of enantioselectivity. We have also established a simple protocol for generating a large library of these BOROXY catalysts with different sterics and electronics at the active site polyborate core. The lowest energy transition structures (TS1 and TS2) leading to the two enantiomeric products have been identified. The transition state model is consistent with the one proposed for our aziridination reaction mediated by the same catalyst,^{1c,d} where the chiral counterion also offers H-bonding sites to orient the substrates at the stereochemistry determining transition state. Attempts to further improve the asymmetric inductions of the present study are ongoing, as are those to apply this novel library of polyborate BOROXY catalysts for other asymmetric reactions.

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Supplementary data

Supplementary data (synthetic procedures, spectral data for all new compounds, coordinates of all calculated structures and

alternative transition structures, pdb files for TS1 and TS2) associated with this article can be found, in the online version, at <http://dx.doi.org/10.1016/j.tetlet.2015.02.063>.

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