Supporting Information

Metallopeptoids as Efficient Biomimetic Catalysts

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Materials. Rink Amide resin was supplied by Novabiochem; (S)-1-phenylethylamine (spe) and benzylamine (pm) were supplied by Acros organics, Israel; [Cu(MeCN)₄]OTf, 4-aminoTEMPO (tempo), 1,10 phenanthroline-5-amine (phen), Amylamine (aa), t-butylmethlamine (tbm), 2-methoxyethyl amine (me), 1-Naphthalenemethylamine (nm), bromoacetic acid and N,N'diisopropylcarbodiimide (DIC) were supplied by Sigma Aldrich. Other reagents and solvents were obtained from commercial sources and used without additional purification.

Instrumentation. Peptoid oligomers were analyzed by reversed-phase HPLC (analytical C18 column, 5 μm, 100 Å, 2.0x50 mm) on a Jasco UV-2075 instrument. A linear gradient of 5–95% ACN in water (0.1% TFA) over 10 min was used at a flow rate of 0.7 mL/min. Preparative HPLC was performed using a phenomenex C18 column (15μm, 100 Å 21.20x100mm) on a Jasco UV-2075 instrument. Peaks were eluted with a linear gradient of 5–95% ACN in water (0.1% TFA) over 50 min at a flow rate of 5 mL/min. Mass spectrometry was performed on an Waters LCT Premier mass and Advion expression mass under electrospray ionization (ESI), direct probe ACN:H₂O (70:30). Reactions were monitored and analyzed by A Varian 3800 Gas Chromatograph using CP-Sil-8 column. CD measurements were performed using a JASCO circular dichroism

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spectropolarimeter Model J-810-150S. The ¹H NMR spectra were recorded on a Bruker 400 MHz instrument. Coupling constants are given in Hz. The following abbreviations are used to indicate the multiplicity: s, singlet; d, doublet; m, multiplet; bs, broad signal.

Preparation of the peptoid oligomers:

Peptoids were synthesized manually on Rink amide resin using the submonomer approach [1]. All peptoid oligomers were synthesized at room temperature. Typically, 100 mg of resin was swollen in DCM for 40 minutes before initiating oligomer synthesis. Multiple washing steps using DMF were performed between each step described below. De-protection of resin was performed by addition of 20% piperidine solution (1.5 ml in DMF) and the reaction was allowed to shake at room temperature for 20 min. Following the reaction, piperidine was washed from the resin using DMF (10 mL g⁻¹ resin) (3 x 1 min). Bromoacetylation was completed by adding 20 eq. bromoacetic acid (1.2 M in DMF, 8.5 mL g⁻¹ resin) and 24 eq. of diisopropylcarbodiimide (2 mL g⁻¹ resin); this reaction was allowed to shake at room temperature for 20 min. Following the reaction, the bromoacetylation reagents were washed from the resin using DMF (10 mL g⁻¹ resin) (3 x 1 min) and 20 eq. of submonomer amine (1.0 M in DMF, 10 mL g⁻¹ resin) were added. The amine displacement reaction was allowed to shake at room temperature for 20 min and was followed by multiple washing steps (DMF, 10 mL g⁻¹ resin) (3 x 1 min). This two-step addition cycle was modified as follows: the last amine displacement with 1,10-Phenanthroline-5-amine (20 equivalents, 0.4 M in DMF, 10 mL g⁻¹ resin) was performed for 24 hr. Bromoacetylations and amine displacement steps were repeated until the peptoids were obtained. To cleave the peptoid oligomers from solid support for analysis, approximately 5 mg of resin was treated with 95% TFA in water (40 mL g⁻¹ resin) for 10 minutes. The cleavage cocktail was evaporated under nitrogen gas and the peptoid oligomers were re-suspended in 0.5 mL HPLC solvent (1:1 HPLC grade acetonitrile:HPLC grade water). To cleave the peptoid oligomers from solid support for preparative HPLC the beads were treated with 5ml of 95% TFA in water for 30 minutes. The cleavage cocktail was evaporated under low pressure, re-suspended in 2 mL HPLC solvent and lyophilized overnight.

Characterization of the peptoid oligomers:

Peptoid **Helix i+3**, **Helix i+1** and **Nonhelix i+3** were characterized by analytical HPLC using a C18 column. The peptoids **DI**, **BT**, **MT**, **RBT**, **PT**, **TT** and **NT** were characterized by preparative HPLC using a C18 column and 1H NMR. Analysis with analytical C18 column was done using a solvent gradient conducted from 5% to 95% solvent B (0.1% TFA in HPLC grade acetonitrile) over solvent A (0.1% TFA in HPLC grade water) in 10 minutes with a flow rate of 0.7 mL min⁻¹. Analysis with preparative C18 column was done using a solvent gradient conducted from 5% to 95% solvent B (0.1% TFA in HPLC grade acetonitrile) over solvent A (0.1% TFA in HPLC grade water) in 50 minutes with a flow rate of 5 mL min⁻¹. Additional characterization was conducted by ESI MS. The peptoids were further purified to >95% by RP-HPLC and lyophilized overnight. HPLC traces of the pure peptoid oligomers are depicted in Figures S1-S9. All peptoids were subjected again to ESI MS characterization. MS spectra of the pure peptoid oligomers are depicted in Figures S10-S19. The two helical peptoids **Helix i+3** and **Helix i+1** were also characterized by CD spectroscopy (Figure S35 and S36). The peptoids **DI**, **BT**, **MT**, **RBT**, **PT**, **TT** and **NT** were characterized further by ¹H NMR (S38-S44).

Catalysis experiments:

In a typical catalytic procedure acetonitrile was distilled over calcium hydride until a water content of 0.05% (w/w) in the solvent was reached. [Cu(MeCN)₄]OTf (0.617 mg, 1.65 μ mol) and the peptoid catalysts (1mg, 1.65 μ mol) were placed in a dry vial, which was then capped with a septum. Dry acetonitrile (0.1 mL) and 1-methyl imidazole (1.8 μ L, 3.3 μ mol) were added, and the mixture was stirred for 5 minutes at rt following the exposure of the solution mixture to air. Benzyl alcohol (34 μ L, 330 μ mol) was then added, and the resulting mixture was stirred for 3hrs under air.

ESI-MS analysis of intermediates 1-6:

Peptoid catalyst **BT** (1.65 μ mol) and [Cu(MeCN)₄]OTf (1.65 μ mol) were placed in a dry vial, which was then capped with a septum and purged with N₂ for 3-4 minutes. Dry acetonitrile 100 μ l was then add as the solvent, followed by these steps (i) a sample was diluted with dry acetonitrile and subjected to MS analysis, (ii) under these conditions,

MMI (3.3 μ mol) was added, stirred and a sample was similarly analyzed. Thereafter, the system was exposed to air, followed by an immediate analysis of a sample by ESI-MS (S32). (iii) Finally, benzyl alcohol (330 μ mol) was added to the mixture, stirred for 10 minutes, and a sample was analyzed by ESI-MS (Fig. S20-S25 respectively). Similar analysis was done with **DI** peptoid catalyst except the stirring time was increased to 30 min upon adding NMI and benzyl alcohol (S26-S31).

References

1. R. N. Zuckermann, J. M. Kerr, S. B. W. Kent, W. H. Moos, *J. Am. Chem. Soc.* **1992**, *114*, 10646.

Table S1. Peptoid oligomer sequences and their corresponding molecular weights.

Peptoid	Oligomer sequence	Oligomer length	Molecular weight		
			Calculated: Found		
Helix i+3	N phen $(N$ spe $)_2N$ tempo $(N$ spe $)_3$	7mer	1269.5:1270.1		
Nonhelix i+3	N phen $(N$ me $)_2N$ tempo $(N$ me $)_3$	7mer	1039.2:1040.0		
Helix i+1	NphenNtempo(Nspe) ₅	7mer 1269.5:1269.7			
DI	NphenNtempo	2mer	463.5: 465.2		
BT	NphenNtempoNpm	3mer	610.7: 612.0		
MT	NphenNtempoNme	3mer 578.6: 580			
RBT	NphenNpmNtempo 3mer 61		610.7: 611.8		
PT	NphenNtempoNaa	3mer 590.7:592.6			
TT	NphenNtempoNtbm	3mer	590.7:592.4		
NT	NphenNtempoNnm	3mer 660.8:662.5			

¹H NMR Data

DI (400 MHz, CD₃OD): δ 1.36 (s, 6H), 1.40 (s, 6H), 1.93-1.96 (m, 2H), 2.09-2.16 (m, 2H), 3.20 (s, 4H), 4.23 (s, 3H), 4.92-4.98 (m, 1H), 6.89 (s, 1H), 7.88-7.94 (m, 2H), 8.64 (d, 1H, J = 5.2 Hz), 8.70 (d, 1H, J = 8.3Hz), 8.77 (d, 1H, J = 8.4 Hz), 9.09 (d, 1H, J = 3.8 Hz).

BT (400 MHz, CD₃OD): δ1.22-1.34 (m, 12H), 1.78-1.88 (m, 2H), 1.97-2.04 (m, 2H), 4.08-4.11 (m, 4H), 4.29-4.33 (m, 1H), 4.48-4.49 (m, 2H), 4.59 (bs, 2H), 6.82-6.89 (m, 1H), 7.13-7.19 (m, 2H), 7.22-7.33 (m, 3H), 7.77-7.83 (m, 2H), 8.47-8.52 (m, 1H), 8.62 (d, 1H, *J* = 4.8 Hz), 8.70-8.73 (m, 1H), 9.04-9.05 (bs, 1H).

MT (400 MHz, CD₃OD): δ1.44-1.49 (m, 12H), 1.95-2.05 (m, 2H), 2.13-2.16 (m, 2H), 3.39-3.47 (m, 3H), 3.55-3.75 (m, 5H), 4.21-4.23 (bs, 2H), 4.35-4.36 (bs, 1H), 4.49-4.54 (bs, 2H), 4.68-4.72 (bs, 3H), 6.96-7.05 (m, 1H), 7.98-7.99 (m, 2H), 8.74-8.86 (m, 3H), 9.12-9.18 (bs, 1H).

RBT(400 MHz, CD₃CN): δ1.36 (s, 6H), 1.47 (s, 6H), 2.08-2.11 (m, 4H), 3.96-4.03 (m, 2H), 4.17-4.25 (m, 3H), 4.34 (bs, 1H), 4.69-4.79 (m, 3H), 6.71 (bs, 2H), 7.00 (s, 1H), 7.34-7.47 (m, 5H), 7.78-7.93 (m, 2H), 8.57-8.67 (m, 2H), 8.80-8.81 (bs, 1 H), 9.07-9.08 (bs, 1H), 12.66 (bs, 1H).

PT (400 MHz, CD₃OD): δ 0.83-0.86 (m, 3H), 1.33-1.34 (m, 4H), 1.42-1.66 (m, 14H), 1.42-1.60 (m, 2H), 1.74-1.76 (m, 1H), 2.02-2.90 (m, 2H), 2.13-2.16 (m, 2H), 3.45-3.53 (m, 2H), 4.18-4.28 (m, 4H), 4.53-4.59 (m, 2H), 6.97-7.02 (bs, 1H), 7.94-7.98 (m, 2H), 8.70-8.75 (m, 2H), 8.82-8.85 (m, 1H), 9.16-9.17 (bs, 1H).

TT (400 MHz, CD₃OD): δ0.80- 0.81 (m, 9H), 1.23-1.30 (m, 12H), 1.79-1.82 (m, 2H), 1.99-2.01 (m, 2H), 3.12-3.13 (m, 4H), 4.01-4.02 (bs, 2H), 4.16-4.18 (bs, 2H), 4.35-4.36 (bs, 2H), 6.76-6.78 (bs, 1H), 7.75-7.78 (m, 2H), 8.52-8.54 (m, 2H), 8.61-8.65 (m, 1H), 8.95 (bs, 1H).

NT(400 MHz, CD₃CN): δ 1.29 (s, δ H), 1.42 (s, δ H), 2.05 (d, δ H, J = 7.6 Hz), 3.96 (d, δ H, J = 5.2 Hz), 4.29 (s, δ H), 4.86-4.94 (m, δ H), 5.01 (s, δ H), 6.20 (s, δ H), 6.49 (s, δ H), 6.79 (s, δ H), 7.21-7.26 (m, δ H), 7.40 (d, δ H), δ HH, δ

HPLC

HPLC solvent A: (0.1 % TFA in water), HPLC solvent B: (0.1 % TFA in acetonitrile)

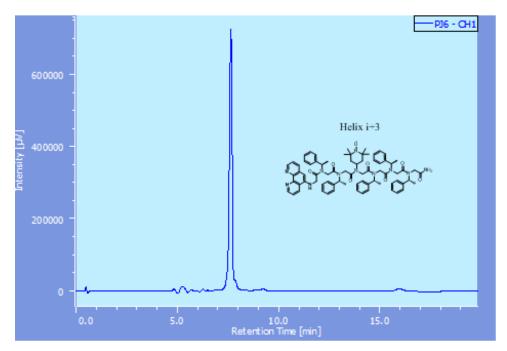


Figure S1. HPLC trace of peptoid Helix i+3

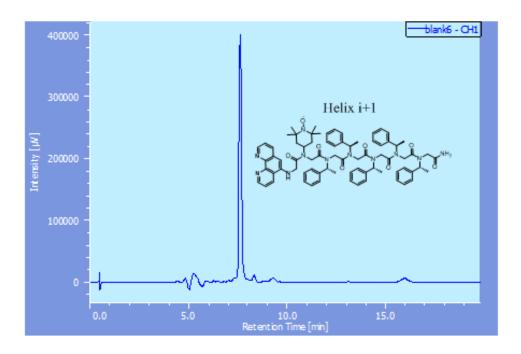


Figure S2. HPLC trace of peptoid Helix i+1

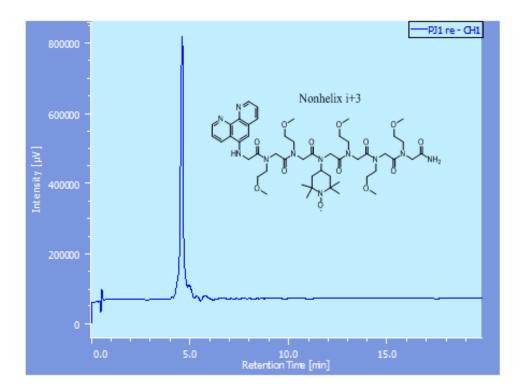


Figure S3. HPLC trace of peptoid Nonhelix i+3

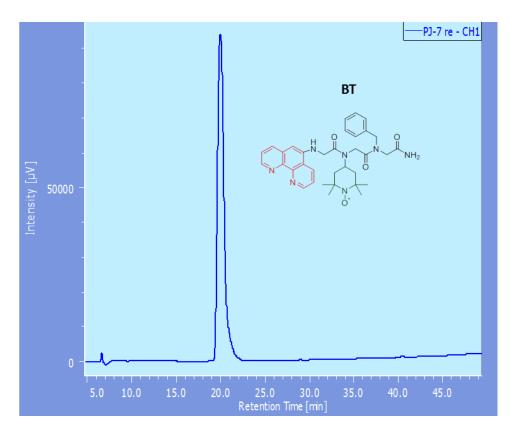


Figure S4. HPLC trace of peptoid BT

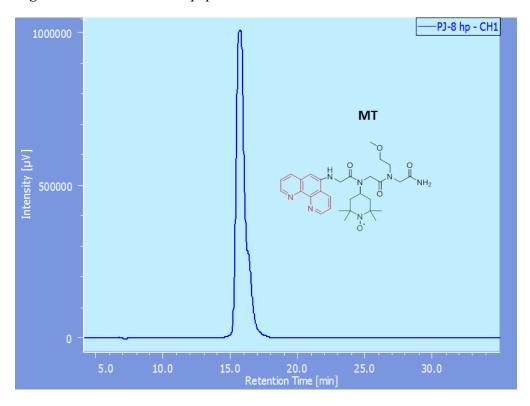


Figure S5. HPLC trace of peptoid MT

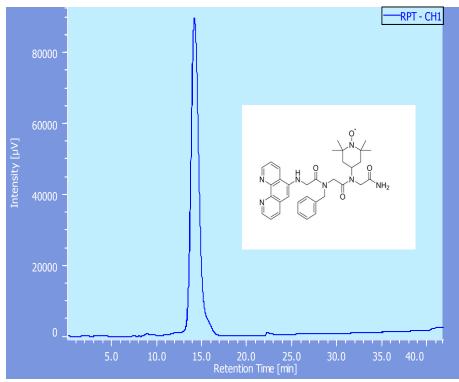


Figure S6. HPLC trace of peptoid RBT

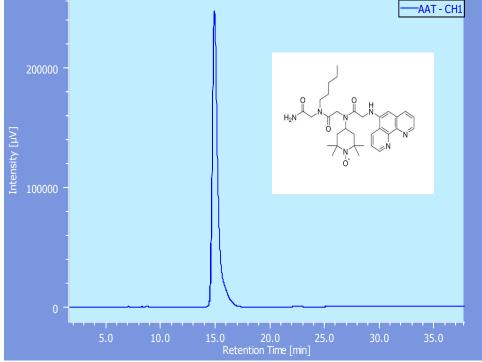


Figure S7. HPLC trace of peptoid PT

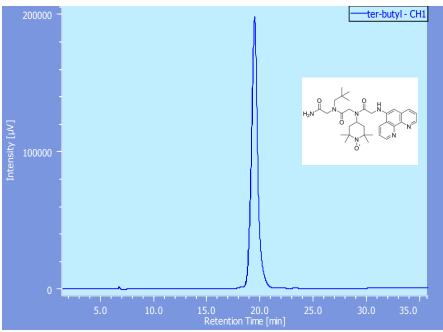


Figure S8. HPLC trace of peptoid TT

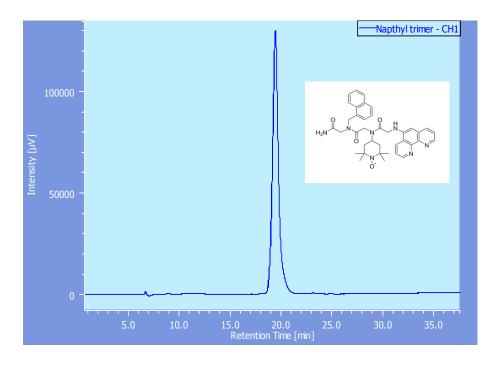


Figure S9. HPLC trace of peptoid NT

ESI MS of peptoid oligomers

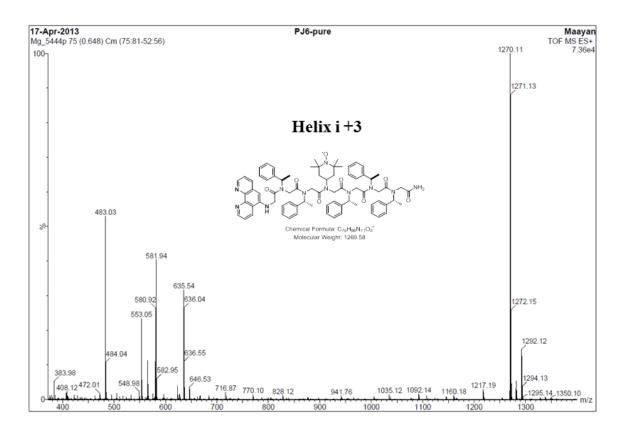


Figure S10. ESI MS of peptoid Helix i+3

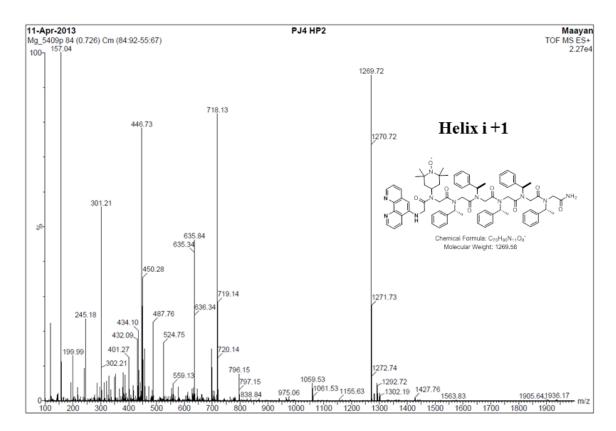


Figure S11. ESI MS of peptoid Helix i+1

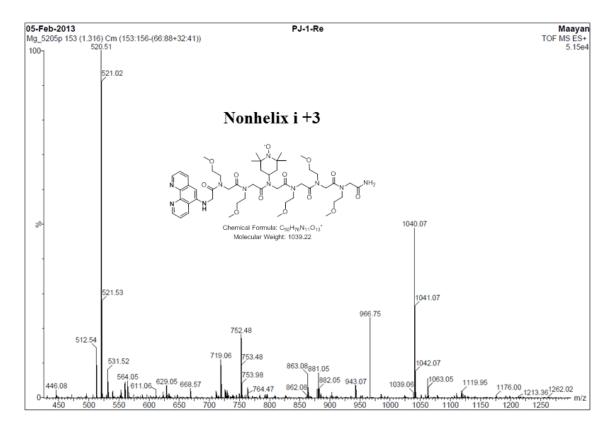


Figure S12. ESI MS of peptoid Nonhelix i+3

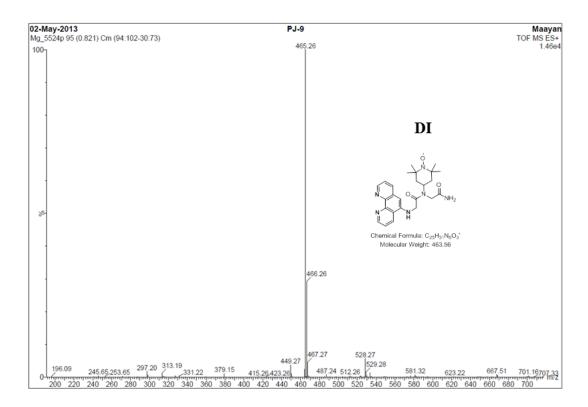


Figure S13. ESI MS of peptoid DI

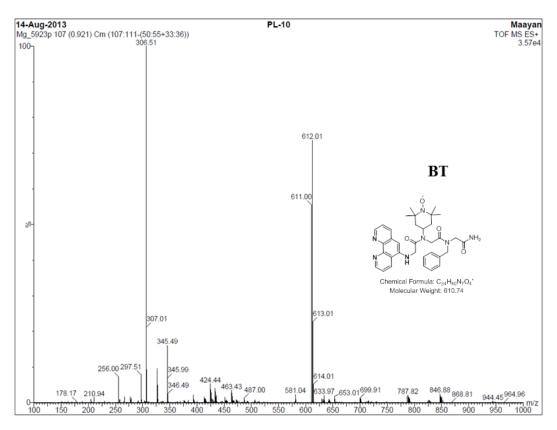


Figure S14. ESI MS of peptoid BT

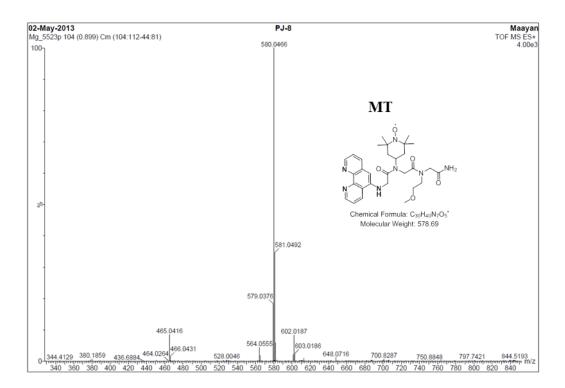


Figure S15. ESI MS of peptoid MT

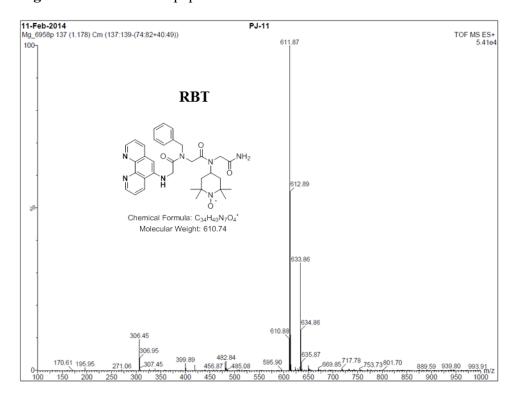


Figure S16. ESI MS of peptoid RBT



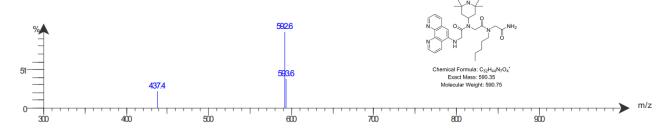
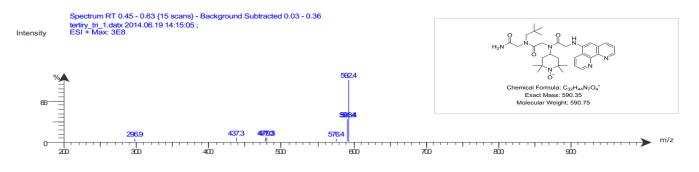


Figure S17. ESI MS of peptoid PT

Figure S18. ESI MS of peptoid TT



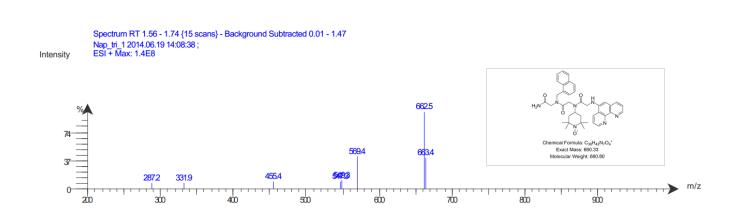


Figure S19. ESI MS of peptoid NT

ESI MS of catalysis intermediates



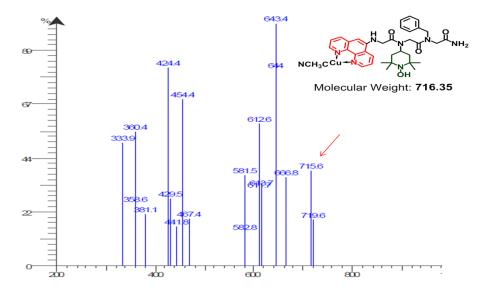


Figure S20. ESI MS of reaction intermediate 1 of BT

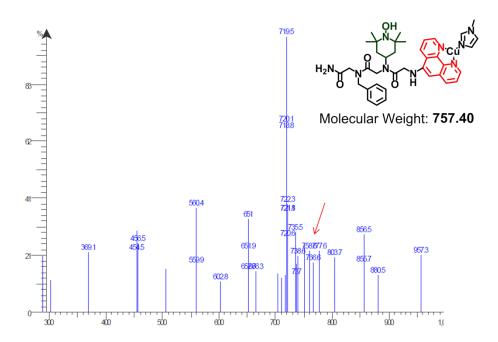


Figure S21. ESI MS of reaction intermediate 2 of BT



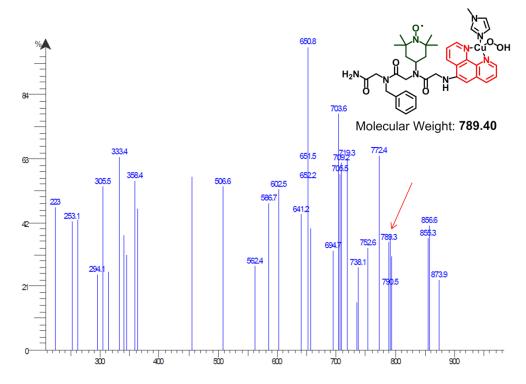


Figure S22. ESI MS of reaction intermediate 4 of BT

 $Spectrum RT 1.14 \{1 scans\} - Background Subtracted 0.97 - 1.16 \\ BT + Cu I nmldatx 2014.11.24 18:20:03 ; \\ ESI + Max: 3.2E6$

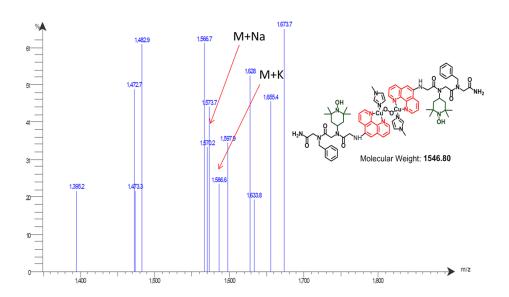


Figure S23. ESI MS of reaction intermediate 3 of BT



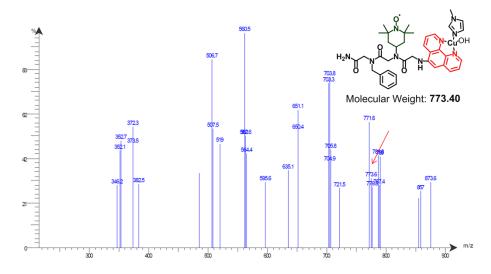


Figure S24. ESI MS of reaction intermediate 5 of BT

```
| Spectrum RT 1.08 - 1.38 {12 scans} - Background Subtracted 1.05 - 1.22 | BT + Cu | nmi_bzoh.datx 2014.11.24 18:23:45 ; | nsity | ESI + Max: 1E6
```

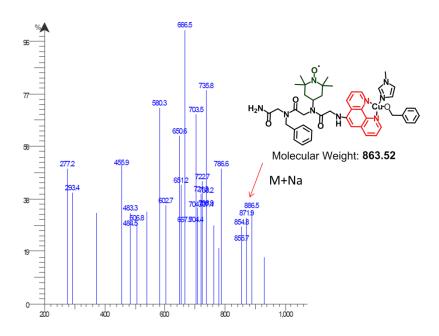


Figure S25. ESI MS of reaction intermediate 6 of BT

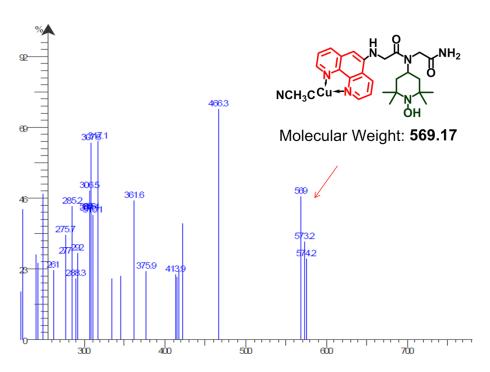


Figure S26. ESI MS of reaction intermediate 1 of DI

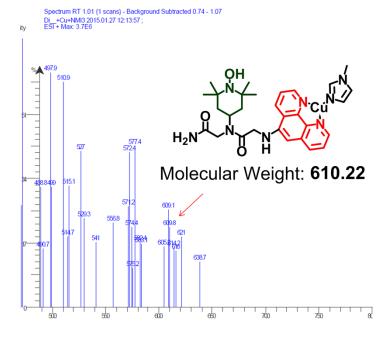


Figure S27. ESI MS of reaction intermediate 2 of DI



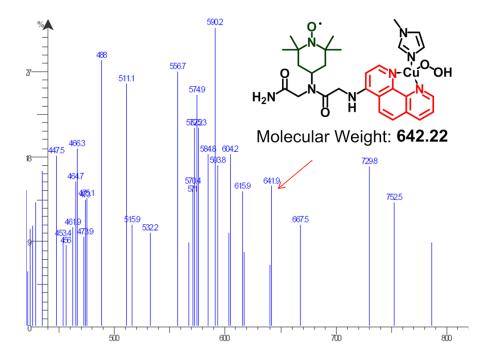


Figure S28. ESI MS of reaction intermediate 4 of DI

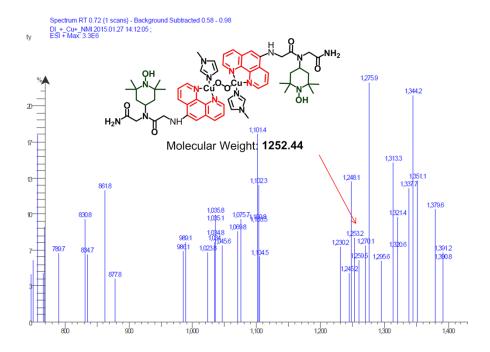


Figure S29. ESI MS of reaction intermediate 3 of DI

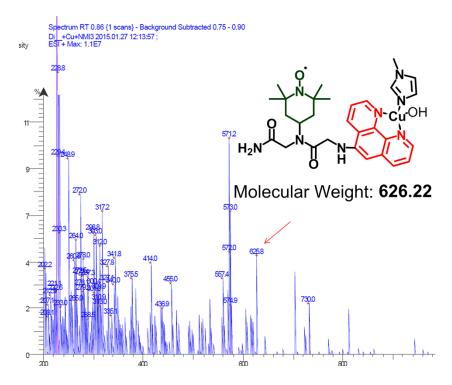


Figure S30. ESI MS of reaction intermediate 5 of DI

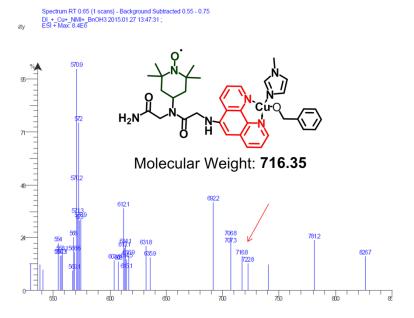


Figure S31. ESI MS of reaction intermediate 6 of DI

Figure S32. ESI-MS analysis of our reaction intermediates as a support for the proposed mechanism suggested by Sthal et al. Reaction conditions: dry acetonitrile (0.1 mL) at room temperature with 330 μ mol benzyl alcohol, 0.5 mol % catalyst(s), 0.5 mol % [Cu(MeCN)₄]OTf and 1 mol % NMI.

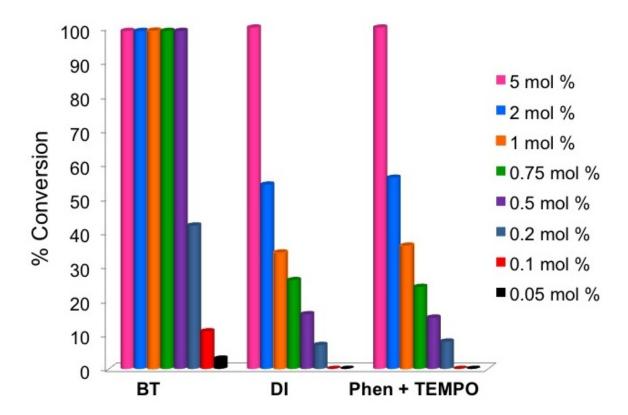


Figure S33. Oxidation of benzyl alcohol with three different catalytic systems in concentrations varied from 5 mol% to 0.05 mol%. All of the reactions were performed in dry acetonitrile (0.1 mL) at room temperature for 3 hrs. with 330 μmol benzyl alcohol, 1 equiv. [Cu(MeCN)₄]OTf and 2 equiv. NMI for each equiv. of catalyst(s). Conversions were determined by gas chromatography

Figure S34. The third set of peptoid sequences that were used to evaluate intramolecular cooperative catalysis in the oxidation of benzyl alcohol.

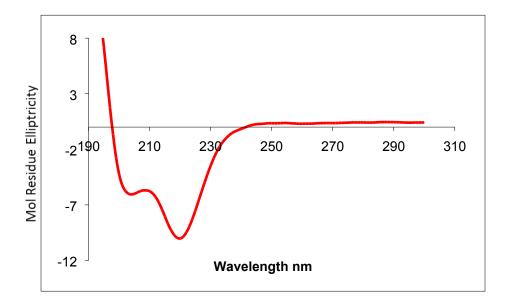


Figure S35. CD spectrum of peptoid **Helix i+1**. The spectrum was recorded at room temperature, in a MeOH/ H_2O (4:1) and peptoid concentration of 5mM.

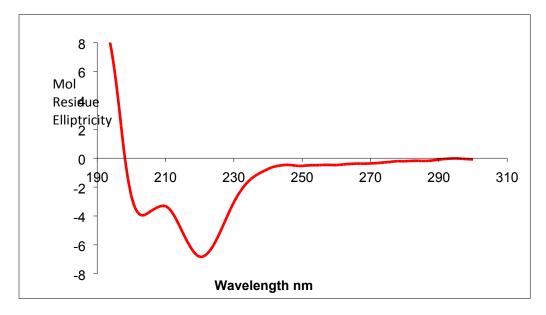


Figure S36. CD spectrum of peptoid **Helix i+3**. The spectrum was recorded at room temperature, in a MeOH/ H_2O (4:1) and peptoid concentration of 5mM.

Catalysis

Table S2.Efficiency of the peptoid catalysts **BT** and **DI** compared to the intermolecular control system in the oxidation of benzyl alcohol, with different catalysts loadings

Entry	Catalyst	% Conversion (TON)[a]						
	loading							
	(mol %)	BT	NT	TT	DI	Phen+TEMPO		
1	0.5	99 (198)	98 (196)	97 (194)	10 (20)	16(32)		
2	0.2	41 (205)	nd	nd	2 (4)	2 (4)		
3[b]	0.2	66 (330)	nd	nd	6 (30)	7 (35)		
4	0.1	11 (110)	nd	nd	nd	nd		
5 ^[b]	0.1	49 (490)	49 (490)	46 (460)	3 (30)	3 (30)		

All of the reactions were conducted in dry acetonitrile as the solvent (0.1ml) at room temperature for 3hrs., using 0.5mol% catalyst,0.5mol% [Cu(MeCN₄]OTf and 1mol% NMI. [a]As determined by gas chromatography.[b] Reaction time was 12 hours. nd = not detected

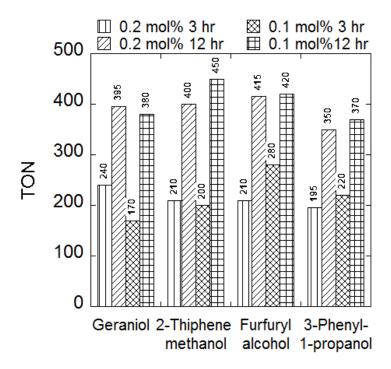


Figure S37. Oxidation of aromatic alcohols catalyzed by BT at low catalyst loadings.

1H NMR Spectra

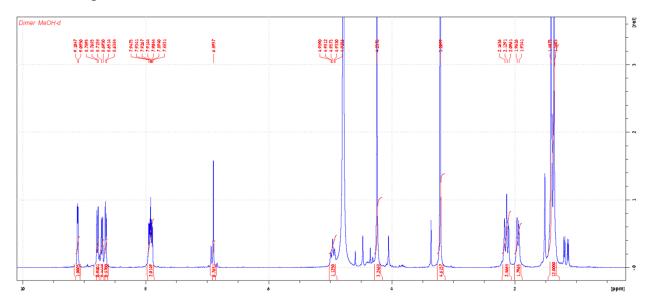


Figure \$38.1H NMR spectra of **DI** in CD₃OD

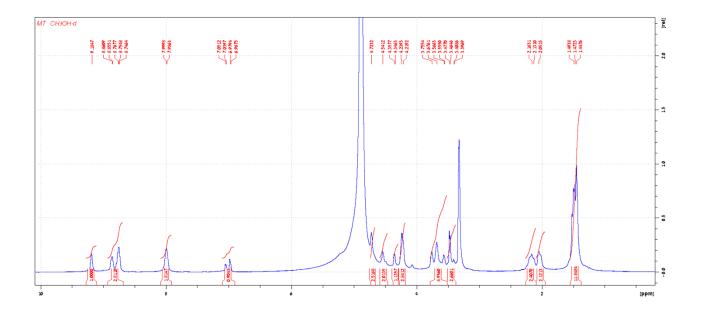


Figure \$39.1H NMR spectra of MT in CD₃OD

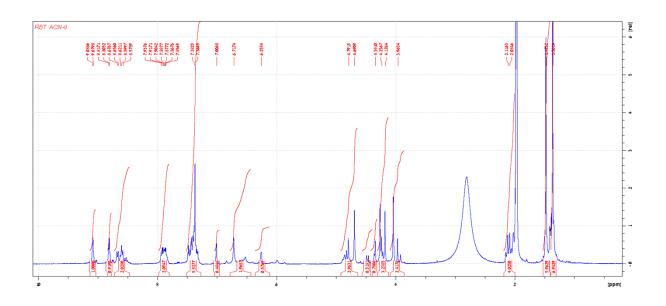


Figure S40. ¹H NMR spectra of RBT in CD₃OD

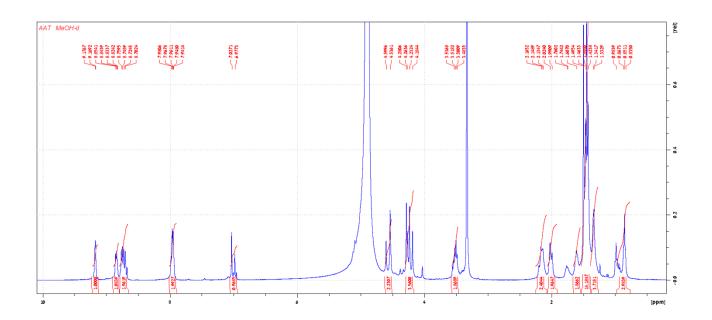


Figure S14. ¹H NMR spectra of PT in CD₃CN

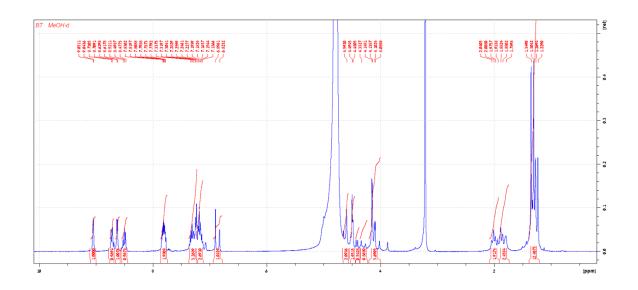


Figure S42. ¹H NMR spectra of BT in CD₃OD

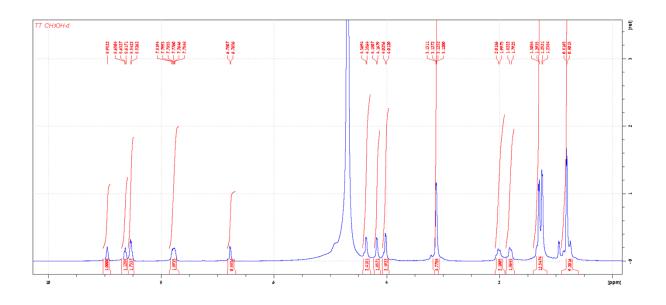


Figure S43. ¹H NMR spectra of TT in CD₃OD

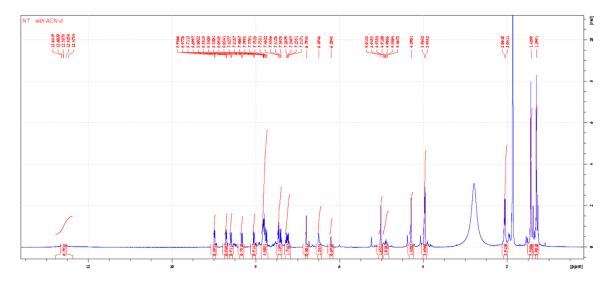


Figure S44. 1H NMR spectra of NT in CD₃CN