

**Development of Chiral DMAP Catalyst for Dynamic Kinetic Resolution of Azole
Hemiaminals**

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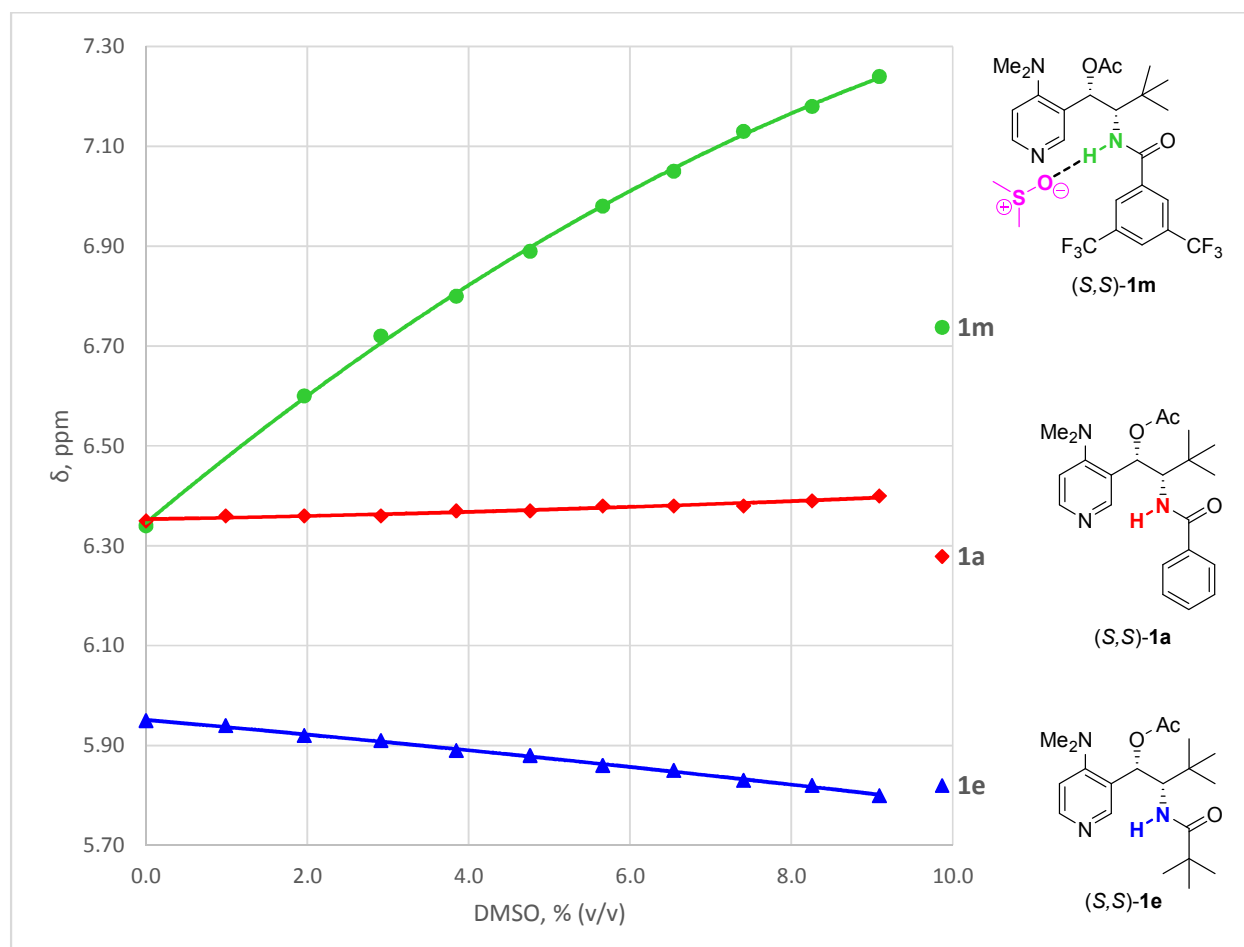
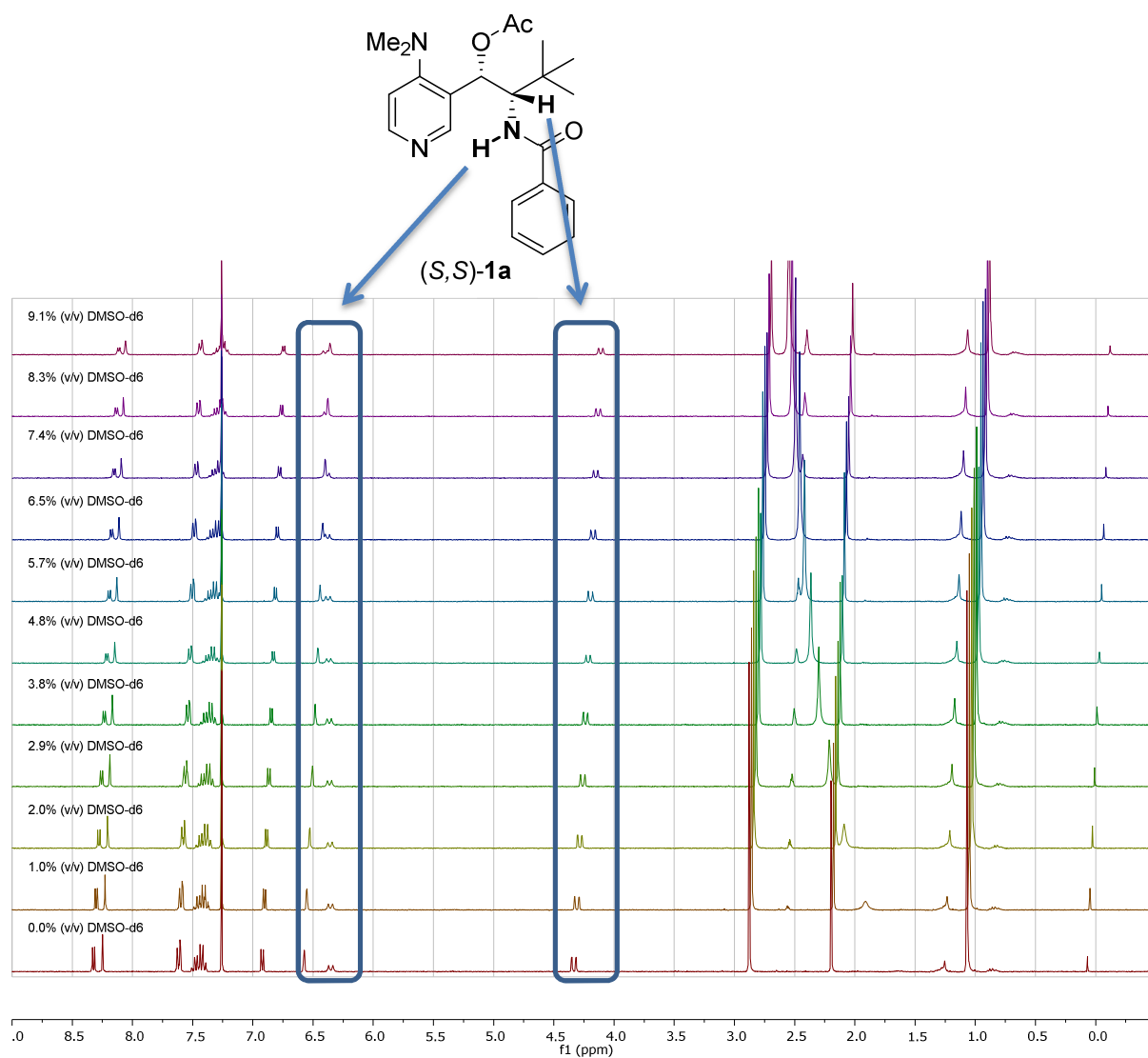
NMR titration of catalysts (*S,S*)-**1a,e,m**

Figure S1. NMR titration of (*S,S*)-**1a,e,m** with DMSO-*d*₆ in CHCl₃-*d* at 298 K.

H-bond formation ability of catalysts (*S,S*)-**1a,e,m** was evaluated by NMR titration of their CHCl₃-*d* solution with DMSO-*d*₆.¹ Accordingly, DMSO-*d*₆ was added portionwise (1 vol% increment) to the CHCl₃-*d* solution of DMAP species (*S,S*)-**1a,e,m** and change of the amide N-H chemical shifts was monitored. The amide ¹H chemical shifts for each of the catalyst (*S,S*)-**1a,e,m** was plotted versus amount of the added DMSO-*d*₆ (Scheme S8). Negligible changes of ¹H chemical shifts were observed for catalysts (*S,S*)-**1a,e** indicating that these DMAP species feature weak H-bonding interactions with DMSO. In contrast, a relatively large downfield shift

(1) (a) Wagner, G.; Pardi, A.; Wuethrich, K. *J. Am. Chem. Soc.* **1983**, *105*, 5948–5949; (b) Jansma, A.; Zhang, Q.; Li, B.; Ding, Q.; Uno, T.; Bursulaya, B.; Liu, Y.; Furet, P.; Gray, N. S.; Geierstanger, B. H. *J. Med. Chem.* **2007**, *50*, 5875–5877.

Figure S3. $^1\text{H-NMR}$ spectra for titration of (S,S)-1a with DMSO- d_6

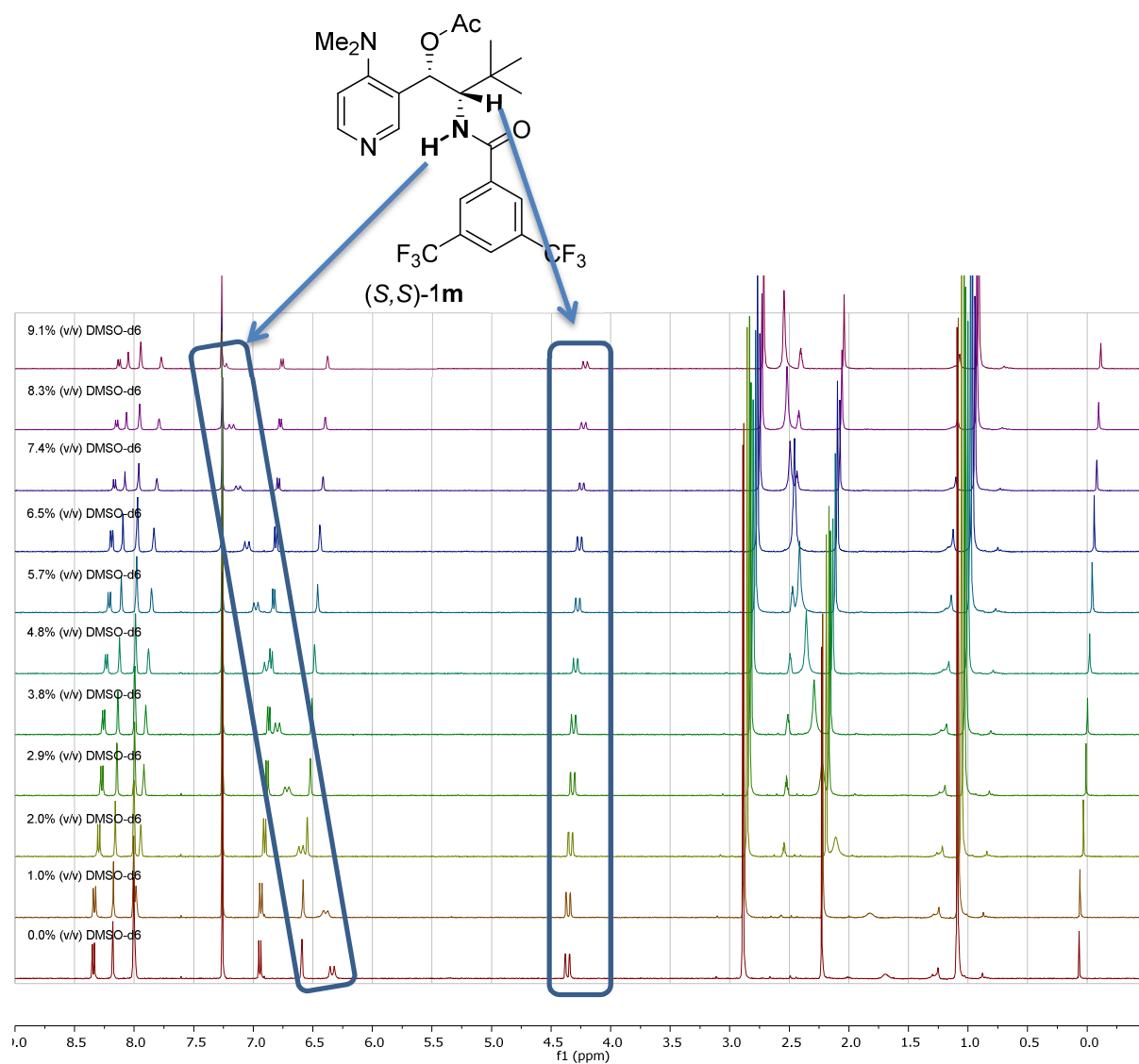
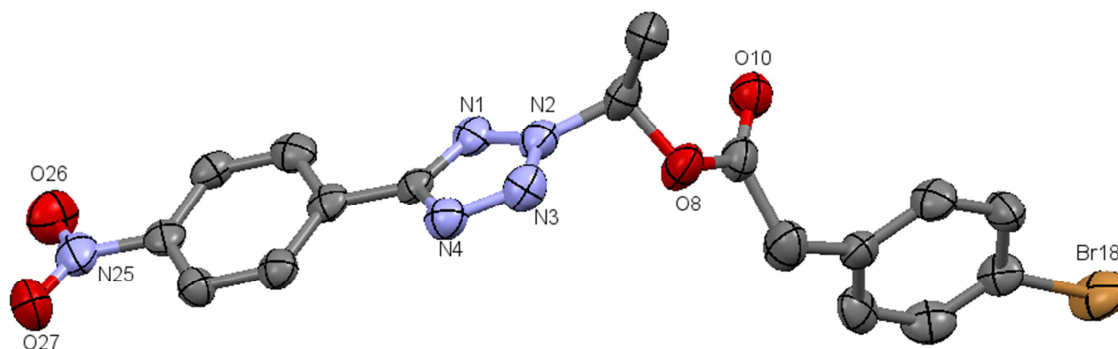


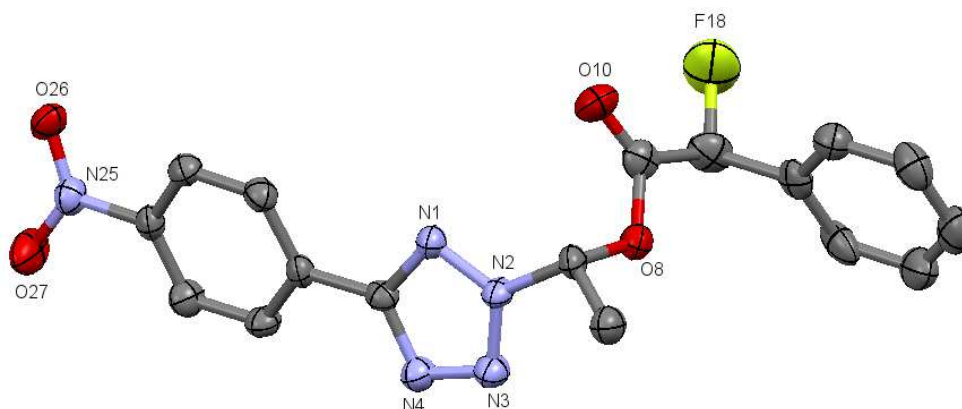
Figure S4. $^1\text{H-NMR}$ spectra for titration of (*S,S*)-**1m** with DMSO- d_6

X-Ray Structure, Crystal Data and Structure Refinements for (*R*)-**12x**

(ellipsoids at 50% probability; hydrogen atoms are omitted for clarity)

Empirical formula	C ₁₇ H ₁₄ BrN ₅ O ₄
Formula weight	432.24
Temperature	193(2) K
Wavelength	0.71073 Å
Crystal system, space group	Monoclinic, P 21
Unit cell dimensions	a = 13.5743(5) Å alpha = 90 deg. b = 7.6627(4) Å beta = 107.8219(17) deg. c = 18.0825(11) Å gamma = 90 deg.
Volume	1790.61(16) Å ³
Z, Calculated density	4, 1.603 Mg/m ³
Absorption coefficient	2.330 mm ⁻¹
F(000)	872
Crystal size	0.24 x 0.16 x 0.11 mm
Two-theta max. for data	56.0 deg.
Limiting indices	-17<=h<=17 -8<=k<=10 -23<=l<=23
Reflections collected / unique	e 7401 / 7394 [R(int) = 0.0127]
Completeness to theta = 28.0	98%
Absorption correction	None
Refinement method	Full-matrix least-squares on F ²
Data / restraints / parameter	s 7394 / 1 / 487
Goodness-of-fit on F ²	1.096
Final R indices [I>2sigma(I)]	R1 = 0.0797, wR2 = 0.1501
R indices (all data)	R1 = 0.1263, wR2 = 0.1697
Absolute structure parameter	0.032(13)
Largest diff. peak and hole	0.848 and -0.797 e.Å ⁻³

X-Ray Structure, Crystal Data and Structure Refinements for (R,S)-14



(ellipsoids at 50% probability; hydrogen atoms are omitted for clarity)

Empirical formula	C ₁₇ H ₁₄ FN ₅ O ₄
Formula weight	371.33
Temperature	173(2) K
Wavelength	0.71073 Å
Crystal system, space group	Monoclinic, P 21
Unit cell dimensions	a = 6.5366(4) Å alpha = 90 deg.
	b = 9.8897(6) Å beta = 96.641(2) deg.
	c = 13.2763(10) Å gamma = 90 deg.
Volume	852.49(10) Å ³
Z, Calculated density	2, 1.447 Mg/m ³
Absorption coefficient	0.113 mm ⁻¹
F(000)	384
Crystal size	0.47 x 0.07 x 0.02 mm
Two-theta max. for data	58.0 deg.
Limiting indices	-8 ≤ h ≤ 8 -12 ≤ k ≤ 13 -17 ≤ l ≤ 14
Reflections collected / unique	6374 / 2316 [R(int) = 0.0945]
Completeness to theta = 29.0	98%
Absorption correction	None
Refinement method	Full-matrix least-squares on F ²
Data / restraints / parameters	2316 / 1 / 244
Goodness-of-fit on F ²	1.008
Final R indices [I > 2σ(I)]	R1 = 0.0662, wR2 = 0.1143
R indices (all data)	R1 = 0.1331, wR2 = 0.1362
Largest diff. peak and hole	0.212 and -0.219 e.Å ⁻³

Spectroscopic Data

