## **Supporting Information for**

# Visualising diastereomeric interactions of chiral amine-chiral copper Salen adducts by EPR spectroscopy and DFT.

Damien M. Murphy,<sup>1</sup> Ignacio Caretti,<sup>2</sup> Emma Carter,<sup>1</sup> Ian A. Fallis,<sup>1</sup>

Marcus C. Göbel,<sup>1</sup> James Landon,<sup>1</sup> Sabine Van Doorslaer,<sup>2</sup> David J. Willock,<sup>1</sup>

<sup>1</sup>School of Chemistry, Cardiff University, Main Building, Park Place, Cardiff CF10

3AT, UK and <sup>2</sup>SIBAC laboratory – Department of Physics, University of Antwerp,

Universiteitsplein 1, B-2610 Wilrijk, Belgium.

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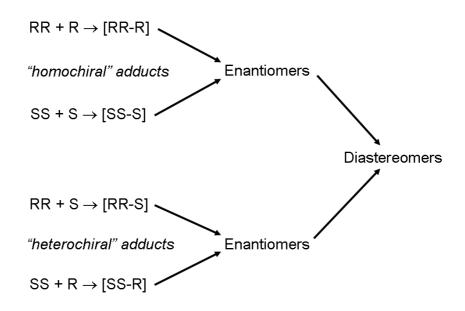
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#### (1) Further Experimental Details:

### Preparation of samples for the EPR measurements

*X-/Q-band CW-EPR/ENDOR*: For CW-EPR/ENDOR measurements, a 1 ml volumetric stock solution of [Cu(1,2)] (0.100 g, 0.16 mmol) in dichloromethane was prepared and 200 µl (Hamilton glass microsyringes) aliquots of this solution were placed in 2 ml vials. The vials were placed in a heating block at 50°C and evaporated to dryness under a slow stream of nitrogen to afford [Cu(1,2)] as a purple glass. The vials were allowed to cool and to each 115 µl of dichloromethane and 115 µl of toluene was added to afford a high quality frozen solution. A number of additional EPR experiments were also performed (under similar conditions) using racemic (or enantiomeric *R-, S-*) methylbenzylamine (MBA). In the presence of amine, the parent complex was dissolved in 100 µl of dichloromethane and 100 µl of amine, with a small amount (30 µL) of toluene added to aid glass formation. The sample volume used in X-band EPR measurements was 150 µL, whereas 20 µL sample volume was used in Q-band ENDOR measurements.

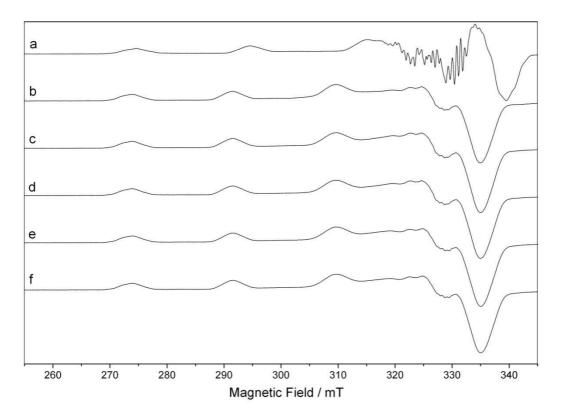
*W-band CW and pulsed EPR:* First, a 32mM stock solution was prepared by dissolving the *R*,*R* or *S*,*S* form of the copper complex under study (Cu[1] or Cu[2]) in 1ml of dichloromethane. Then, 100µl aliquots of this solution were placed in 4ml vials and evaporated to dryness at 50°C under nitrogen atmosphere inside a glovebox. Finally, 230µl of toluene (in case of Cu[1]) or dichloromethane (in case of Cu[2]) was added to the vials. To study amine coordination, 200µl of either *R-*,*S-* or *rac*-methylbenzylamine (MBA) was added together with 30µl of toluene.



Scheme S1: Summary of the configurational stereoisomers examined:  $\mathbf{RR} = R, R$ -[Cu(1,2)],  $\mathbf{SS} = S, S$ -[Cu(1,2)],  $\mathbf{R} = R$ -MBA,  $\mathbf{S} = S$ -MBA. [**RR-R**] therefore represents the specific complex-amine adduct of R, R-[Cu(1,2)] + R-MBA.

## (2) X-band EPR spectra of [Cu(2)] in MBA

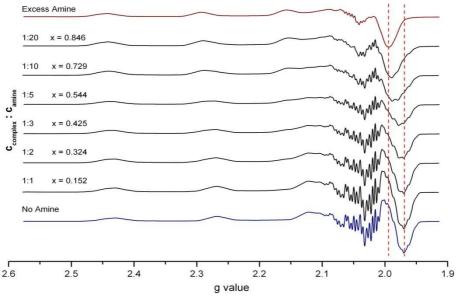
**Figure S1**: X-band CW-EPR spectra recorded at 140K of (a) the non-coordinated [Cu(2)] complex in toluene:dichloromethane together with the different [Cu(2)]-MBA combinations of (b) *rac-rac*, (c) *RR-S*, (d) *SS-R*, (e) *RR-R* and (f) *SS-S*.



The equivalent series of spectra for [Cu(1)] were given in Fig.1 of main paper.

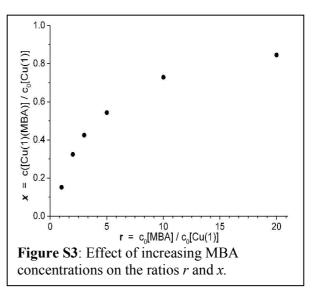
#### (3) Determination of the number of coordinated MBA molecules (*n*) to [Cu(1)]

A series of *rac-rac* samples, i.e., [Cu(1)]+MBA, were prepared using a constant concentration of [Cu(1)] (c = 3.3205 mmol dm<sup>-3</sup>) and the ratio of MBA was varied from 1 to 20. The solvent used throughout was toluene. The resulting EPR spectra are shown in Fig.S2 below. For comparison, the EPR spectra of [Cu(1)] recorded in neat toluene (no amine; blue trace) and in an excess of amine (1:1200; red trace) are also shown.



**Figure S2**: X-band CW-EPR spectra recorded at 130K of *rac*-[Cu(1)] dissolved in toluene with increasing concentrations of *rac*-MBA.

The spectra in Fig.S2 can be seen to progressively shift from the extreme cases of *no-amine present*, a typical non-coordinated spectrum (blue trace), to the *excess amine* case giving a coordinated spectrum (red trace). These shifts are most notable in the overshoot region of the spectra. The intervening EPR spectra (with Cu:MBA ratios ranging from 1:1 to 1:20) can then be treated as a linear combination of the blue and red base spectra. These intervening spectra were thus fitted using an algorithm based on combinations of red and blue spectra giving the coefficient x (the ratio of bound



to unbound MBA): as expected x increases as the amount of MBA increases (Fig.S3). The data can then be analysed as follows to find n (the number of coordinated MBA's):

The equilibrium for MBA binding can be written as:

$$RR-[Cu(ligand)] + n R-MBA \leftrightarrow [RR-Cu(ligand)(R-MBA)_n]$$
 (eq.S1)

$$K = \frac{c([Cu(ligand)(MBA)_n])}{c([Cu(ligand)]) \cdot c(MBA)^n}$$
(eq.S2)

Where Cu(ligand) represents either Cu(1) or Cu(2). If *r* represents the Cu:MBA ratio and c<sub>0</sub> the initial concentration, then:

$$c_0(MBA) = r \cdot c_0([Cu(ligand)])$$
(eq.S3)

and

$$c([Cu(ligand)(MBA)_n]) = x \cdot c_0([Cu(ligand)])$$
(eq.S4)

$$c(MBA) = c_0(MBA) - n \cdot c([Cu(ligand)(MBA)_n])$$
  

$$c(MBA) = r \cdot c_0([Cu(ligand)]) - n \cdot x \cdot c_0([Cu(ligand)]) \qquad (eq.S5)$$
  

$$c(MBA) = (r - n \cdot x) \cdot c_0([Cu(ligand)])$$

while the amount of non-coordinated Cu(ligand) is given as:

$$c([Cu(ligand)]) = c_0([Cu(ligand)]) - c([Cu(ligand)(MBA)_n]) \quad (eq.S6)$$
$$c([Cu(ligand)]) = (1-x) \cdot c_0([Cu(ligand)])$$

Inserting eq.S5 and S6 into S2 gives:

$$K = \frac{x \cdot c_0([Cu(ligand)])}{(1-x) \cdot c_0([Cu(ligand)]) \cdot ((r-n \cdot x) \cdot c_0([Cu(ligand)]))^n}$$
(eq.S7)

$$K = \frac{x}{1-x} \cdot \left( \left( r - n \cdot x \right) \cdot c_0 \left( \left[ Cu(ligand) \right] \right) \right)^{-n}$$
(eq.S8)

$$\frac{x}{1-x} = K \cdot \left( \left( r - n \cdot x \right) \cdot c_0 \left( \left[ Cu(ligand) \right] \right) \right)^n$$
 (eq.S9)

$$\ln\left(\frac{x}{1-x}\right) = \ln(K) + n \cdot \ln(r - n \cdot x) + n \cdot \ln(c_0([Cu(ligand)]))$$
  

$$A = \ln(K) + n \cdot \ln(c_0([Cu(ligand)]))$$
  

$$\ln\left(\frac{x}{1-x}\right) = n \cdot \ln(r - n \cdot x) + A$$
 (eq.S10)

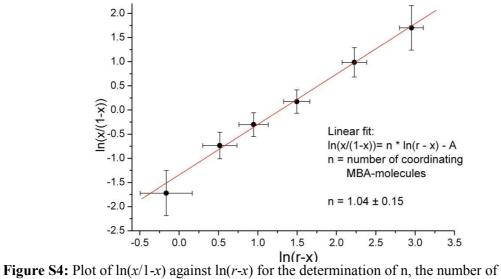
Since *r* ranges from 1 to 20 and is much larger than x (ranges from 0 to 1), the following simplification may be implemented:  $r \gg x$ 

$$n \in \{1; 2\}$$
(eq.S11)  
$$r - n \cdot x \approx r - x$$

Hence:

$$\ln\left(\frac{\mathbf{x}}{\mathbf{1}-\mathbf{x}}\right) = n \cdot \ln(r - \mathbf{x}) + \mathbf{A}$$
(eq.S11)

Thus a plot of  $\ln(x/1-x)$  against  $\ln(r-x)$  should give a straight line of slope *n*. The resulting graph is shown in Fig. S4 with  $n = 1.04 \pm 0.15$ .

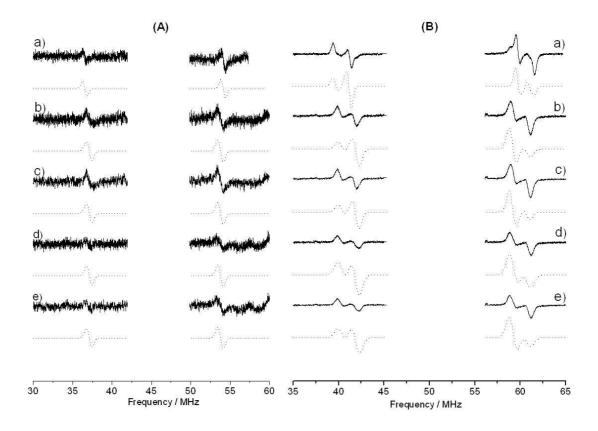


**Figure S4:** Plot of  $\ln(x/1-x)$  against  $\ln(r-x)$  for the determination of n, the number of coordinated MBA molecules in the [Cu(1)] complex.

Thus we can conclude that only 1 MBA molecule coordinates to [Cu(1)]. A similar treatment was also performed for [Cu(2)], where n = 1 was also found. These results are expected for such as bulky square planar Cu-salen complex interacting with a bulky MBA substrate.

## (4) Supporting <sup>1</sup>H and <sup>14</sup>N ENDOR spectra

**Figure S5**; Q-band <sup>1</sup>H CW-ENDOR spectra (recorded at 10K) showing the H<sup>iminc</sup> couplings of (a) *rac*-[Cu(1)] dissolved in d<sup>8</sup>-toluene/d<sup>6</sup>-dichloromethane, and (b) *SS-R*, (c) *RR-S*, (d) *SS-S* and (e) *RR-R*. Spectra recorded at  $g = g_{\parallel} = g_3$  and  $g = g_{\perp} = (g_1 + g_2)/2$ . For clarity, the central part of the spectra, containing proton couplings from the remaining ligand nuclei and coordinated MBA, have been removed.



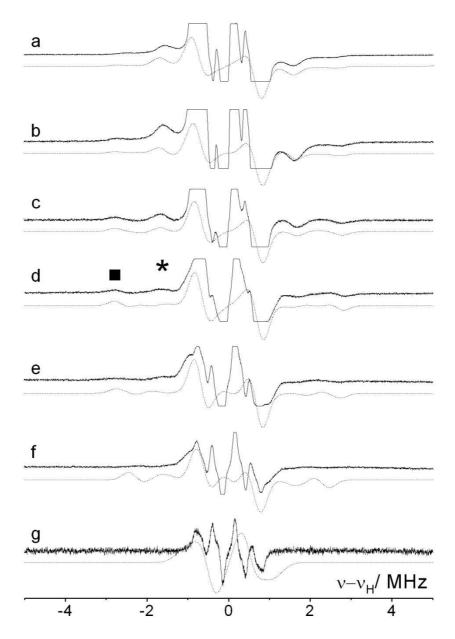
The equivalent series of spectra for the [Cu(2)] complex were given in Fig.4 of the main paper.

Proton	$A_1$ /	$A_2$ /	$A_{3}$ /	$a_{\rm iso}$ /	$^{(c)} heta_{ m H}$ / °	A <sub>dipolar</sub> /	R / Å	
	MHz	MHz	MHz	MHz		MHz		
Single Crystal <sup>(41)</sup>							ENDOR	X-ray
<sup>(a)</sup> [Cu(Salen)] - H <sup>imine</sup>	18.43	19.38	22.80	20.22	90		3.97	3.78
	18.64	19.45	23.62	20.57			3.76	3.73
<sup>(a)</sup> [Cu(Salen)] - H <sup>methine</sup>	-1.43	-1.43	5.54	0.90	90		3.30	3.19
	-3.47	-0.90	5.13	0.76			3.37	3.36
<sup>(b)</sup> tol/dcm								
$[Cu(1)]$ - $H^{\text{methine}}$	-1.41	-1.43	5.54	0.91	50	4.63	3.30	
$[Cu(1)] - H^{tert-butyl}$	-1.64	-1.50	3.30	0.05	70	3.25	3.69	
$[Cu(1)] - H^{imine}$	18.5	17.2	22.1	19.27	90	2.83	3.84	
[Cu(2)] - H <sup>imine</sup>	18.4	17.7	22.45	19.52	90	2.93	3.80	
<sup>(b)</sup> MBA								
$[Cu(1)] - H^{imine}$	17.2	16.6	21.0	18.26	90	2.73	3.90	
$[Cu(2)] - H_a^{imine}$	16.6	17.2	21.3	18.37	90	2.93	3.81	
- $H_b^{imine}$	16.6	16.1	21.3	18.00	90	3.30	3.66	
[Cu(1)] - H <sup>NH2</sup>	-2.52	-4.87	3.62	-1.26	10	4.88	3.22	
$[Cu(2)] - H^{NH2}$	-2.62	-5.2	3.62	-1.40	10	5.02	3.18	

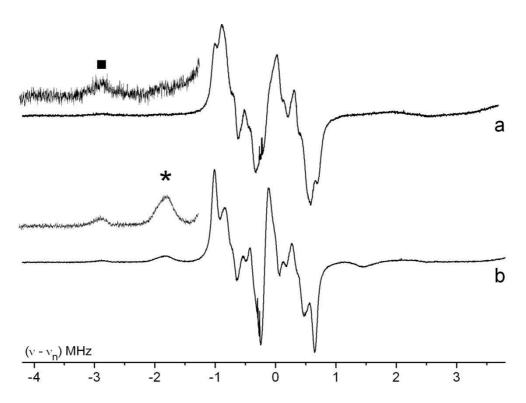
**Table 1;** <sup>1</sup>H principal hyperfine values for R, R-[Cu(1)] and R, R-[Cu(2)] dissolved in toluene/dichloromethane or MBA. For comparison the ENDOR data for the structurally related [Cu(Salen)] complex<sup>(42)</sup> is also shown.

<sup>(a)</sup>Two crystallographically distinct sites identified in the single crystal ENDOR spectrum of [Cu(Salen)]. <sup>(b)</sup>The frozen solution ENDOR spectra of [Cu(**1**,**2**)] were recorded either using toluene/dicholomethane or neat MBA. <sup>(c)</sup> $\theta_{\rm H}$  is defined as the angle between  $g_z$  and  $A_3$ .

**Figure S6**; Q-band <sup>1</sup>H CW-ENDOR spectra (10K) of *rac*-[Cu(1)] dissolved in d<sup>8</sup>-toluene/d<sup>6</sup>-dichloromethane, recorded at (a) 1204.6, (b) 1197.7, (c) 1186.2, (d) 1165.7, (e) 1140.5, (f) 1117.7 and (g) 1086.8 mT. (experimental = solid, simulated = dashed line). Only the strongly coupled protons from the methine and *tert*-butyl groups are shown in the simulations (the remaining weak/remote ligand protons responsible for the inner intense peaks were not simulated, as the error in their assignment is significant). \* = *tert*-butyl peak,  $\blacksquare$  = methine peak.

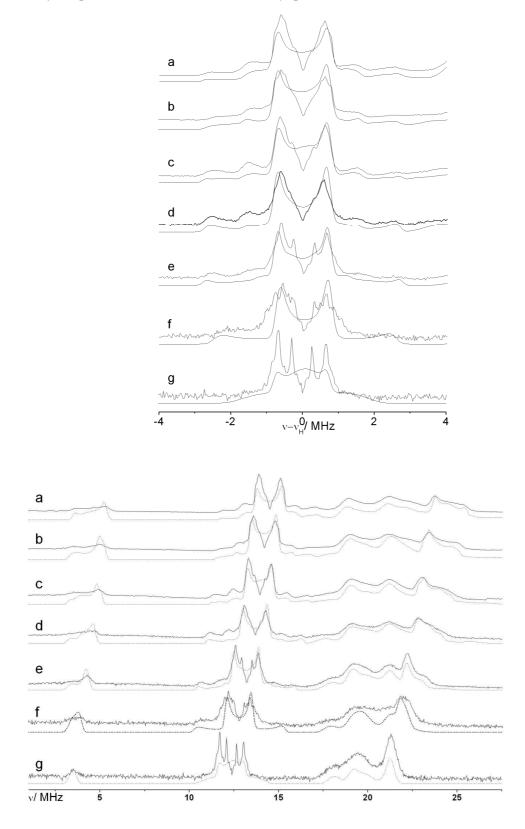


**Figure S7**; X-band <sup>1</sup>H CW-ENDOR spectra (10K) of (a) *rac*-[Cu(1)] and (b) *rac*-[Cu(2)] dissolved in d<sup>8</sup>-toluene/d<sup>6</sup>-dichloromethane, recorded at the field position corresponding to g =  $g_{\parallel}$ . The absence of the *tert*-butyl peak (\*) in [Cu(2)] is very obvious compared to [Cu(1)]. The methine protons are present in both samples ( $\blacksquare$ ).

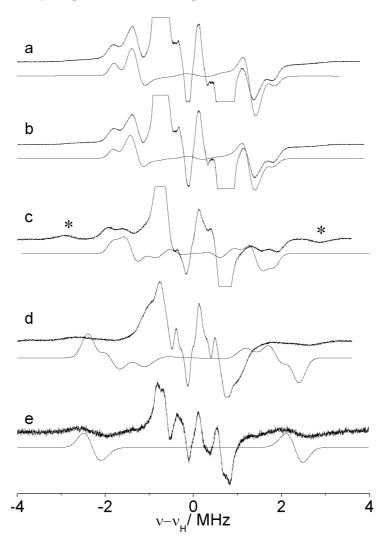


These spectra were used in the identification and assignment of the *tert*-butyl groups in [Cu(1)].

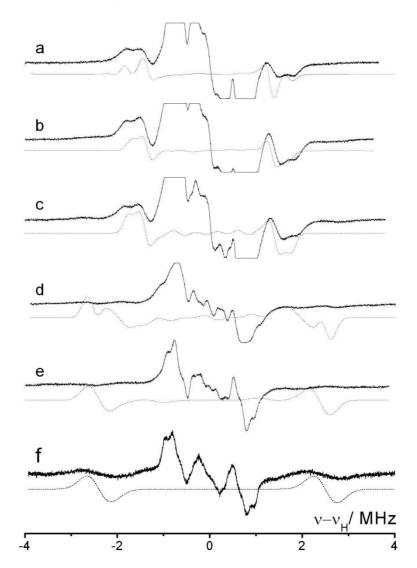
**Figure S8**: X-band Davies ENDOR spectra (recorded at 10K) of *rac*-[Cu(1)] dissolved in  $d^8$ -toluene/ $d^6$ -dichloromethane, recorded at the field positions (a) 340.0, (b) 333.0, (c) 327.0, (d) 320.6, (e) 310.0, (f) 300.0 and (g) 290.0 mT (experimental = solid, simulated = dashed line). Only the peaks for the methine and *tert*-butyl protons are included in the simulations.



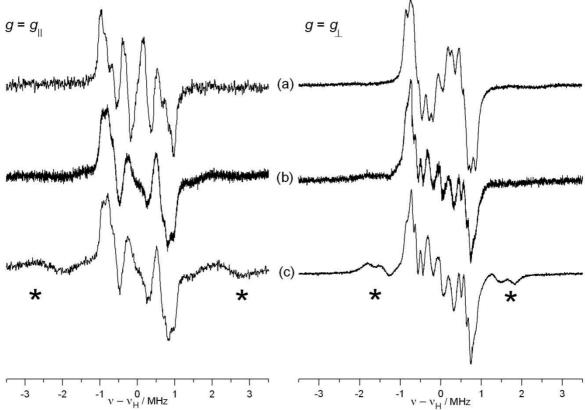
**Figure S9**: Q-band CW <sup>1</sup>H ENDOR spectra of *rac*-[Cu(1)] dissolved in *rac*-MBA recorded at the field positions of (a) 1191.0 (b) 1188.8 (c) 1171.7 (d) 1125.0 and (e) 1069.0 mT (experimental = solid, simulated = dashed line). The peaks labelled \* are due to the ligand methine protons. Only the peaks for the amine protons are included in the simulations.



**Figure S10**; Q-band CW-<sup>1</sup>H ENDOR spectra of *rac*-[Cu(**2**)] dissolved in *rac*-MBA recorded at (a) 1201.1, (b) 1191.9, (c) 1186.2, (d) 1142.8, (e) 1101.6 and (f) 1073.1 mT. (experimental = solid, simulated = dashed line).

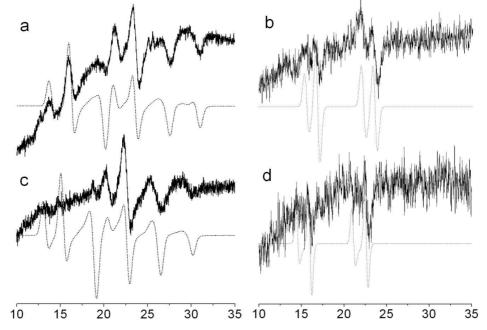


**Figure S11**; Q-band <sup>1</sup>H CW-ENDOR spectra (recorded at 10K) of *rac*-[Cu(1)] dissolved in d<sup>8</sup>-toluene/d<sup>6</sup>-dichloromethane, (b) *rac*-[Cu(1)] in *racemic* d<sup>2</sup>-MBA, and (c) *rac*-[Cu(1)] in *rac*-MBA (protic). Spectra recorded at  $g = g_{\parallel} = g_3$  and  $g = g_{\perp} = (g_1+g_2)/2$ .

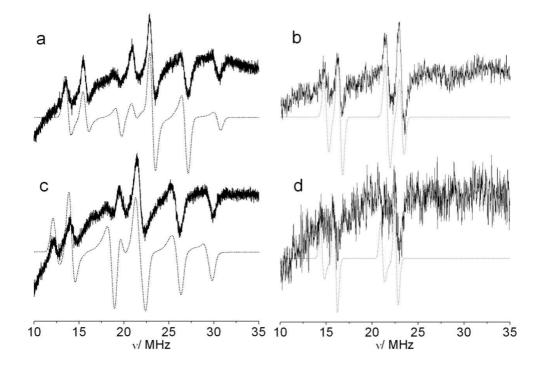


The  $-NH_2$  protons from the coordinated MBA are labelled \* in the Figure. These are clearly visible in (c), obtained using protic MBA, but largely absent in (b), recorded using d<sup>2</sup> deuterated-MBA. (Note; deuteration of MBA by D<sub>2</sub>O does not give 100% exchange, hence the presence of residual MBA-NH<sub>2</sub> proton peaks in (b).

**Figure S12**; Q-band <sup>14</sup>N CW-ENDOR spectra of [Cu(1)] dissolved in (a,b) toluene and (c,d) MBA respectively. The spectra were recorded at the field positions (a,c)  $g = g_{\perp}$  and (b,d)  $g = g_{\parallel}$ . Solid lines = experimental, dashed lines = simulated.



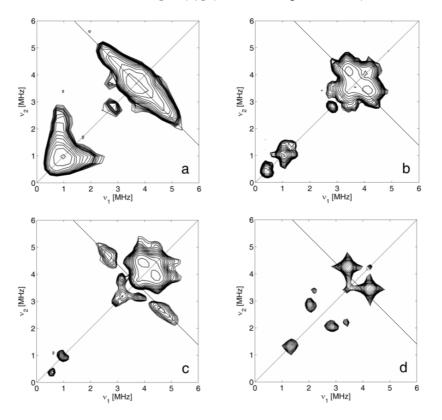
**Figure S13**: Q-band <sup>14</sup>N CW ENDOR spectra of [Cu(2)] dissolved in (a,b) toluene and (c,d) MBA respectively. The spectra were recorded at the field positions (a,c)  $g = g_{\perp}$  and (b,d)  $g = g_{\parallel}$ . Solid lines = experimental, dashed lines = simulated.



NOTE: simulations assuming the contributions of both nitrogen nuclei (with the hyperfine tensor in-plane rotated over 90°) are identical to those assuming only one nitrogen, because of the near axiality of the g tensor.

## (5) HYSCORE spectra

**Figure S14:** X-band HYSCORE spectra of *S*,*S*-[Cu(1)] dissolved in (a) toluene, (b) *R*-MBA and (c) pyridine, recorded at 15K around the  $g = g_{\perp}$  field position (340 mT) with  $\tau = 96$ ns and  $\tau = 176$  ns. (d) Simulation of spectrum (b) showing the weak interaction of the amine <sup>14</sup>N with [Cu(1)] (see text for parameters).

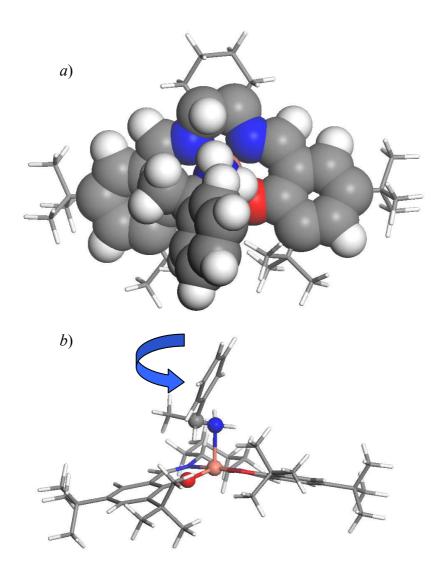


This Figure (a) shows the low-frequency region of the HYSCORE spectrum of [Cu(1)] in toluene, recorded at  $g = g_{\perp}$ . In the low-field area of the spectrum, a ridge due to interactions with the  ${}^{13}C$  nuclei of ligand (1) is observed ( ${}^{13}C$  in natural abundance). The corresponding HYSCORE spectrum of [Cu(1)] in MBA is shown in (b). Two distinct peaks at ~(3.4, 4.2) MHz are observed, centred around the  $^{13}C$ Larmor frequency and could thus stem from nearby <sup>13</sup>C nuclei in altered arrangement. However, in principle, an appropriate combination of <sup>14</sup>N hyperfine and nuclear quadrupole couplings could also lead to double-quantum cross-peaks at this position. To test this, the HYSCORE spectrum of [Cu(1)] dissolved in pyridine (Py) was recorded. The sp<sup>2</sup>-hybridized nitrogen of pyridine is expected to have different nuclear quadrupole values compared to the sp<sup>3</sup>-hybridized nitrogen of MBA. The resulting HYSCORE spectrum of [Cu(1)] in pyridine is given in (c). A clear shift of the double-quantum cross peaks, consistent with a larger nuclear quadrupole coupling of the <sup>14</sup>N nucleus, can be seen. The <sup>13</sup>C ridge found in (a) is now also visible, where earlier it was masked by the intense cross peaks in (b). The HYSCORE spectrum of [Cu(1)]+Py therefore proved that the cross peaks observed for [Cu(1)]+MBA are indeed due to the weak interaction with the amine nitrogen. The simulated parameters are given in Table 3.

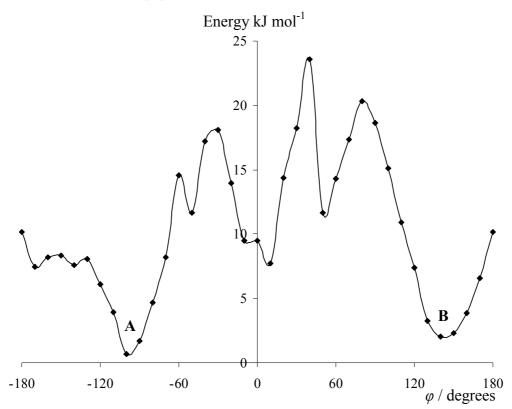
## (6) Additional DFT details

**Figure S15**: The starting model of [Cu(1)]+MBA used in QMMM calculations. *a*) Space fill atoms at QM level (BHandH), stick atoms at MM level (UFF). *b*) The torsion angle,  $\varphi$ , used to scan the orientation of MBA relative to the [Cu(1)] complex; this angle is defined with respect to  $O(Salen) \cdots Cu \cdots N(MBA \cdots C^*(MBA))$  where the defining atoms are shown as spheres in b).

Thus an angle of  $\varphi = 0^\circ$  is shown in b) where the  $\mathbb{C}^*$  (of MBA) atom is above the O(of Salen) atom. For clarity, the MBA-phenyl ring is shown oriented away from the complex. Clockwise turn is represented as -ve degrees and anticlockwise as +ve degrees.



**Figure S16**: The relative energy *vs* torsion angle,  $\varphi$  (see Figure S15*b* for definition and value of  $\varphi = 0^{\circ}$ ) for *R*-MBA with *R*,*R*-[Cu(1)]. For this scan of the torsion angle Cu···N(MBA) is constrained to 2.2 Å and each point in the plot represents an optimisation with the additional constraint of torsion angle value. Structures from minima **A** and **B** were further optimised without constraints to produce the structures given in Figure 5 (main paper).



Full author details for reference (32) are:

[32] Gaussian 03, Revision C.02, Frisch, M.J.; Trucks, G.W.; Schlegel, H.B.; Scuseria, G.E.;
Robb, M.A.; Cheeseman, J.R.; Montgomery, Jr. J.A.; Vreven, T.; Kudin, K.N.; Burant, J.C.;
Millam, J.M.; Iyengar, S.S.; Tomasi, T.; Barone, V.; Mennucci, B.; Cossi, M.; Scalmani, G.;
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P.Y.; Morokuma, K.; Koth, J.; Salvador, P.; Dannenberg, J.J.; Zakrzewski, V.G.; Dapprich,
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Chen, W.; Wong, M.W.; Gonzalez, C. and Pople, J.A.; Gaussian, Inc., Wallingford CT, 2004.