

Supplementary Information for:

**Self-assembly of Zn(salphen) complexes: Steric regulation,
stability studies and crystallographic analysis revealing an
unexpected dimeric 3,3'-*t*-Bu-substituted Zn(salphen)
complex**

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Crystallographic comments

Complex 2:

The unit cell of this compound contains a CH₃CN molecule and one of the benzene rings at the main molecule is showing some disorder. The completeness of the data is 95 %. The structure has an R1 value of 4.0 %.

Complex 5:

The unit cell contains two half independent molecules of the complex. The unit cell contains also three independent molecules of CH₃CN. Some of the tert-butyl groups are disordered. The completeness of the data is 91 %. The structure has an R1 value of 4.42%.

Complex 6:

The unit cell of this compound contains one molecule of CH₃CN and non relevant disorder was observed. The completeness of the data is 91 %. The structure has an R1 value of 2.67 %.

Complex 9:

The structure contains three molecules of CH₃CN and one of the tert-butyl groups is disordered in two positions. The structure has an R1 value of 8.28 %.

Complex 11:

The structure is a solvate of 2,6-lutidine (2,6-dimethylpyridine). The 2,6-lutidine molecule is disordered in two overlaid positions (50:50) which are related through a C2-operation. The structure has an R1 value of 4.76 %.

The residual electron density near the Zn atom is a residual artifact due to the presence of a second crystal that was not enough intense to perform a combined data reduction and absorption. This is also shown in the unusually large second term of the weighting scheme: 5.9979

Complex 18:

The unit cell does not contain solvent molecules and only diminutive disorder was observed. The completeness of the data is of 94 %. This structure has an R1 value of 3.22 %.

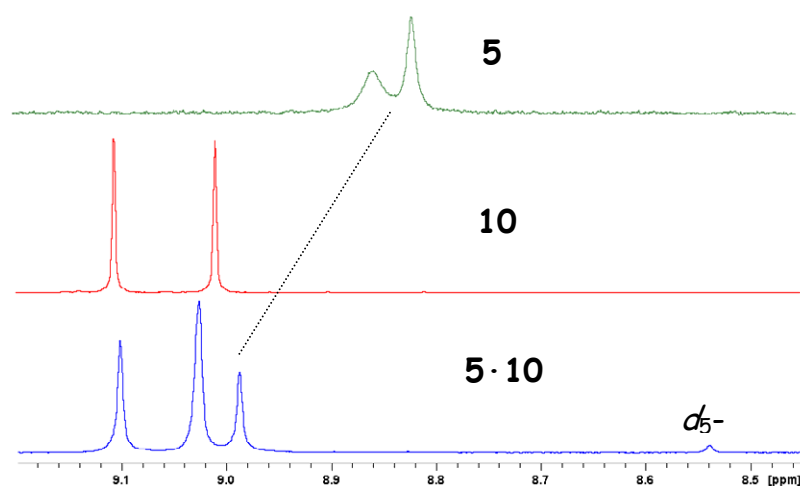
Complex 19:

The unit cell of this compound contains one molecule of CH₃CN and non relevant disorder was observed. The completeness of the data is 87 %. The crystal measured was a twin (crystal ratio of 51:49) and for the absorption correction TWINABS was used

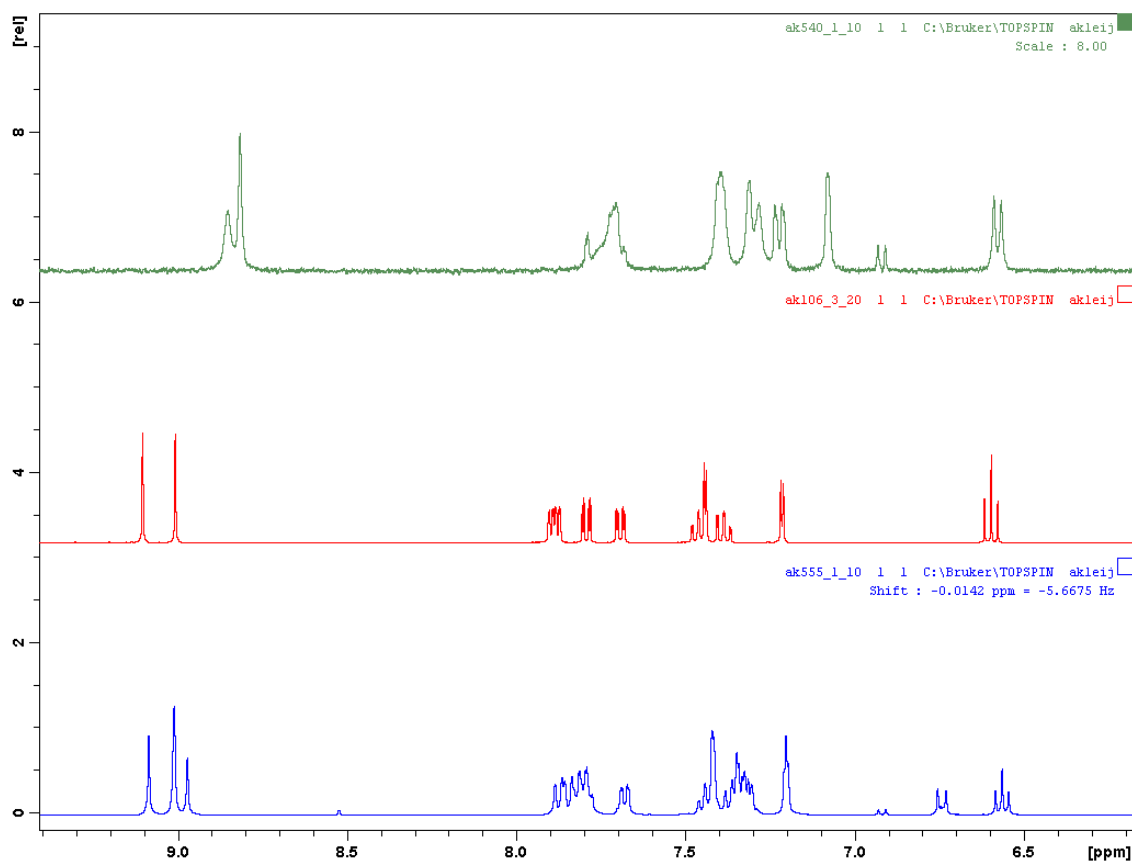
(TWINABS Version 2008/4 Bruker AXS; Blessing, Acta Cryst. (1995) A51 33-38).
The structure has a R1 value of 5.7 %.

^1H NMR comparison between dimers (**5**)₂ and (**10**)₂ and hetero-dimeric assembly **5**·**10**

Imine region of the ^1H NMR (d_6 -acetone) spectra of the respective species:

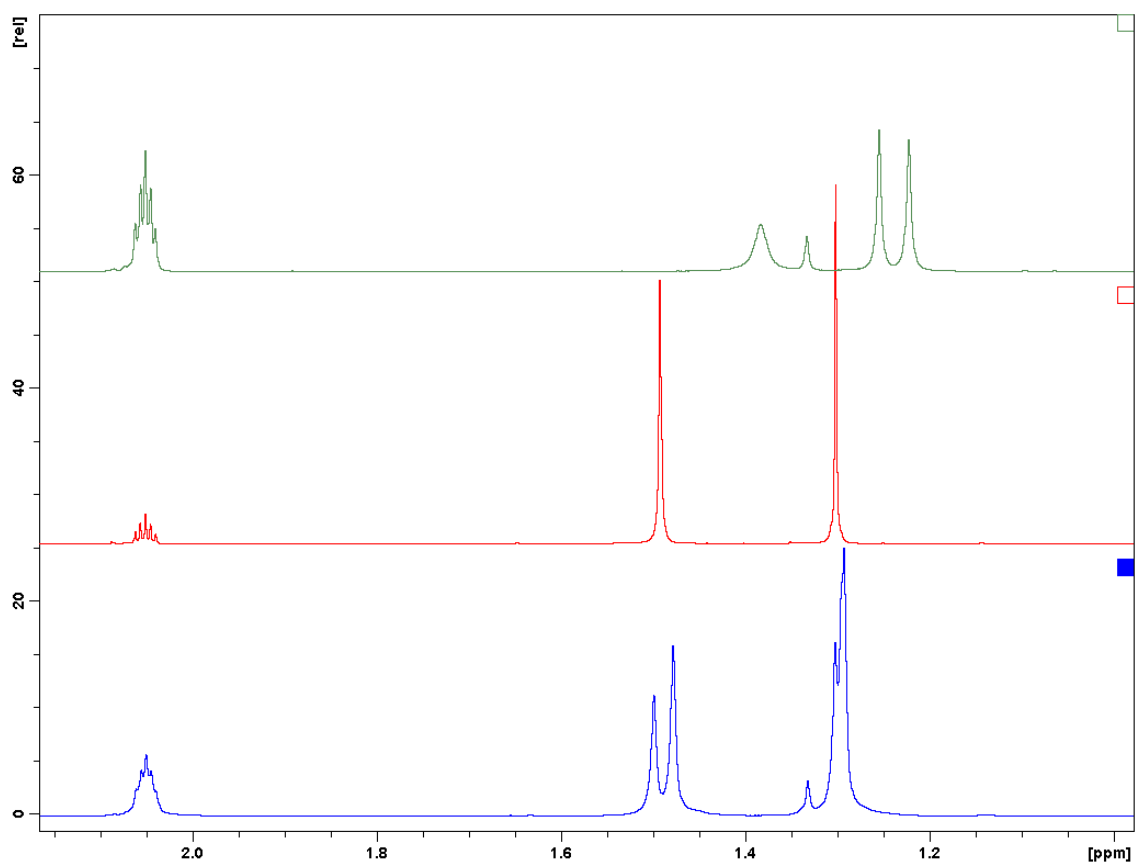


Full aromatic region:



In **GREEN**: complex **5**, in **RED**: complex **10**, and in **BLUE** hetero-complex **5**·**10**.

t-Bu region:



In **GREEN**: complex 5, in **RED**: complex 10, and in **BLUE** hetero-complex 5·10.

MALDI-TOF mass spectrometric analysis of **5·10**.

Calcd masses (molecular ions) of the DIMERS:

See: <http://winter.group.shef.ac.uk/chemputer/isotopes.html>

For **(5)₂**: $m/z = 1096$ (100%)

For **(10)₂**: $m/z = 1074$ (100%)

For **5·10**: $m/z = 1085$ (100%)

