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- State of the art:
 - \rightarrow role of the thioether in the active site is still unclear
 - \rightarrow Investigation of ligands with N₂S and N₃S systems to mimic the active site of PHM^[4,5]

See the Peptidylglycine α hydroxylating monooxygenase in VR





You can see the active site of PHM also on YouTube here!

copper(I) complexes with all ligands:

$Lx + Cu(CH_3CH)_4CF_3SO_3 \longrightarrow [CuLx(CH_3CN)_x]CF_3SO_3 + x CH_3CN$

- ligands are dissolved in acetonitril
- copper(I)-salts ($CF_3SO_3^-$, PF_6^- or $[BAr_4^-]^-$) are added in solution
- solvent is removed \rightarrow light green solid







LCu^OCuL



UV/VIS



Ligands L2, L7 + O_2 :

band at 389 nm

 \implies bis (μ -oxo)dicopper(III) complex





Ligands L1, L4 + O_2 :

band at 525 nm and 600 nm

trans-1,2 peroxodicopper(II) complex





Summary

The tripodal ligands (L2, L7) form bis (μ -oxo)dicopper complexes as predicted from literature^[4]. Two of the tetrapodal ligands with thioether ligand system (L1, L4) show *trans-1,2* peroxodicopper complexes. The ligand L3 (tetrapodal ligand) is able to form both copper complexes.



Crystal structure of a copper complex with L2



JUSTUS-LIEBIG-



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