Meeting 1: Enumeration

Stefan Gugler

Massachusetts Institute of Technology sgugler@mit.edu

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Mono-Heavy-Atomic Isoelectronic Ligand Distribution

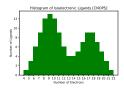


Figure: CNOPS: 15625 ligands

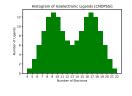


Figure: CNOPSSi: 22500 (?) ligands

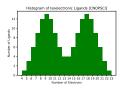


Figure: CNOPSCI: 22500 (?) ligands

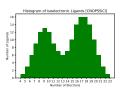


Figure: CNOPSSiCI: 30625 ligands

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Mono-Heavy-Atomic Iso-VE Ligand Distribution

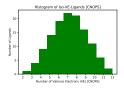


Figure: CNOPS: 15625 ligands

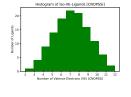


Figure: CNOPSSi: 22500 (?) ligands

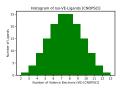


Figure: CNOPSCI: 22500 (?) ligands

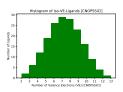


Figure: CNOPSSiCI: 30625 ligands

Now we go to Di-Heavy-Atomic Ligands ...

- Exhaustive combinatorial combination from the Mono-Heavy-Atomic Ligands to get Di-Heavy-Ligands
- Over heavy atoms identities $a_i \in \{C, N, O, P, S, (Si, Cl)\}$, charges $c_i \in \{-2, +2\}$, number of H atoms $h_i \in \{0, 4\}$, with heavy atoms $i \in 1, 2$, where i = 1 is the coordinating atom.
- Pruned by means of a health function, u, of ligand j, to encoding heuristics.

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Now we go to Di-Heavy-Atomic Ligands ...

• Only ligands with $c_1 \leq 0 \land c_2 \leq 0$ are considered.

$$u_{\text{octet,i}} = \begin{cases} 10 + 2 \cdot (8 - VE_i) & \text{if } 8 - VE_i < 0\\ 10 - 1 \cdot (8 - VE_i) & \text{if } 8 - VE_i \ge 0 \end{cases}$$
(1)

$$u_{\text{charge}} = \begin{cases} 0 & \text{if } c_1 + c_2 > 0 \\ 3 & \text{if } 0 \ge c_1 + c_2 \ge -2 \\ 1 & \text{if } c_1 + c_2 = -3 \\ 0 & \text{if } c_1 + c_2 = -4 \end{cases}$$
 (2)

$$u_{\rm VSEPR} = 5 - |\underbrace{(VE_1 - 2 \cdot LP_1 + c_1 - 2 \cdot h_1)}_{ready\ electrons} - \underbrace{(VE_2 - 2 \cdot LP_2 + c_2 - 2 \cdot h_2)}_{ready\ electrons}|$$

(3)

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Now we go to Di-Heavy-Atomic Ligands ...

$$u_{\rm CA} = \begin{cases} 1 & \text{if } h_1 = 4 \\ 2 & \text{if } h_1 = 3 \\ 3 & \text{if else} \end{cases}$$
 (4)

$$u_{\text{shell}} = \begin{cases} 1 & \text{if } \neg (VE_1 + VE_2)\%2\\ 0 & \text{if else} \end{cases}$$
 (5)

$$u_{\text{total}} = u_{\text{shell}} \cdot \left(\frac{1}{2}(u_{\text{octet},1} + u_{\text{octet},i}) + u_{\text{charge}} + u_{\text{VSEPR}} + u_{\text{CA}}\right)$$
 (6)

where VE_i denotes the number of valence electrons, LP_i the number of lone pairs.

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Distribution of the Score

Comparison of edge cases. Mayer B.O. with multiwfn.

1	2	Mayer B.O.	score	comment
СНННН++	CHHHH++	< 0.05	12	rather high
СНННН–	CHHHH–	0.928	-	$c_{tot} > 0$
CHHH	NH	1.308	0	open shell
CH4++	-	0.577	-	Mono-Heavy-Atom
C-	0+	-	16	dummy

Distribution of the Score

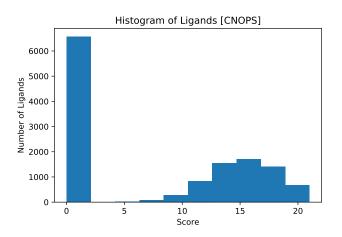


Figure: A histogram describing the score distribution. There are 30 ligands with score 21 and 180 ligands with score 20. CO is has a score of 16. There are 2917 ligands at least as good as CO.

Code Output

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 \begin{bmatrix} [\lceil [NHH-] \rceil, 10, 8, 1, -1, 2, 0, 1, 10, \rceil [CHHH-] \rceil, 10, 8, 0, -1, 3, 0, 1, 10], 21.0] \\ [\lceil [NHH-], 10, 8, 1, -1, 2, 0, 1, 10, \rceil [NHH-] \rceil, 10, 8, 1, -1, 2, 0, 1, 10], 21.0] \\ [\lceil [NHH-], 10, 8, 1, -1, 2, 0, 1, 10, \rceil [NHH-] \rceil, 10, 8, 2, -1, 1, 0, 1, 10], 21.0] \\ [\lceil [NHH-], 10, 8, 1, -1, 2, 0, 1, 10, \rceil [PHH-] \rceil, 18, 8, 1, -1, 2, 0, 1, 10], 21.0] \\ [\lceil [NHH-], 10, 8, 1, -1, 2, 0, 1, 10, \rceil [SH-] \rceil, 18, 8, 2, -1, 1, 0, 1, 10], 21.0] \\ [\lceil [OH-], 10, 8, 2, -1, 1, 0, 1, 10, \rceil [CHHH-] \rceil, 10, 8, 0, -1, 3, 0, 1, 10], 21.0] \\ [\lceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, \rceil [CHHH-] \rceil, 10, 8, 0, -1, 3, 0, 1, 10], 21.0] \\ [\lceil [OH-], 10, 8, 2, -1, 1, 0, 1, 10, \rceil [NHH-] \rceil, 10, 8, 1, -1, 2, 0, 1, 10], 21.0] \\ [\lceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, \rceil [NHH-] \rceil, 10, 8, 1, -1, 2, 0, 1, 10], 21.0] \\ [\lceil [OH-], 10, 8, 2, -1, 1, 0, 1, 10, \rceil [NHH-] \rceil, 10, 8, 2, -1, 1, 0, 1, 10], 21.0] \\ [\lceil [OH-], 10, 8, 2, -1, 1, 0, 1, 10, \rceil [NHH-] \rceil, 10, 8, 2, -1, 1, 0, 1, 10], 21.0] \\ [\lceil [OH-], 10, 8, 2, -1, 1, 0, 1, 10, \rceil [OHH] \rceil, 10, 8, 2, -1, 1, 0, 1, 10], 21.0] \\ [\lceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, \rceil [OHH] \rceil, 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, \rceil [OHH] \rceil, 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, \rceil [OHH] \rceil, 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, [OHH]], 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, [OHH]], 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, [OHH]], 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, [OHH]], 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, [OHH]], 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, [OHH]], 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, [OHH]], 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, [OHH]], 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, [OHH]], 10, 8, 2, 0, 2, 0, 1, 10], 21.0] \\ [\rceil [OH-], 10, 8, 2, 0, 2, 0, 1, 10, [OHH]], 10, 10, 2, 2,
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