The Valence Multipole Model: Using small molecules to create a new bond energy expression for Molecular Mechanics



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Traditional Molecular Dynamics



+ $\sum_{i < j} \sum \frac{q_i q_j}{4\pi\varepsilon_0 r_{ij}}$ U + $\sum_{bonds} \frac{1}{2} k_b (r - r_0)^2 \qquad \longrightarrow \qquad r_0$ + $\sum_{angles} \frac{1}{2} k_a (\theta - \theta_0)^2$ + $\sum_{torsions} k_{\phi} [1 + \cos(n\phi - \delta)]$

Problems with MM

- There isn't really a single ideal bond length or bond angle value for a given bond type.
 - MM structures tend to be too symmetrical
 - Usually can't do coordination number changes
- The number of "adjustable parameters" in a MM model goes up AT LEAST proportionally to the number of atom types squared (often n³ or n⁴).
 - Especially ridiculous in models that allow chemical reactions
 - Exception: Rule-based models, which can do the entire periodic table, but generally suck.
- Models tuned to reproduce experimental data in a VERY NARROW range of chemical scenarios.



The Bond-Valence Model

Definitions:

- Atomic Valence (V_i): a measure of the number of valence electron states available for bonding.
 - Usually taken as the oxidation number
- Bond Valence (s_{ij}): the number of electron pairs involved in a given bond (by sharing and/or electron transfer).
 - Equivalent to "bond order", related to bond length
 - Does not have to be an integer!



Problems With the BVM

- 1. Only addresses bond lengths, not the angular distribution of bonds.
- 2. Since its calibrated for crystals molecules have had big misfits.
- 3. Only addresses polar bonds, not 100% covalent bonds.
- 4. Nobody has a clear idea how bond valence relates to energy.



Wander et al., The use of cation-cation and anion-anion bonds to augment the bond-valence model, Am. Min. 2015.



Johnson's Force Constant Model of Bonding



Valence Multipole Model

 Breaks energy into total bonding (S_{Total}): monopolar, no directionality, Vector (VVS): dipolar, noncentrosymmetric, quadrupole(QVS): centrosymmetric....



• Bickmore et al., Electronic structure effects in the vectorial bondvalence model, Am. Min. 2013

A More Flexible Equation

• One of a family of forms (exp-exp, pow-pow, geometric).

 $|s_{ij}| = (1 - w)(R/R_0)^{-1/B_{pow}} + we^{(R_0 - R)/B_{exp}}$

- Can flexibly and accurately fit anything from triple bonds to bonds as small as a few hundredths of a bond order.
- Suggests ionic-covalent character change around 1v.u.

• Wander et al., AIM analysis and the form of the bond-valence equation, Am. Min. 2015

Energy Expression

 Many body energy (atom centered) composed of only two body terms (Bond Valence Vectors).

$$E_{Total} = \frac{1}{2} V_I D_{E1} \left(\left(\frac{S_{Tot}}{V_I} \right)^B \sqrt{\frac{F_1}{2D_{E1}}} - 1 \right)^2 + k_{VVS} (VVS - VVS_{Ideal})^2 + E_{QV} + E_{Charge}$$

- Bond energy only definable in context of other bonds reaching an atom.
- VVS_{Ideal} depends on number and row number (electronegativities) of atoms surrounding it.

New challenges

 Quality bond energy per unit valence curves essential to getting high accuracy predictions.



AlSiHO Fitting Set Results

• Fitting Set Contains Both Molecules and Crystals



AlHSiO Check Set

- Some outliers: Unsatisfied Valences (Radicals), Bad Vector Sums (Poor H Positions).
- Otherwise errors less than 5kJ/mol/atom.
- Results oxidation state dependent. Reduced Ai Si different force field.





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