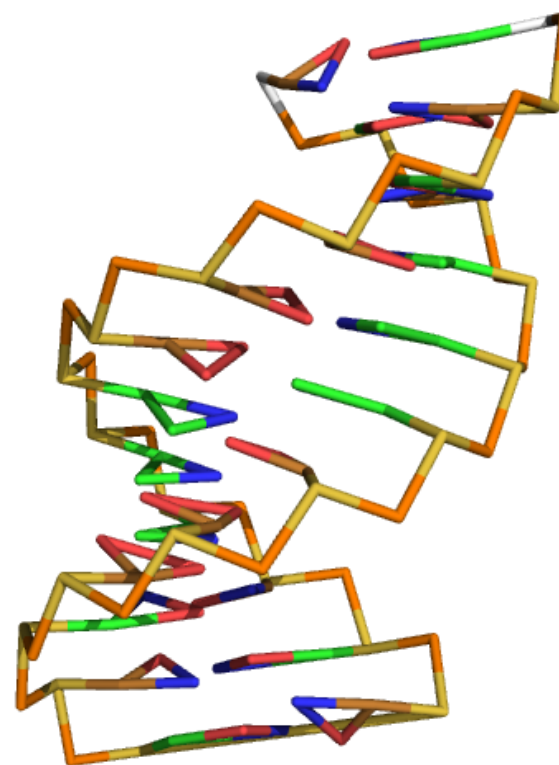
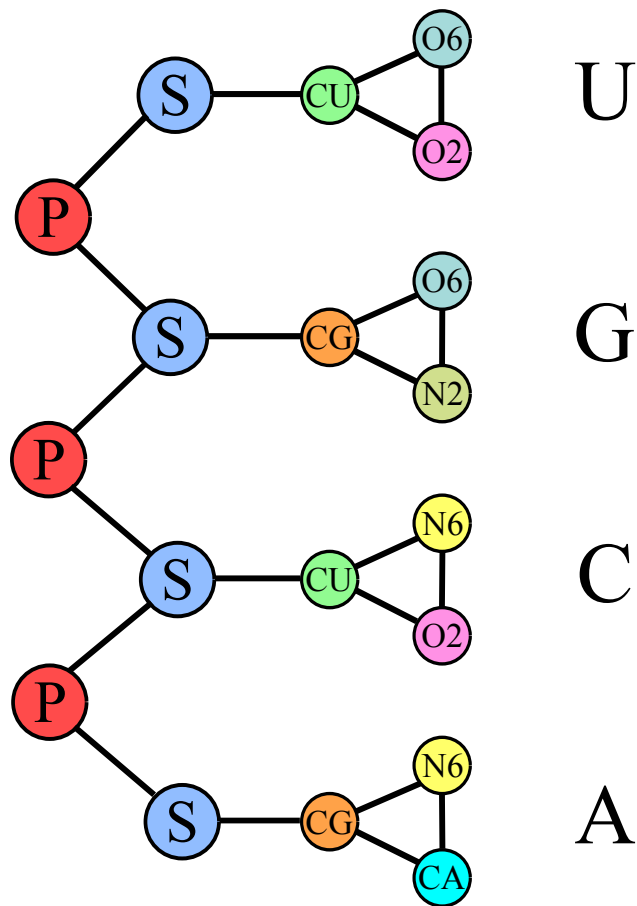


RNA Model Energy Optimization

11/14/14

Coarse-Grained Model



PDB ID: 1AL5

Force-Field

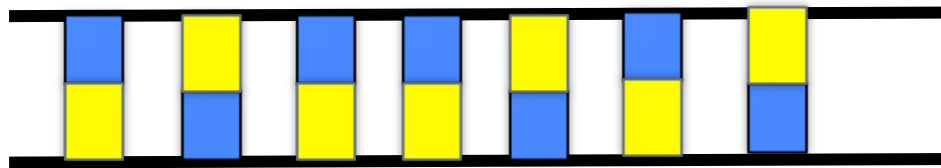
- Standard bond stretching, angle bending and torsion
- Debye-Hückel electrostatics and Buckingham vdW

$$E_{ele} = \sum_{i>j} \frac{q_i q_j}{4\pi D r_{ij}} e^{-\frac{r_{ij}}{\xi}}$$

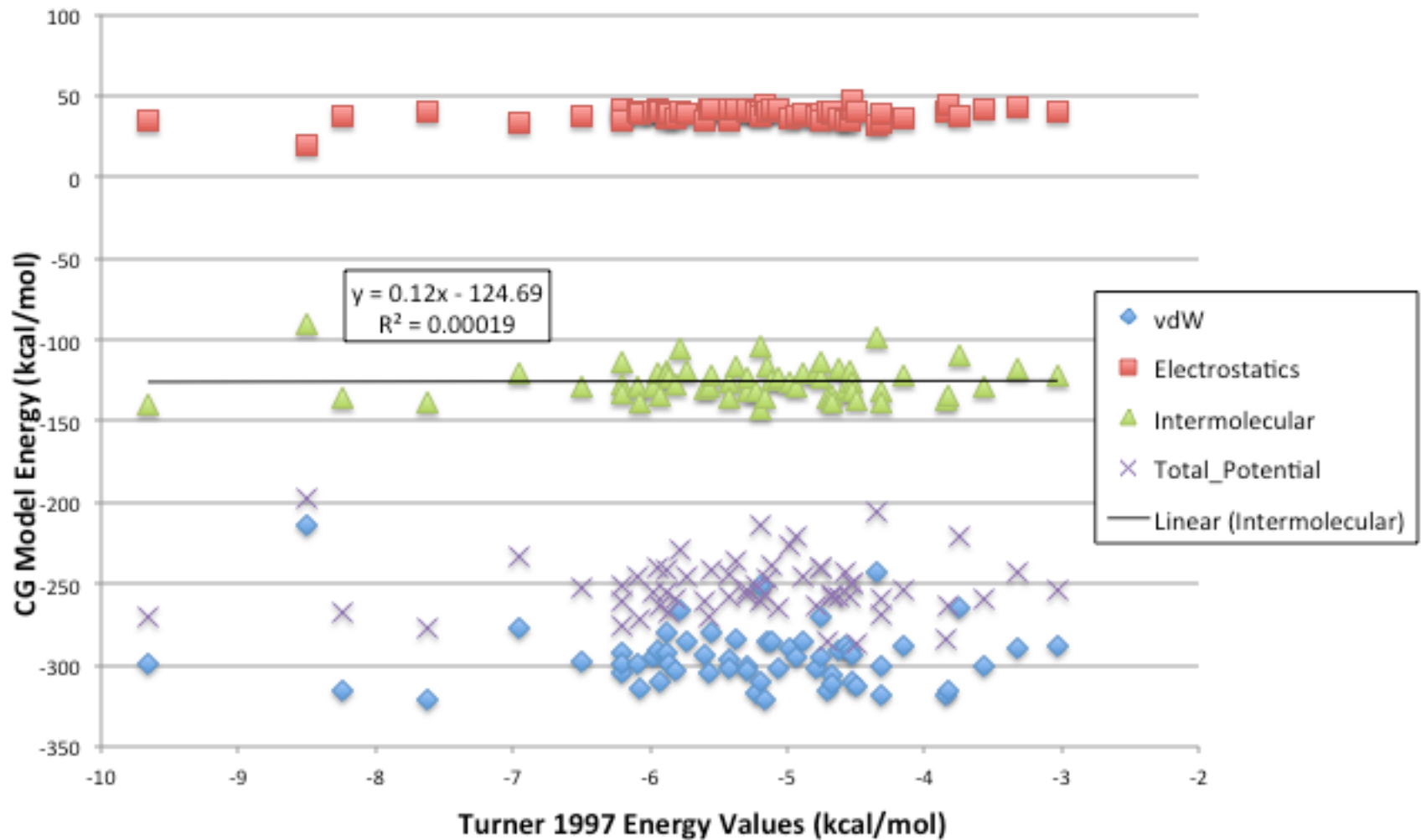
$$E_{vdW} = \sum_{i,j} \varepsilon_{ij} \left[-2.25 \left(\frac{\sigma_{ij}}{r_{ij}} \right)^6 + 1.84 \times 10^5 e^{-\left(\frac{12r_{ij}}{\sigma_{ij}} \right)} \right]$$

Optimization

- Original model parameterized to statistical potentials from 668 RNA structures (PDB, NDB, CRW)
- RNA secondary structure field optimizes to free energy
- Optimize our model to Turner's 1997 melting free energies for 60 duplexes

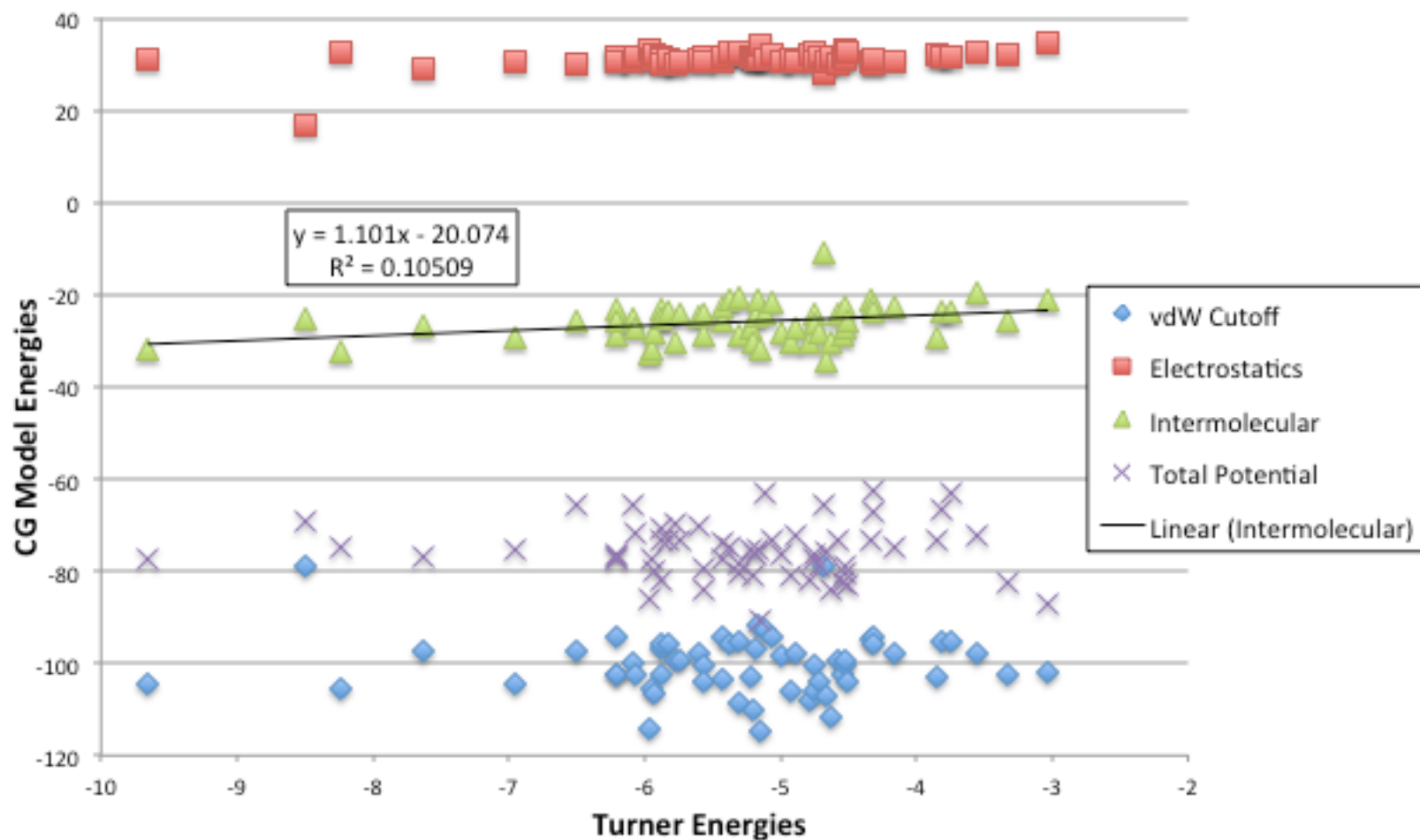


Helix Energy Correlation Initial Model



Interplay between vdW and electrostatics needs to be adjusted:
-Dielectric and vdW/vdWpr

Energy Correlation vdW Cutoff 5 Angstroms



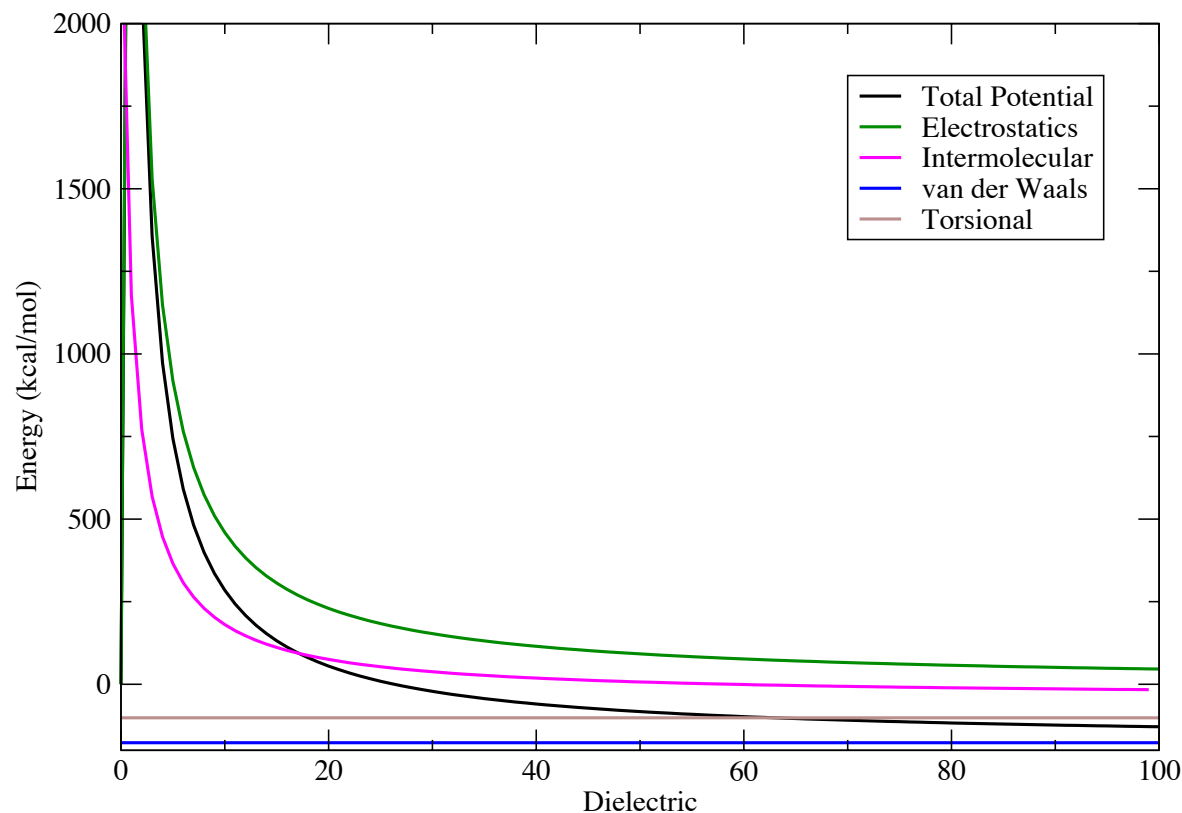
- Using modest vdW cutoff, vdW energy increases by 200 kcal/mol.
- Long-range interactions are captured from model
- Interplay between vdW and electrostatics needs to be adjusted: Dielectric and vdW/vdWpr

Dielectric Optimization

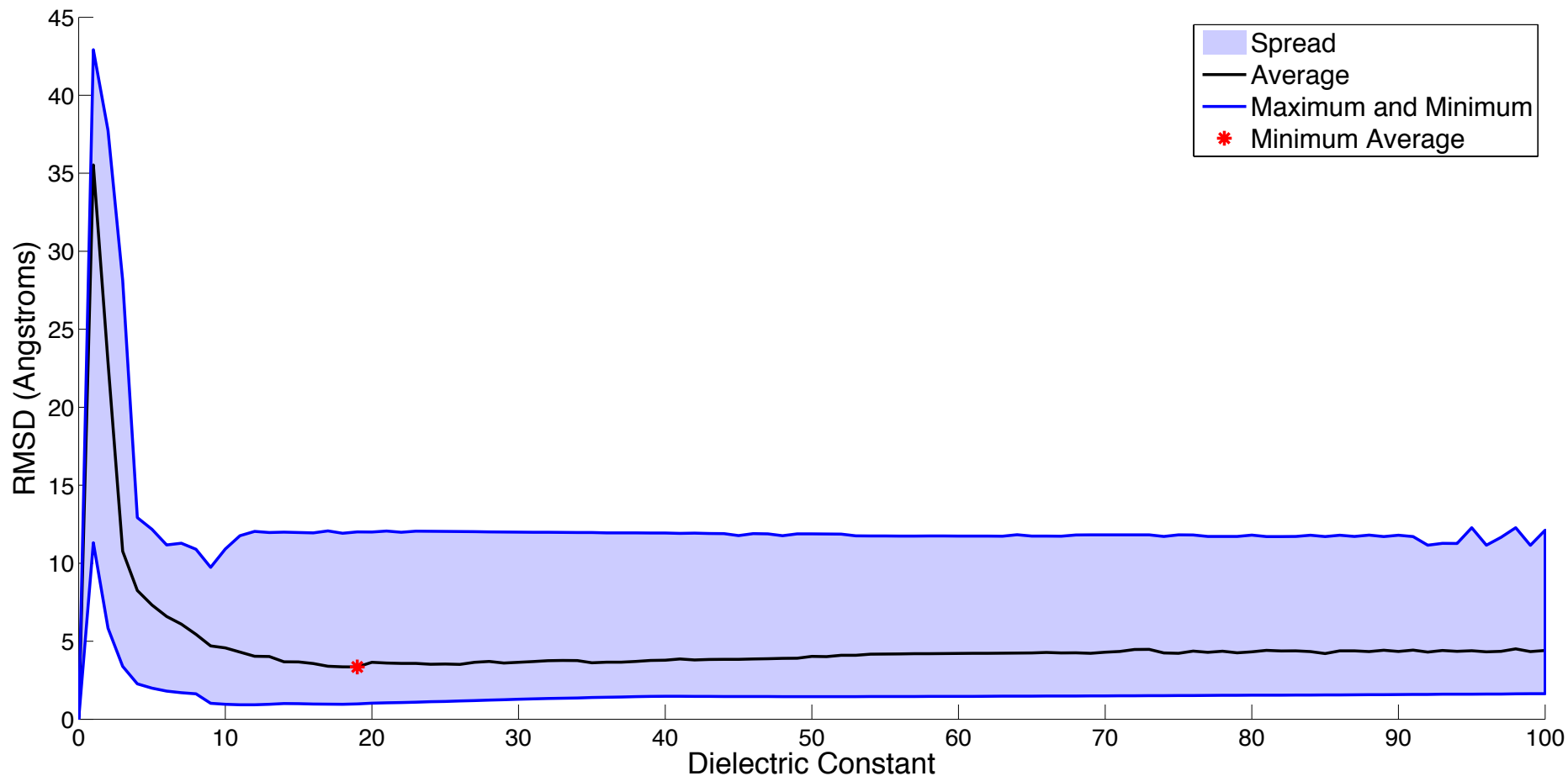
$$E_{ele} = \sum_{i>j} \frac{q_i q_j}{4\pi D r_{ij}} e^{-\frac{r_{ij}}{\xi}}$$

Previous Value: 78

Energy Composition as a function of Dielectric 10_23_14



RMSD Over Dielectric Values



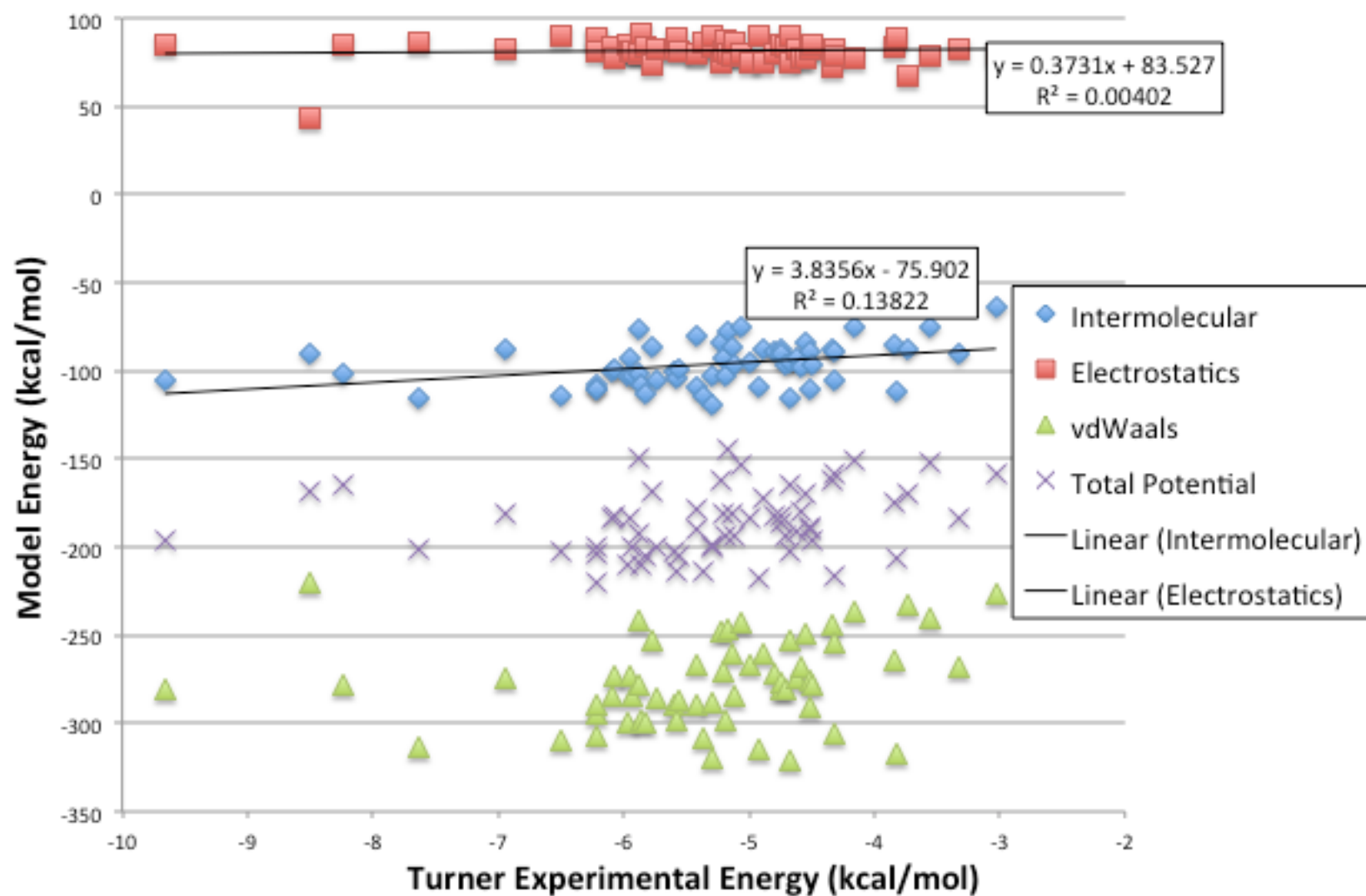
Electrostatics Interaction:

$q = -1.0$

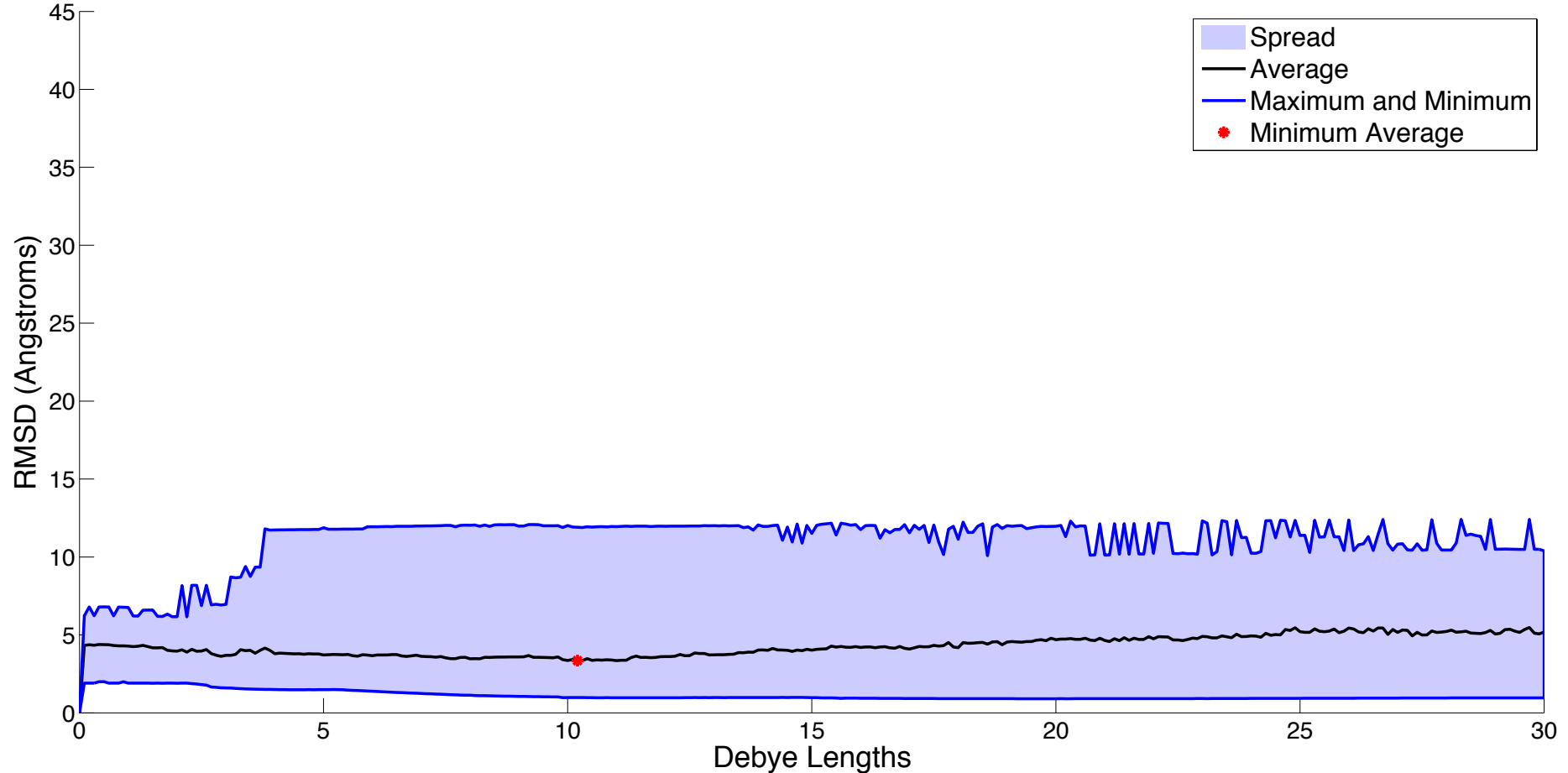
Debye length = 10.0 Å (default)

Minimum Average: 19

Energy Composition Dielectric 19



RMSD Over Debye Values



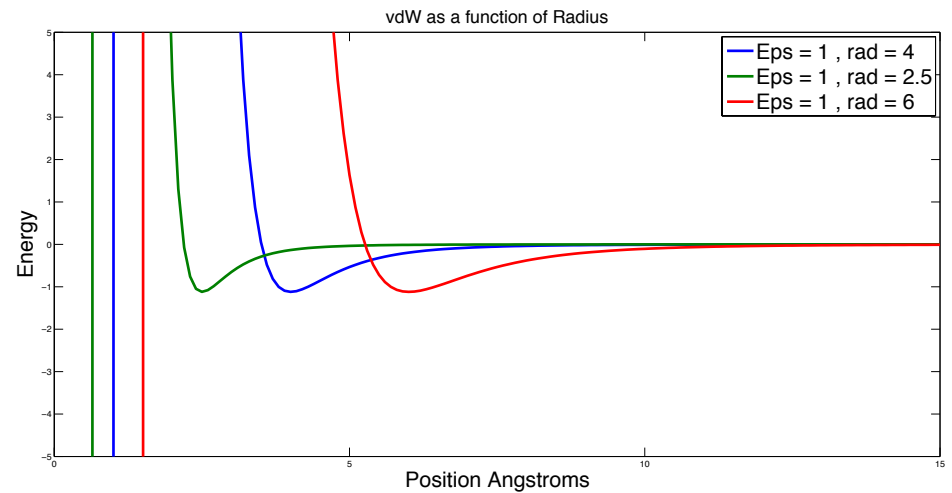
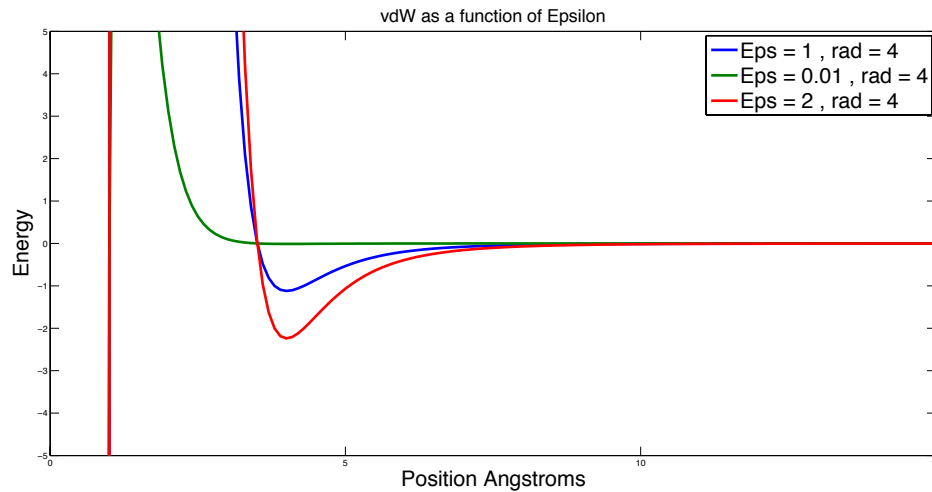
Electrostatic Interactions:

$q = -1.0$

Dielectric = 19

Minimum Debye: 10.2

$$E_{vdW} = \sum_{i,j} \varepsilon_{ij} \left[-2.25 \left(\frac{\sigma_{ij}}{r_{ij}} \right)^6 + 1.84 \times 10^5 e^{-\left(\frac{12r_{ij}}{\sigma_{ij}} \right)} \right]$$



Previous Parameters

Current vdW parameters

