

### Electrostatics in Biomacromolecules

Electrostatics important in (at least)

binding for

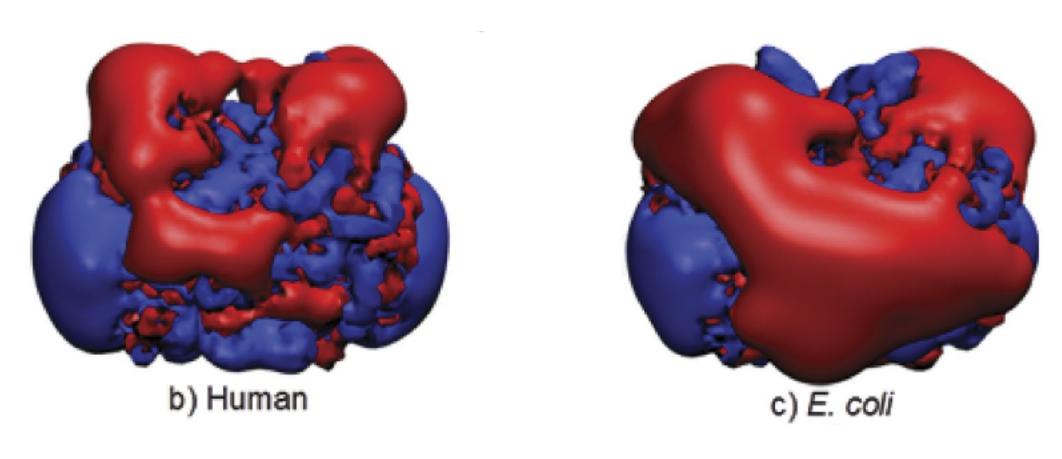
 steering, facilitating approach of species

 complexation, as complementarity means lower potential energy

 enzyme catalysis, as electric potential stabilizes transition state

 Thus, electrostatic potential usually conserved near functional sites

#### Thymidylate Synthase

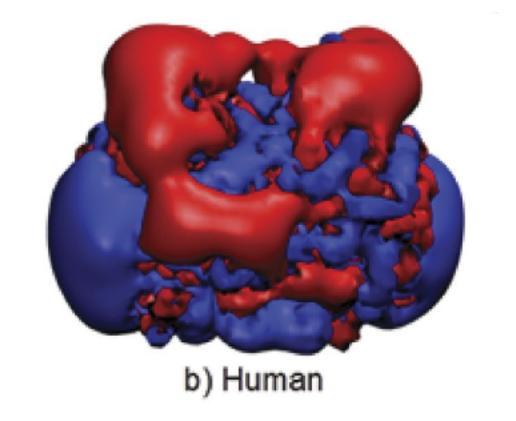


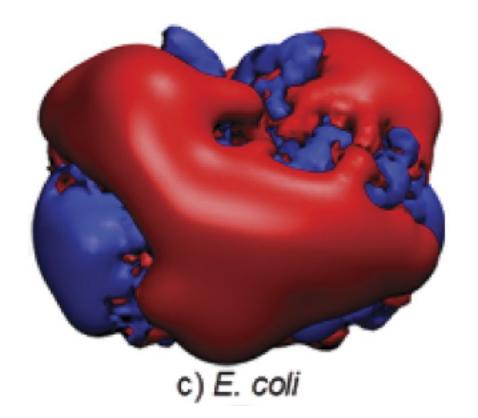
blue isocontours +1 kT/e, red isocontours -1 kT/e From Fig 1 of Garg *et al.* (2015).

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# Modeling Electrostatics

- In biological macromolecules, the electrostatic potential is usually calculated based on the Poisson-Boltzmann equation
  - The Poisson equation describes the potential field due to a given charge distribution. Atoms in the biomolecule are assumed to have a fixed charge.
  - The Poisson-Boltzmann equation assumes that (infinitely small) ions surround a biomolecule in accordance with the Boltzmann distribution
- The PB equation is a partial differential equation that is solved numerically
- The equation is often linearized to be more numerically stable
- Chun Liu in Applied Math has worked on versions
  - that are time-dependent
  - account for the finite size of ions