



# *Finite Element Analysis of Rydberg States*

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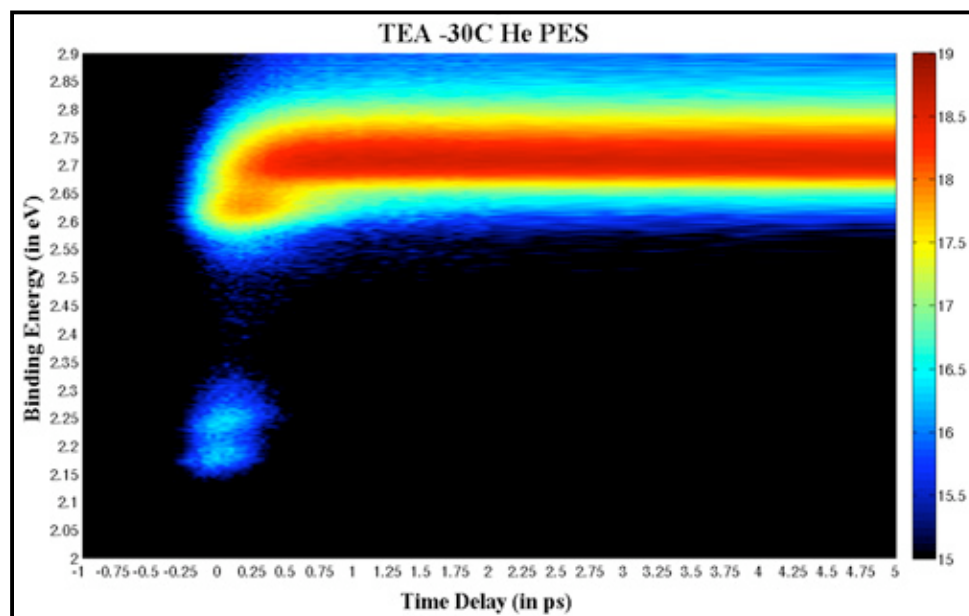
## **Funding**

- Army Research Office
- Department of Energy



## Challenge: Ultrafast Structural Dynamics

- Observe molecules while they do chemistry
- Requires  $<100$  fs ( $10^{-15}$  s) time resolution
- Requires new spectroscopic tools

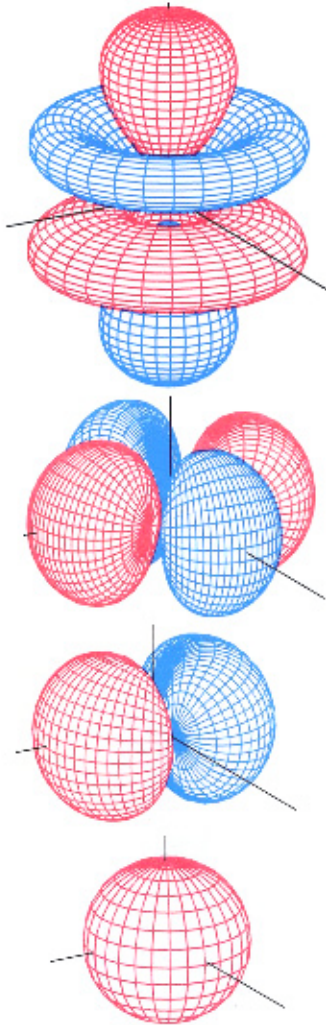


### Recent Discovery:

- Rydberg spectra encode molecular structure
- Technology developed to measure with fs time resolution
- Needed: Computational link spectrum  $\longleftrightarrow$  structure

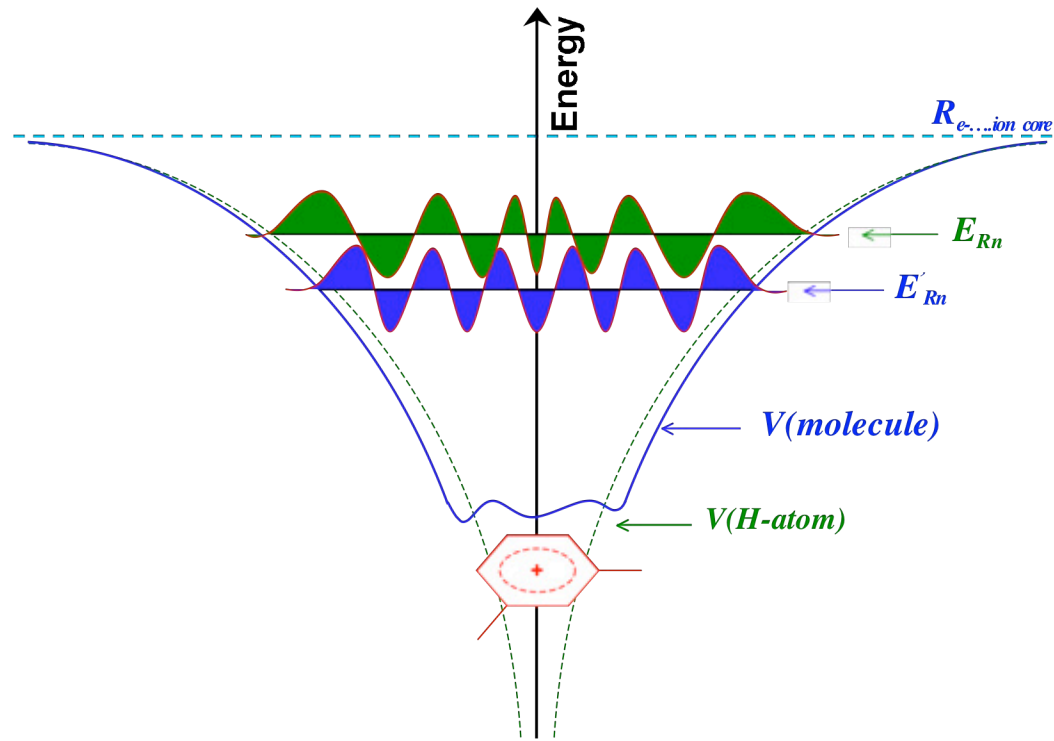


# Quantum Mechanics of Rydberg States



$$-\frac{\hbar^2}{2m} \nabla^2 u + Vu = \lambda u$$

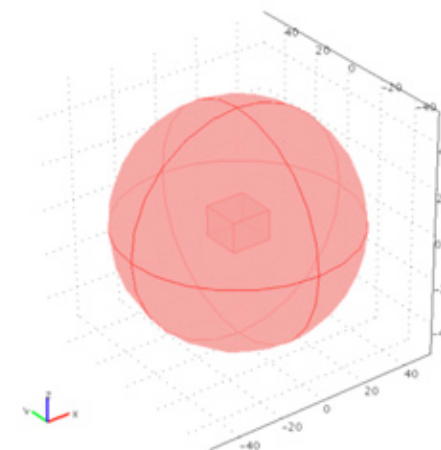
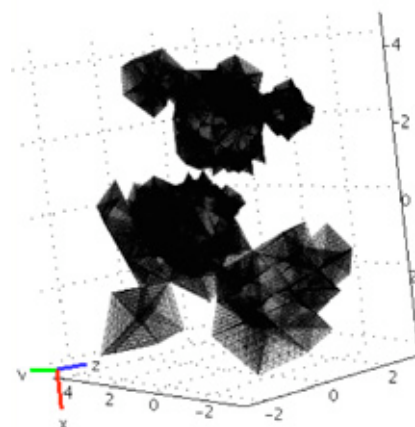
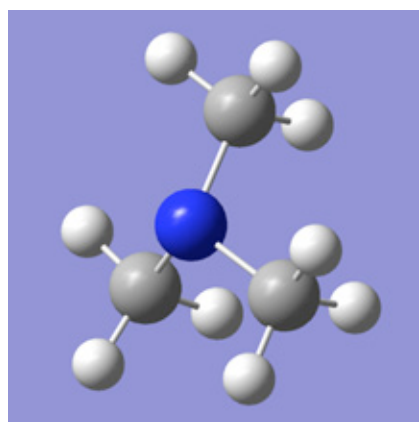
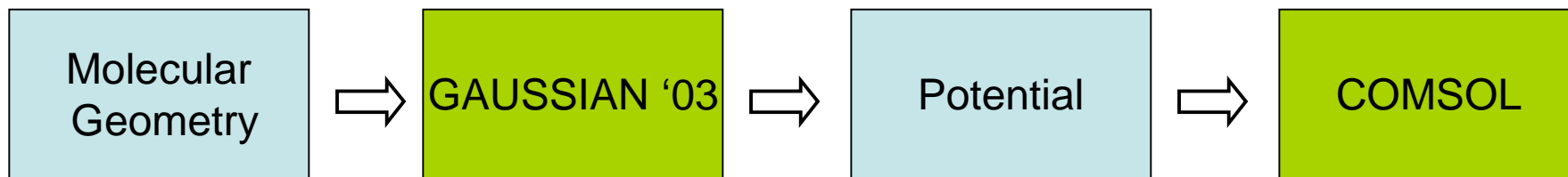
Schrödinger equation





# Calculation Scheme

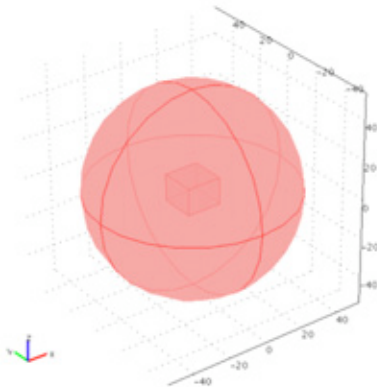
$$-\frac{\hbar^2}{2m}\nabla^2 u + Vu = \lambda u$$





## Inside COMSOL

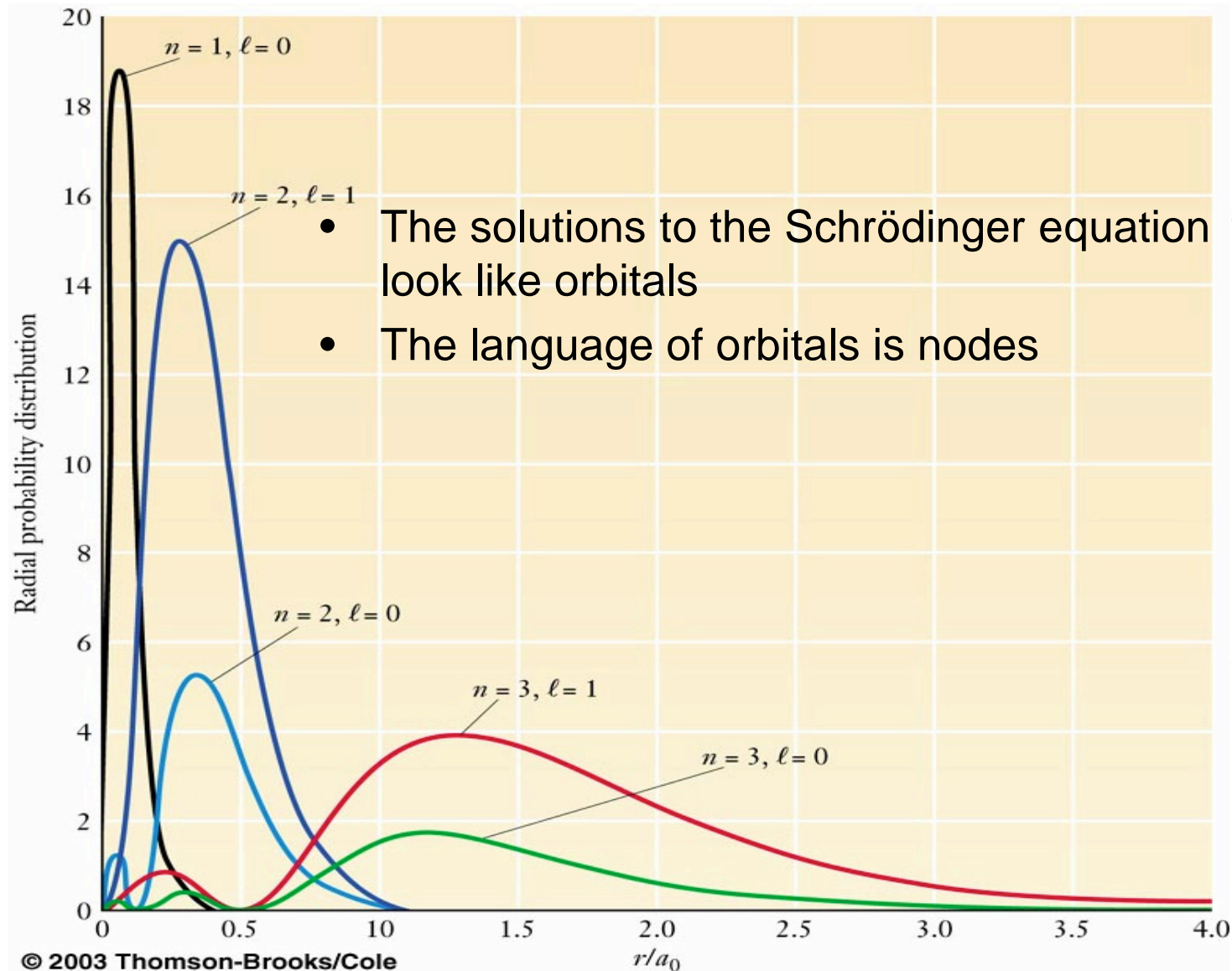
$$-\frac{\hbar^2}{2m}\nabla^2 u + Vu = \lambda u$$



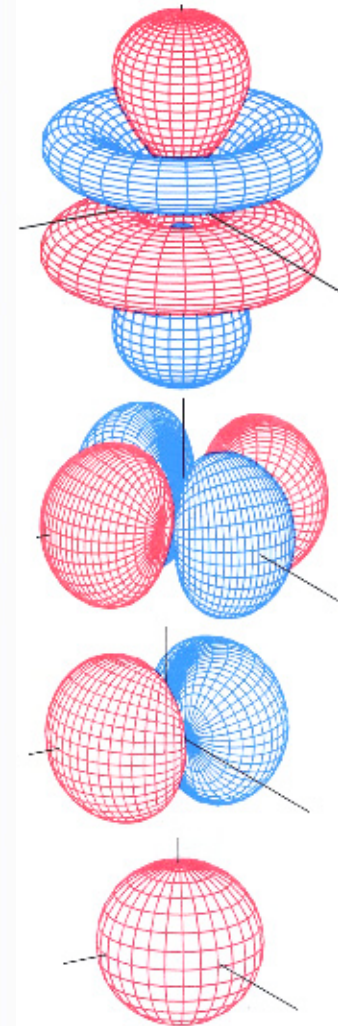
- *PARDISO solver*
- *20 eigenvalues around expected value*
- *Dirichlet Boundary Condition*
- *Matrices set to Hermitian*
- *Fine Potential Resolution in Inner Cube*
- *Coarser Potential Resolution Outside*
- *Mesh Density Proportional to Potential Resolution*



# Orbitals

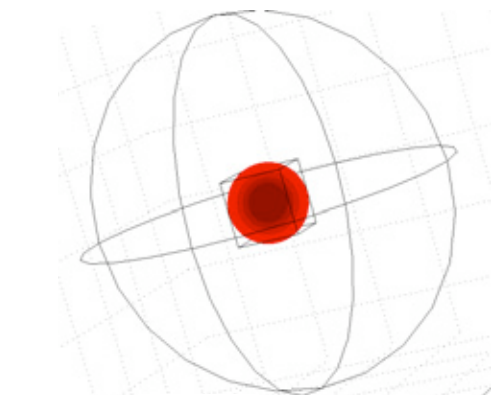


- The solutions to the Schrödinger equation look like orbitals
- The language of orbitals is nodes

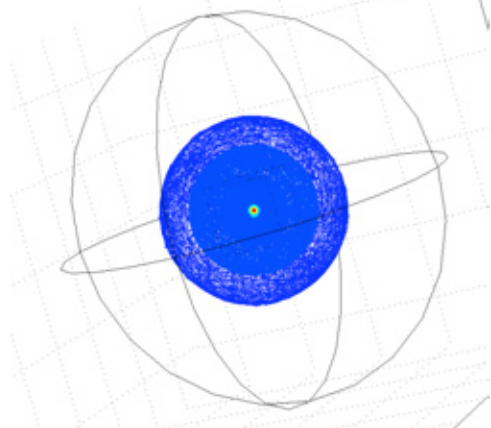




# Hydrogen in Cartesian Coordinates

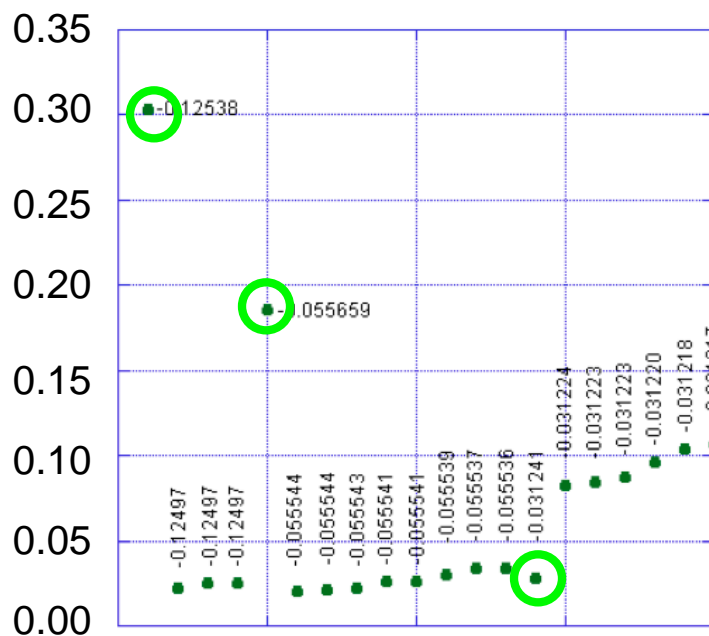


$$\sqrt{\langle r^2 \rangle} = 6.46 \text{ au}$$



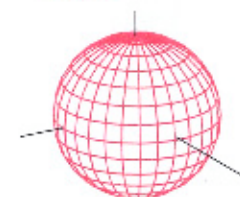
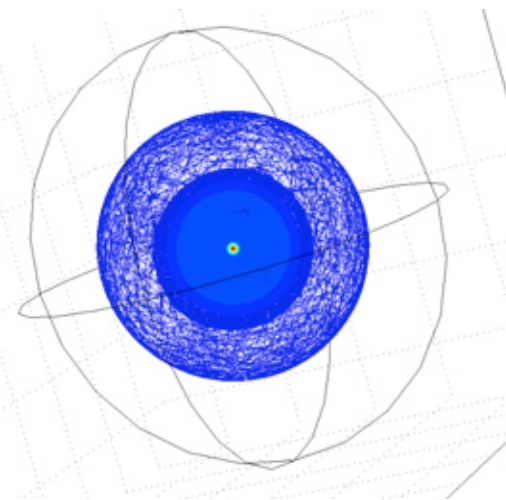
$$\sqrt{\langle r^2 \rangle} = 14.36 \text{ au}$$

Percent Error



Eigenvalue Number

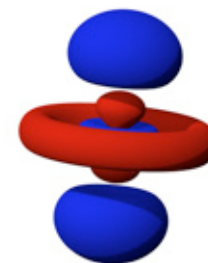
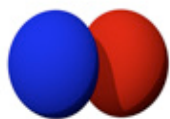
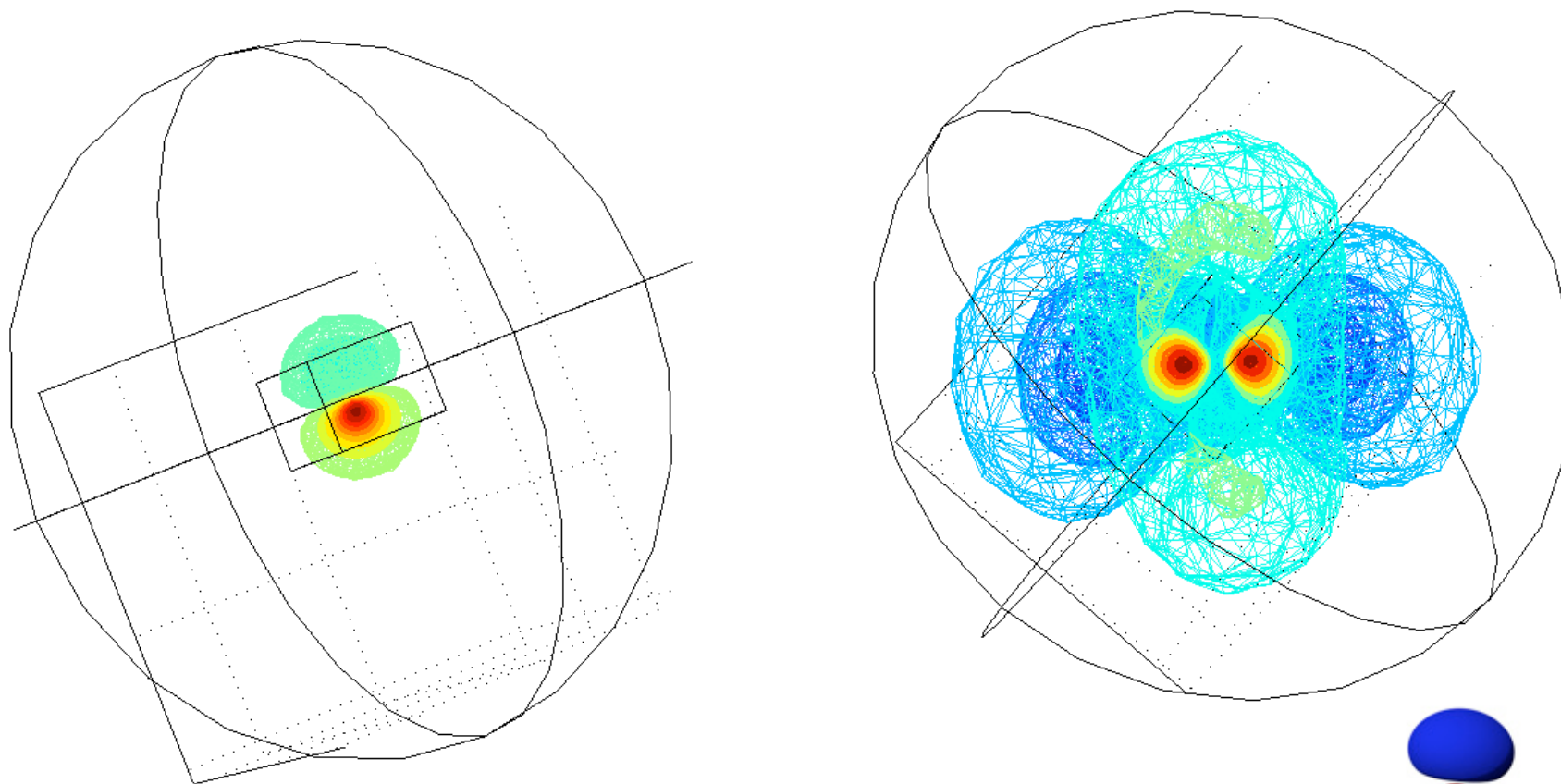
$$\sqrt{\langle r^2 \rangle} = 25.14 \text{ au}$$







# *Hydrogen Interpolated Solution*

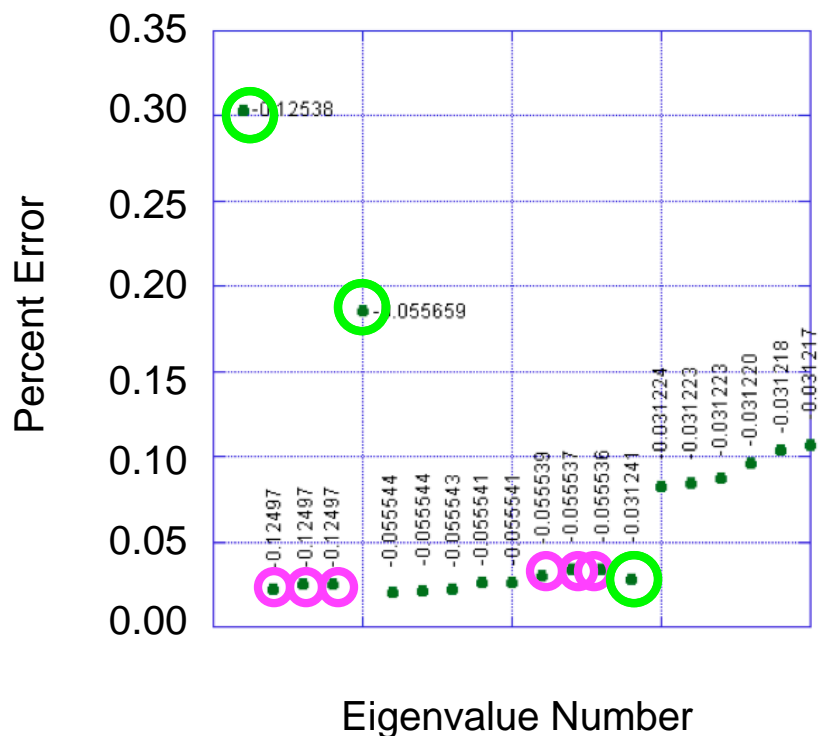




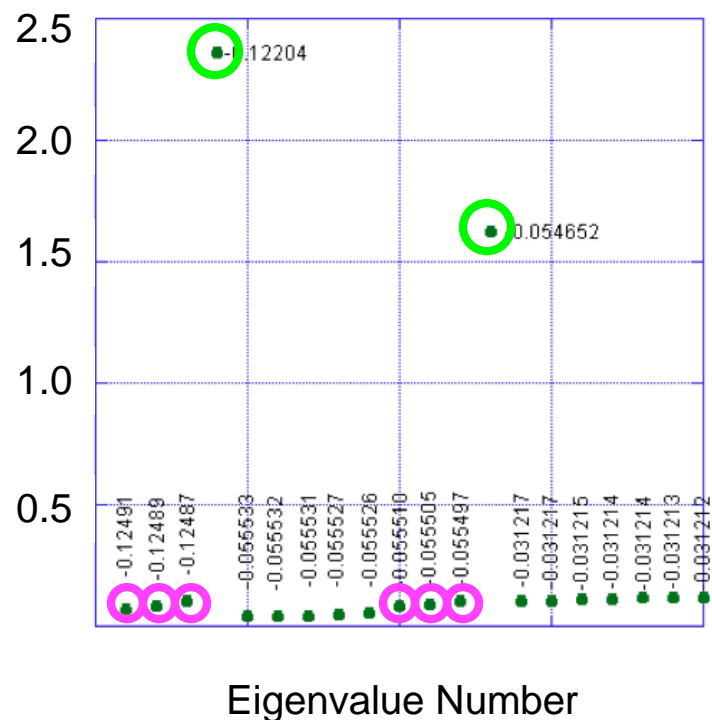


# Hydrogen in Cartesian Coordinates

Using analytical potential



Using imported potential

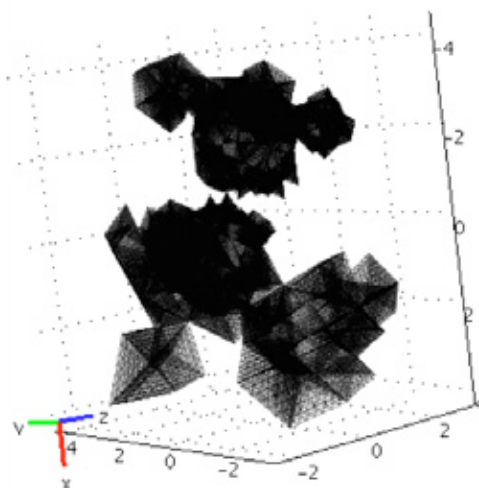
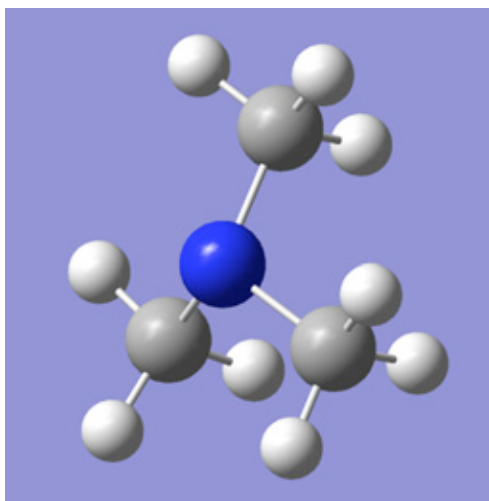


Conclude:

- Errors are in the 1% range for s orbitals using interpolated potentials
- For higher angular momentum states, errors remain  $\leq .1$  %

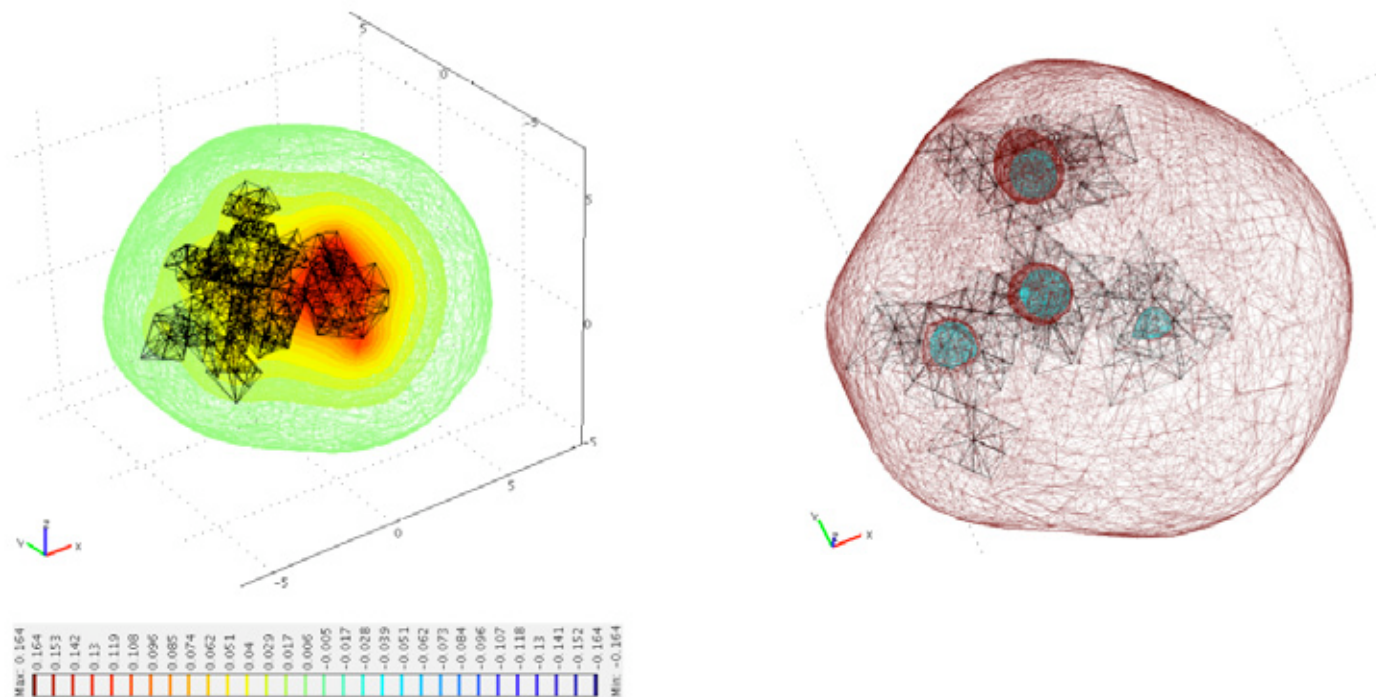


## *Trimethylamine (TMA) - Introduction*





# 3s Eigenvalue



$$E_{\text{This work}} = -3.323 \text{ eV}$$

$$E_{\text{Experiment}} = -3.087 \text{ eV} \quad \% \text{ error} = 7.6\%$$

$$E_{\text{Previous ab initio}} = -3.202 \text{ eV} \quad \% \text{ error} = 3.4\%$$

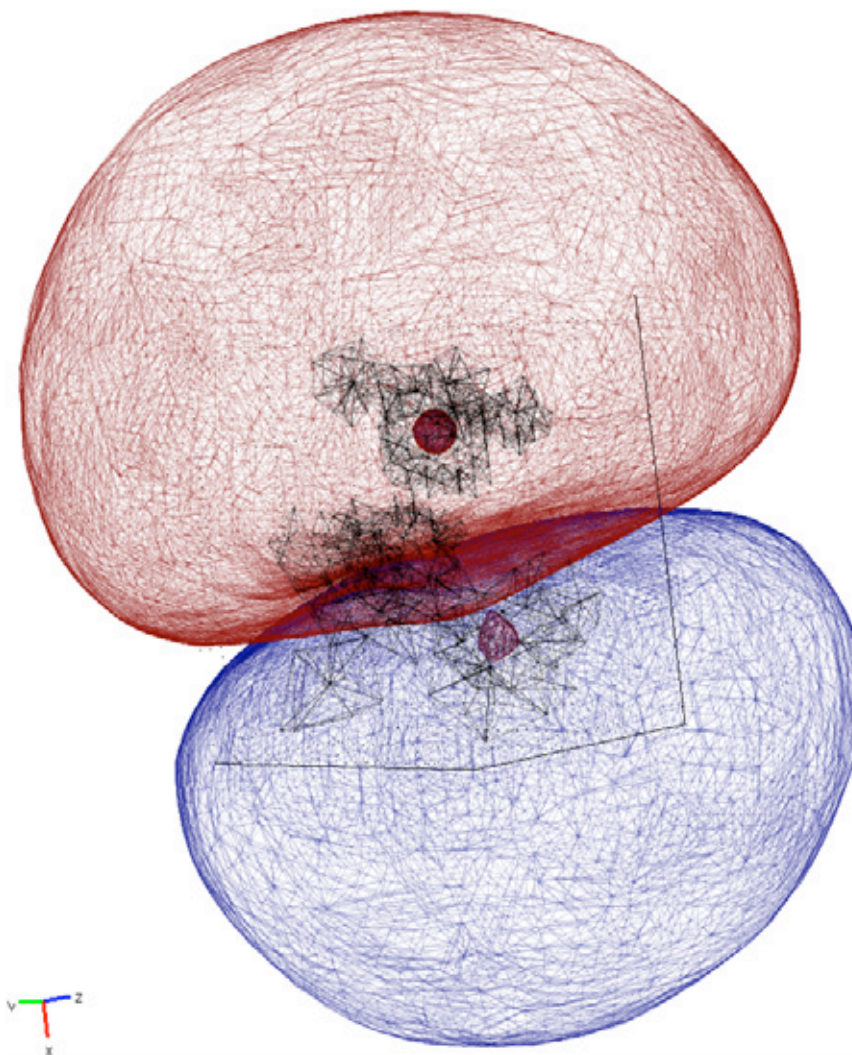
$$\sqrt{\langle r^2 \rangle} = 3.92 \text{ au}$$



## 3p Eigenvalue

$E_{\text{this work}} = -2.217 \text{ eV}$	
$E_{x,y \text{ experiment}} = -2.251 \text{ eV}$	%error = 1.5%
$E_{z \text{ experiment}} = -2.204 \text{ eV}$	%error = 0.6%
$E_{x,y \text{ ab initio}} = -2.517 \text{ eV}$	%error = 12 %
$E_{z \text{ ab initio}} = -2.317 \text{ eV}$	%error = 6.4%

$$\sqrt{\langle r^2 \rangle} = 4.038 \text{ au}$$





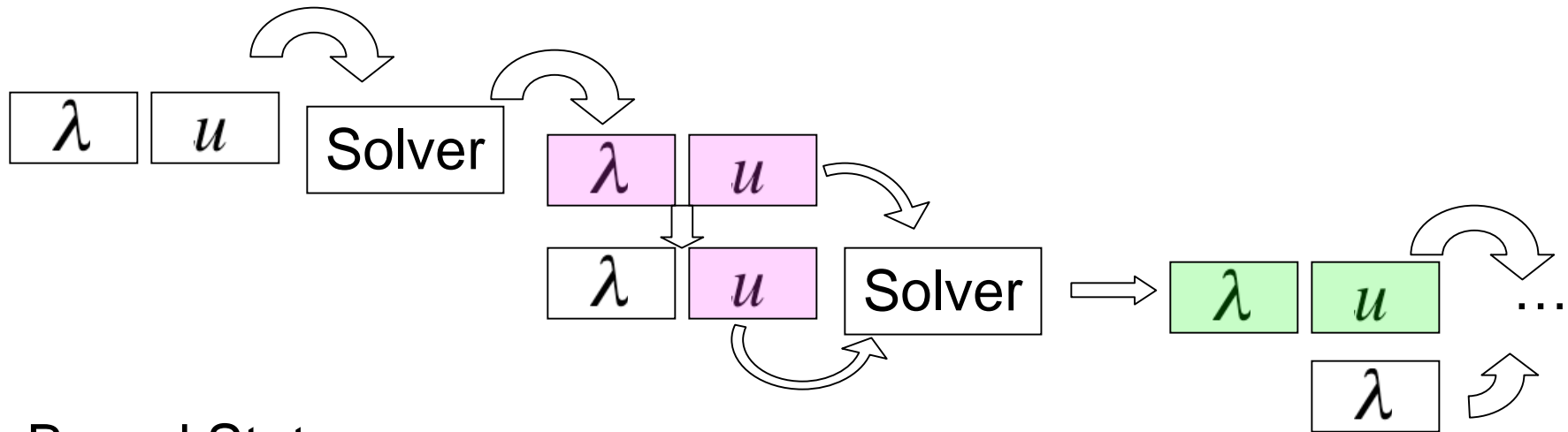
## *TMA - Conclusion*

- We have demonstrated a new conceptual approach to calculate Rydberg orbitals
- We have calculated the most accurate TMA 3p eigenvalue ever.
- Our method shows promise of scaling well for eigenvalue computation of large molecules, opening up new venues of computation.



# Algorithmic Issues

Initial Conditions



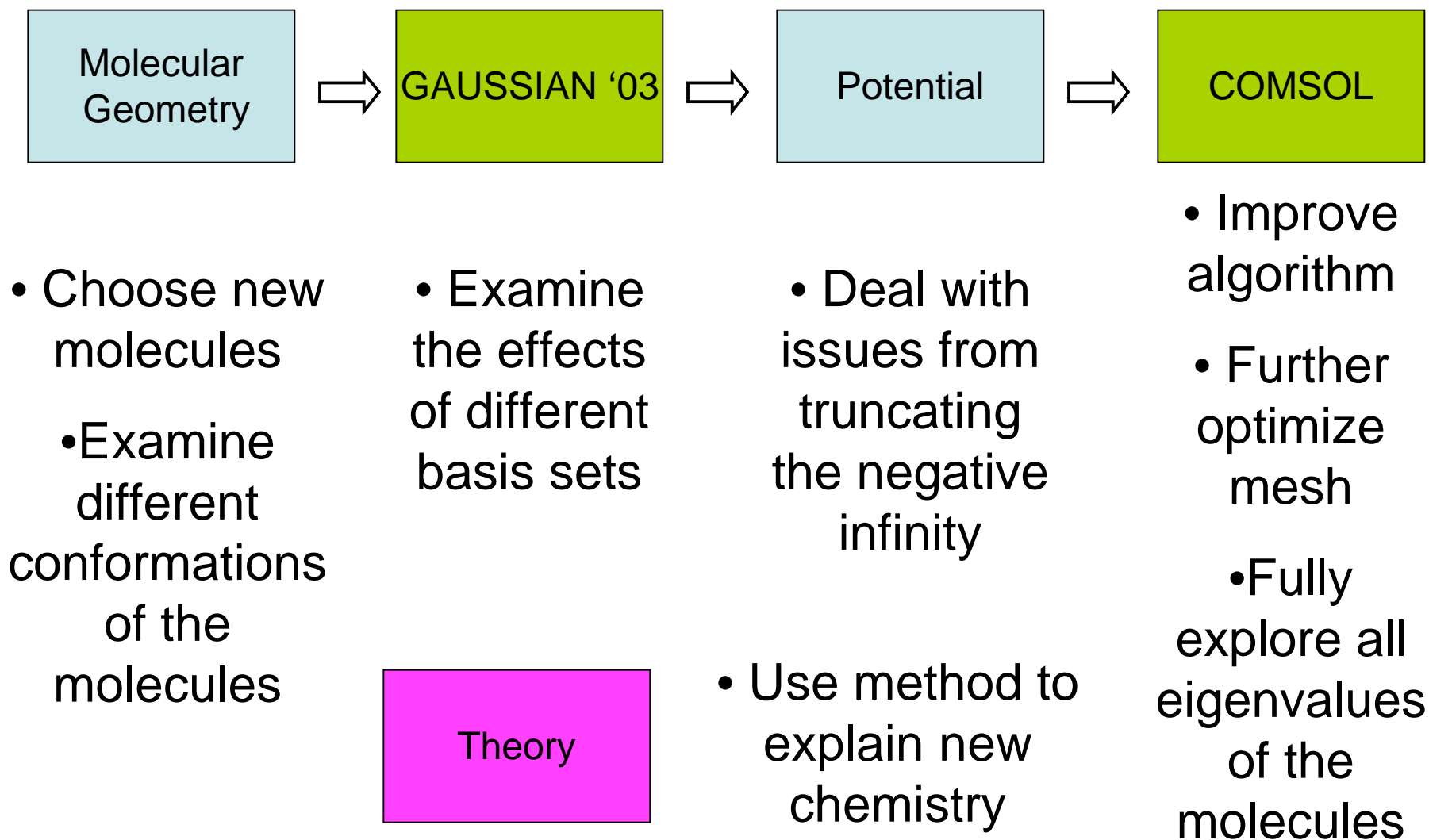
Bound States

Or, how can I bias the solver to return only negative eigenvalues?



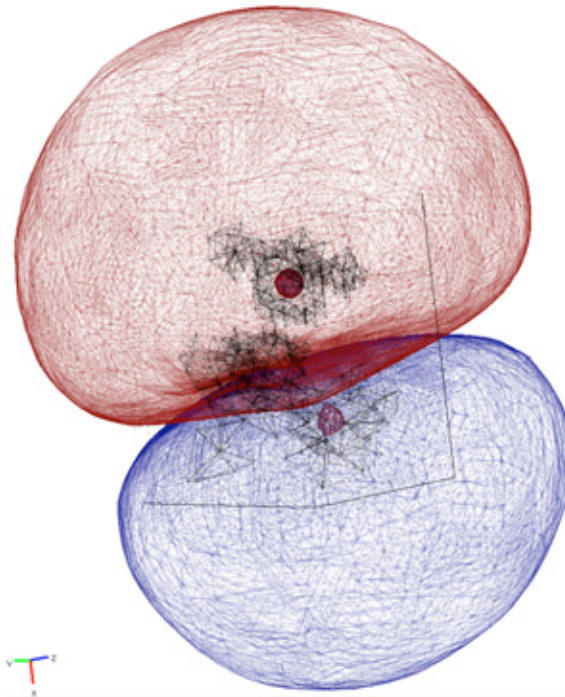


## Next Steps





*Thank You!*



*Any Questions? Comments? Ideas?*