

Cyclodextrins: From Physics to the Pharmacy

Kacey Meaker

NIST Computational Chemistry Group

AIP/SPS Internship Program 2006

Randolph-Macon Woman's College



Purpose

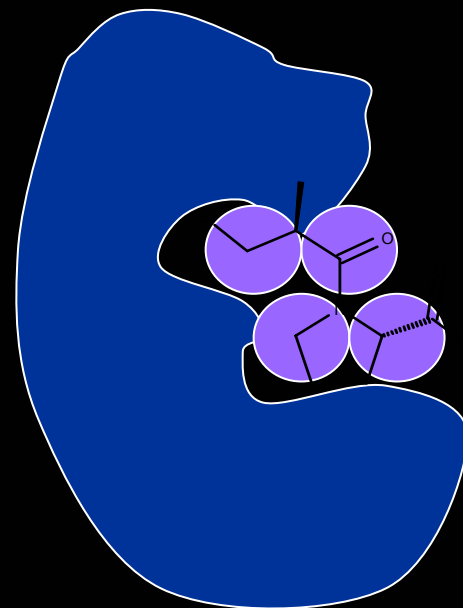


- Improve physical models for use in drug design
- Extend the medicinal applications of cyclodextrins



Computational Methods

- Merck Molecular Force Field (MMFFs)
- Monte Carlo Simulations
- Cluster Analysis
- Docking and Scoring



Jensen, F. *Introduction to Computational Chemistry*. John Wiley & Sons (1999).

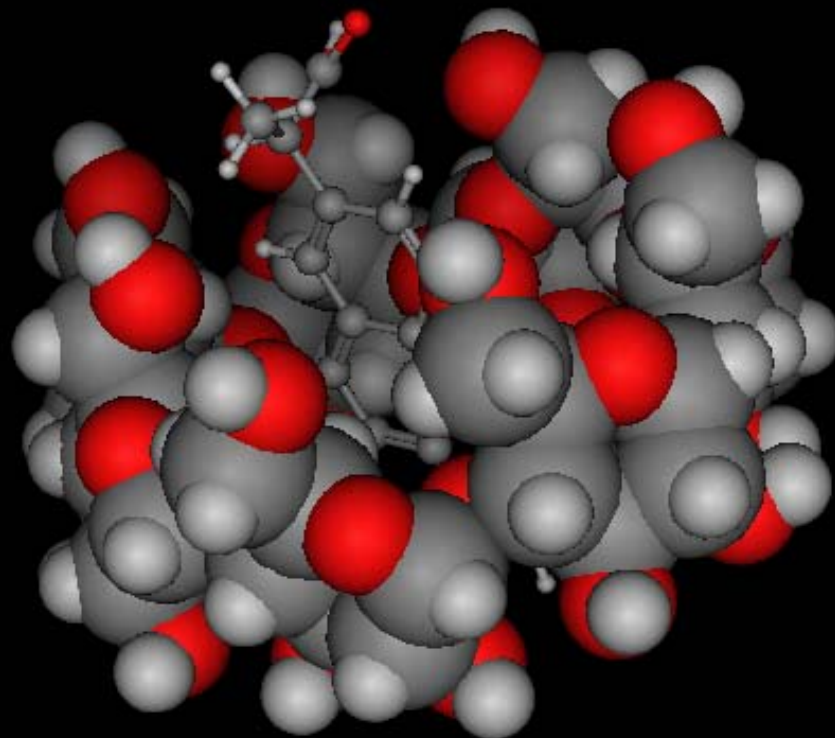
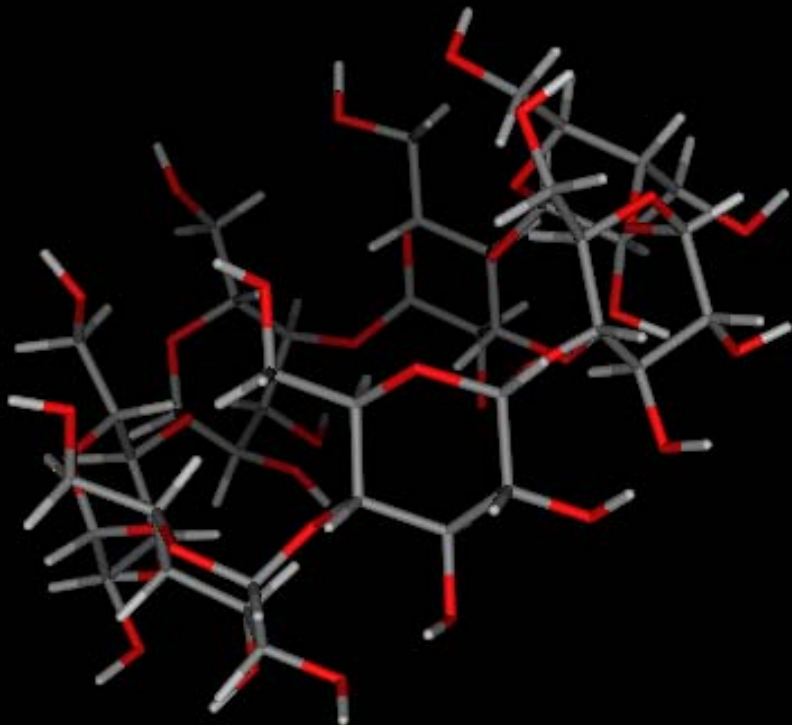
Halgren, T. A. *J. Comput. Chem.* **17**, 490-519 (1996).

Leach, A. R. *Molecular Modelling: Principles and Applications*. Addison Wesley Longman Limited (1996)

Shenkin, P.S. and McDonald, D. Q. *J. Comput. Chem.* **15**, 899-916 (1994).



Cyclodextrins 101



The Problem with Current Screening



- **Current Screening**: docking and scoring receptors and test ligands
- **Problem**: inconsistencies due to complexity of biological systems and incomplete physical models
- **Solution**: use a less complicated system like cyclodextrins that can be more easily understood



Training Wheel Molecules

- Use cyclodextrins as training wheels to improve the model
- Calculate the free energy of binding with increasingly sophisticated methods
- Compare to experimental data to find where the errors are



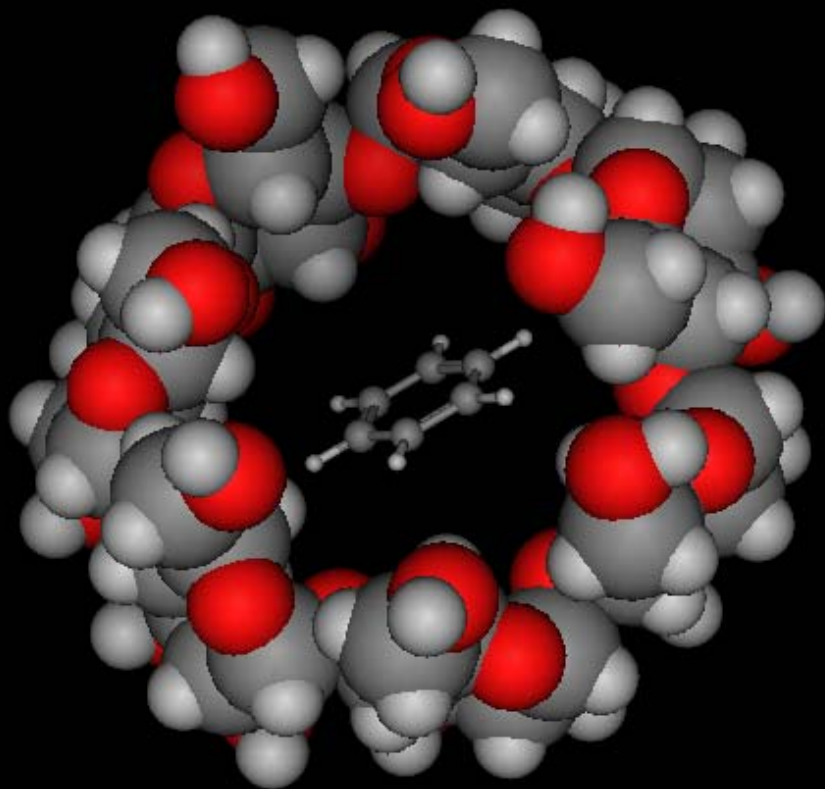
Progress toward Improving Models



- Calculated the free energy of binding for 22 derivatized β -cyclodextrins with 3 guest molecules for later experimental verification
- Possible future collaboration with University of Maryland to obtain experimental data



The Problem with β -cyclodextrin



Heat of Solvation of Derivatized β -CD's

| Attached Group | Heat of Solvation (kcal/mol) | Attached Group | Heat of Solvation (kcal/mol) |
|----------------|------------------------------|---------------------|------------------------------|
| Naked | -475.39 | Ala-ala-cys-glu | -714.33 |
| Hydroxypropyl | -500.81 | Ala-ala-cys-arg | -403.88 |
| Ala | -512.16 | Ala-ala-cys-cys-glu | -695.62 |
| Cys | -515.80 | Ala-ala-cys-cys-arg | -475.91 |
| Glu | -612.69 | Bradykinin | -503.95 |
| Arg | -349.32 | Zonulin 1 | -551.52 |
| Ala-ala-glu | -646.51 | Zonulin 3 | -674.57 |
| Ala-ala-arg | -375.20 | | |



Amino Acids and Drug Interactions



- **Hypothesis:** Positive amino acids will cause negative drugs to bind more strongly and negative amino acids will cause negative drugs to bind less strongly
- This will allow us to choose different amino acids for different drugs and also be able to target drugs to different proteins



Docking: Benzene



| Attached Group | Free Energy of Binding (kcal/mol) | Change from Derivaterizing (kcal/mol) | Attached Group | Free Energy of Binding (kcal/mol) | Change from Derivaterizing (kcal/mol) |
|----------------|-----------------------------------|---------------------------------------|---------------------|-----------------------------------|---------------------------------------|
| Naked | -1.6868 | | Ala-ala-cys-glu | -1.6283 | 0.0585 |
| Hydroxypropyl | -1.7324 | -0.0456 | Ala-ala-cys-arg | -1.7554 | -0.0686 |
| Ala | -1.7818 | -0.0950 | Ala-ala-cys-cys-glu | -1.7953 | -0.1085 |
| Cys | -1.7648 | -0.0780 | Ala-ala-cys-cys-arg | -1.8979 | -0.2111 |
| Glu | -1.0936 | 0.5932 | Bradykinin | -1.6725 | 0.0143 |
| Arg | -1.7815 | -0.0947 | Zonulin 1 | -1.7458 | -0.0590 |
| Ala-ala-glu | -1.6963 | -0.0095 | Zonulin 3 | -2.0955 | -0.4087 |
| Ala-ala-arg | -1.7457 | -0.0589 | | | |



Docking: Naproxen



| Attached Group | Free Energy of Binding (kcal/mol) | Change from Derivaterizing (kcal/mol) | Attached Group | Free Energy of Binding (kcal/mol) | Change from Derivaterizing (kcal/mol) |
|----------------|-----------------------------------|---------------------------------------|---------------------|-----------------------------------|---------------------------------------|
| Naked | -4.3237 | | Ala-ala-cys-glu | -4.2644 | 0.0593 |
| Hydroxypropyl | -5.2387 | -0.9150 | Ala-ala-cys-arg | -4.4746 | -0.1509 |
| Ala | -5.1545 | -0.8308 | Ala-ala-cys-cys-glu | -3.9905 | 0.3332 |
| Cys | -4.3915 | -0.0678 | Ala-ala-cys-cys-arg | -4.4381 | -0.1144 |
| Glu | -4.7334 | -0.4097 | Bradykinin | -3.7227 | 0.6010 |
| Arg | -4.3931 | -0.0694 | Zonulin 1 | -2.8666 | 1.4571 |
| Ala-ala-glu | -4.6726 | -0.3489 | Zonulin 3 | -3.3238 | 0.9999 |
| Ala-ala-arg | -4.7032 | -0.3795 | | | |



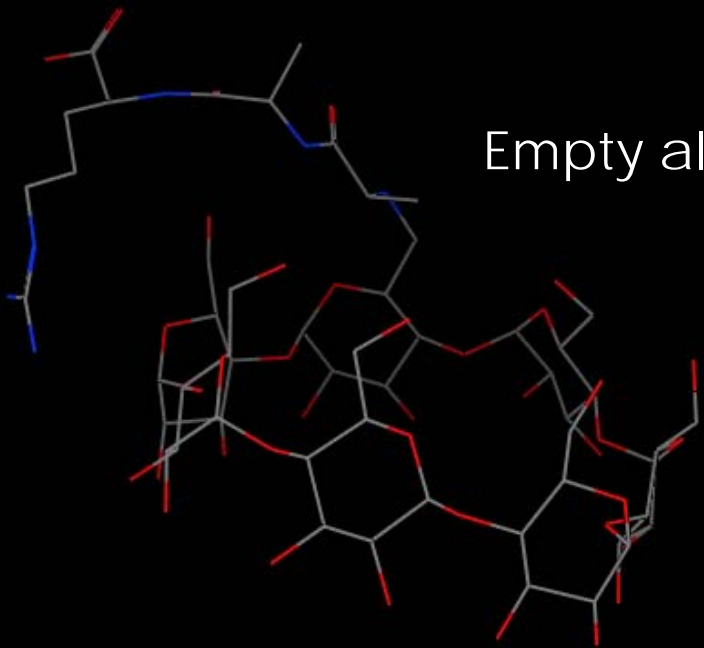
Docking: Flurbiprofen

| Attached Group | Free Energy of Binding (kcal/mol) | Change from Derivatizing (kcal/mol) | Attached Group | Free Energy of Binding (kcal/mol) | Change from Derivatizing (kcal/mol) |
|----------------|-----------------------------------|-------------------------------------|---------------------|-----------------------------------|-------------------------------------|
| Naked | -4.1109 | | Ala-ala-cys-glu | -3.8418 | 0.2691 |
| Hydroxypropyl | -3.9512 | 0.1597 | Ala-ala-cys-arg | -3.5591 | 0.5518 |
| Ala | -4.5858 | -0.4749 | Ala-ala-cys-cys-glu | -4.1750 | -0.0641 |
| Cys* | -2.2175 | 1.8934 | Ala-ala-cys-cys-arg | -3.5683 | 0.5426 |
| Glu | -4.4139 | -0.3030 | Bradykinin | -4.1060 | 0.0049 |
| Arg | -4.3971 | -0.2862 | Zonulin 1 | -5.5171 | -1.4062 |
| Ala-ala-glu | -4.4936 | -0.3827 | Zonulin 3 | -5.2514 | -1.1405 |
| Ala-ala-arg | -4.3563 | -0.2454 | | | |



* Conformation of flurbiprofen is a false, very high energy conformation which leads to such a low binding energy. Calculation needs to be re-optimized and re-docked.

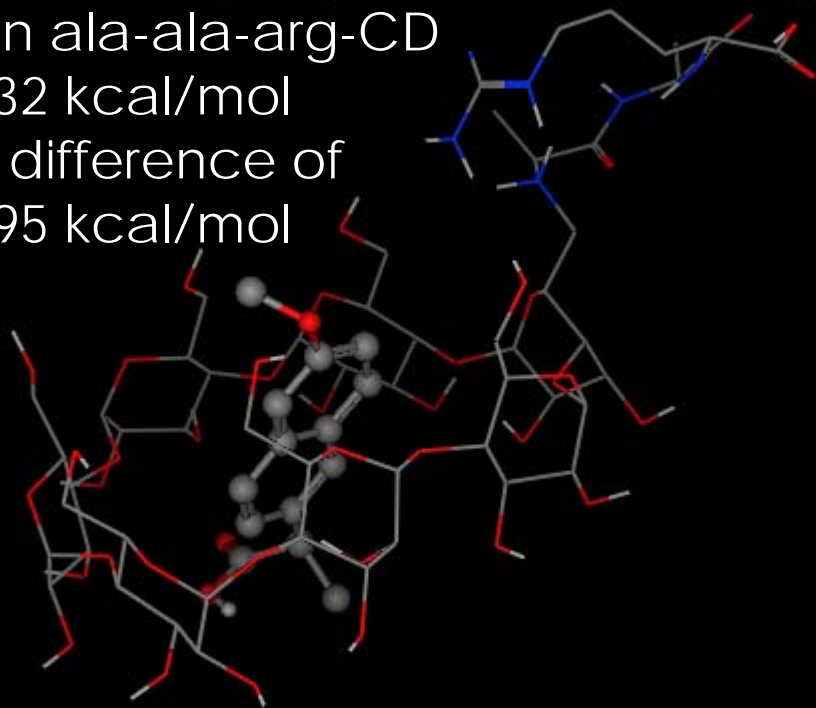
Empty ala-ala-arg-CD



Naproxen in ala-ala-arg-CD

-4.7032 kcal/mol

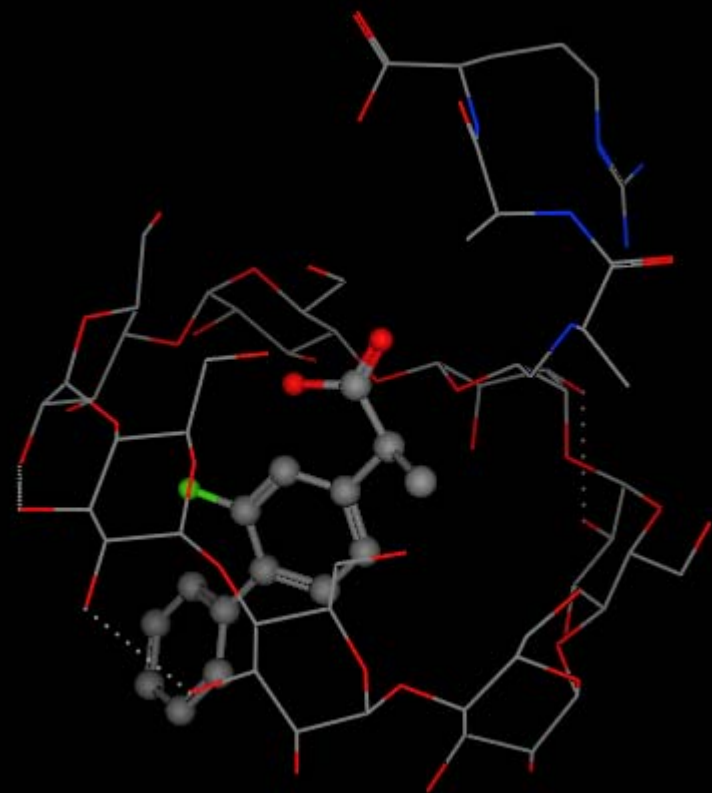
Energy difference of
-0.3795 kcal/mol

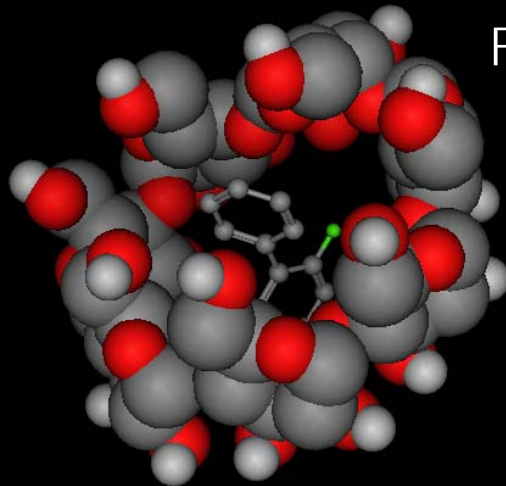


Flurbiprofen in ala-ala-arg-CD

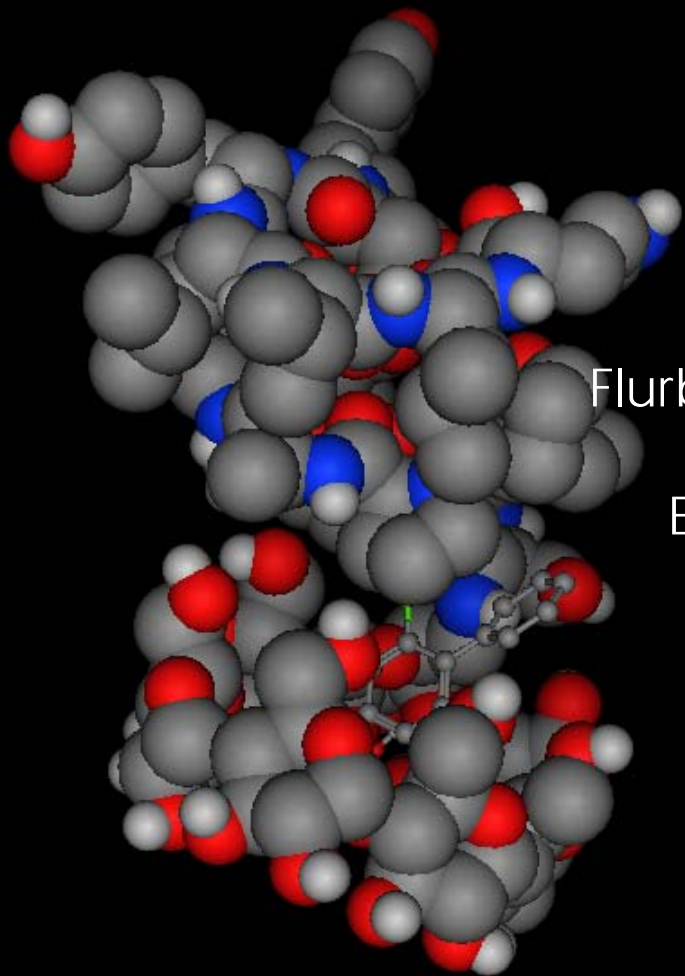
-4.3563 kcal/mol

Energy difference of
-0.2454 kcal/mol

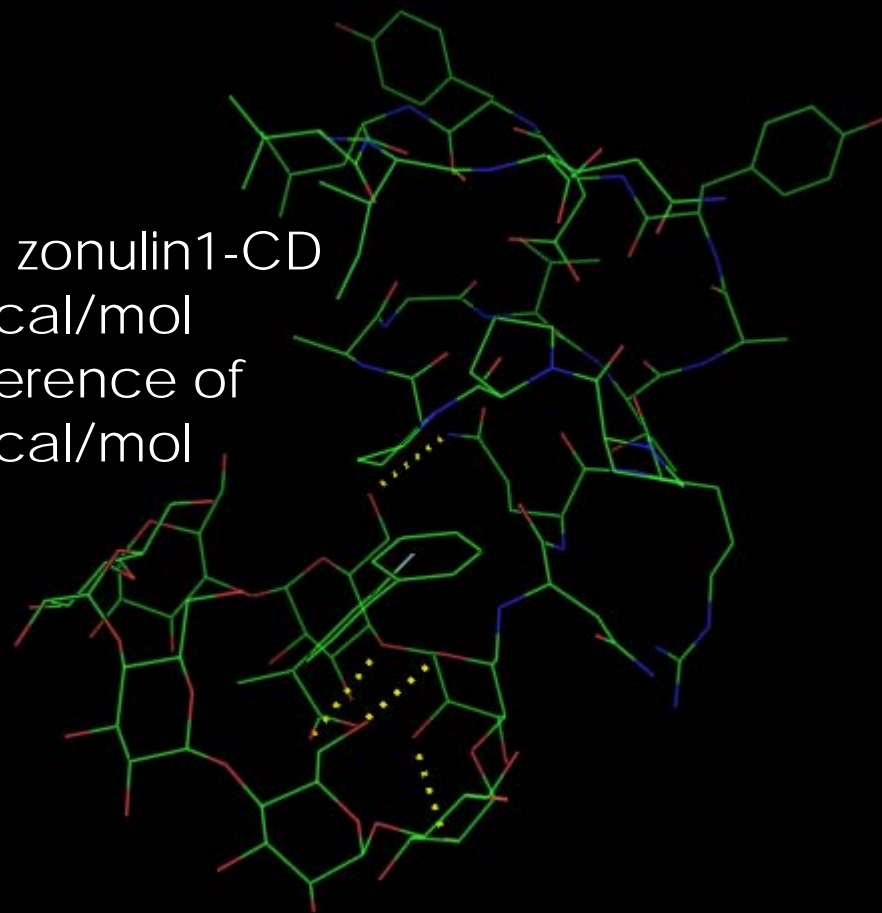




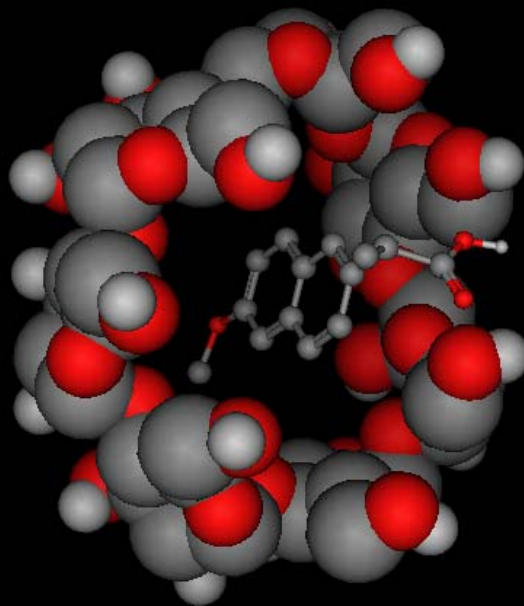
Flurbiprofen in Naked CD
-4.1109 kcal/mol



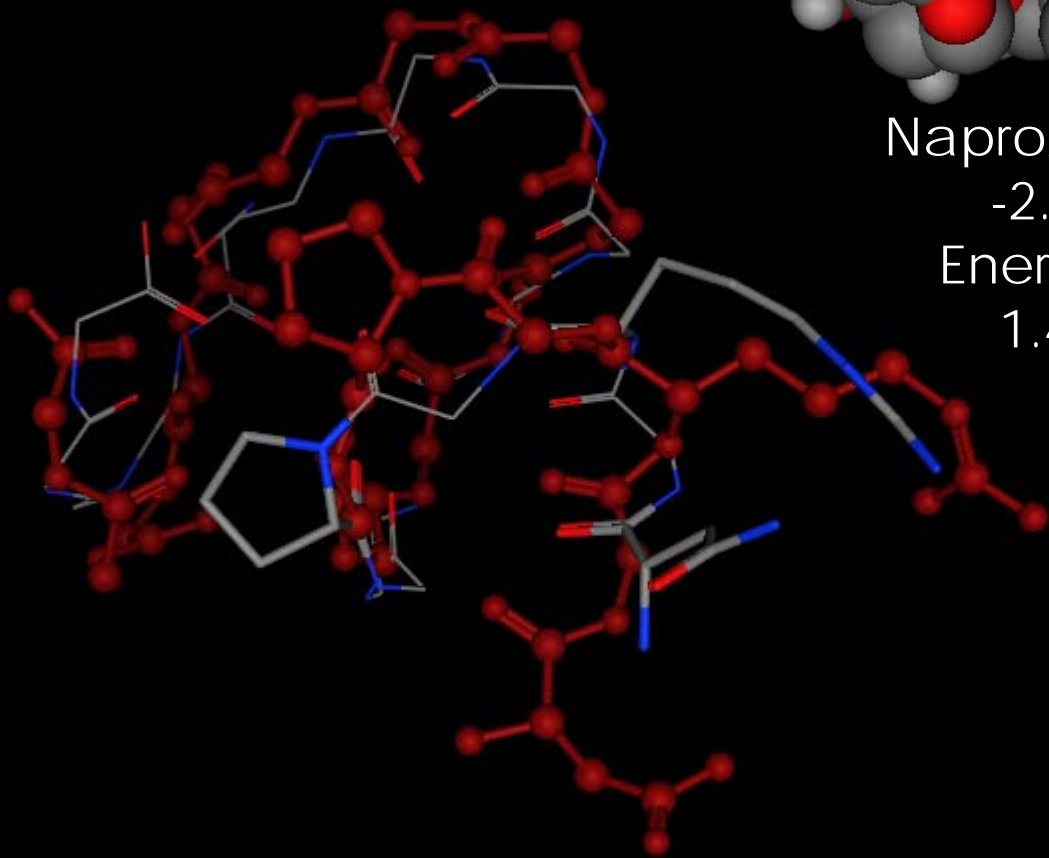
Flurbiprofen in zonulin1-CD
-5.5171 kcal/mol
Energy difference of
-1.4062 kcal/mol



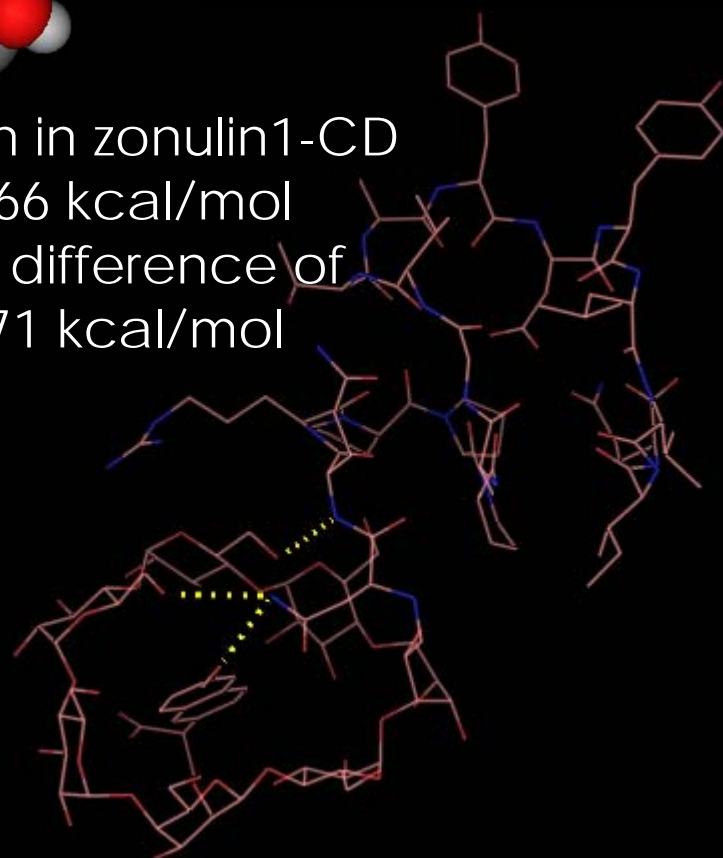
Zonulin1 peptide from
naproxen (red)
Zonulin1 peptide from
flurbiprofen (multi-colored)

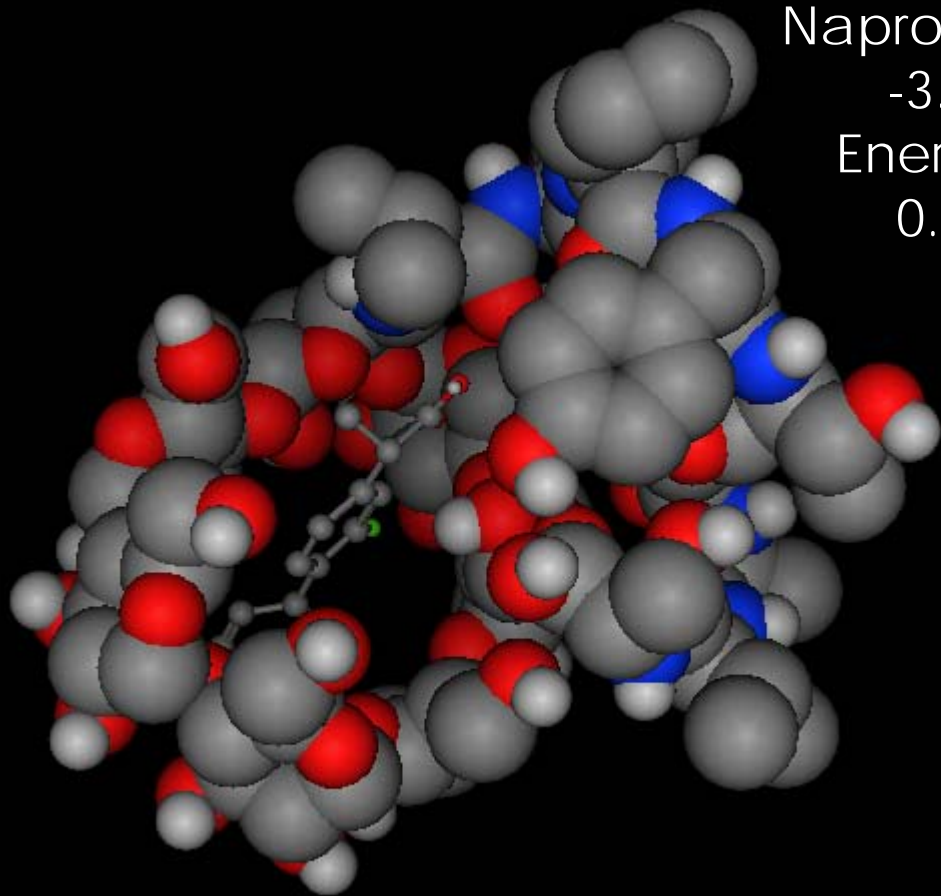


Naproxen in Naked CD
-4.3237 kcal/mol



Naproxen in zonulin1-CD
-2.8666 kcal/mol
Energy difference of
1.4571 kcal/mol



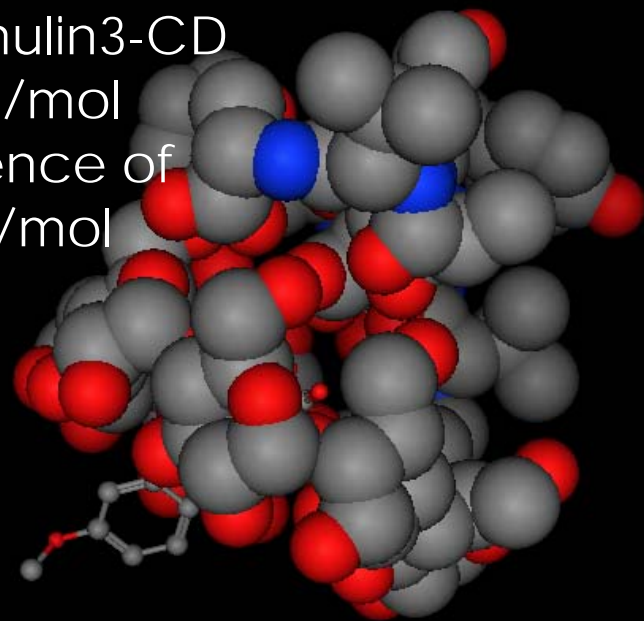


Naproxen in zonulin3-CD

-3.3238 kcal/mol

Energy difference of

0.9999 kcal/mol

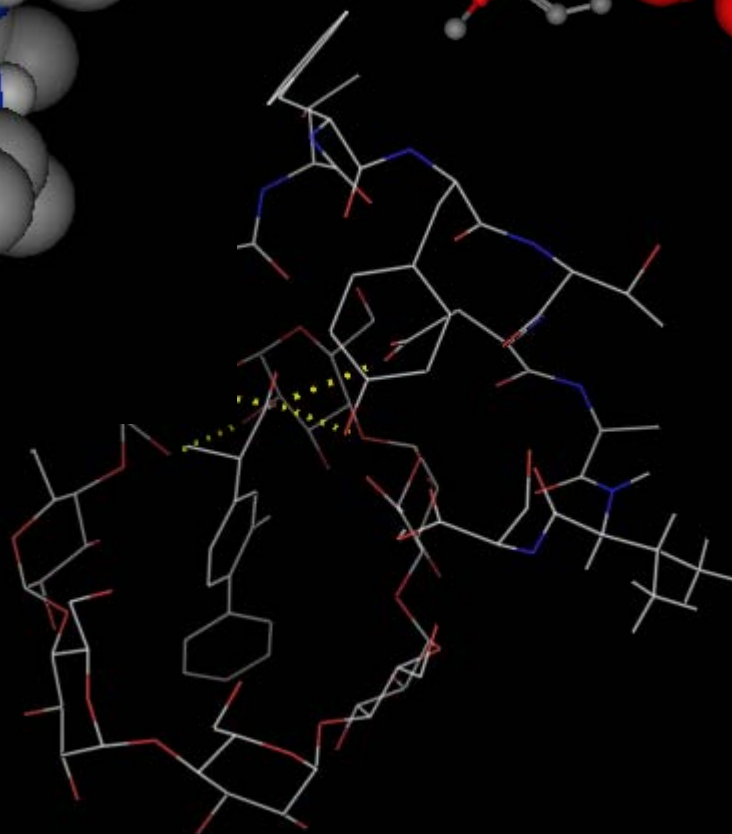


Flurbiprofen in zonulin3-CD

-5.2514 kcal/mol

Energy difference of

-1.1405 kcal/mol



Results from Derivatization



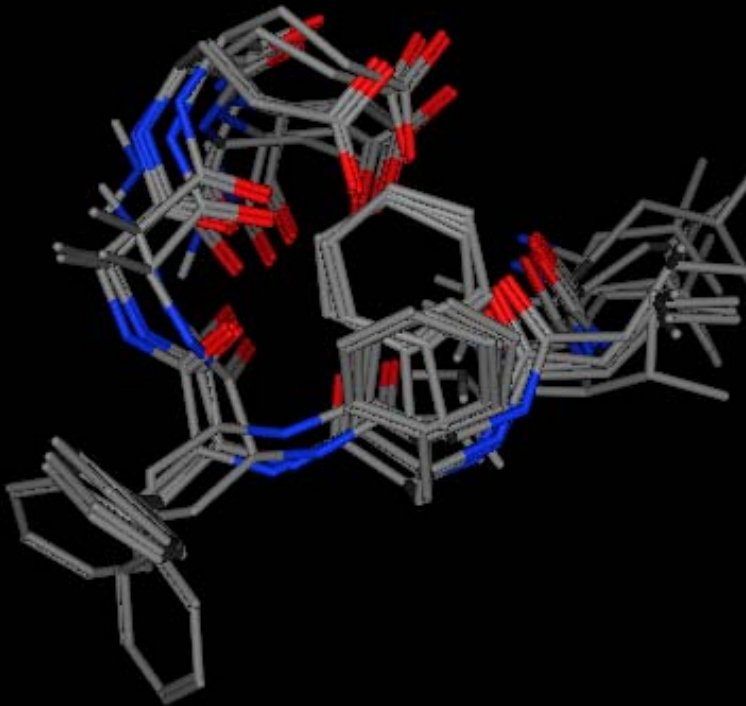
- Changes due to derivatization with amino acids have more to do with changes in pocket characteristics than a simple positive-negative interaction
- Need to do more analysis
- Derivatization with zonulin peptides results in a very stable docking with flurbiprofen



Alzheimer's: A Real Life Example

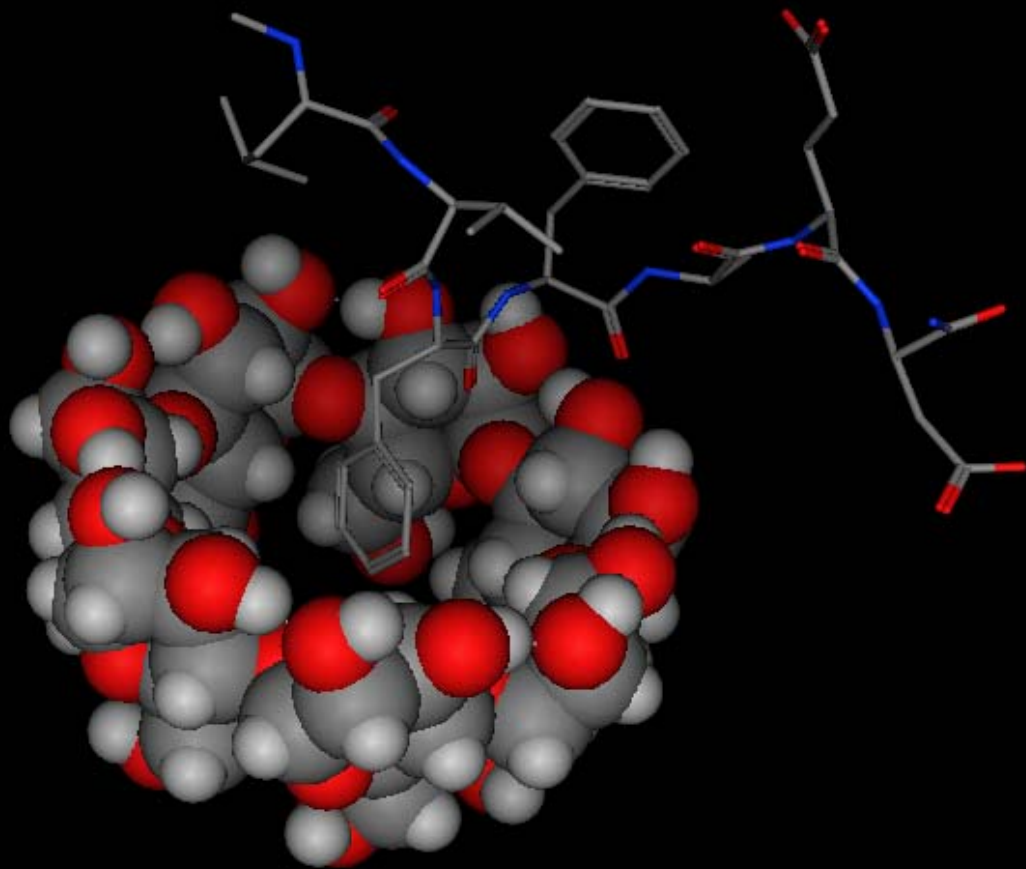
Conformational Analysis after 6000 steps

| Peptide Residues | Minimum Energy (kcal/mol) | Conformations within 3 kcal/mol |
|------------------|---------------------------|---------------------------------|
| 16-23 | -1373.73 | 90 |
| 12-28 | -6318.39 | 56 |
| 1-40 | -15341.57 | 30 |



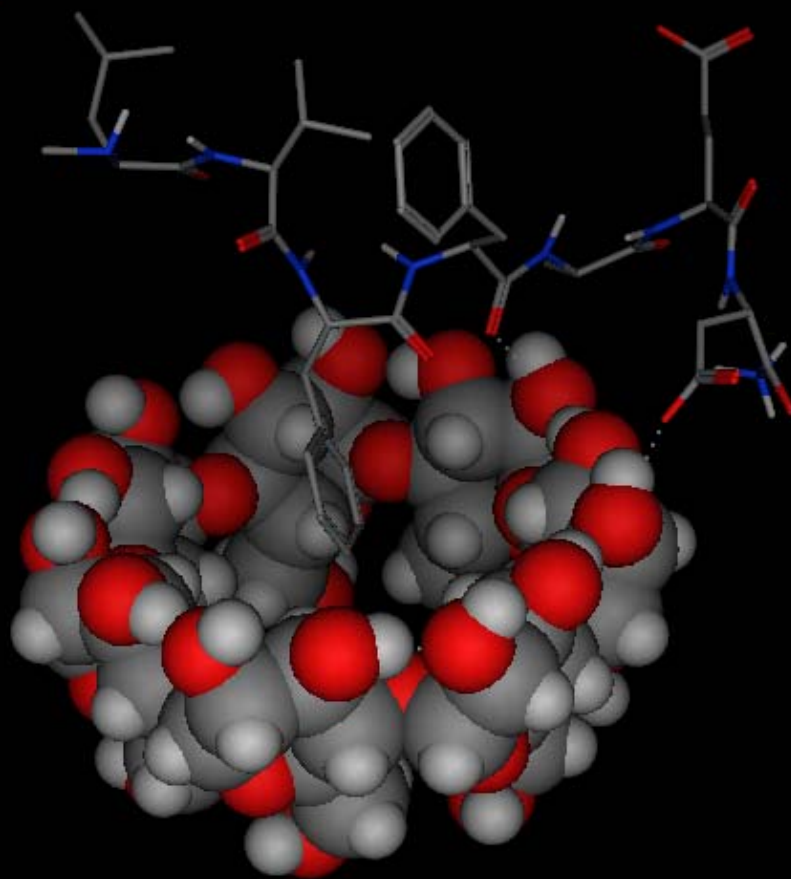
Sticht, H., Bayer, P. et al. *Eur. J. Biochem.* **233**, 293-298 (1995).


Danielson, J., Jarvet, J. et al. *Biochemistry.* **43**, 6261-6269 (2004).



Residues 16-23
Phe-19 docked into CD
0.60 kcal/mol

Residues 16-23
Phe-20 docked into CD
-2.37 kcal/mol



- 
- Possible to derivatize a cyclodextrin so that it is an enzyme that acts on phenyl rings
 - This would prevent aggregation in the brain

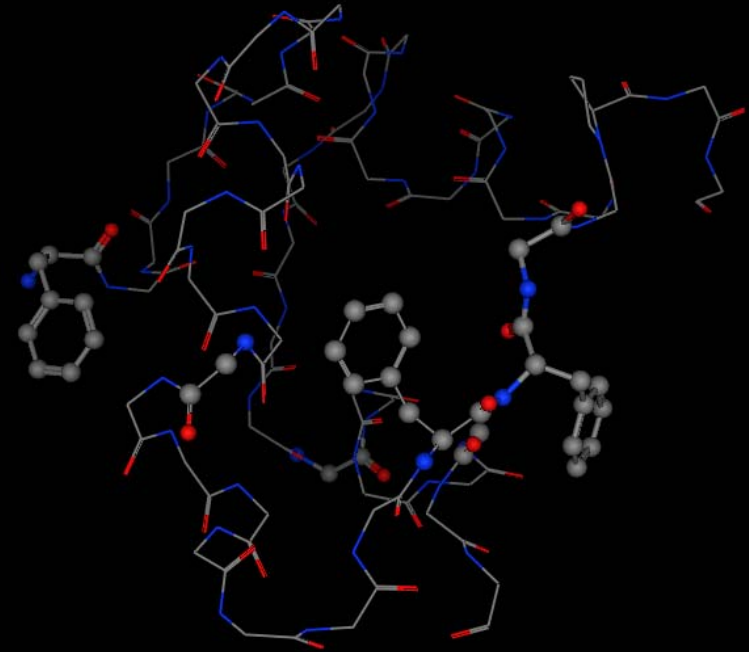


Breslow, R. *Acc. Chem. Res.* **28**, 146-153 (1995).

Danielson, J., Jarvet, J. et al. *Biochemistry.* **43**, 6261-6269 (2004).

Insulin

- Similar structure is found in insulin
- May also be possible to sequester insulin using cyclodextrins



Conclusion



- Constructed models of derivatized cyclodextrins and calculated energies with force fields
- Derivatization with zonulin peptides results in stable binding with flurbiprofen
- A cyclodextrin binds quite well to the Phe-20 residue of the amyloid beta protein



Acknowledgements



- Dr. Jayne Kapur
- NIST Computational Chemistry Group
- AIP/SPS Internship Program



Notes



- Solubilities
 - β -CD: 18 mg/mL
 - HP- β -CD: 500 mg/mL
- Bradykinin
 - arg-pro-pro-gly-phe-ser-pro-phe-arg
 - Dilates blood vessels in brain






- Zonulin peptides

- Open tight junctions in brain
- Closely related to zonula occludens toxin from cholera
- Zonulin 1: asn-gln-arg-pro-pro-pro-ala-gly-val-thr-ala-tyr-asp-tyr-leu-val-ile-gln
- Zonulin 3: val-thr-phe-tyr-thr-asp-ala-val-ser



- 
- Amyloid precursor protein (APP) → β -secretase → γ -secretase → $A\beta(1-40)$ peptide
 - Flurbiprofen
 - Interferes with γ -secretase to reduce $A\beta$ peptide levels
 - In phase 3 of clinical trials

