Kolloquium Visualization of molecule surface dynamics

Masterarbeit Raphael Menges

Contents

- Motivation
- Surface Extraction
- Ascension
- Layers
- Residue Surface-Proximity
- Evaluation
- Conclusion

Motivation

- Surface of molecules defines binding behavior
- Folding simulation data hard to interpret
- Available software does not support scientists in surface dynamics visualization and analysis

Surface Extraction Methods



Solvent Accessible Surface



Surface Atom Extraction based on Extended Spheres





SAEES

Optimization of Cutting Face List



```
SAEES
```

Check Endpoint of Faces Intersections





Compute Shader Program

Per Molecule (OpenGL)



Per Atom (GLSL)

SAEES Results







Ascension **Results**



Ascension Results





Layers **Results**



Residue Surface-Proximity

- Proteins consist of amino acid residues
- Scientists are interested in their movement
- Average layer membership over folding simulation tracked
- Surface-proximity or coreproximity per amino acid computed



Evaluation

Evaluation Ascension





Evaluation Residue Surface-Proximity



Evaluation Residue Surface-Proximity



Conclusion

- Supports scientists in molecule design
- Surface dynamics visualization and analysis is novel research area
- Interdisciplinary field with potential

Thank you for your attention!