Source:

1 | Preliminary Research

1.1 | Sources

`https://www.frontiersin.org/articles/10.3389/fchem.2019.00540/full`

1.2 | Notes

1.2.1 | Folding Simulation Methods

1. all-atome molecular dynamics (MD)

    - Obtains all desired information regarding the kinetics and thermodynamics
    - very slow (supercomputers -> microseconds of simulation)