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Biocomplexity Faculty Search
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To Whom it May Concern:

I am writing this letter in strong support of Chenglong Li. In the past two years, I have had the pleasure of advising on his work on the design of inhibitors for AICAR transformylase. With this work, he has greatly extended the functionality of our docking method AutoDock, demonstrating that it is an effective tool for virtual database screening. His hierarchical approach will become a standard for use of AutoDock in computer-aided drug design.

In the two years that I have worked with him, Dr. Li has proven himself to be a thorough and efficient researcher. He has worked effectively in a collaboration between chemists, experimental biologists and computational chemists, combining diverse information from chemistry and from computational simulation in the search for better inhibitors. He has been very successful in his past work, showing a good facility for finding the appropriate methods for each new challenge. I fully expect that he will show continued success in the field of computational chemistry.

Sincerely,

David S. Goodsell