

## Editorial: Special Issue on “Evaluation of Computational Methods”

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This issue comprises one of a continuing series of “special” issues of JCAMD which target a topic of particular interest. In this issue, guest editors Ajay Jain and Anthony Nicholls tackle a topic in which they are well versed: how can we tell whether or not one method of ligand/protein docking or ligand based virtual screening is “better” than another. They’ve assembled an impressive list of authors to provide 11 papers dealing with different aspects of the problem. Ajay and Anthony also provide a guest editorial in which they summarize the results of the authors and make a series

of recommendations on how to conduct and report those studies.

How do you see this problem? Do you feel it has been adequately addressed? Do you have a conflicting point of view? If so, please let us know. We’ve instituted “letters to the editor” to allow our readers to make their viewpoint known. Simply go to the JCAMD submission web page, follow directions for submission of a new contribution and select “Letters to the Editor” as the “article type.” We’ll do our best to let your viewpoint be heard.

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Guest Editors: Ajay Jain, Anthony Nicholls

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