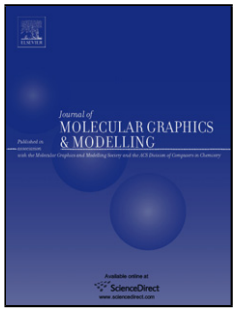

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Homology modeling of a Class A GPCR in the inactive conformation: a quantitative analysis of the correlation between model/template sequence identity and model accuracy

Stefano Costanzi,[a,b*] Matthew Skorski,[a] Alessandro Deplano,[a] Brett Habermehl,[a] Mary Mendoza,[a] Keyun Wang,[a] Michelle Biederman,[a] Jessica Dawson,[a] and Jia Gao[a]

[a]*Department of Chemistry, American University, Washington, DC 20016, USA*

[b]*Center for Behavioral Neuroscience, American University, Washington, DC 20016, USA*

*Address correspondence to:

Stefano Costanzi, Ph.D.

Department of Chemistry,

American University

4400 Massachusetts Ave, NW, 20016 Washington, DC, USA

Tel: +1-202-885-1722

Email: costanzi@american.edu