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Author: Stefano Costanzi Matthew Skorski Alessandro Deplano Brett Habermehl Mary Mendoza Keyun Wang Michelle Biederman Jessica Dawson Jia Gao

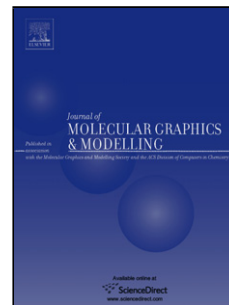
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**Homology modeling of a Class A GPCR in the inactive conformation:  
a quantitative analysis of the correlation between model/template  
sequence identity and model accuracy**

Stefano Costanzi,<sup>a,b\*</sup> Matthew Skorski,<sup>a</sup> Alessandro Deplano,<sup>a</sup> Brett Habermehl,<sup>a</sup> Mary Mendoza,<sup>a</sup> Keyun Wang,<sup>a</sup> Michelle Biederman,<sup>a</sup> Jessica Dawson,<sup>a</sup> and Jia Gao<sup>a</sup>

<sup>a</sup>*Department of Chemistry, American University, Washington, DC 20016, USA*

<sup>b</sup>*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

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