A multi-scale approach for deciphering HIV infection

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Publication date
2011

Citation for published version (APA):

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Publications

Published:


Conferences:


143
G. Ertaylan and P.M.A. Sloot: A complex automata model of HIV-1 co-receptor tropism: Understanding mutation rate pressure, in Reviews in Antiretroviral Therapy, Washington D.C., USA, December 2007

Submitted:

Ilkay Altintas; Manish Kumar Anand; Adam Belloumi; Gokhan Ertaylan; Bartosz Balis; Marian Bubak; Peter M.A. Sloot: Collaborative Provenance for Workflow-Driven Science - A Position Paper. Future Generation Computer Systems

In preparation:

G. Ertaylan; P.M.A. Sloot: An evolutionary model on HIV’s co-receptor switch.