

## Descriptor- and Fragment-based QSAR Models for a Series of *Schistosoma mansoni* Purine Nucleoside Inhibitors

Humberto F. Freitas,<sup>a</sup> Matheus P. Postigo,<sup>b</sup> Adriano D. Andricopulo<sup>\*,b</sup> and  
Marcelo S. Castilho<sup>\*,a</sup>

<sup>a</sup>Departamento do Medicamento, Faculdade de Farmácia, Universidade Federal da Bahia,  
40170-115 Salvador-BA, Brazil

<sup>b</sup>Laboratório de Química Medicinal e Computacional, Instituto de Física de São Carlos,  
Universidade de São Paulo, 13566-970 São Carlos-SP, Brazil

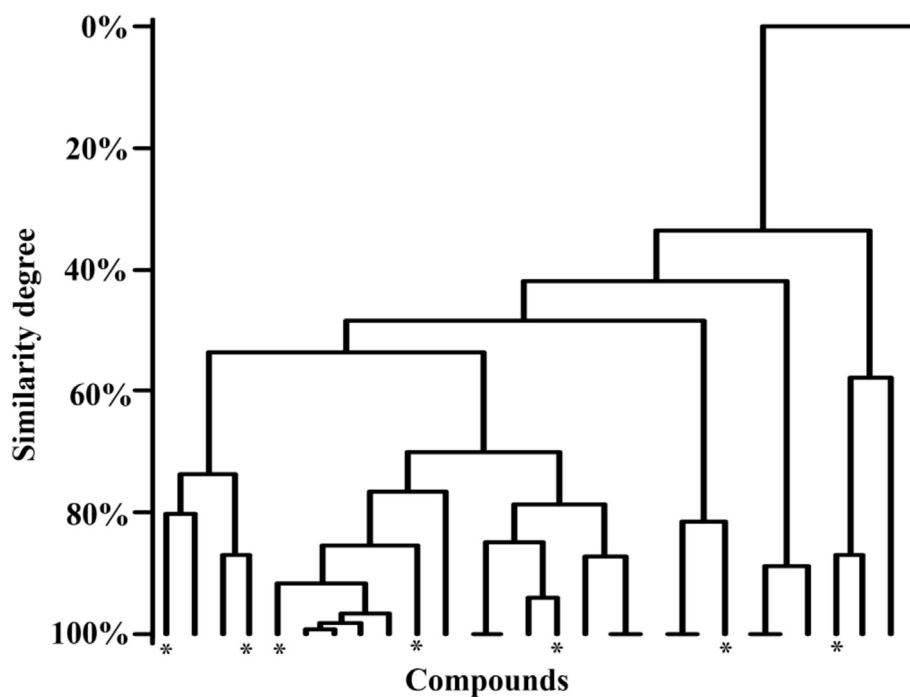


Figure S1. Hierarchical cluster analysis of all deazapurine analogs employed in QSAR model development. Test set compounds are marked with \*.