A QSAR Study of Artemisinin Analogs

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What is Artemisinin?



- Treatment for drug resistant P. falciparum
- Rapid clearance of cerebral malaria
- Some analogs are neurotoxic

Mode of Action

- Two modes are theorized
 - The active conformation is simply the energetically minimized conformation
 - Artemisinin complexes with hemin via the peroxy moiety to generate bio active radicals^a

^aS.R. Meshnick, Trans. R. Soc. Trop. Med. Hyg. 1994, 88, 31-32

Backbone Structures



Backbone Structures



Miscellaneous Structures



Only 2 acyclic stuctures present

Handling the Compounds

The dataset contained several enantiomers:

The dataset also contained charged species

Previous Study

- CoMFA & Hologram QSAR^a (seems similar to atom pair fragment descriptors)
- ho r^2 & q^2 was used to assess quality of models
- Model goodness of fit was characterized by the ration of standard error to activity range (s/AR)
- Part of the study also included racemates.

Endpoint Details

- All the analogs are assumed to act via similar mechanisms
- All were tested using the same assay in vitro against chloroquine resistant & mefloquine sensitive *P. falciparum* W-2 clone.
- Due to interday variations of IC₅₀ for artemisinin, relative activity (RA) was taken as the property under study

Relative Activity

RA is calculated using

 ${\rm RA} = \frac{{\rm IC}_{50} \text{ of artemisinin}}{{\rm IC}_{50} \text{ of analog}} \times \frac{{\rm MW} \text{ of artemisinin}}{{\rm MW} \text{ of analog}}$

log RA is used in the study

Plan of Action

- Structures were drawn in with Corina
- Optimization with PM3 was skipped
- Ignore enantiomeric pairs and cap charged species with hydrogens
- MPOLR, MRFRAC & GEOWIND could not be used

Plan of Action

- Initially carry out a study with topological indices
- Atom pair fragments may be something to look into
- Would hydrophobicity be useful in this situation?