

## 一种新的De Novo生物活性分子设计方法

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收稿日期 修回日期 网络版发布日期 接受日期

**摘要** 由药效团进行虚拟活性结构生成与3D-QSAR模型相结合, 筛选出有前途的结构多样性的化合物, 并从中寻找活性先导化合物,

是一种新的分子设计方法。采用这种方法对抗小麦赤霉病类含氟农药进行了研究, 共生成了53

个虚拟活性结构, 通过3D-QSAR模型筛选出其中10个活性较高的结构,

在活性最高的化合物基础上进行了结构修饰,

得到了活性更高且毒性较低的理想化合物。研究表明这种方法能突破原模型化合物结构模式的局限, 可以找到结构新颖的活性先导化合物, 是一种非常有前途的分子设计方法, 而且具有较高的筛选效率。

**关键词** [药效团](#) [分子设计](#) [生物活性](#) [筛选](#) [结构生成](#) [计算机应用](#)

分类号 [06-39](#)

## A new method for De Novo bio-active molecular design

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**Abstract** A new De Novo method for bio-active molecular design is presented, which combines virtual bio-active structural generation with 3D-QSAR study. This method could generate a lot of highly diverse molecules and find bio-active lead compounds. the method is illustrated through a study on a set of fluorine -containing pesticides for anti-gibberella. With the constraints of the pharmacophore obtained by DISCO, 53 virtual bio-active structures were generated, and their anti-gibberella activities were predicted by CoMFA. The 10 most active compounds were selected and screened by 3D-QSAR. The first one was investigated in depth by modifying and simplifying its structure. the results showed that the method was a feasible means for bio- active molecular design. It had high screen efficiency.

**Key words** [MOLECULAR DESIGN](#) [BIOLOGICAL ACTIVITY](#) [SELECTION](#) [COMPUTER APPLICATIONS](#)

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