

## CASE STUDY BASED PUZZLE

BP 807 ET: COMPUTER AIDED DRUG DESIGN

## UNIT 2: COMFA AND COMSIA

## Case Study Puzzle: 3D-QSAR in Drug Design

Topic: CoMFA and CoMSIA Approaches

## Case Scenario:

A pharmaceutical research team is working on a new series of benzimidazole derivatives as potential anti-inflammatory agents.

Ten compounds (A–J) were synthesized with different substitutions at R<sub>1</sub>, R<sub>2</sub>, and R<sub>3</sub> positions. Biological activity was measured as IC<sub>50</sub> (μM) against a target enzyme.

The researchers decided to apply 3D-QSAR techniques to understand the structure–activity relationship (SAR) and guide further optimization.

They used:

- CoMFA to analyze steric and electrostatic fields
- CoMSIA to analyze steric, electrostatic, hydrophobic, H-bond donor, and H-bond acceptor fields

## Puzzle 1: Method Selection Challenge

## 🔍 Clue:

One compound (Compound F) showed excellent activity despite having a bulky hydrophobic group at R<sub>2</sub>.

## Puzzle Question:

Which 3D-QSAR method is more suitable to explain the contribution of hydrophobic interactions in this case?

Options:

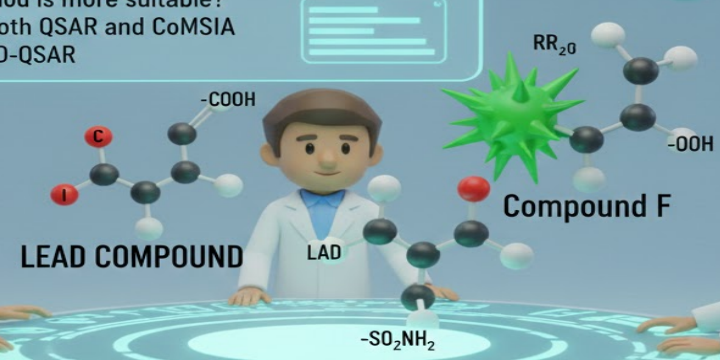
- A. CoMFA only
- B. CoMSIA only
- C. Both CoMFA and CoMSIA
- D. 2D-QSAR

📖 Hint: Think about which method includes hydrophobic field descriptors.

### Puzzle 1: Method Selection Challenge

**Clue:** Compound F showed excellent activity despite bulky hydrophobic group at R<sub>2</sub>

**Puzzle Question:** Which 3D-QSAR method is more suitable?  
 C. Both QSAR and CoMSIA  
 D. 2D-QSAR



**LEAD COMPOUND**      **LAD**      **Compound F**

Puzzle Questions	Solutions
1. Drug Design Approach? 2. Type of the Bioisoterism? 3. CoMSIA only 4. One Advantage?	1. 3D-QSAR Method? 2. CoMSIA only C. CoMSIA only 3. Includes Hydrophobic Field

## Puzzle 2: Grid & Alignment Mystery

### Clue:

During model development, the prediction accuracy dropped significantly when compound alignment was changed.

### Puzzle Question:

Which factor is most critical in CoMFA and CoMSIA for obtaining a reliable model?

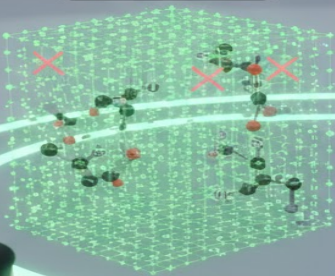
Options:

- A. Molecular weight
- B. Grid spacing
- C. Molecular alignment
- D. Number of compounds


☞ Hint: 3D-QSAR models depend heavily on spatial consistency.

# Grid & Alignment Mystery

**Bad Alignment**



**Good Alignment**



Which factor is most critical in CoMFA and CoMSIA?

A. Molecular weight

B. Grid spacing

C. Molecular

D. Number of compounds

**C. Molecular Alignment**

## Puzzle 3: Contour Map Decoder

### Clue:

The CoMFA steric contour map shows:

- Green contours near R<sub>1</sub>
- Yellow contours near R<sub>3</sub>

### Puzzle Question:

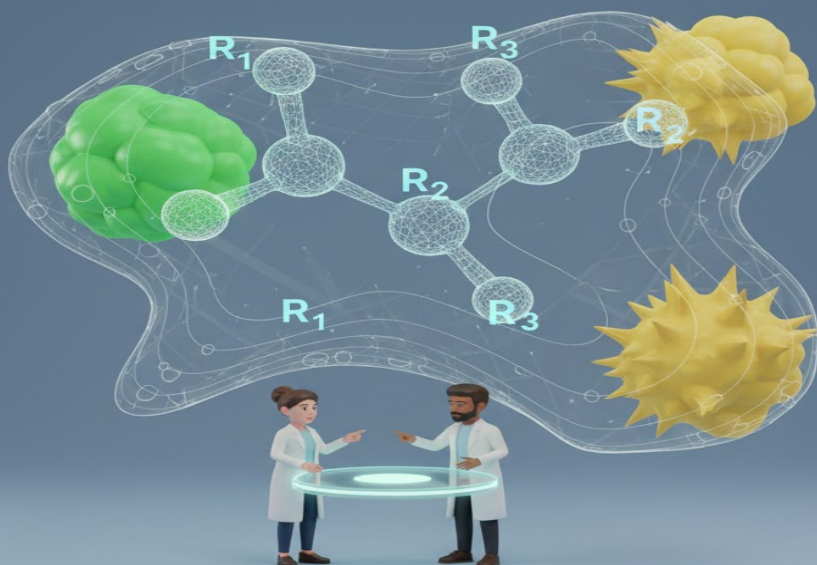
What modification is most likely to increase activity?

Options:

- A. Add bulky group at R<sub>3</sub>
- B. Remove substituent at R<sub>1</sub>
- C. Add bulky group at R<sub>1</sub>
- D. Add electronegative group at R<sub>3</sub>

☞ Hint: Green = favorable steric bulk, Yellow = unfavorable steric bulk.

### Puzzle 3: Contour Map Decoder



 **Puzzle Question** is most likely to increase activity?

- A. Add bulky group at  $R_3$  ☐
- B. Remove substituent at  $R_1$  ☐
- C. Add bulky group at  $R_1$  ☐
- D. Add electronegative group at  $R_3$  ☐



#### Hint

Green = favorable steric bulk,  
Yellow, unfavorable steric bulk