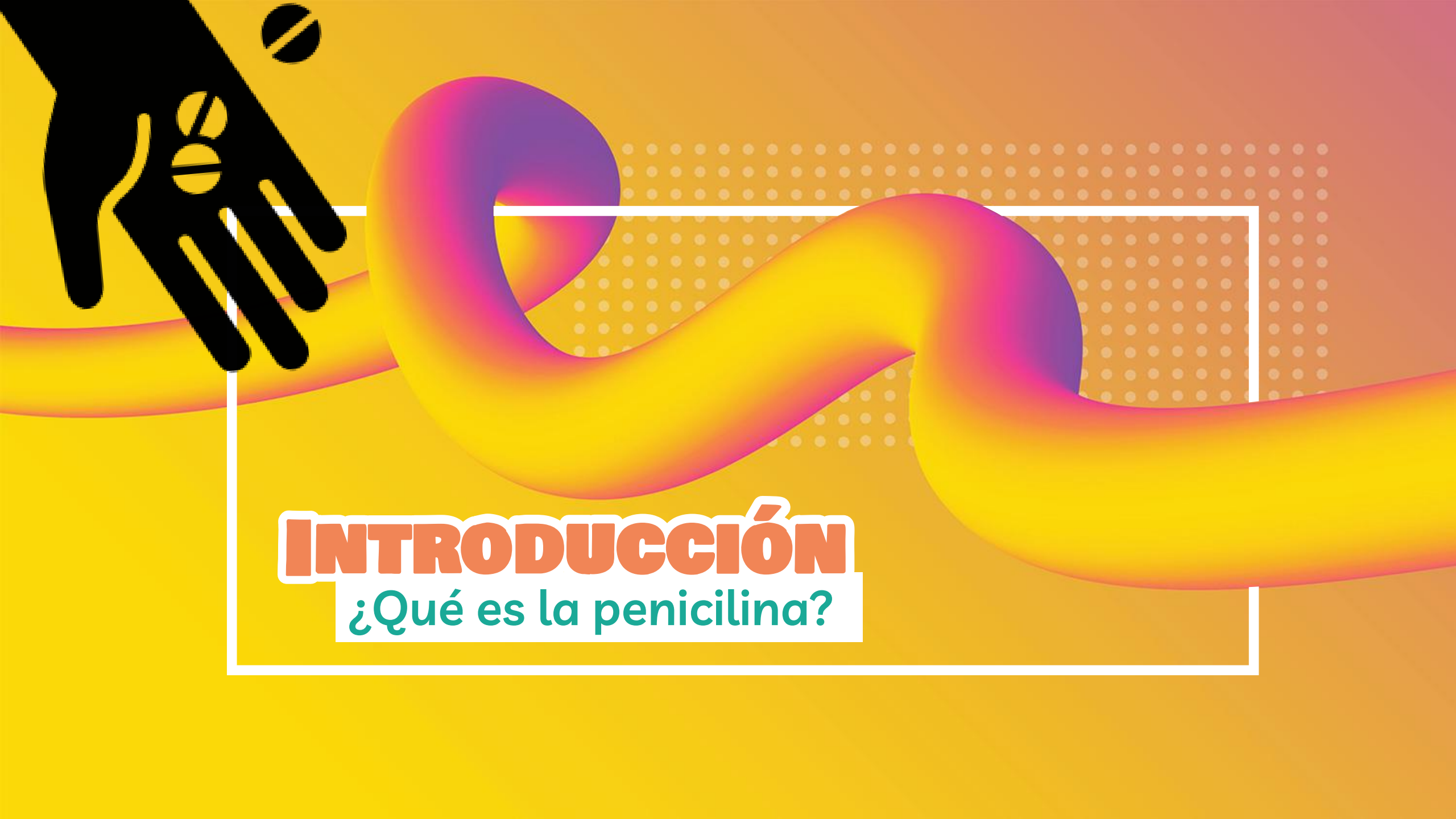


# Complemento contra resistencia bacteriana a la meticilina



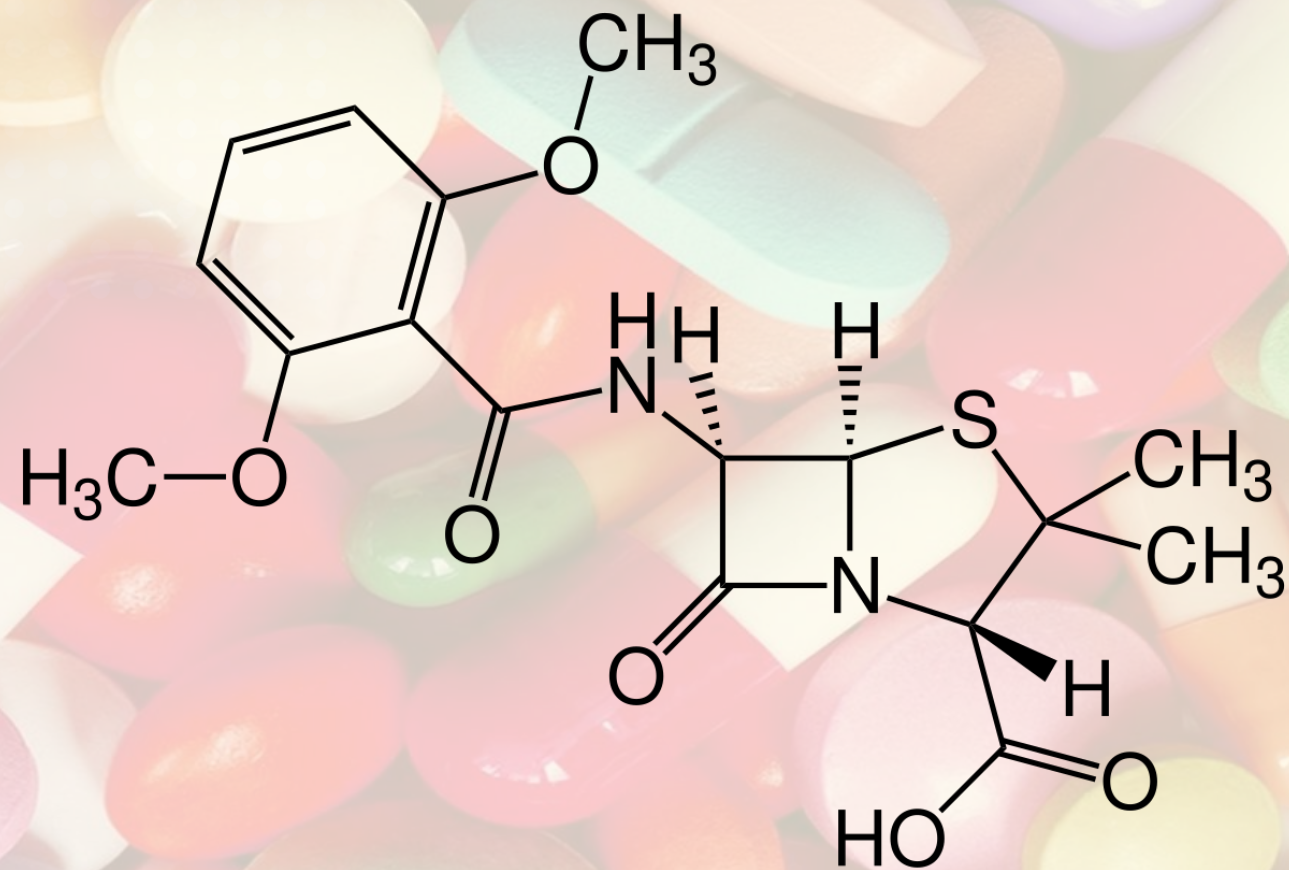
Clubes de Ciencia  
México



# **INTRODUCCIÓN**

¿Qué es la penicilina?





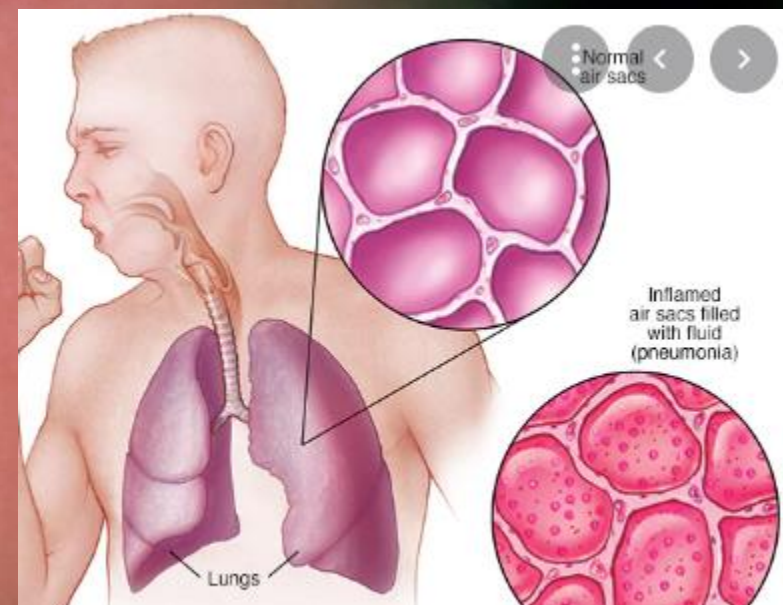


# PROBLEMÁTICA



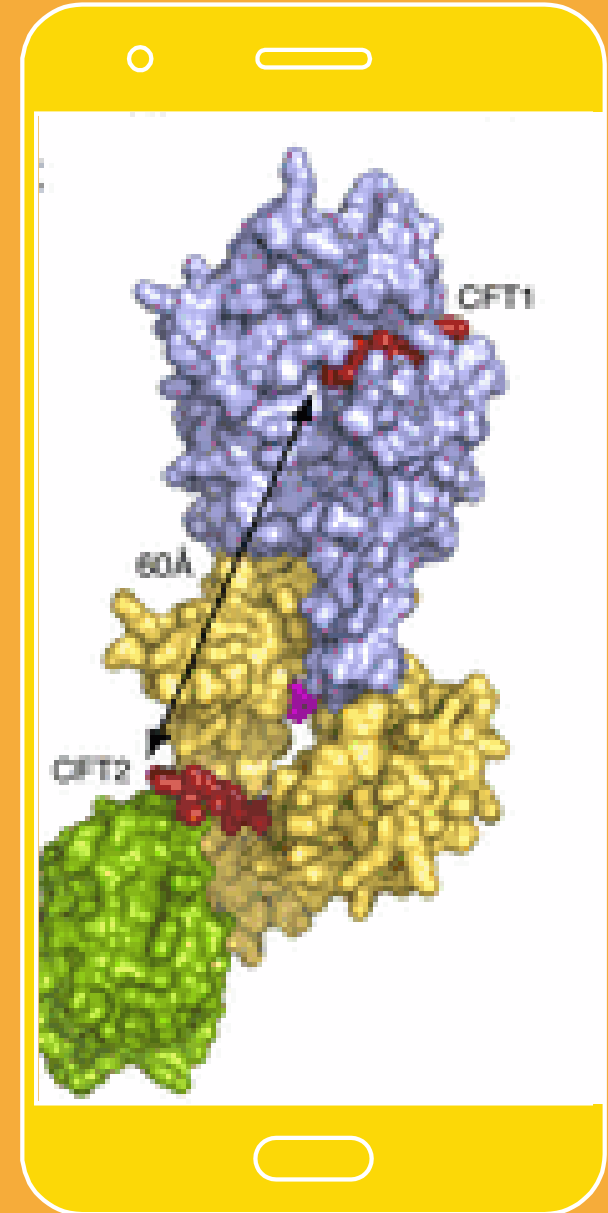
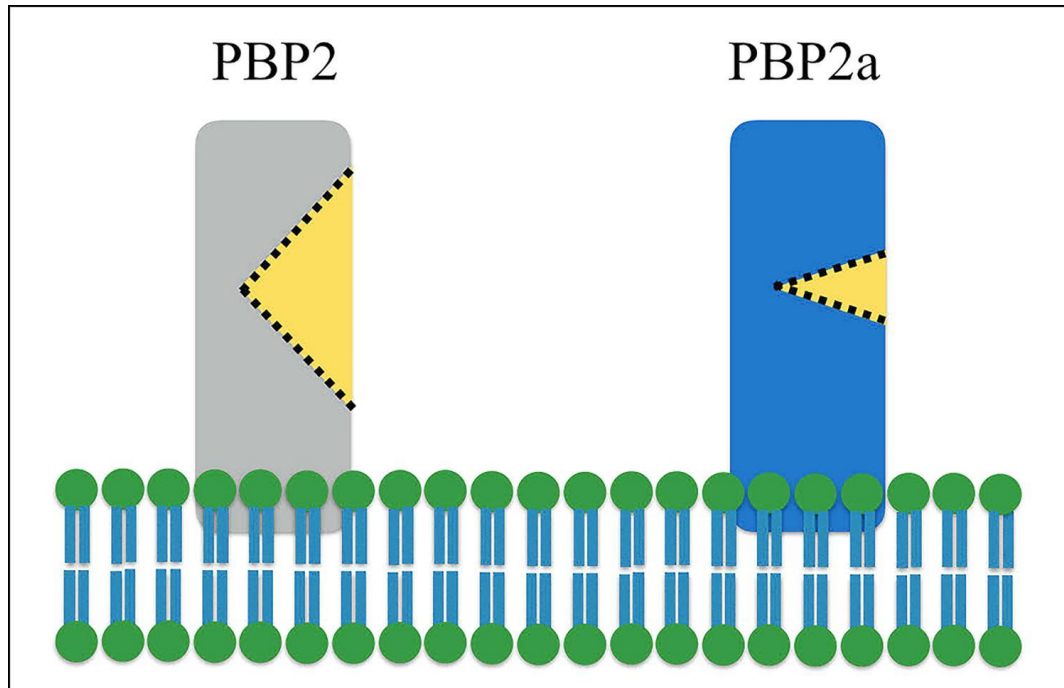
¿Y cuál es el problema con esta bacteria?





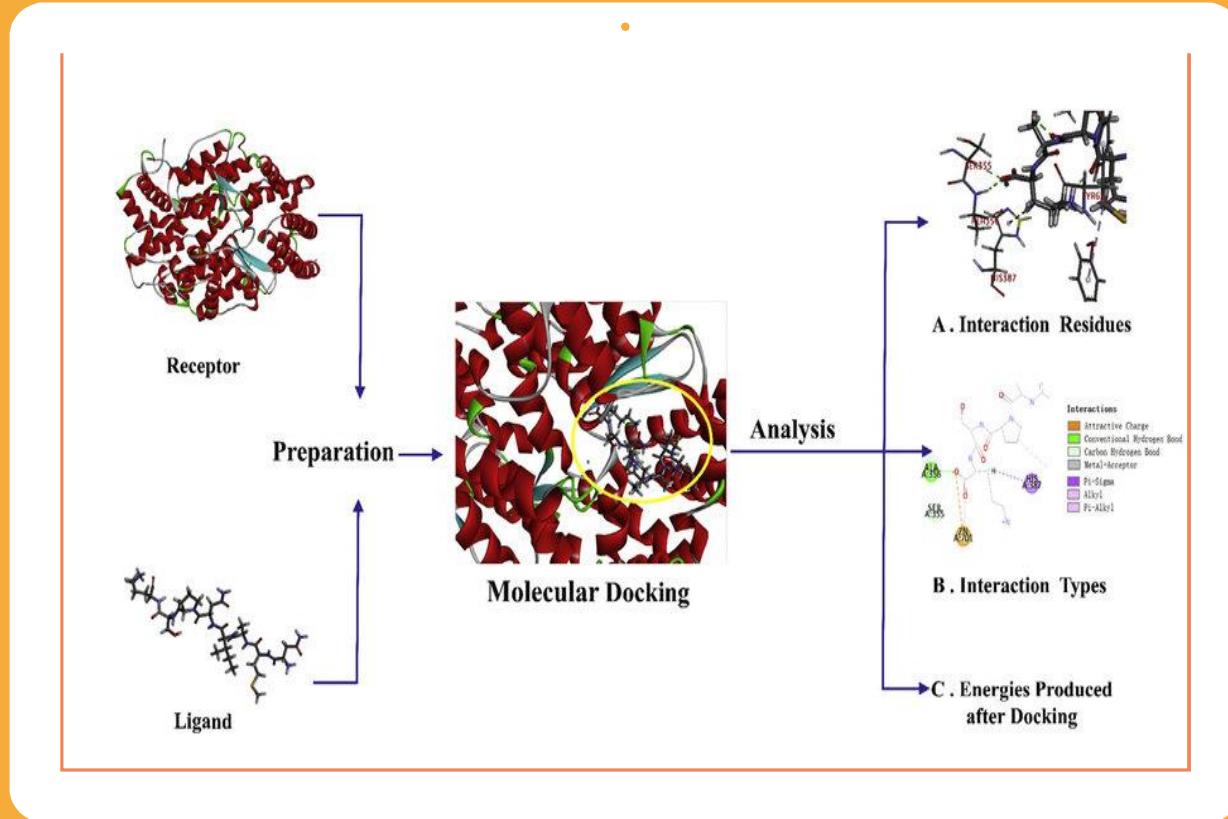
# SOBRE LA PBP2A

¿Qué es la proteína PBD2a?

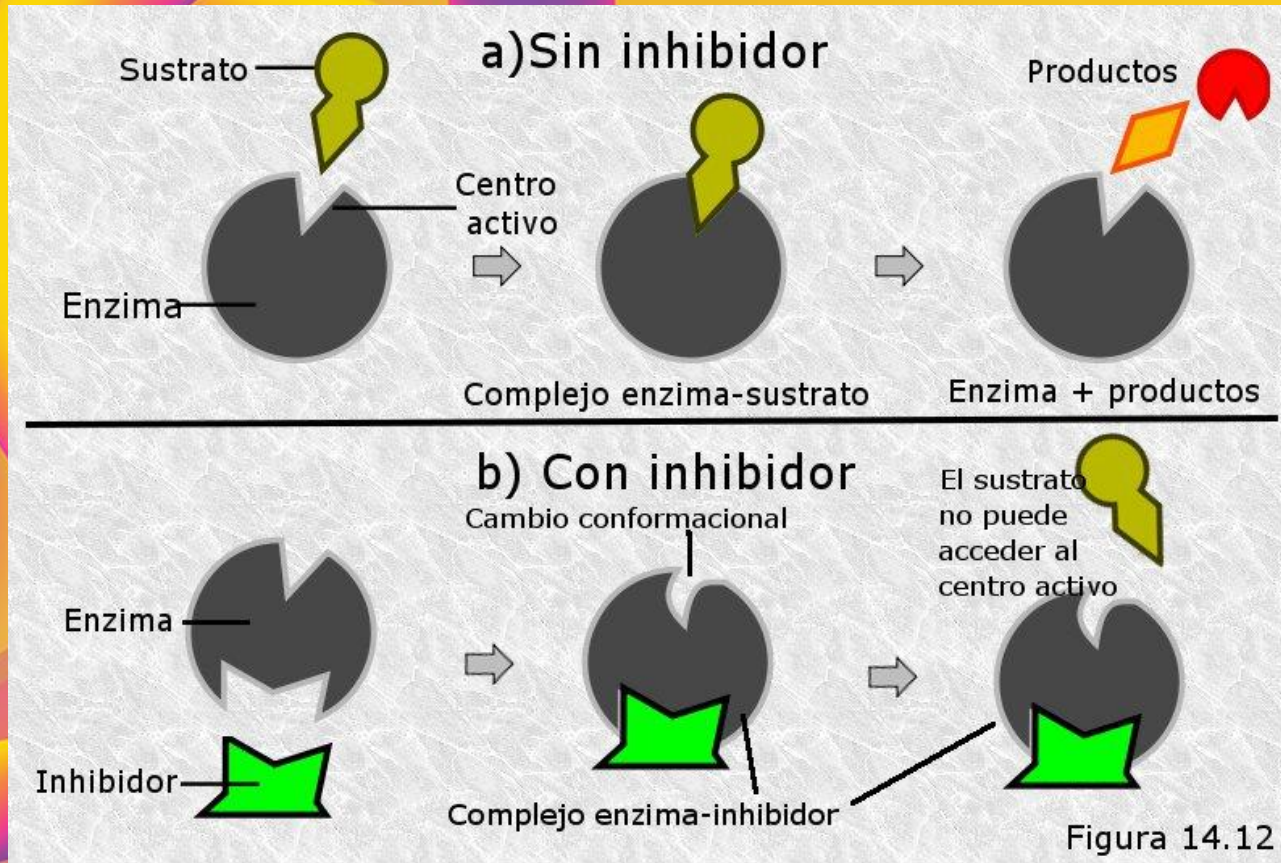


# SOBRE EL DOCKING DE PBP2A CON INHIBIDORES

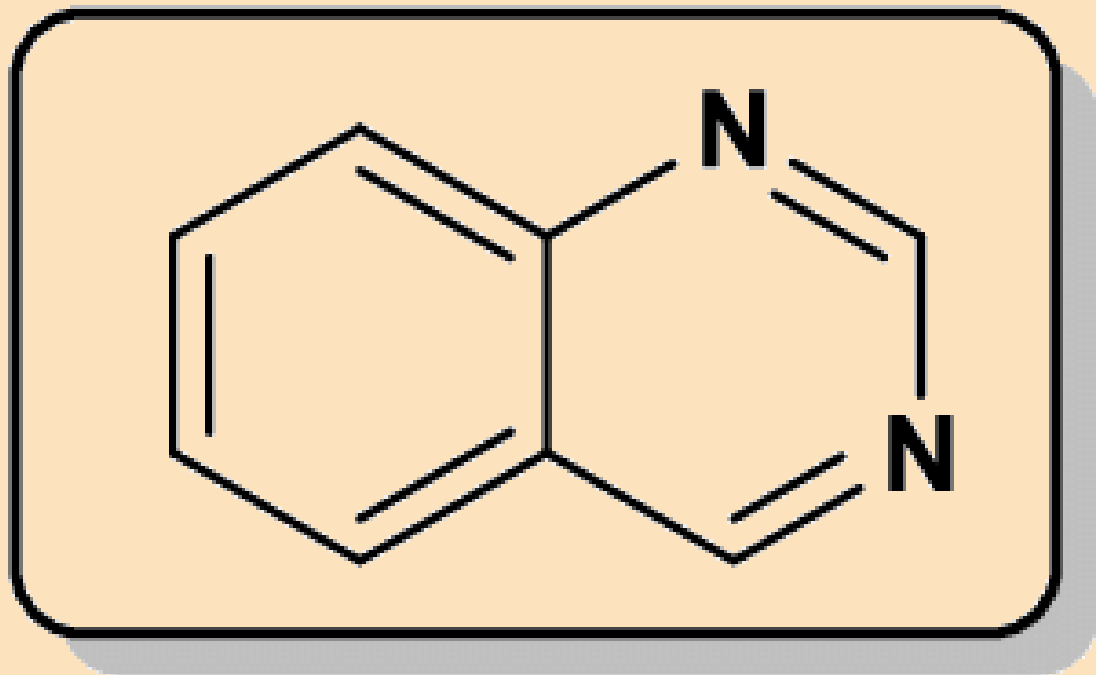
¿Inhibidores...?











AutoDock  
Tools-1.5.

6



CADD-1.5.

6



PMV-1.5.6



Vision-1.5.

6

¿Qué  
proponemos  
nosotros?

- Proponemos emplear el inhibidor Quinazolinona alostérica como inhibidor de la expresión de la enzima.
- Se utiliza pyMol para la representación gráfica de las enzimas y moléculas y para la simulación del docking AutoDockTools

Python Shell

File Edit Debug Options Windows Help

ADT4.2 Ligand Flexible Residues Grid Docking Run Analyze

DashBoard AniMol Tools

Sel: CMD

All Molecules  
Current Selection  
6q9n (1)  
1mw1 (2)

Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB13:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB14:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB15:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB16:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB17:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB18:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB19:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB20:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB21:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB22:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB23:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB24:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB25:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB26:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB27:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB28:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB29:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB30:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB31:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB32:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB33:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB34:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB35:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB36:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB37:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB38:0  
Sorry, there are no Gasteiger parameters available for atom Inset (2):E:BOB39:0

Exception in Tkinter callback  
Traceback (most recent call last):  
File "D:\Autodock\lib\lib-tk\Tkinter.py", line 1403, in \_\_call\_\_  
return self.func(\*args)  
File "D:\Autodock\lib\site-packages\AutodockTools\autocglCommands.py", line 44  
79, in chooseMolecule\_cb  
mol = self chooser.getMolSet()  
File "D:\Autodock\lib\site-packages\AutodockTools\guiTools.py", line 178, in getMolSet  
if self.mode=='single': return mols[0]  
File "D:\Autodock\lib\site-packages\AutodockTools\guiTools.py", line 28, in \_\_getitem\_\_  
def \_\_getitem\_\_(self, i): return self.data[i]  
IndexError: list index out of range

Mod: None Time: 3.127 Selected: 1mw1 (2) Done 100% Spin off FR: 47.6

ADT4.2 Ligand Flexible Residues Grid Docking Run Analyze

DashBoard AniMol Tools

Sel: CMD

All Molecules  
Current Selection  
6q9n (1)  
1mw1 (2)

Mod: None Time: 51.305 Selected: 1mw1 (2) Done 100% Spin off FR: 43.5

AutoDockTools

File 3D Graphics Edit Select Display Color Compute Hydrogen Bonds Grid3D Help

ADT4.2 Ligand Flexible Residues Grid Docking Run Analyze

DashBoard AniMol Tools

Sel: CMD

All Molecules  
Current Selection  
6q9n (1)  
1mw1 (2)

computeGasteigerGC WARNING

Total gasteiger charge added = 19.9646

Acceptar

Mod: None Time: 3.127 Selected: 1mw1 (2) Done 100% Spin off FR: 47.6

AutoDockTools

File 3D Graphics Edit Select Display Color Compute Hydrogen Bonds Grid3D Help

ADT4.2 Ligand Flexible Residues Grid Docking Run Analyze

DashBoard AniMol Tools

Sel: CMD

All Molecules  
Current Selection  
6q9n (1)  
1mw1 (2)

addKollmanChargesGC WARNING

Total Kollman charge added = -124.366

Acceptar

Mod: None Time: 0.334 Selected: 1mw1 (2) Done 100% Spin off FR: 47.6