

*XIII Escola de Verão em Química Farmacêutica  
e Química Medicinal*

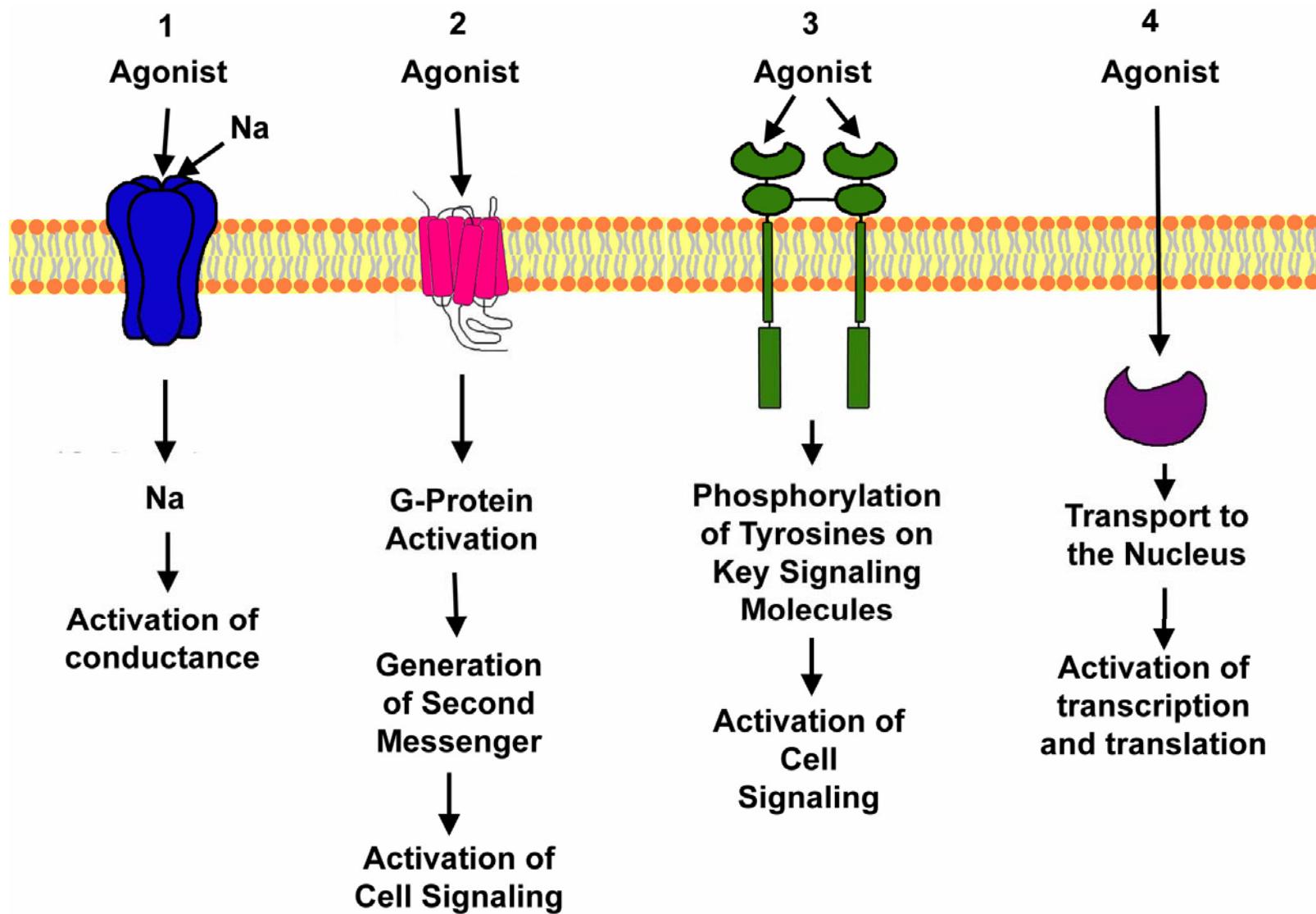
# Princípios Básicos de Farmacologia

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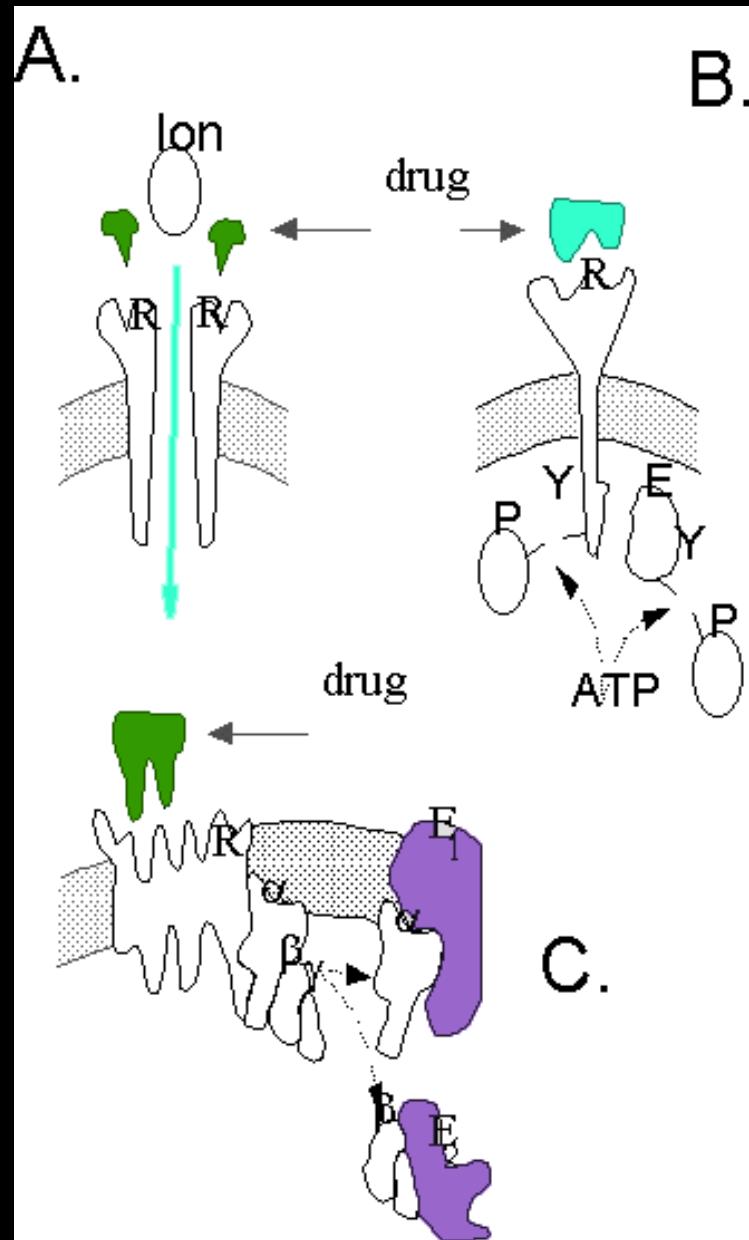
Fevereiro, 2007.

# Receptor Biológico

- Saturabilidade
- Especificidade
- Reversibilidade
- Restabelecimento da função por reconstituição
- Identificação do gene associado



# The pharmacologist's dream!!!!



Droga ou  
fármaco

Sítio de ligação

mecanismo de  
sinalização

ativação/inibição enzimática  
modulação de canais iônicos  
transcrição de DNA...

resposta  
celular

contração, secreção,  
metabolismo...

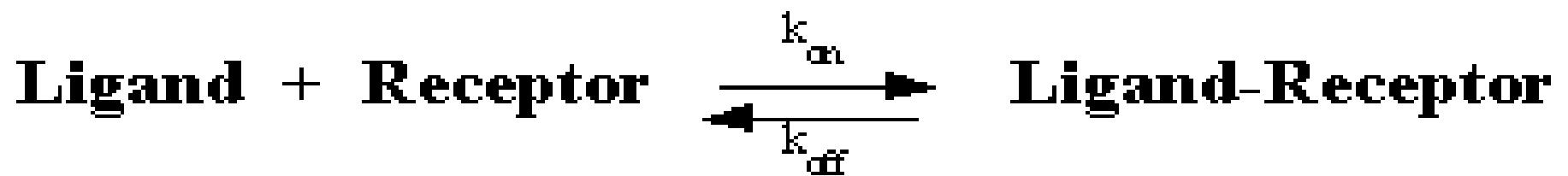
modificação do funcionamento  
do organismo

modificação do funcionamento  
de um órgão

Hill, 1909

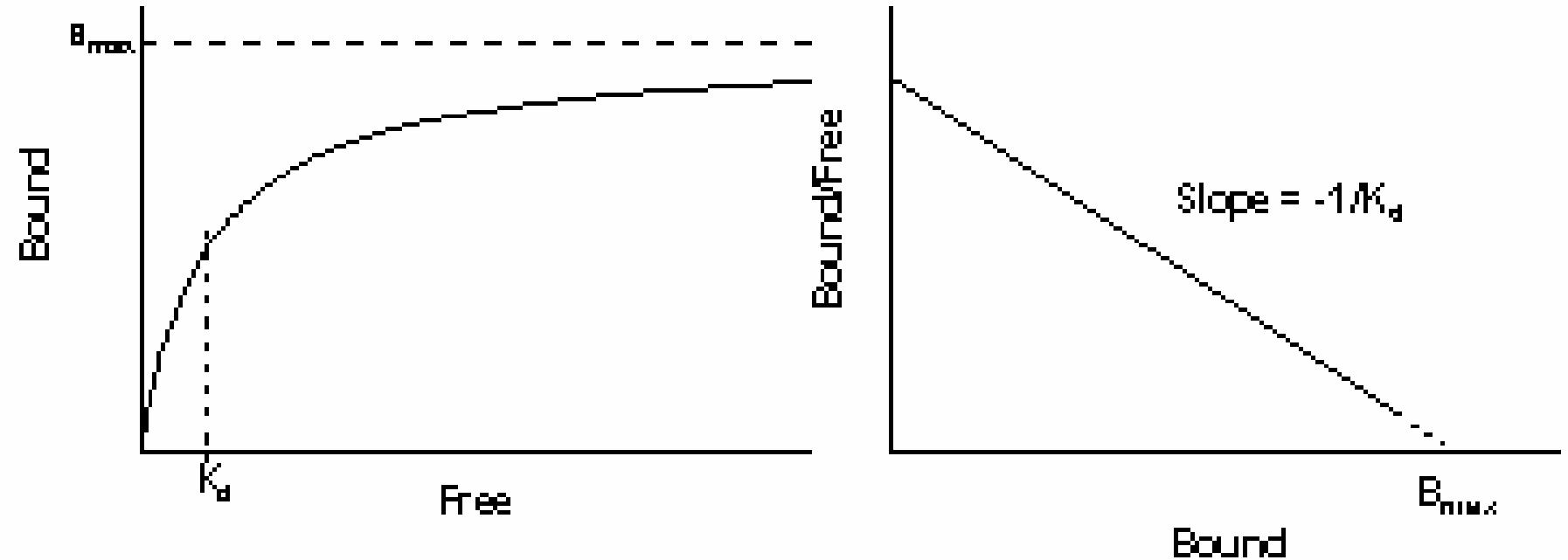
$$\frac{E}{E_{\max}} = \frac{[A]^{nH}}{[A]^{nH} + [A]_{50}^{nH}}.$$

## **Lei da Ação das Massas ( Clark,1926)**



**modelo chave fechadura**

# Modelo do Efeito Máximo

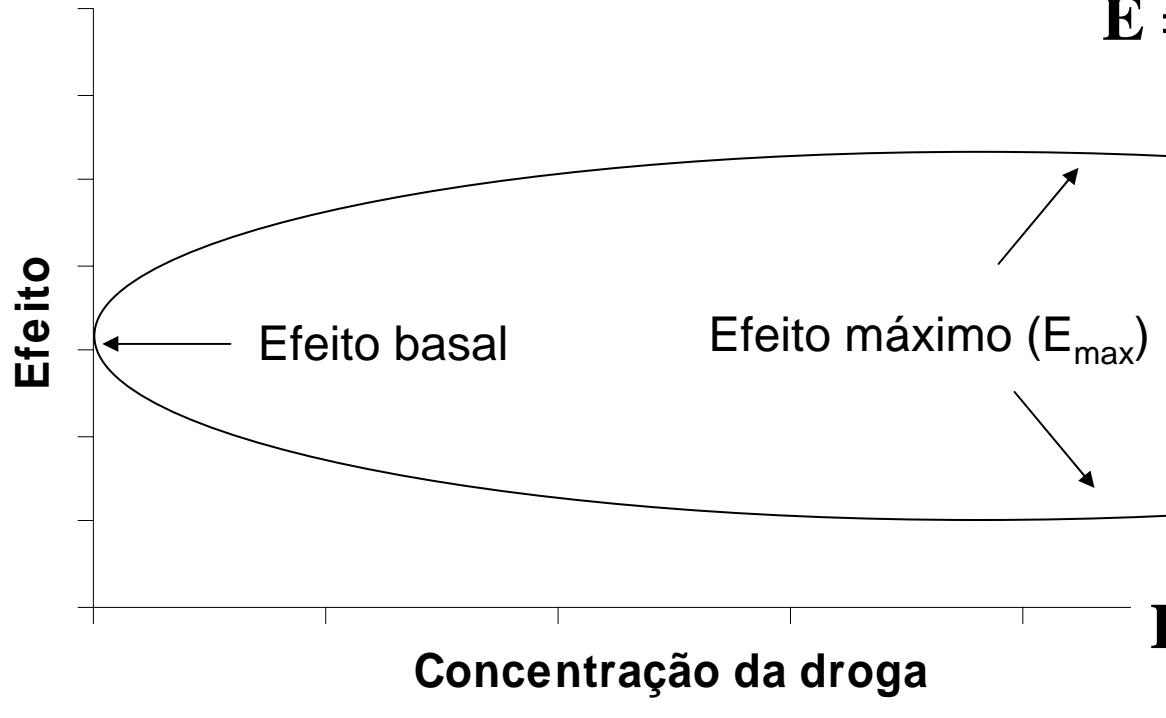


$$E = \frac{[C] \cdot E_{\max}}{CE_{50} + [C]}$$

$$r = \frac{[L]}{K_d + [L]}$$

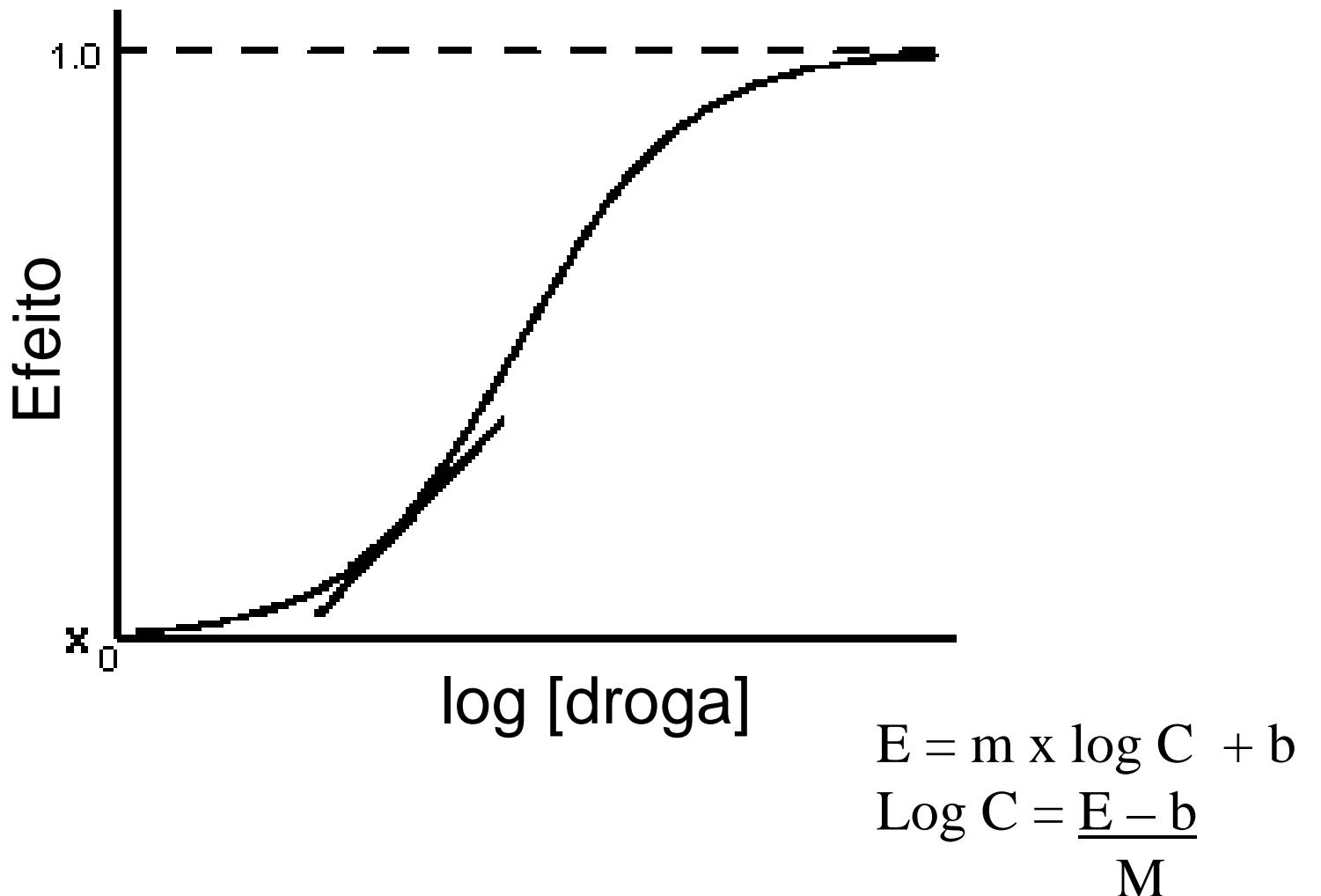
$$E = m \times C + E_0$$

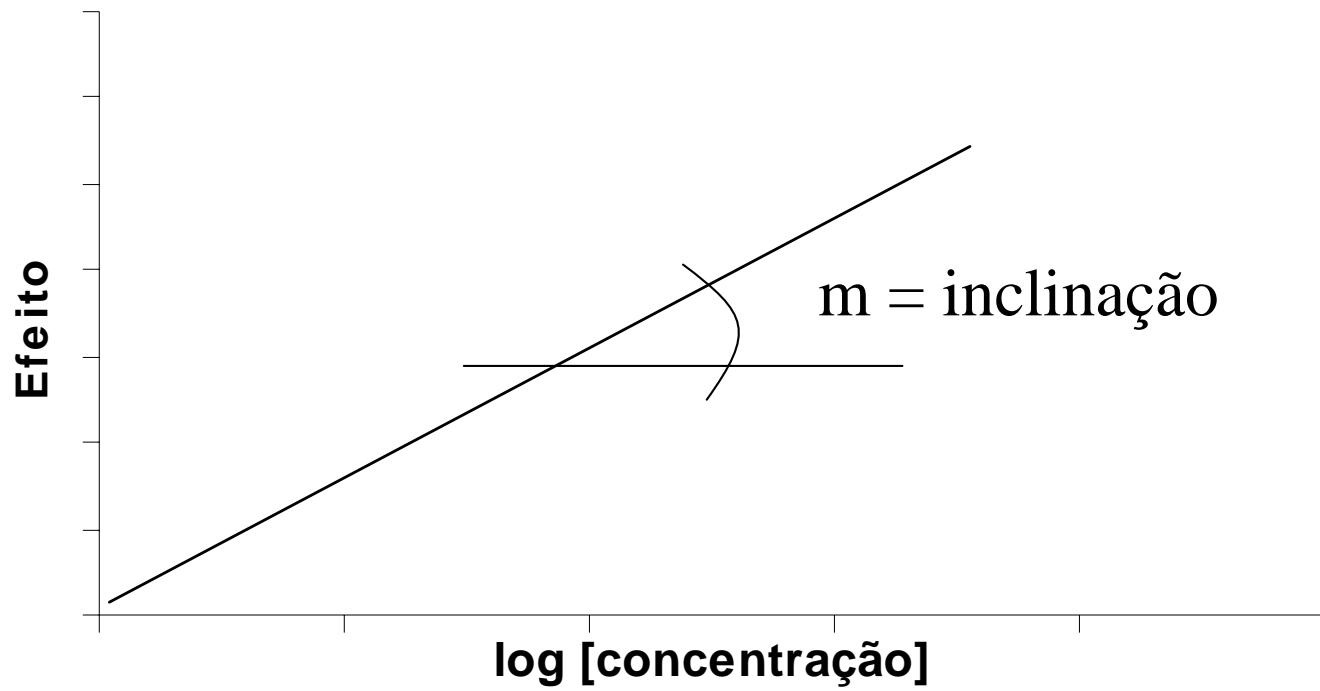
$$E = \frac{E_b - E_{\max} \cdot C}{CE_{50} + [C]}$$



$$E = \frac{E_b - E_{\max} \cdot C}{CE_{50} + [C]}$$

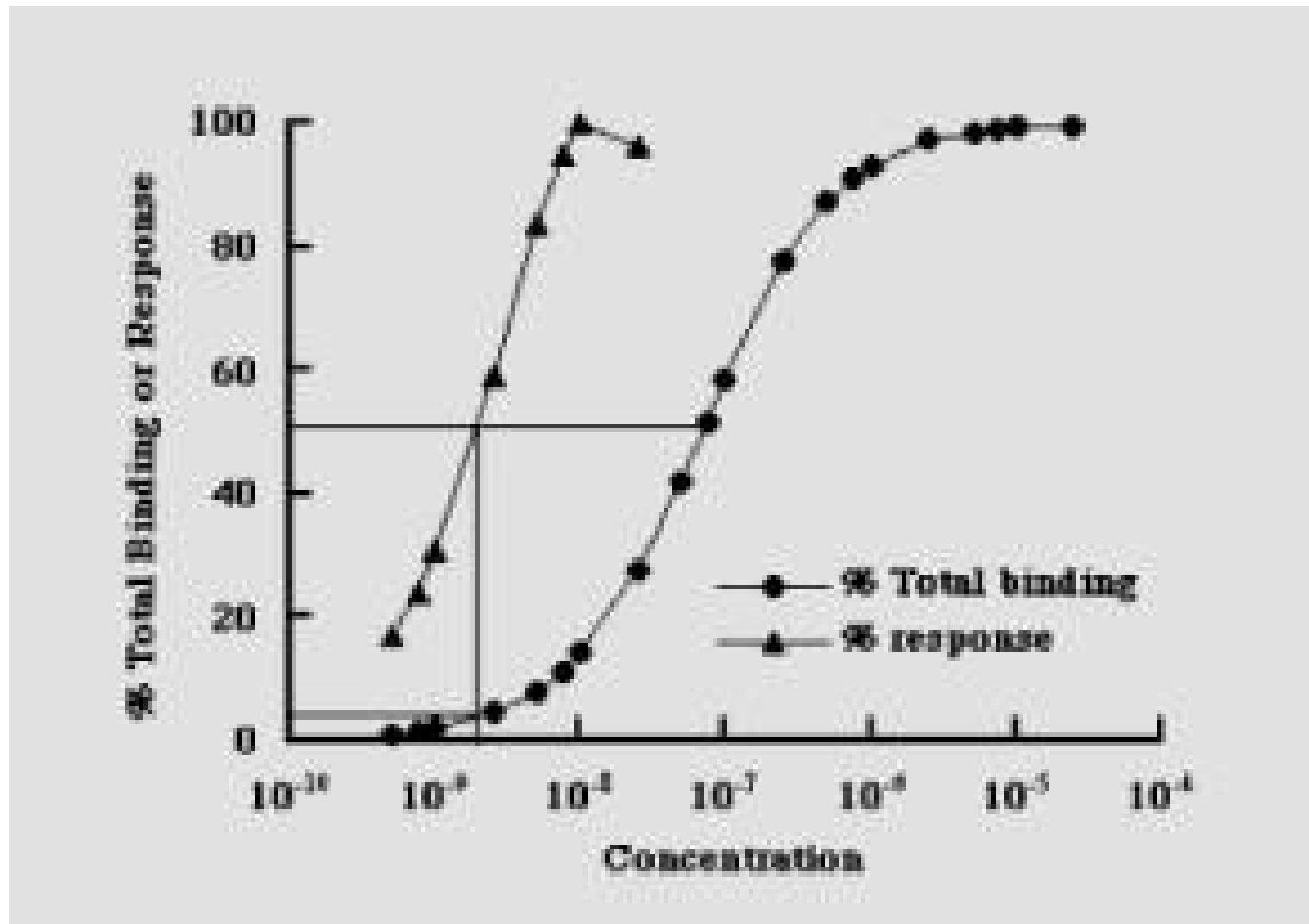
## Log - Sigmoidal





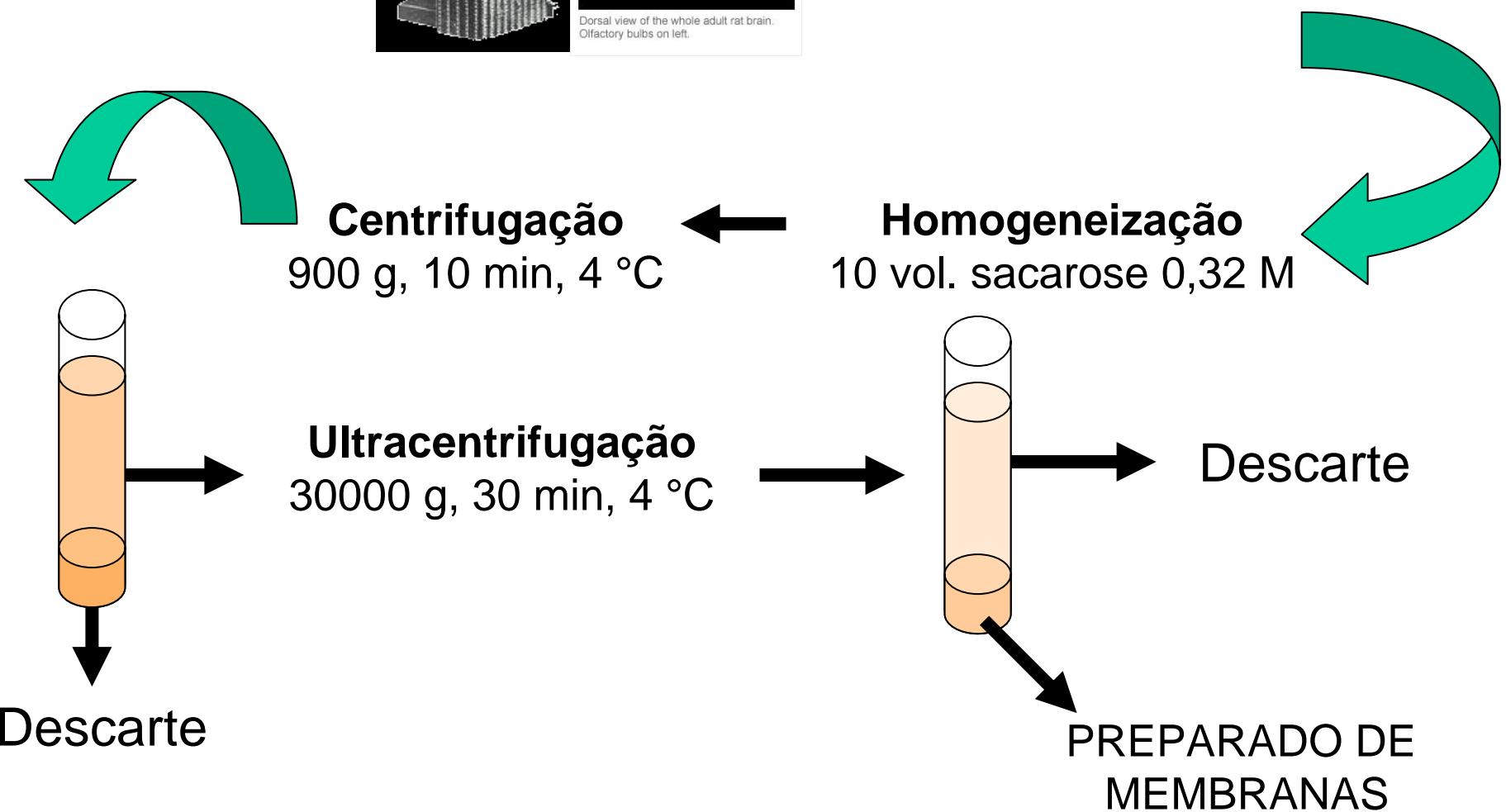
$$E = m \times \log C + b$$

# SPARE RECEPTORS

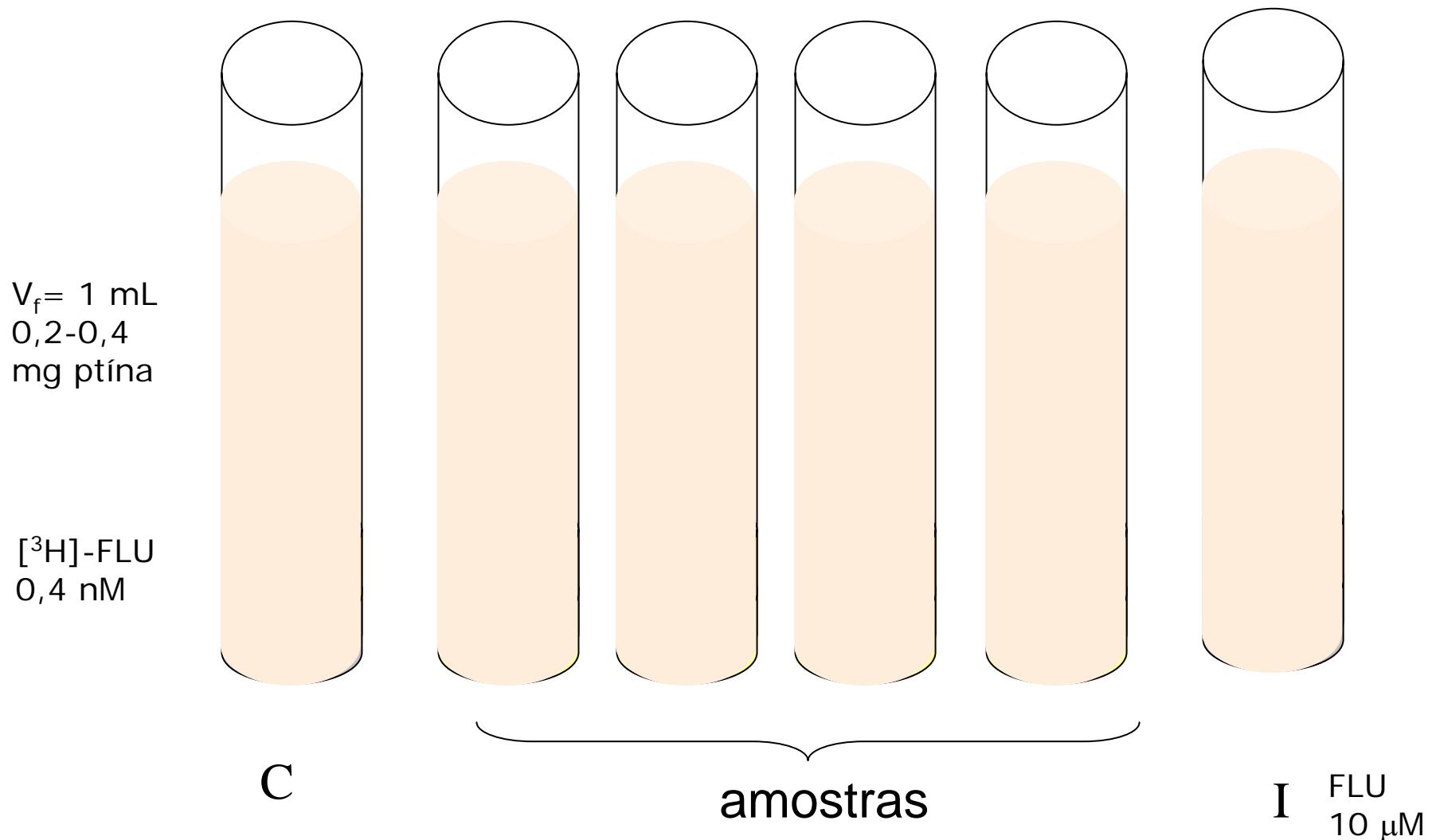




Estruturas de interesse



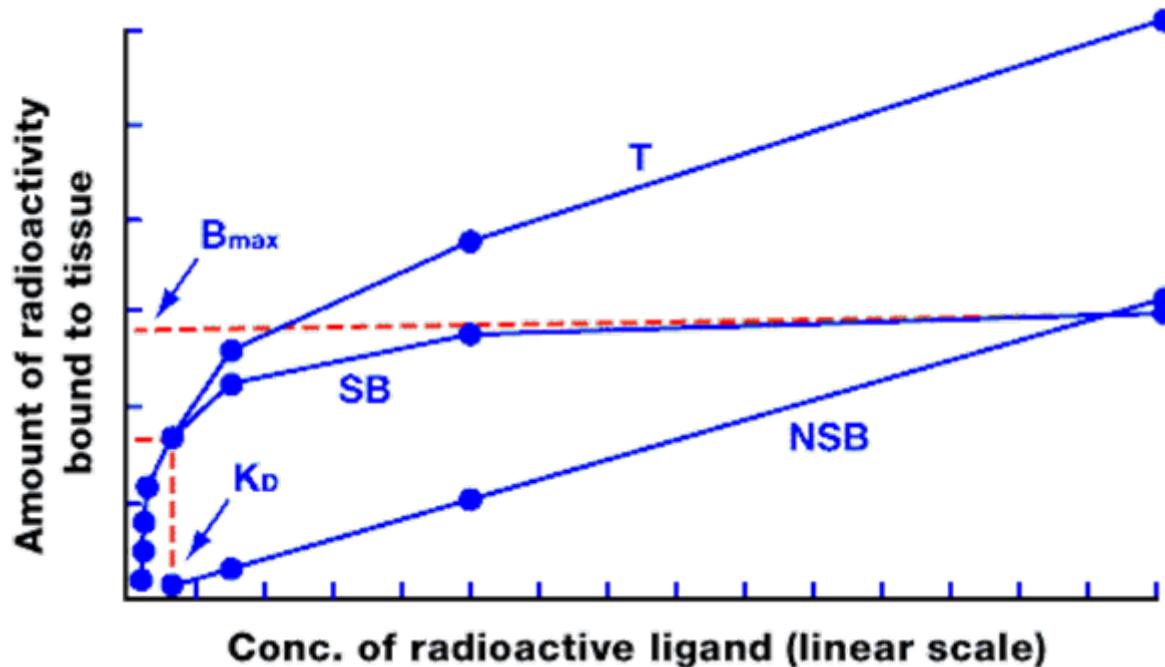
# Binding $[^3\text{H}]\text{-flunitrazepam}$





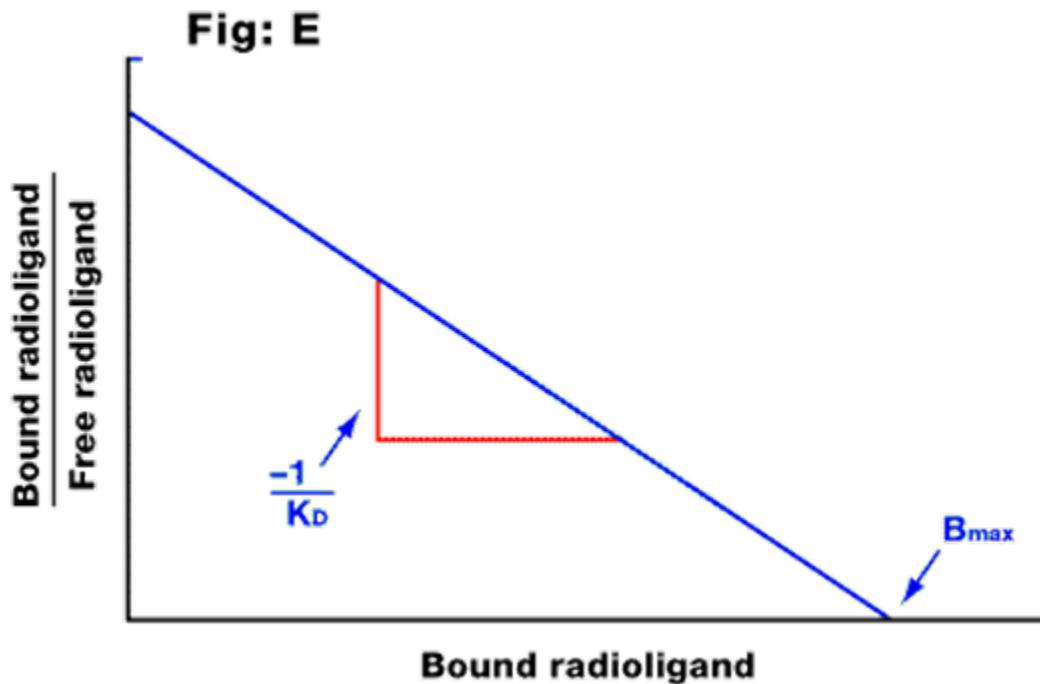


## Curvas de saturação



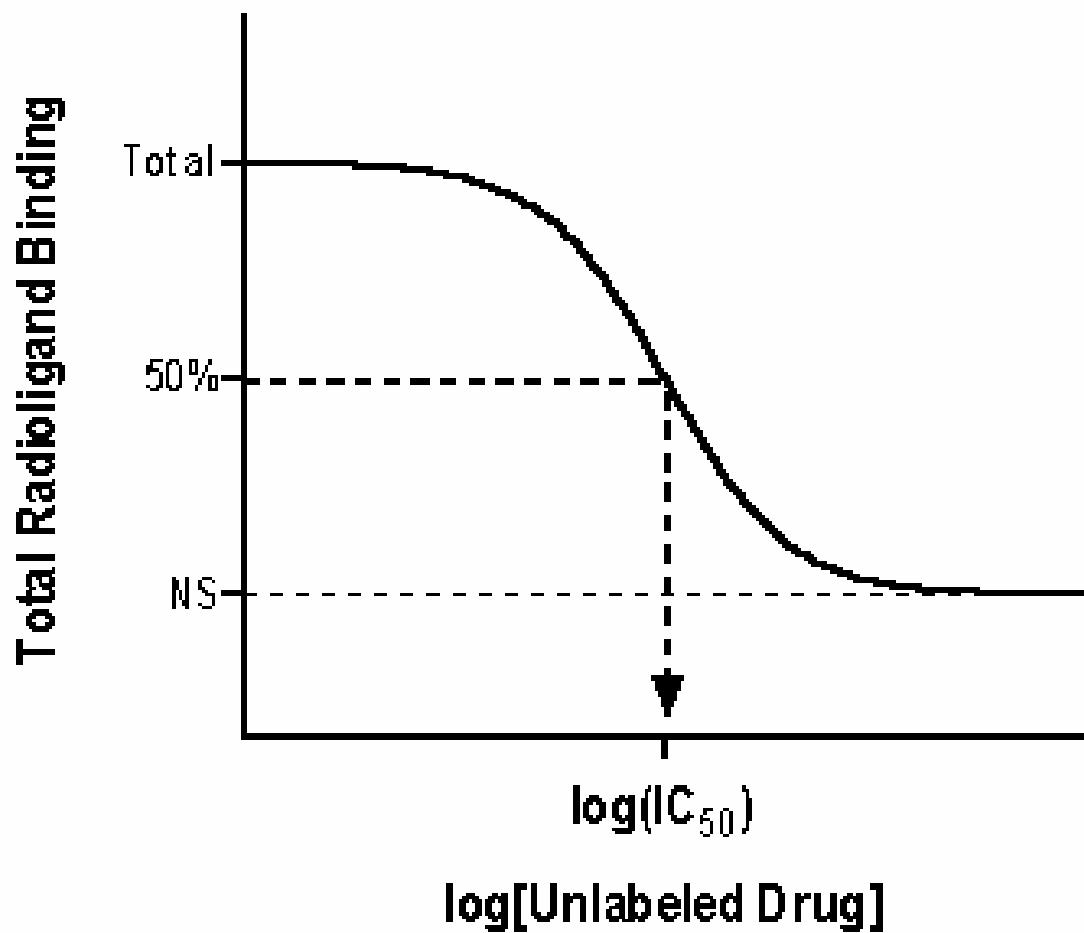
Saturation curve for a ligand binding to a homogeneous receptor population. The total binding (T) includes a component of non-specific binding (NSB), which is non-saturable, and the remainder is specific binding (SB) which saturates at  $B_{max}$ . Note that if the x-axis were logarithmic, the SB curve would be sigmoidal (compare with the log-concentration - response curve). The  $K_D$  of the ligand is the concentration which occupies 50% of the receptors, and can be calculated from a Scatchard plot.

# Scatchard Plot

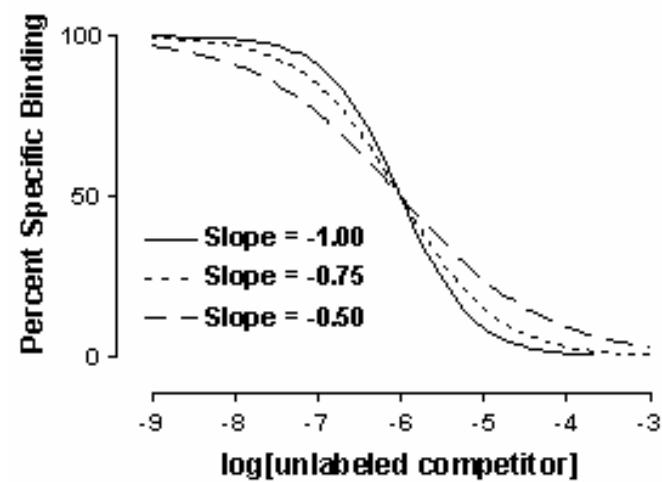
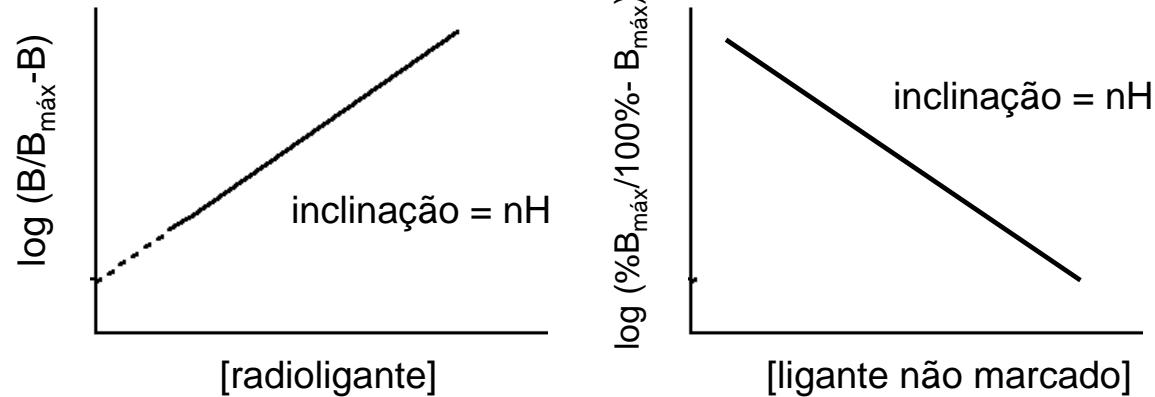


Scatchard plot of the saturable component of the data in the previous figure. The intercept on the x axis is equal to the  $B_{max}$ ; the slope is equal to  $-1/K_D$ .

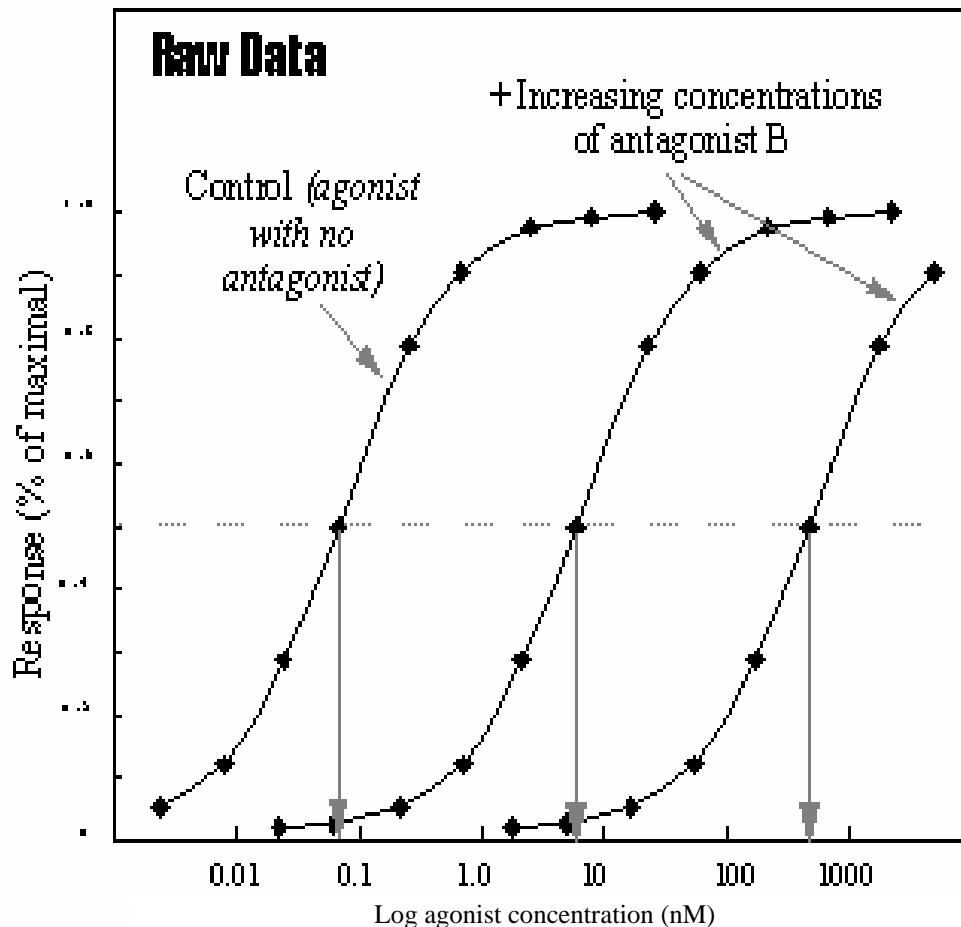
## Curvas de Deslocamento

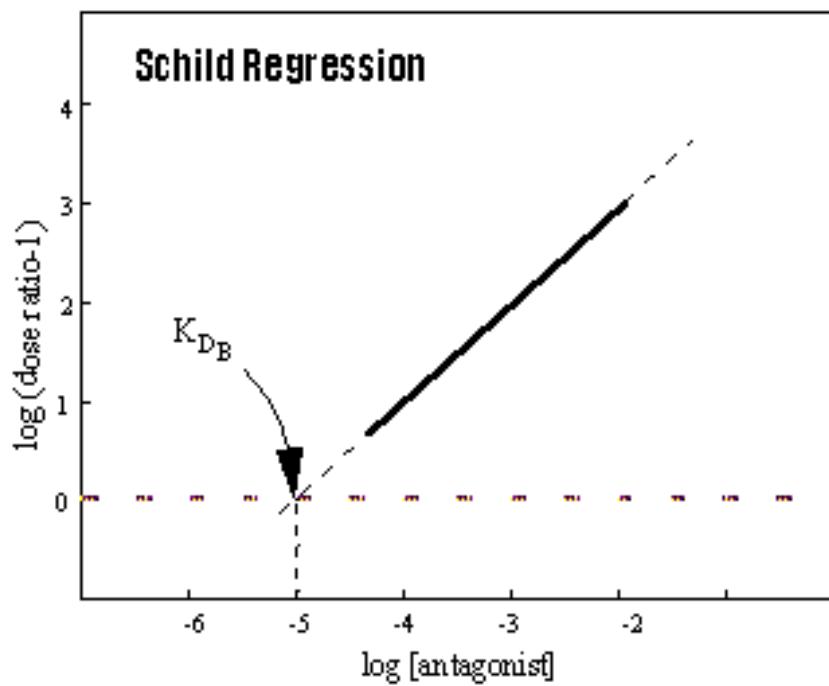
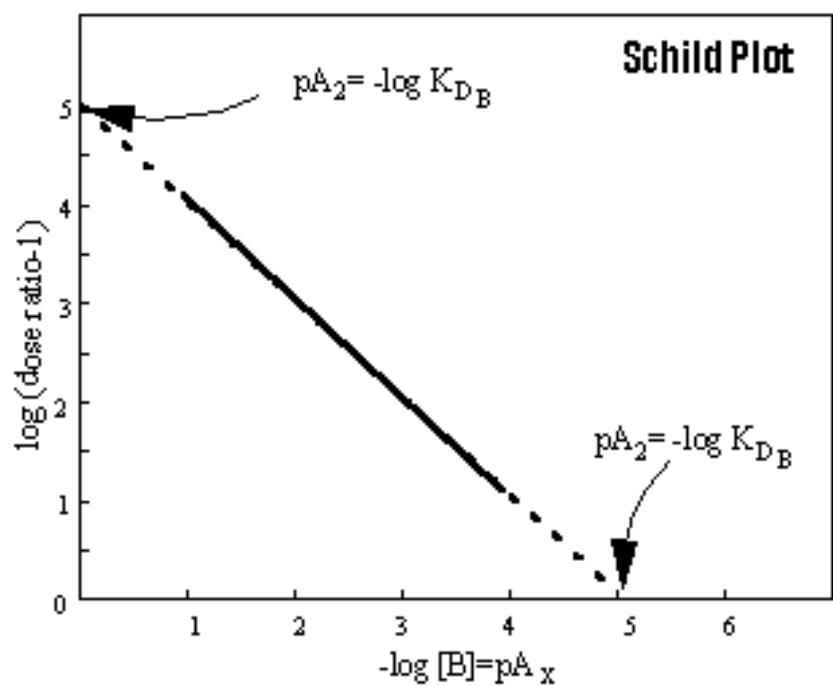


# Plot de Hill

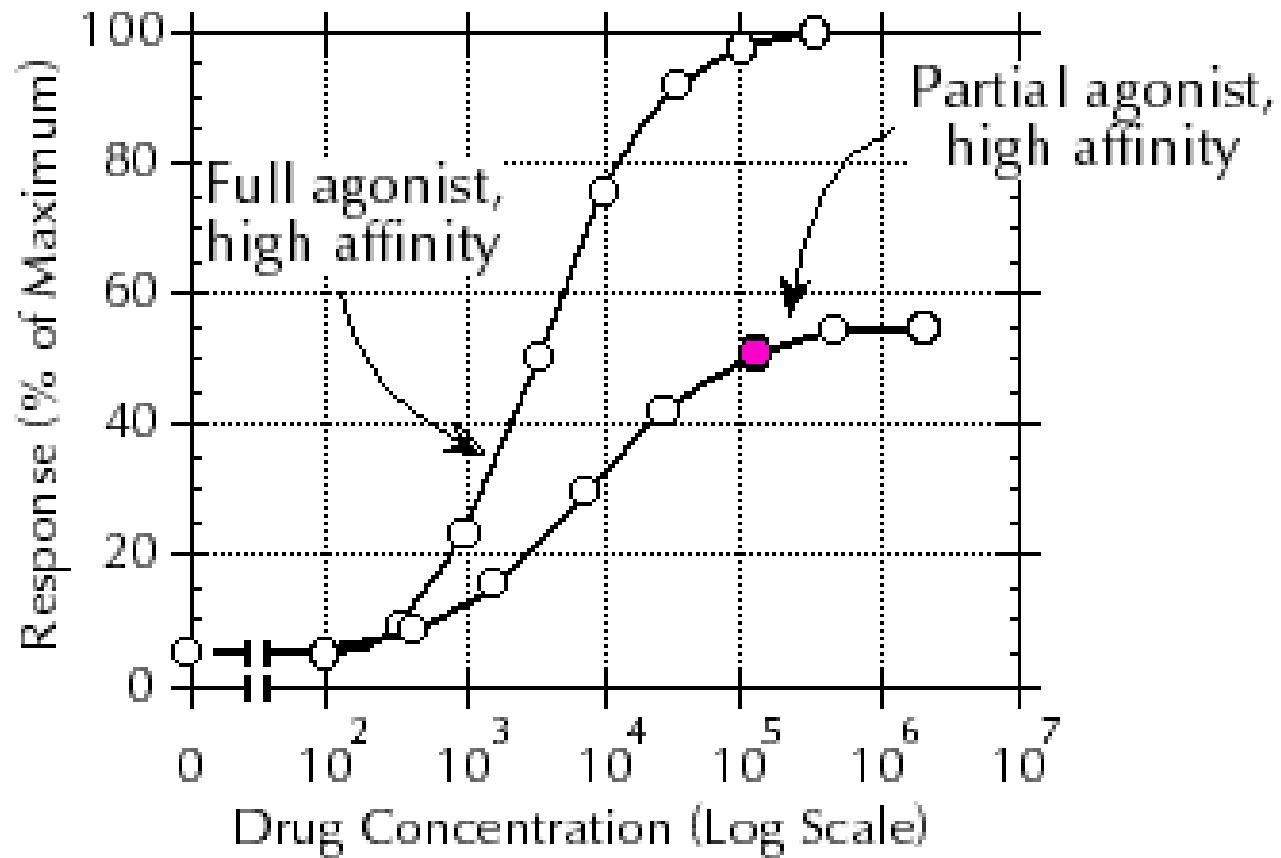


# Antagonistas

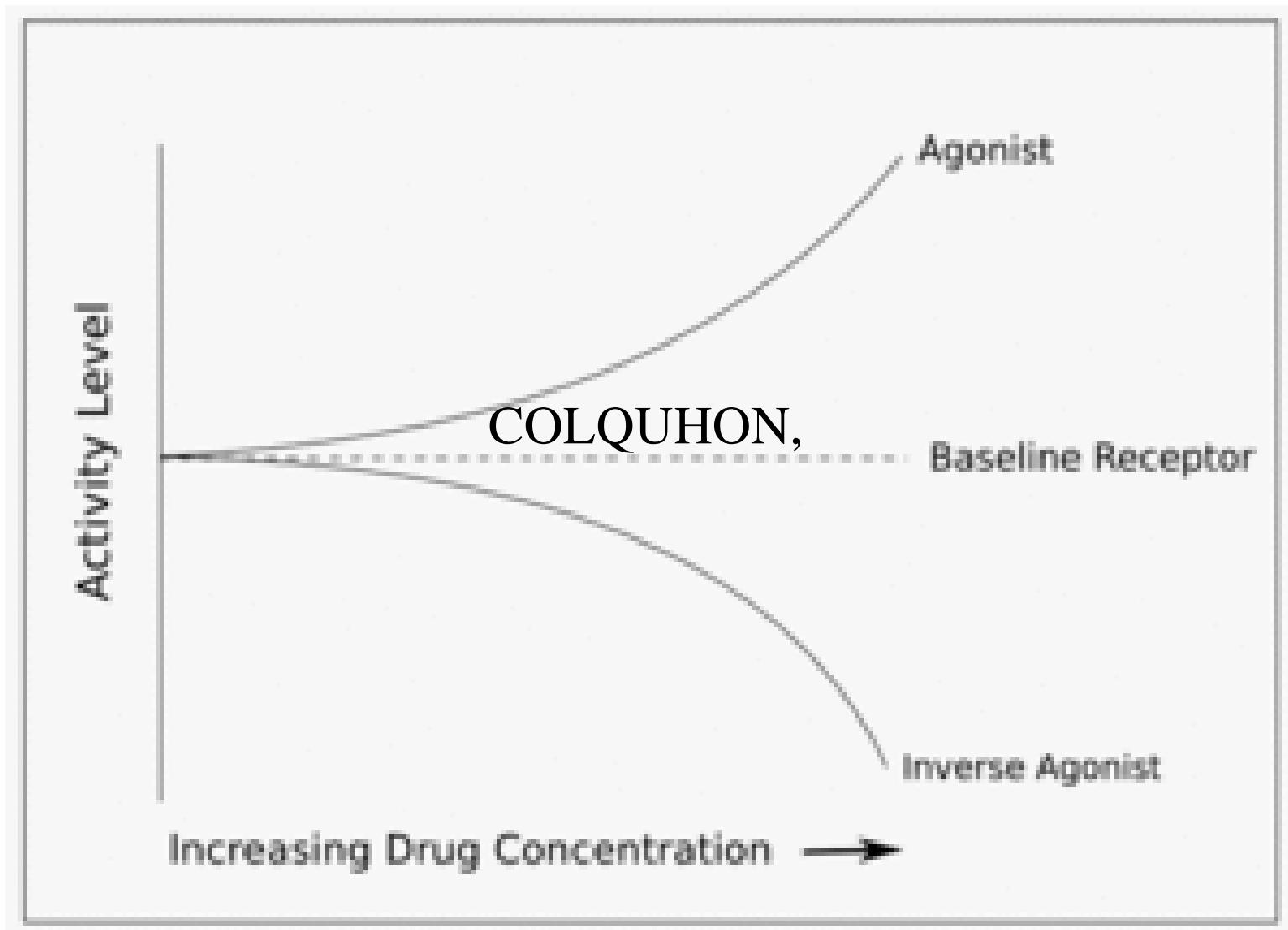




# Agonistas parciais



# Agonistas inversos





Gilda Neves  
(doutorado)



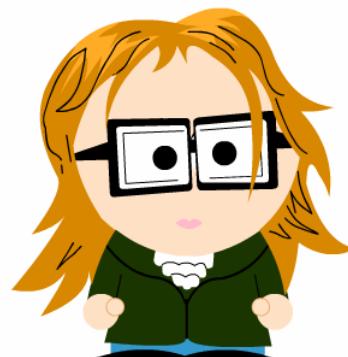
Scheila Valério  
(BIC)



Raquel Fenner  
(doutorado)



Gustavo Prove  
(mestrado)



Mariana Pranke  
(BIC)



Camila Boque  
(BIC)



Andresa Betti  
(BIC)



Alice Viana  
(doutorado)