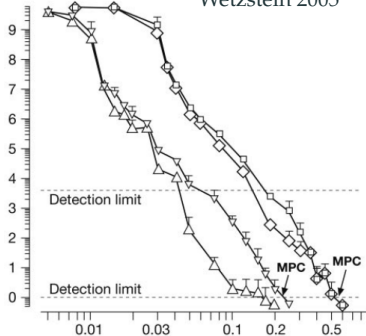


Number of
recovered cells

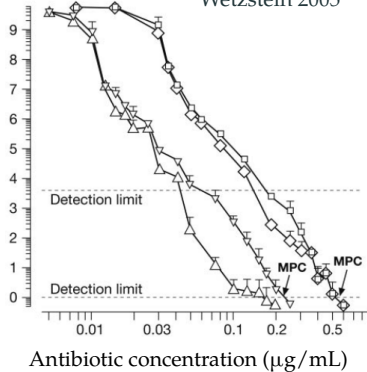
Wetzstein 2005



Antibiotic concentration ($\mu\text{g/mL}$)

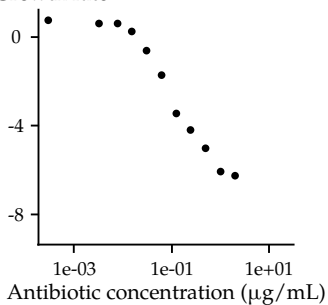
Number of
recovered cells

Wetzstein 2005



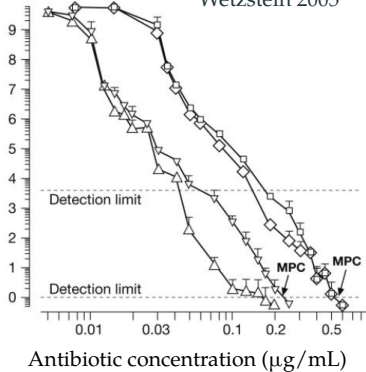
Growth rate

Regoes et al. 2004



Number of
recovered cells

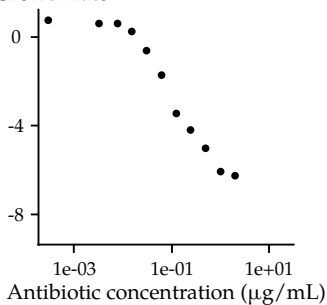
Wetzstein 2005



E.coli ATCC8739
Pradofloxacin
Medium USP

Growth rate

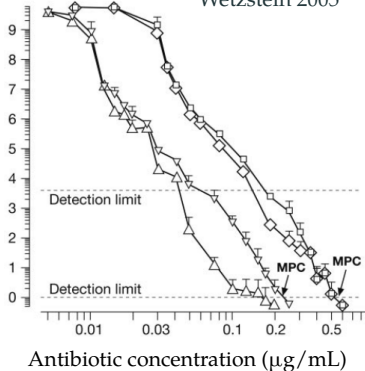
Regoes et al. 2004



E.coli CAB1
Ciprofloxacin
Medium LB

Number of
recovered cells

Wetzstein 2005

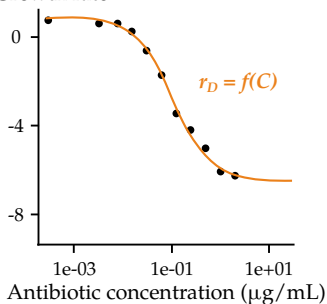


E.coli ATCC8739
Pradofloxacin
Medium USP

Relationship between **antibiotic
concentration C** and **decay rates r_D**

Regoes et al. 2004

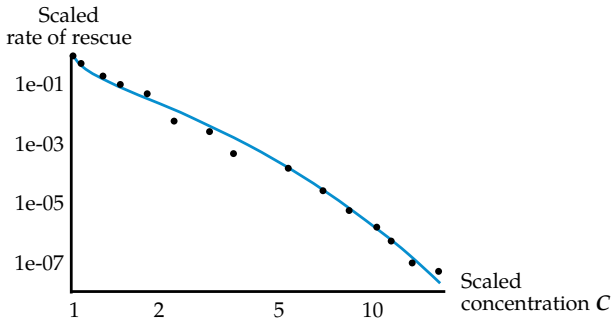
Growth rate



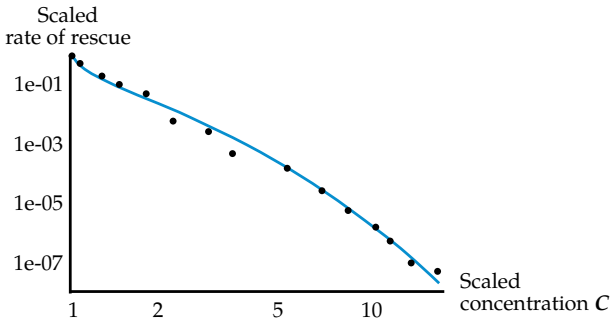
E.coli CAB1
Ciprofloxacin
Medium LB

Model "Weak U" $r_D = f(C)$

Model "Weak U" $r_D = f(C)$



Model "Weak U" $r_D = f(C)$



The model combining the "Weak U" model and the function $r_D = f(C)$ has **too many parameters**.

The model gives a **better visual fit than a model without context-dependence**