

Pharmacokinetic-Pharmacodynamic (PKPD) Modeling Characterizing Resistance for Predictions of Bacterial Kill *in vivo*

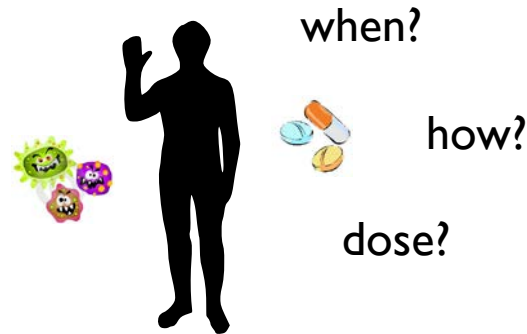
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Pascale David- Pierson, Neil John Parrott, Olaf Kuhlmann,
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Pharmacometrics group Uppsala Universitet, Uppsala, Sweden
Hoffmann LaRoche Ltd., Basel, Switzerland

Introduction

Dose selection antibiotics



Difficulties:

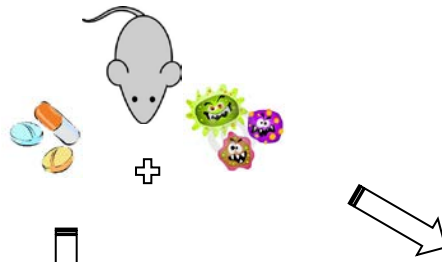
- Bacterial load
- Variable susceptibility
- Short term
- Sudden onset
- Emergence of resistance
- Typically highly efficient comparators

Advantages:

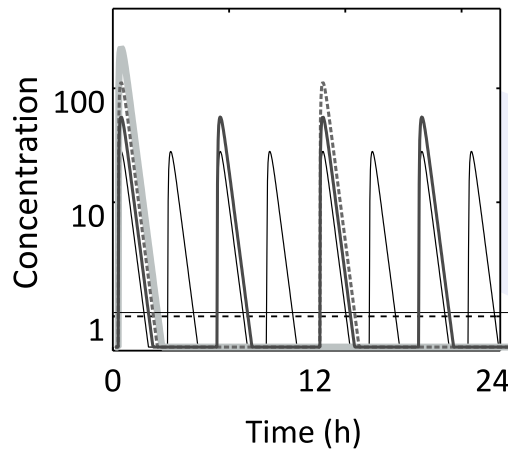
- Ability to study PD preclinically
 - in vitro*
 - in vivo* animal models

Introduction

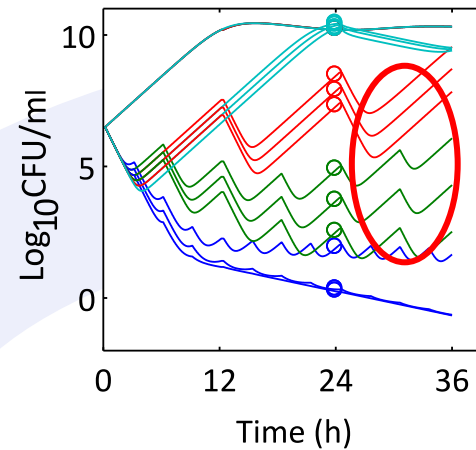
PK/PD indices



PK

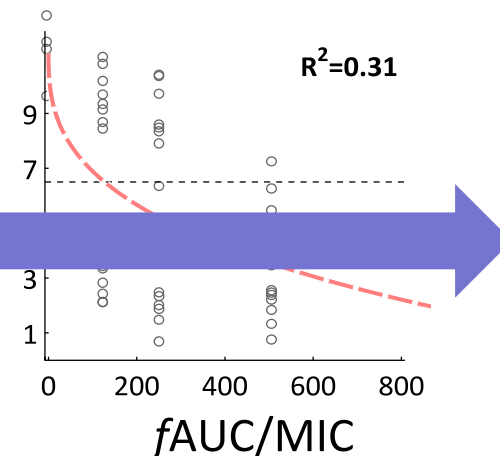
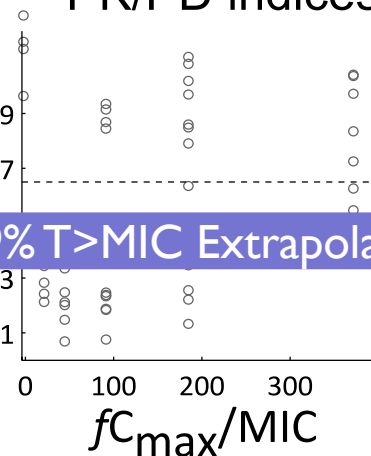
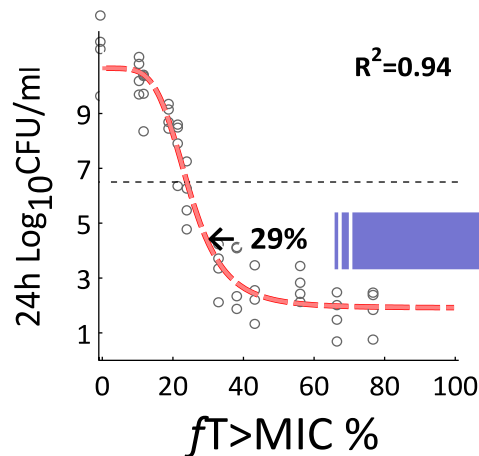


PD



Regrowth?
Resistance?

PK/PD indices

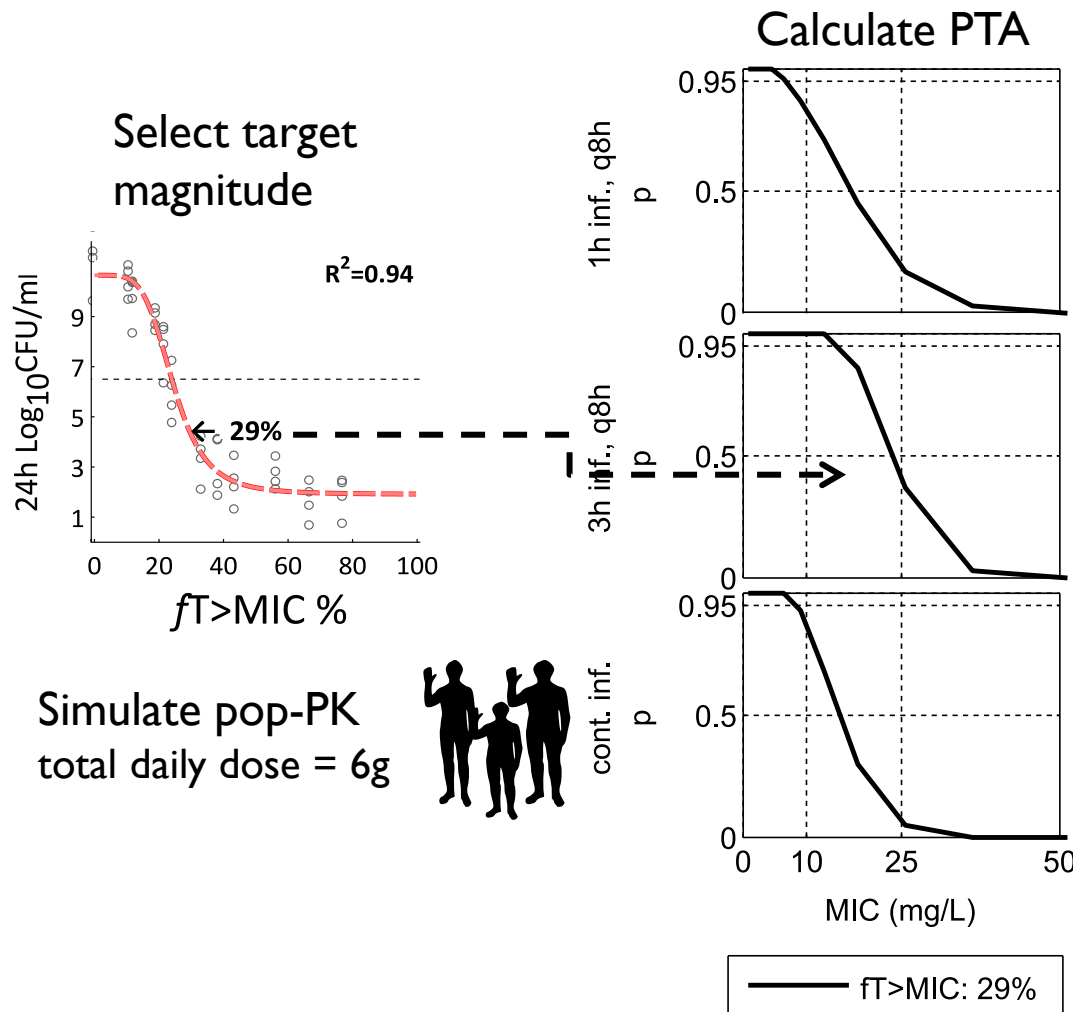




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Introduction

Probability of target attainment (PTA)



Same target and magnitude for different patient populations?

Same magnitude for bacterial strains with different susceptibility?

3h infusion q8h
→ PTA 95%
MIC < 14mg/L

Dose



- Describe bacterial kill and resistance using a previously developed PKPD-model
- Replicate an *in vivo* dose finding study (PK/PD indices)
 - Evaluate the predictability of the PKPD model
 - Compare dose predictions for a resistant and a susceptible strain

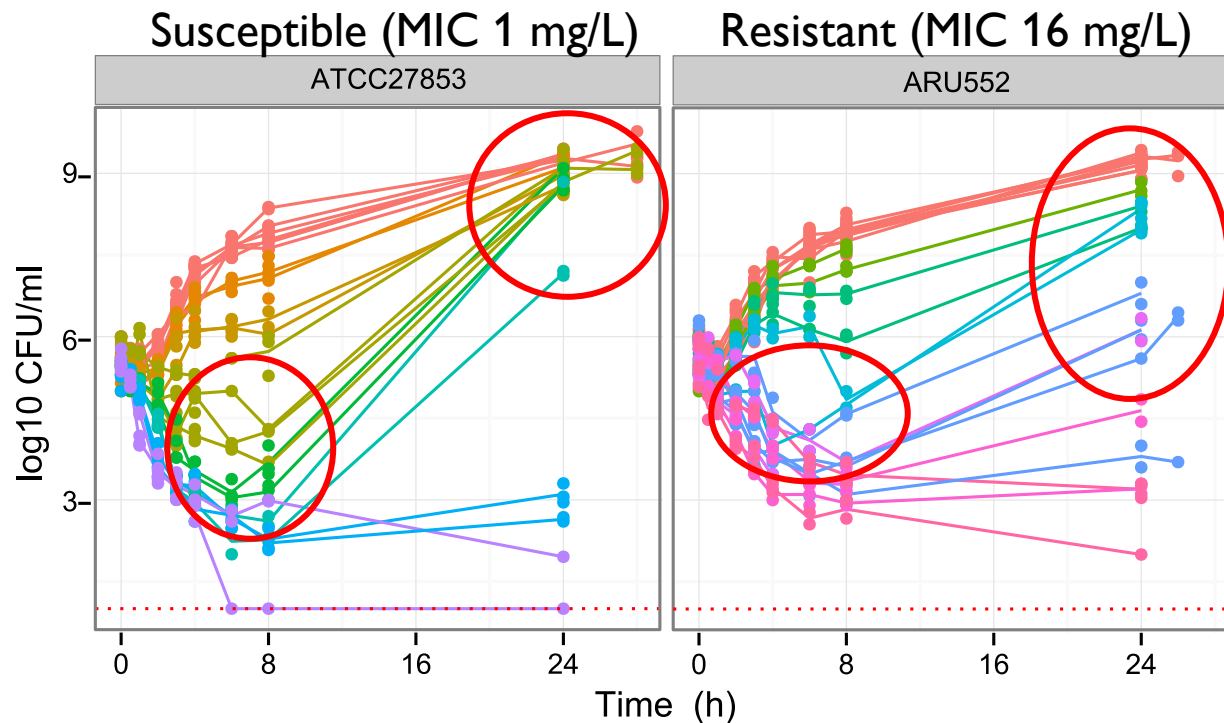
For *Pseudomonas aeruginosa* and meropenem

Meropenem *P. aeruginosa* PKPD model

In vitro time-kill data

2 strains *Pseudomonas aeruginosa*:
ATCC27853 (Susceptible), ARU552 (Resistant)

Meropenem: 0.05-3.2 × MIC (ATCC27853), 0.25-32 × MIC (ARU552)



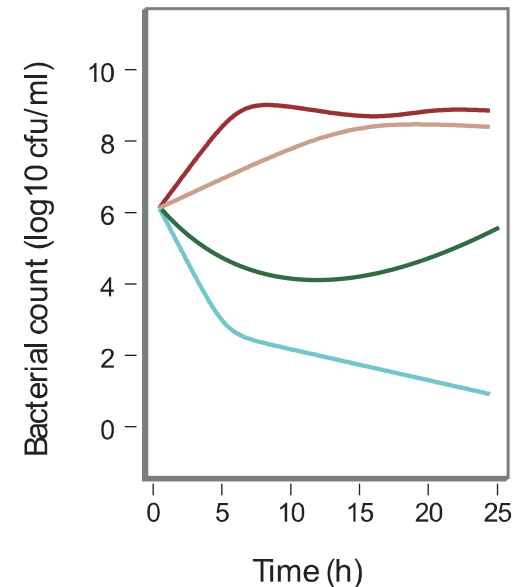
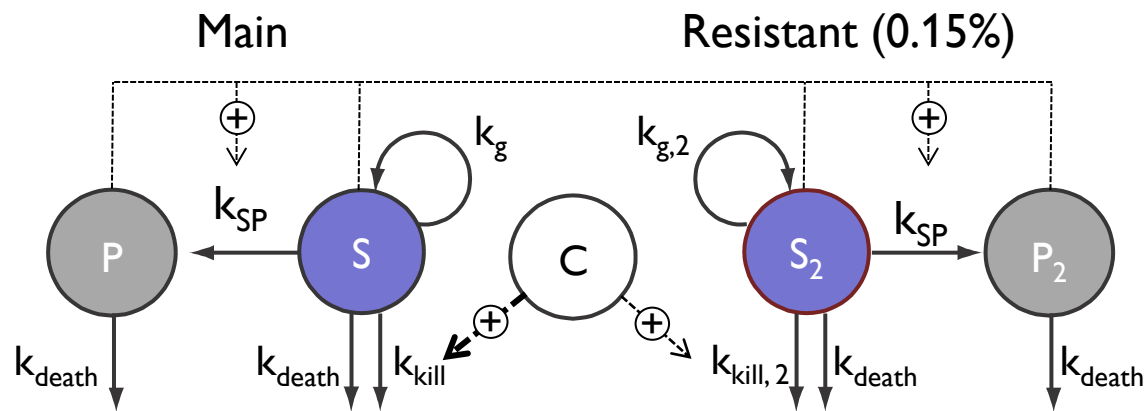
2 types of resistance:

MIC classification
MIC 16 vs 1 mg/L

Selection by exposure
Regrowth during
experiment due to
resistance

Meropenem *P. aeruginosa* PKPD model

Structure

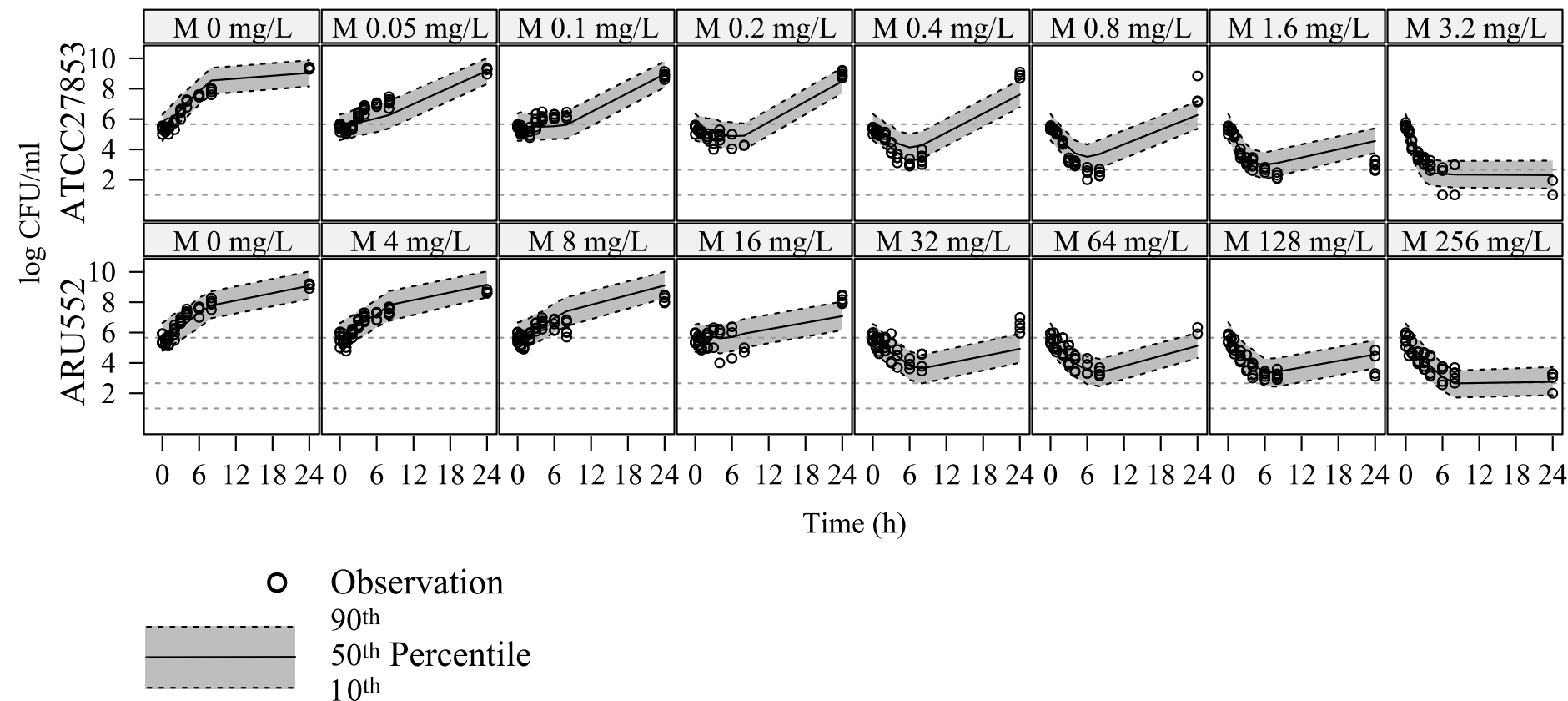


Concentration to suppress growth of main population:
ATCC 0.16 mg/L, ARU552 17 mg/L

Resistant subpopulation:
~20 times higher concentration for same effect
~50% reduced growth rate ARU552

Meropenem *P. aeruginosa* PKPD model

VPC

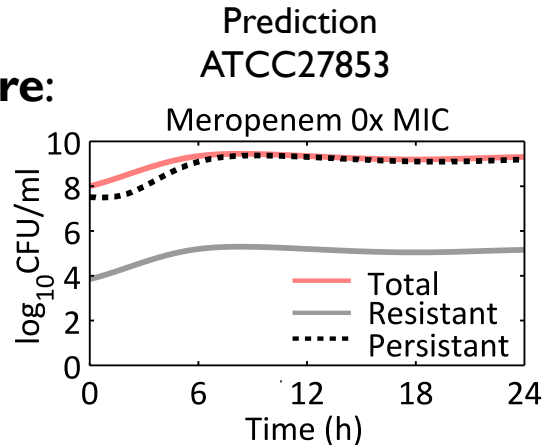


Meropenem *P. aeruginosa* PKPD model

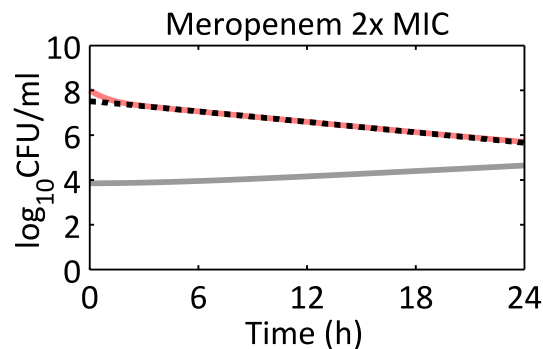
Predictions

Selective pressure:

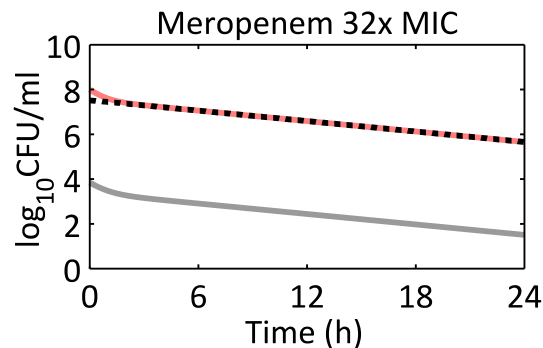
None
(0 x MIC)



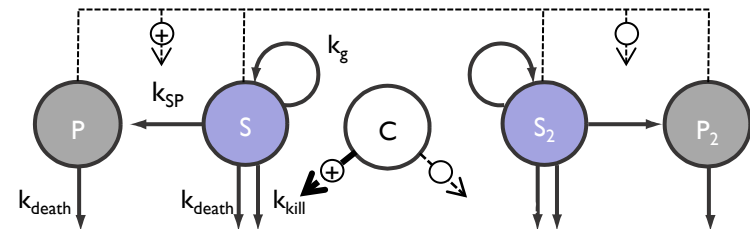
Mutant selective
(2 x MIC)



Mutant preventive
(32 x MIC)



Total = Main + Resistant
Persistent = P + P₂



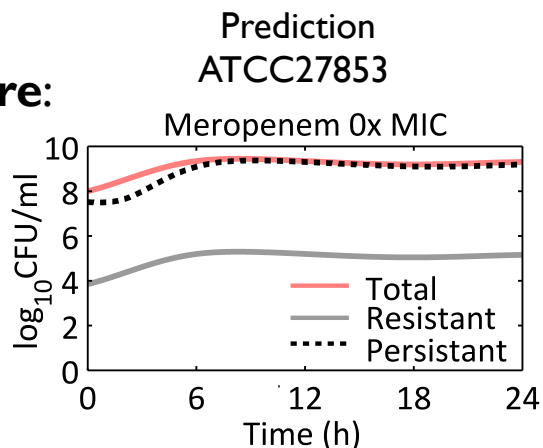


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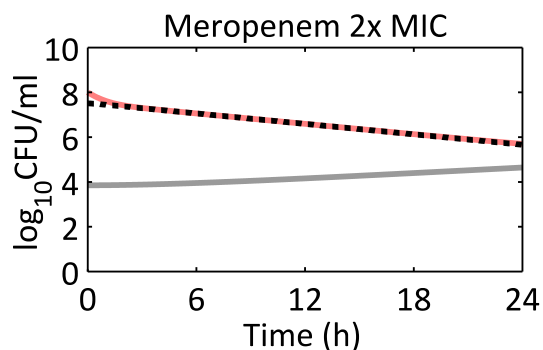
Meropenem *P. aeruginosa* PKPD model Predictions

Selective pressure:

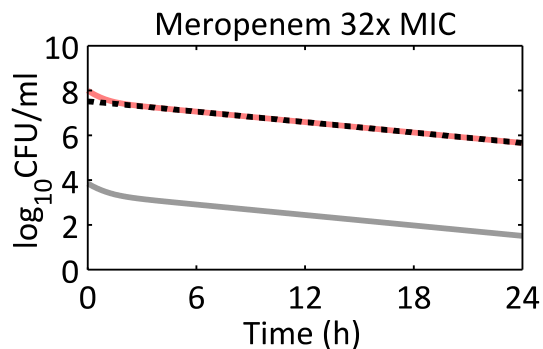
None
(0 x MIC)



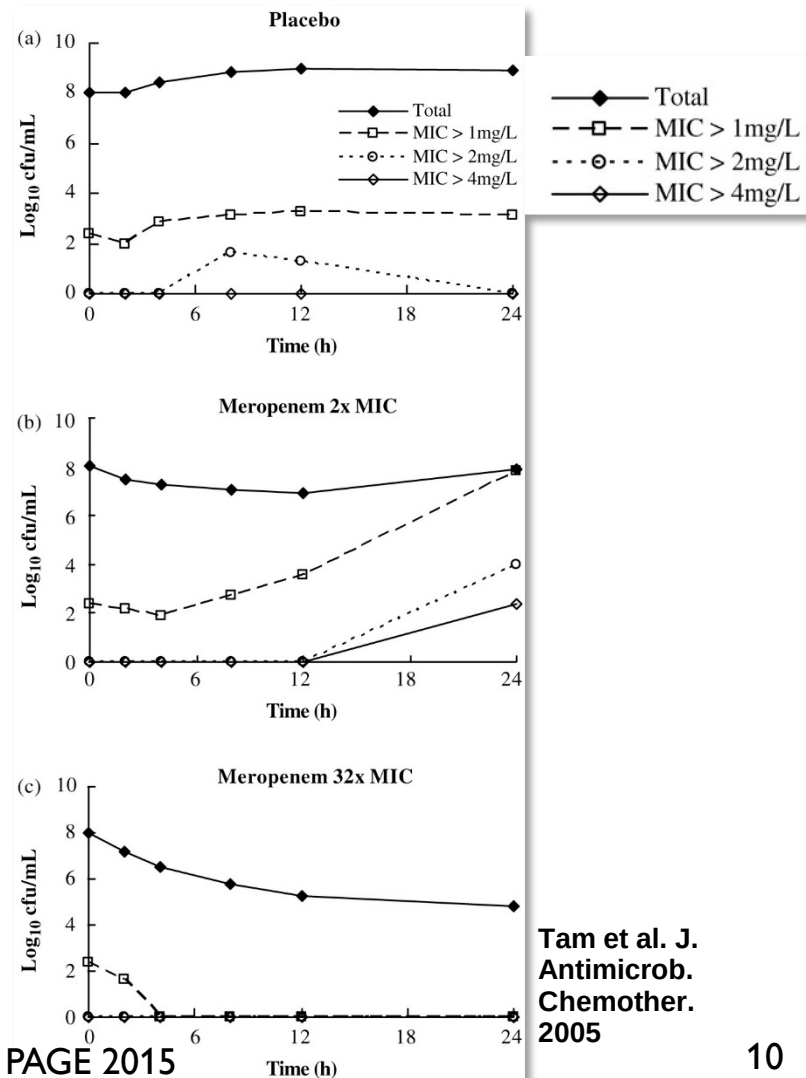
Mutant selective
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(32 x MIC)

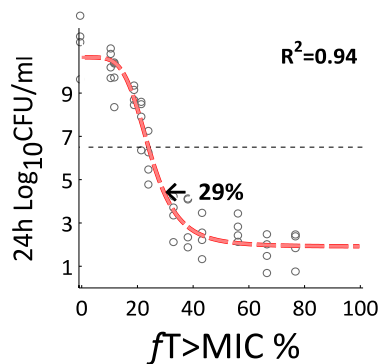
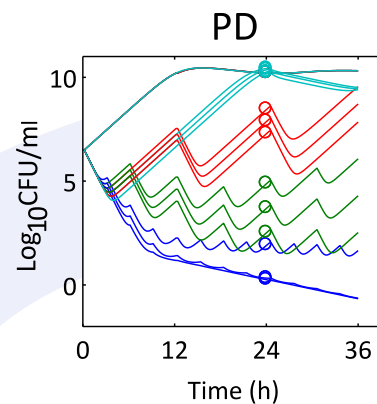
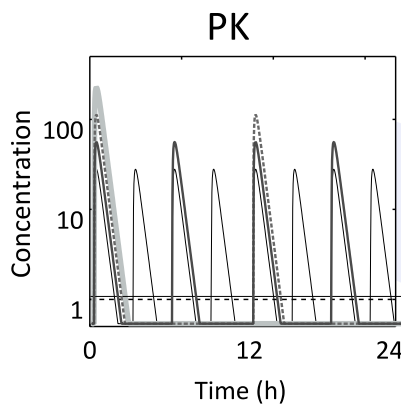
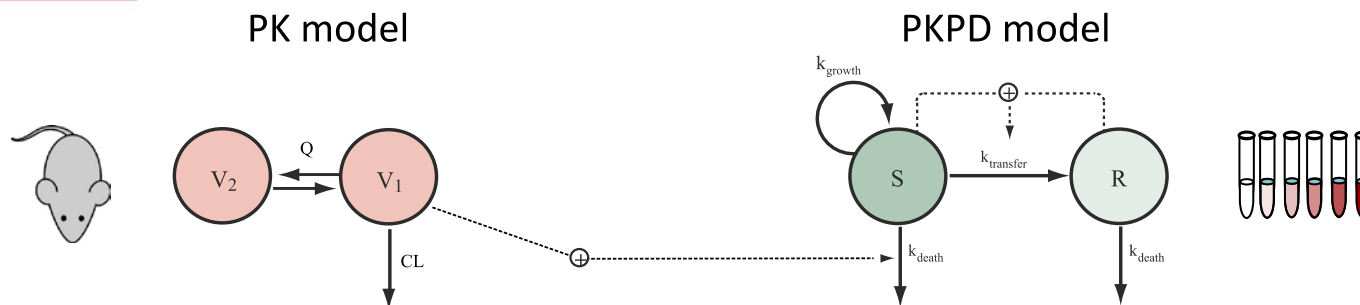


Tam et al 2005
ATCC27853

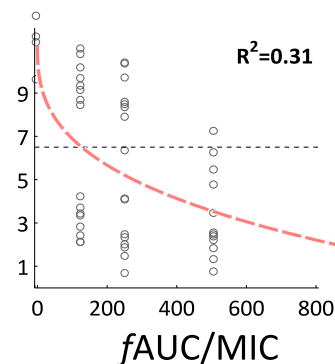
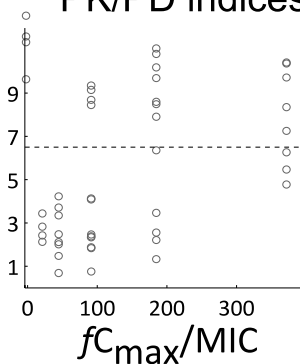




In silico, PK/PD indices



PK/PD indices



Predicting PK/PD indices Mouse



In vivo

(Sugihara et al,
AAC 2010)

P. aeruginosa 12467
MIC=2mg/L



Simulation

1-comp PK

(Katsube et al,

J. Pharm. Sci. 2008)

P. aeruginosa ATCC27853

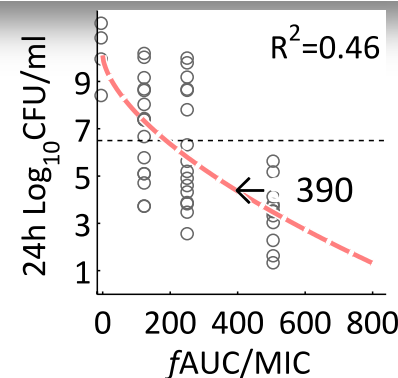
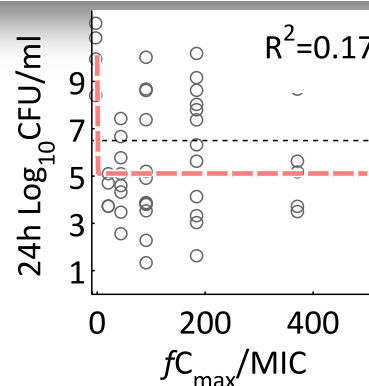
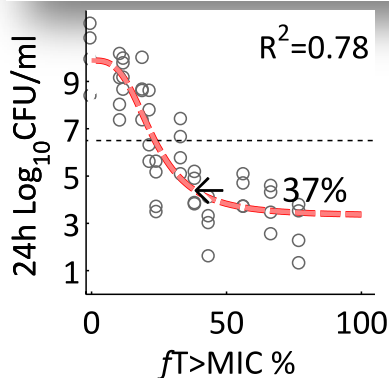
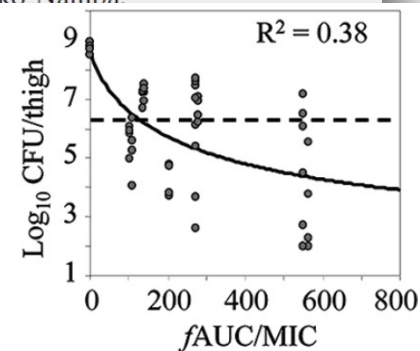
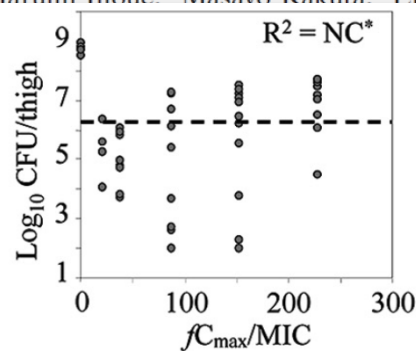
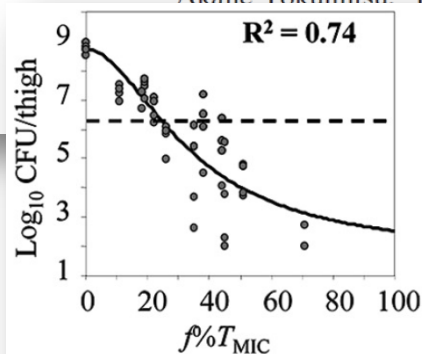
MIC=1mg/L

ANTIMICROBIAL AGENTS AND CHEMOTHERAPY, Dec. 2010, p. 5298–5302
0066-4804/10/\$12.00 doi:10.1128/AAC.00267-10
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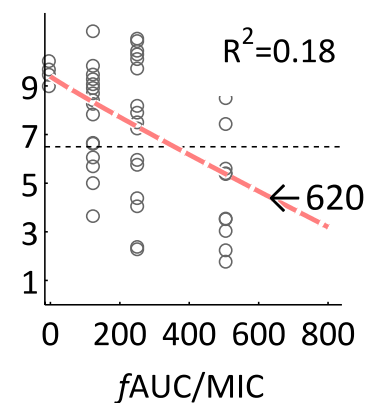
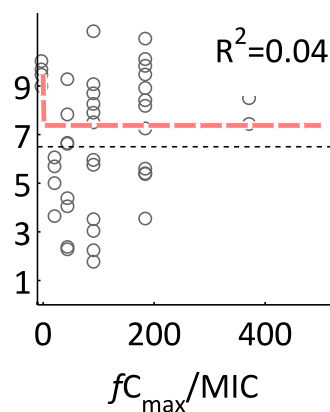
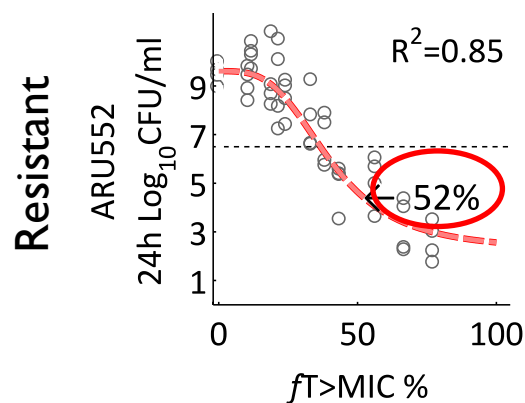
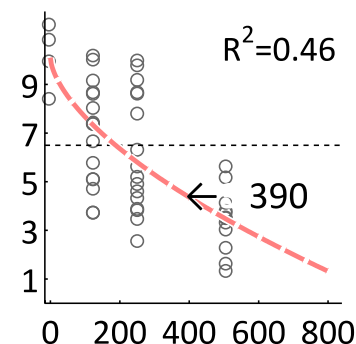
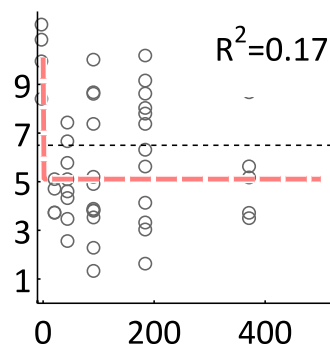
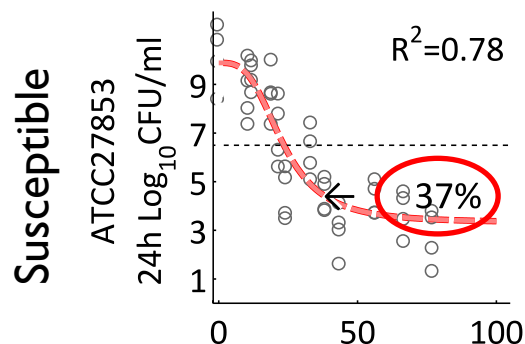
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In Vivo Pharmacodynamic Activity of Tomopenem (formerly CS-023) against *Pseudomonas aeruginosa* and Methicillin-Resistant *Staphylococcus aureus* in a Murine Thigh Infection Model[∇]

Kiyoshi Sugihara,^{1*} Chika Sugihara,¹ Yoko Matsushita,² Naotoshi Yamamura,² Mitsutoshi Uemori,³
Akane Tokumitsu,¹ Harumi Inoue,¹ Masayo Kakuta,¹ Eiko Namba,¹

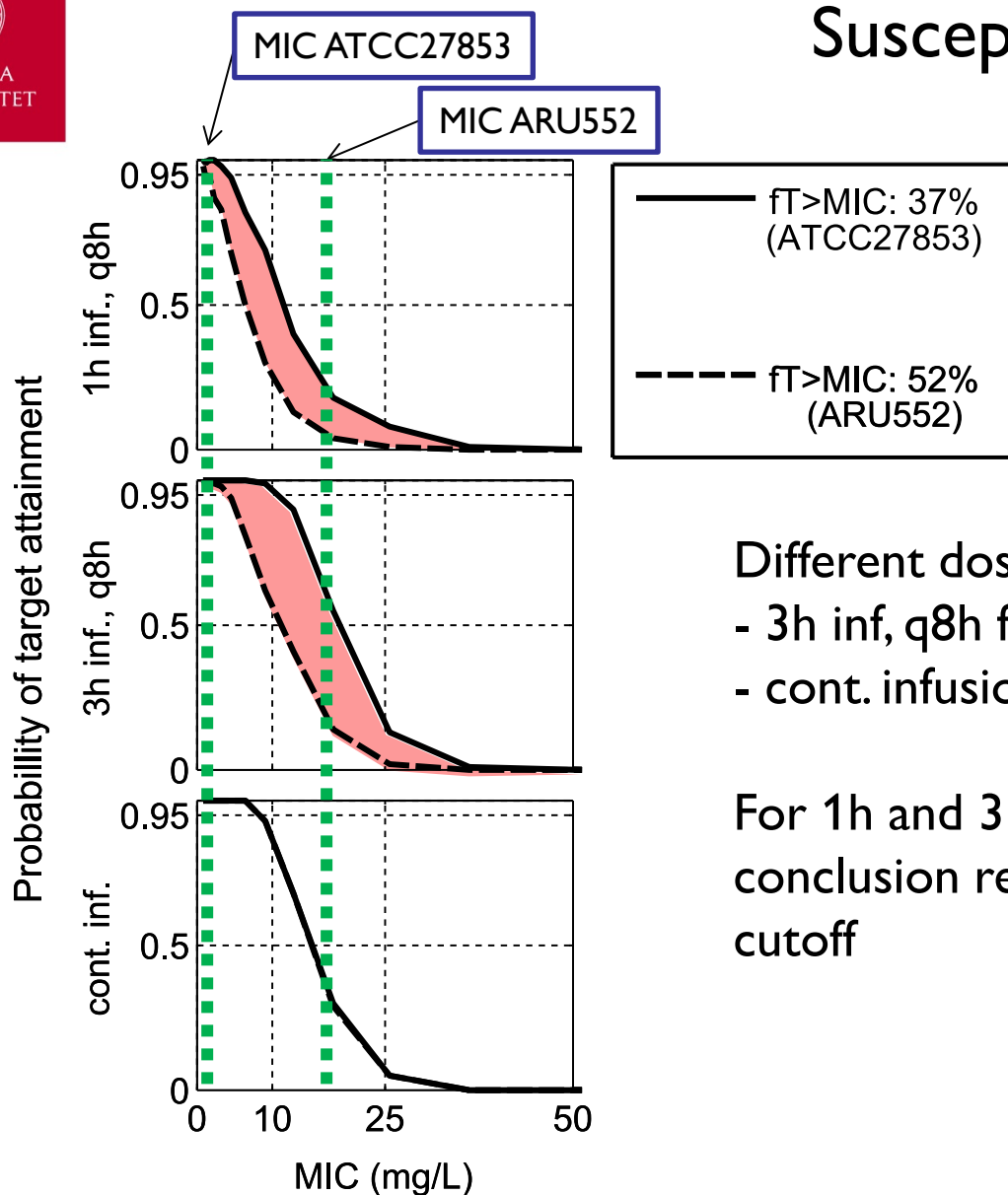


Predicting PK/PD indices Susceptible vs. Resistant



37% to 52% → 2-fold dose/MIC difference for q8h dosing regimen!

Probability of target attainment (PTA) Susceptible vs. Resistant



Total daily
dose: 6g

Different dose regimen selected

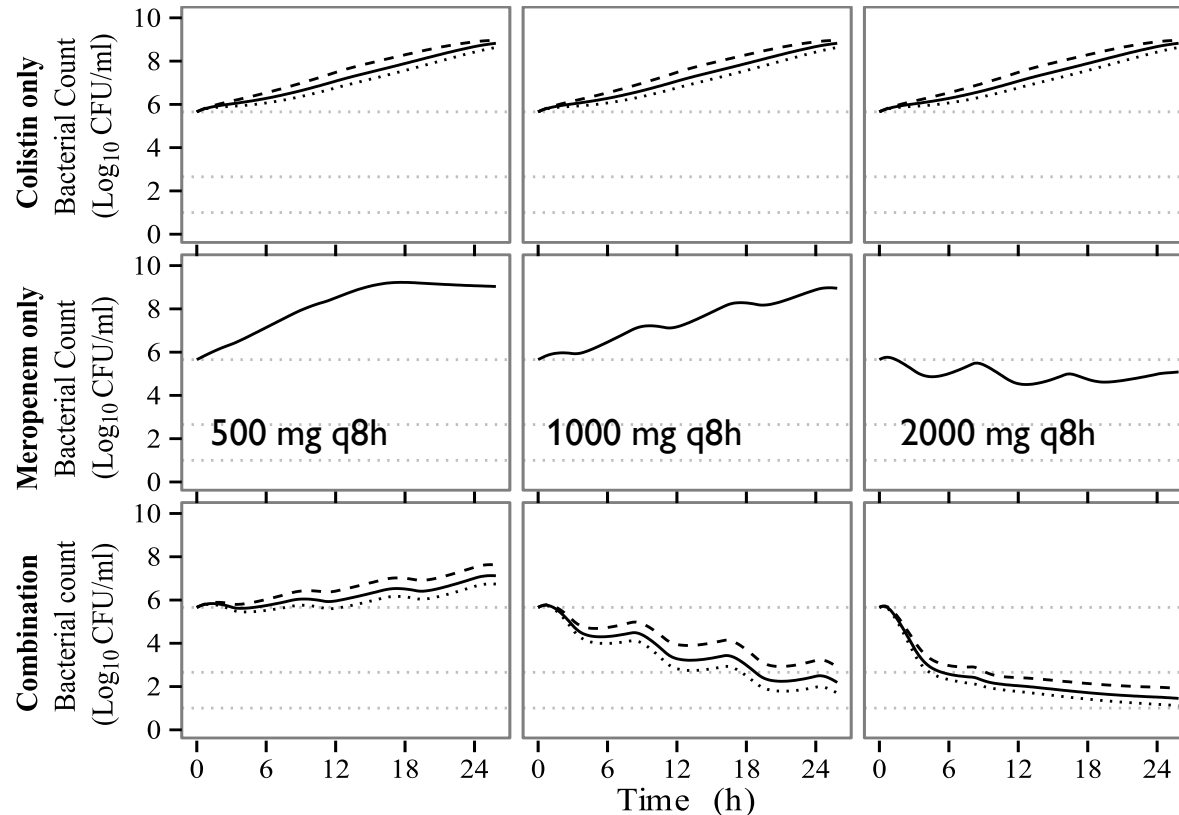
- 3h inf, q8h for 37% T>MIC
- cont. infusion for 52% T>MIC

For 1h and 3h infusion different
conclusion regarding treatment
cutoff

Predict full time course combination dose simulation

Meropenem resistant ARU552

(MIC meropenem 16mg/L, MIC colistin 1.5 mg/L)



PKPD model of colistin and meropenem in combination against *P. aeruginosa*

Combination is predicted to achieve improved kill against meropenem resistant strain

Conclusions

- The PKPD model describes two types of resistance:
 - MIC classification
 - Regrowth from resistant subpopulation
- An *in vivo* dose finding study was replicated in simulation
 - The PK/PD index magnitude was MIC dependent
 - Dose selection differed between strains with different susceptibility

PKPD modeling and dose simulation offers a more powerful dose finding alternative!

Acknowledgments

Friends and colleges at the Pharmacometrics group, Uppsala university

Co-authors Meropenem-Colistin PKPD model:

Ami F Mohamed, Matti Karvanen, Elisabet I Nielsen, Otto Cars,
Lena E Friberg