

TI Pharma mechanism-based PKPD modeling platform the objective

Development and implementation of a mechanism-based PKPD modeling platform as the scientific basis for rational drug discovery and innovation

- Mechanism-based PKPD model library
- Database of 'biological system specific' information

# **TI Pharma mechanism-based PKPD modeling platform the organization**

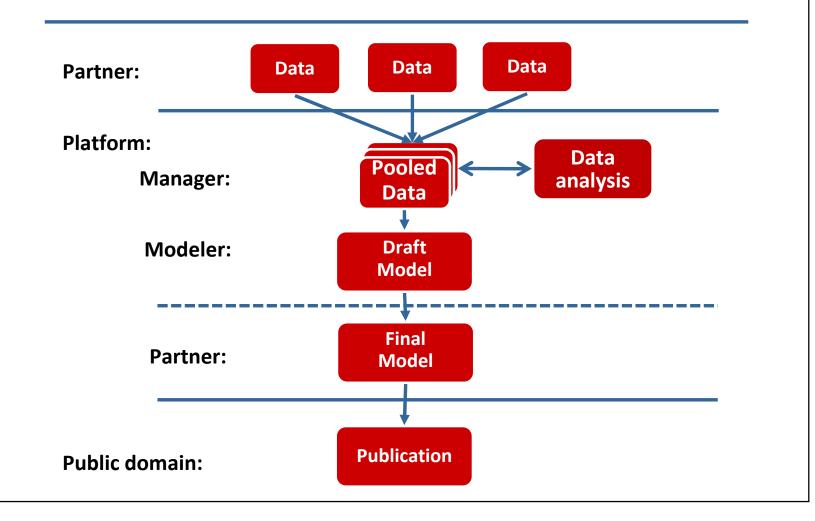
- University-industry consortium with 4 academic and 8 industrial partners
- Dedicated infrastructure for data management, data analysis and reporting: sharing of data, models and biological system specific information
- Emphasis on key factors in the discovery/development and the clinical application of novel drugs
  - Translational pharmacology (efficacy and safety)
  - Developmental pharmacology (pediatrics, elderly)
  - Disease system analysis (osteoporosis, COPD)

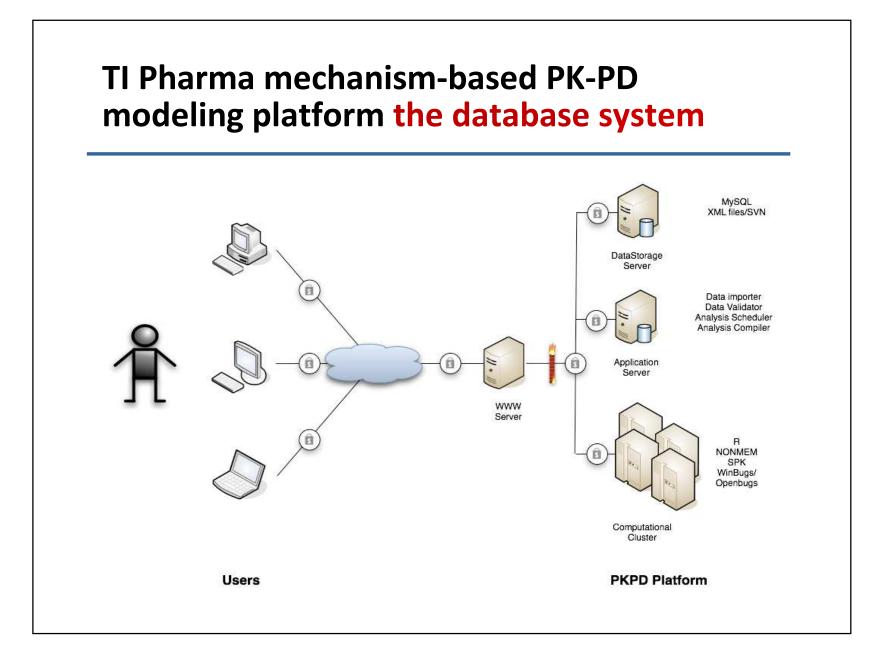


## **TI Pharma mechanism-based PKPD modeling platform the operation**

- Development of mechanism-based PK-PD models on basis of existing data
- Strict data access restrictions
- Centralized computing network facility for data management and analysis
- Model library interface for users

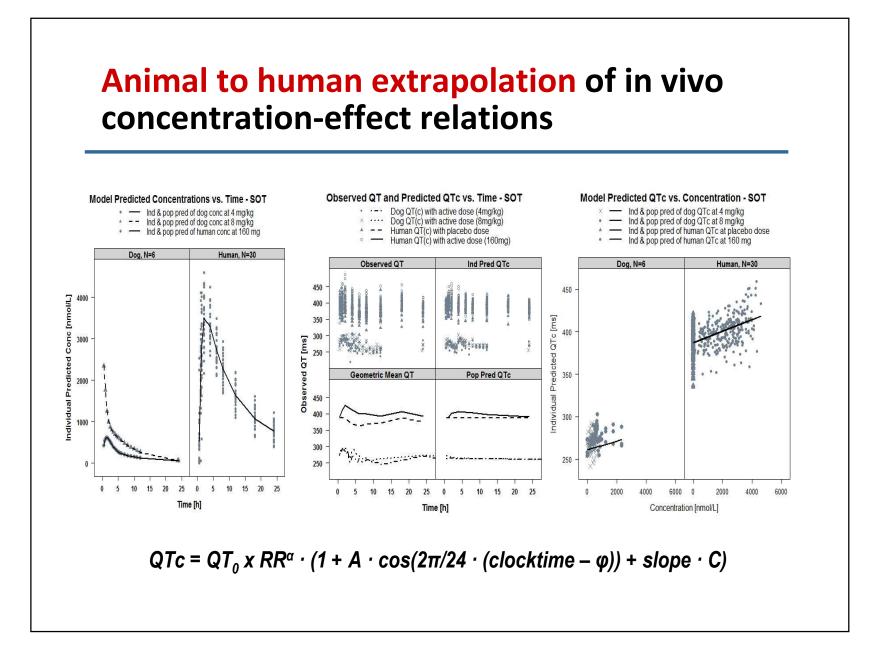
#### TI Pharma mechanism-based PK-PD modeling platform the information flow



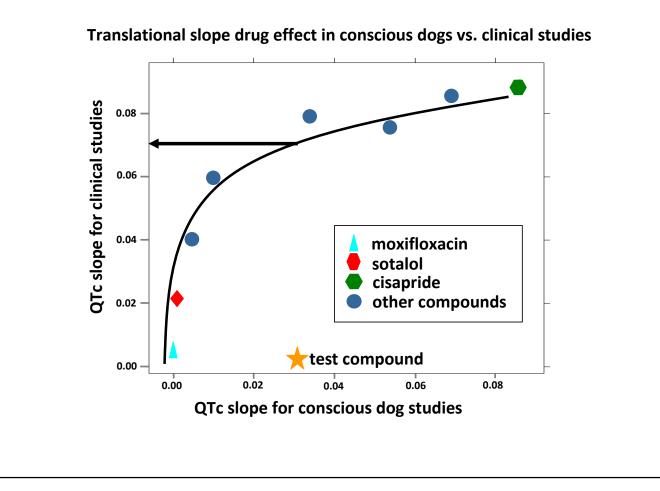


## Prediction of pharmacology in man cardiovascular safety

| Drug         |          | N   | Dose              | Variables used for modelling & simulation and sampling scheme         |
|--------------|----------|-----|-------------------|---|
| Moxifloxacin | *        | 4   | 3, 10, 30 mg/kg   | Clock time, RR, QT over 24 h  |
|              | 775      |     |                   | plasma PK from literature   |
|              | <b>İ</b> | 137 | 400mg             | Clock time, RR, QT, plasma PK over<br>24 h                            |
| Sotalol      | 沅        | 4   | 4, 8 mg/kg        | Clock time, RR, QT over 48h, plasma<br>PK literature                  |
|              | <b>İ</b> | 30  | 160 mg            | Clock time, RR, QT, plasma PK over<br>24 h                            |
| Cisapride    | 沅        | 4   | 0.6, 2, 6 mg/kg   | Clock time, RR, QT, plasma PK over<br>24 h, plasma PK from literature |
|              | <b>İ</b> | 24  | 10, 20, 40, 80 mg | Clock time, RR, QT, plasma PK over<br>24 h                            |
| NCE          | 沅        | 4   | 1.5 µg            | Clock time, RR, QT, plasma PK over 24 h,                              |
|              | <b>İ</b> | 24  | 1.5 µg            | Clock time, RR, QT, plasma PK over<br>24 h                            |



# Indentifying the animal to human translation function for QTc interval prolongation



# Prediction of cardiovascular risk in real life situations not in trial simulation

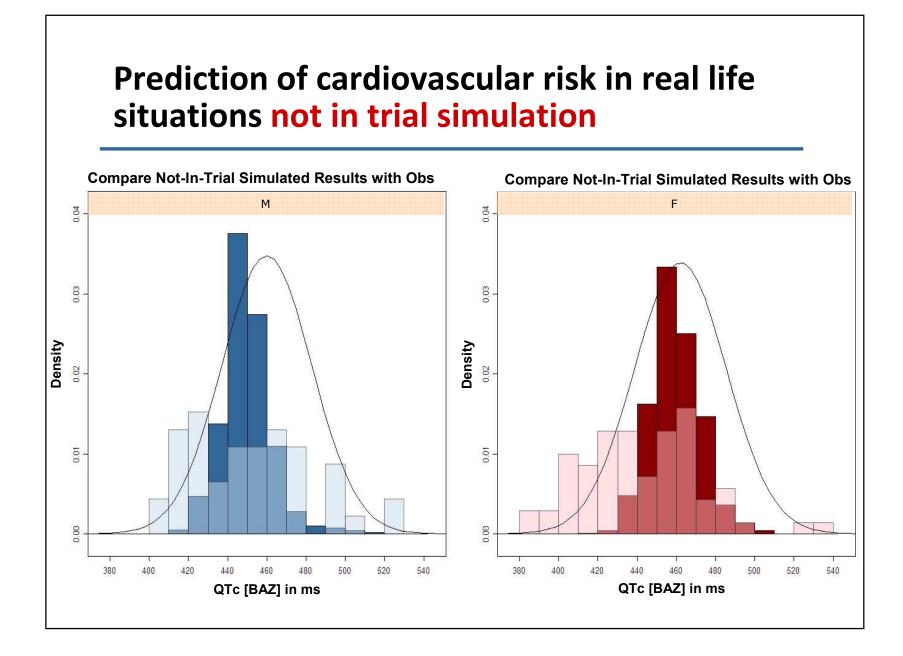
QTc = baseline + drug effect + co-morbidities + co-medications +  $\varepsilon$ 

"Rotterdam study"

- Baseline = sex; linear increase with age
- Co-morbidities = heart failure, MI, diabetes
- Co-medication = anti-arrhythmics
- Between subject variability

#### **Drug effect**

 $QT_c = QT_0 \times RR^{\alpha} \cdot (1 + A \cdot cos(2\pi/24 \cdot (clocktime - \varphi)) + slope \cdot C)$ 



#### Prediction of cardiovascular risk from ECG findings to sudden cardiac death

- Not in trial simulation to predict natural variation in QTc in the target population
- Analyze relationship between variation in QTc and cardiovascular risk in the target population
  - Delta analysis
  - Threshold analysis
- Develop and incorporate cardiovascular risk prediction model

**Prediction of cardiovascular risk** 

