

Overview of the e-LICO Project

e-LICO Kick-Off Meeting
Geneva, February 12-13, 2009



Main project deliverable

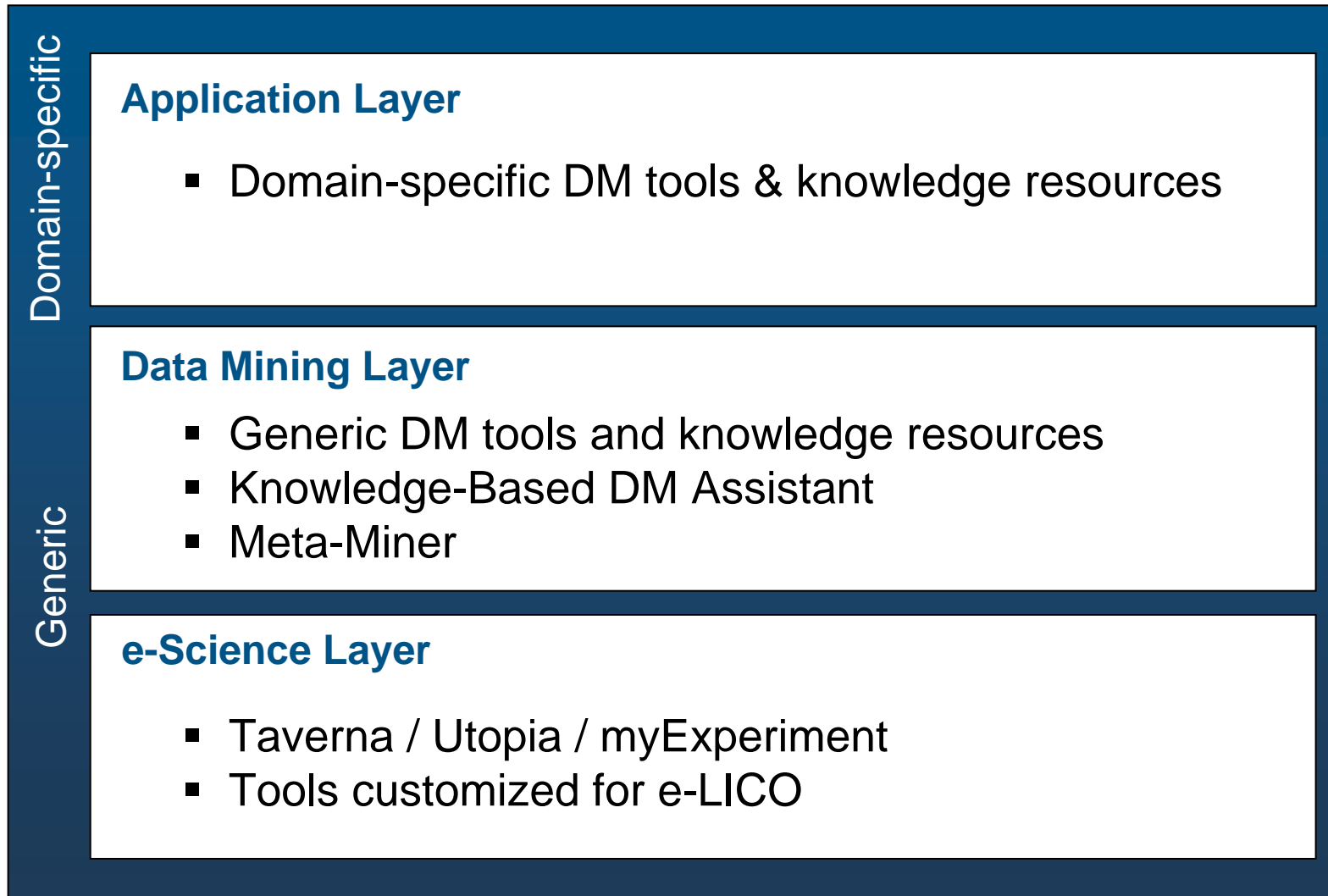
A virtual data mining laboratory that features

- a collaborative e-science infrastructure
- a self-improving DM assistant
- adaptability to different sciences

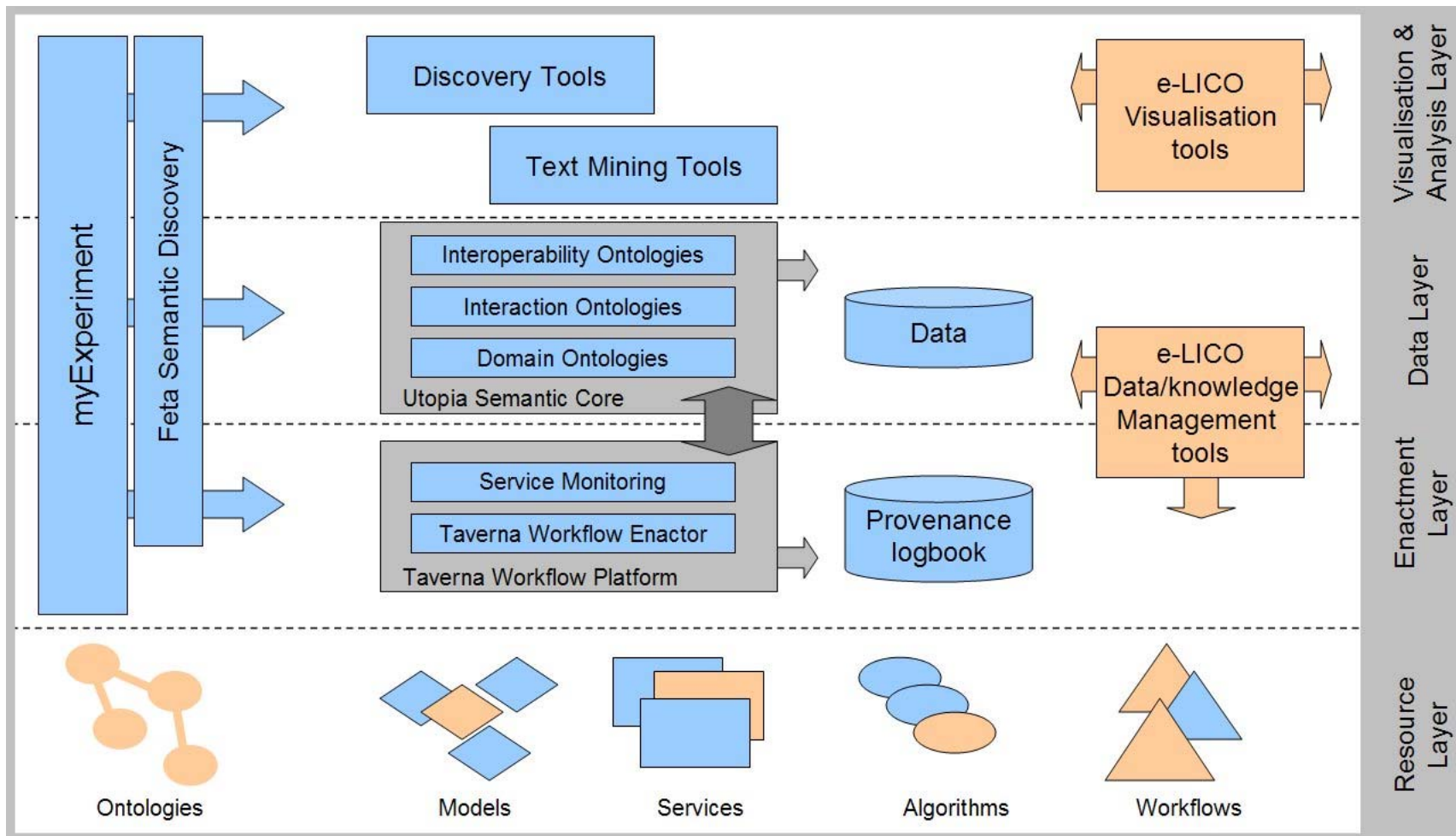
Key ideas

- ontology-based planning to build DM workflows
- incremental self-improvement through meta-mining
- a novel meta-learning technique that blends
 - probabilistic reasoning
 - kernel-based learning from complex structures
- reliance on social networks of committed scientists

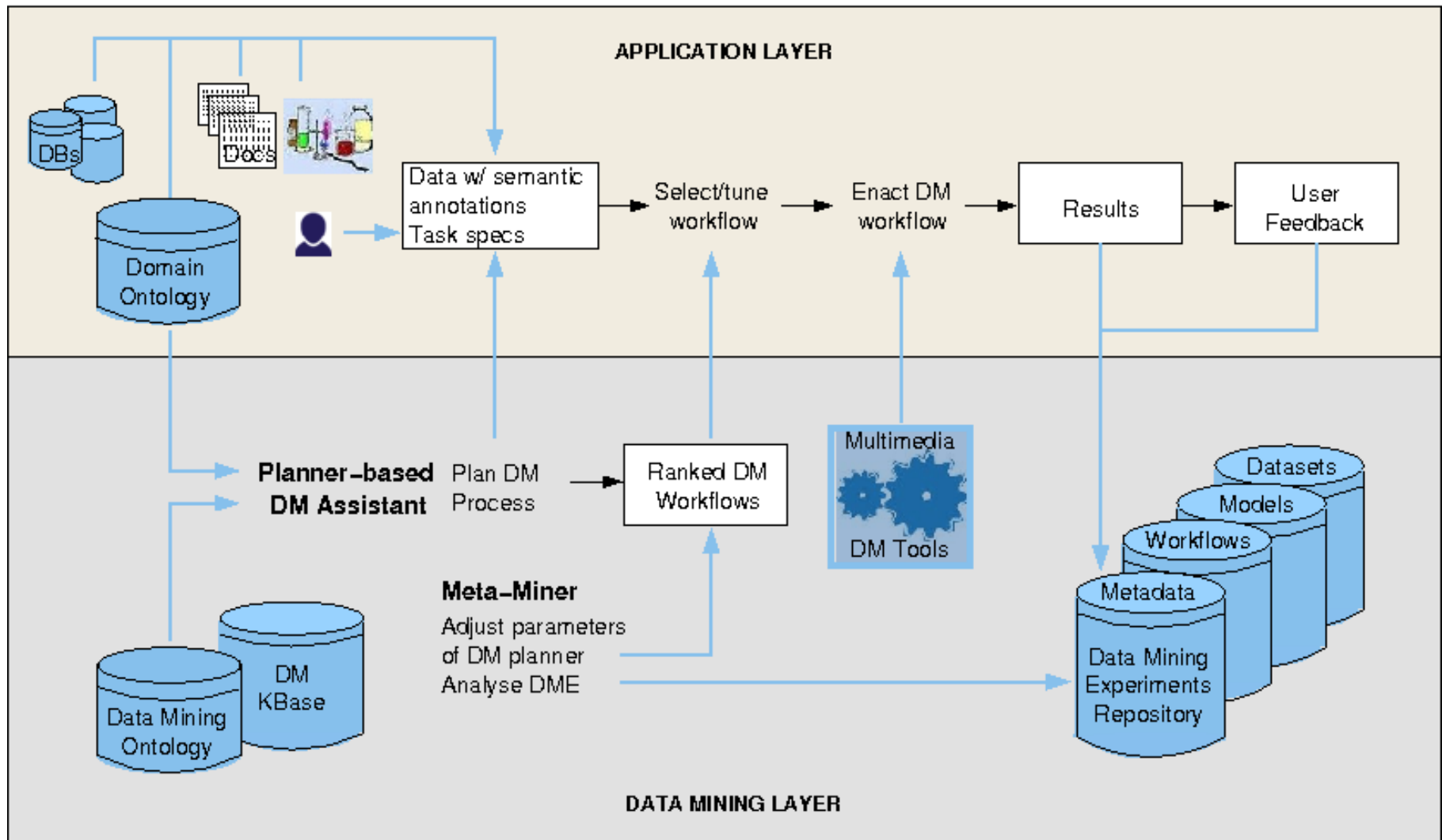
System Architecture



The e-science infrastructure



The DM and application layers



The pilot application

- The application layer is an empty shell
- Each instance of the DM lab comes from filling this shell
- Pilot domain: systems biology approach to the kidney and urinary pathways (KUP)
- Close links with COST Action EuroKUP (2008-2011)

Concrete project outputs

- **Major output: generic e-LICO**
 - Generic software
 - Generic DM knowledge sources
- **By-product: KUP application layer**
 - Biological software
 - Biological knowledge resources

Software

- e-LICO infrastructure and tools for collaborative authoring and annotation
- DM, TM and IM tools developed in the project
- DP, DM, TM and IM workflows
- Intelligent discovery assistant
- Probabilistic, kernel-based meta-miner

Content

- DM ontology
- DM KDB
- DM Experiment Repository (potential counterpart of UCI ML Repository for meta-data)

Software

- Biological data, text and image mining tools

Content

- KUP ontology
- KUP knowledge/data base
- Predictive models for KUP diseases
- Repository of KUP data mining experiments