



Decision Support for Antibiotic Prescription

Dr. Guy Tsafnat,

University of New South Wales, Centre for Health Informatics, UNSW

Line R. Sanden and Ninna Kæseler

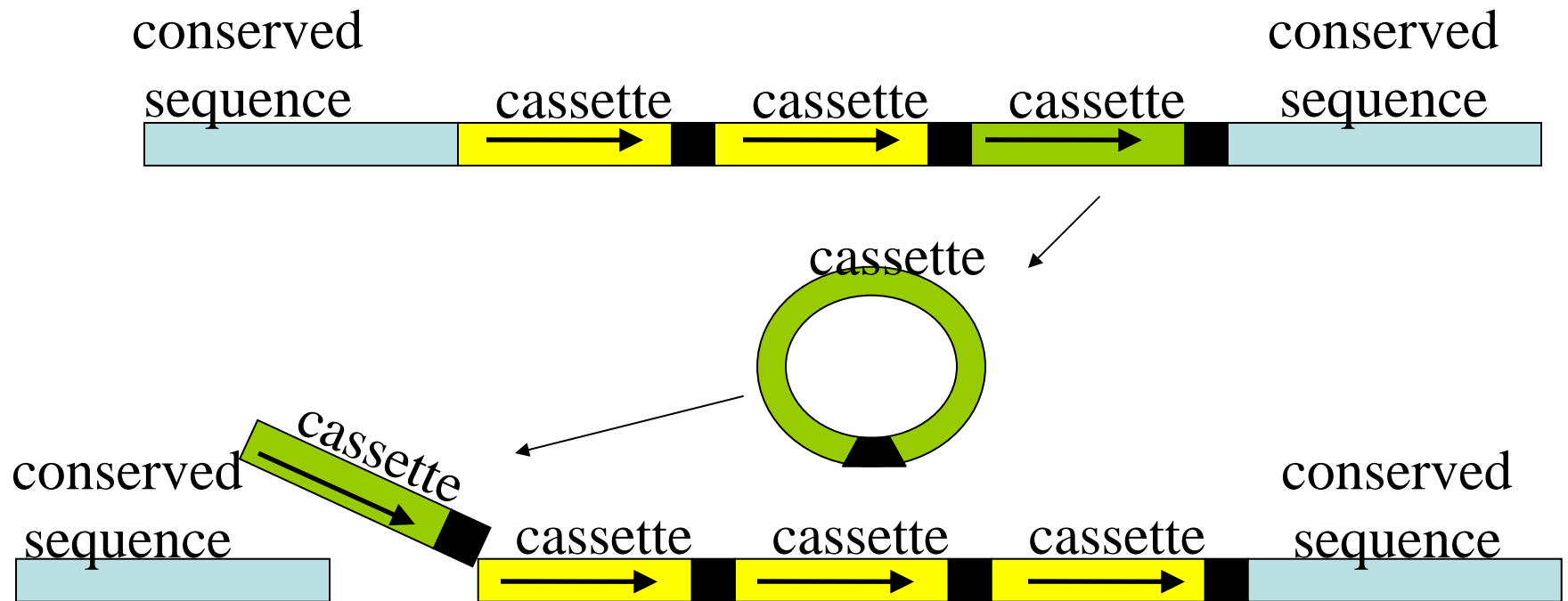
Dept. of Health Science & Technology, Aalborg University, Denmark

The ABR Problem

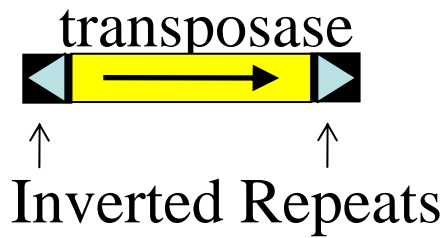
- Some bacteria are resistant to some antibiotics
- Antibiotic resistance spreads (horizontal transfer) and aggregates →
 - More multi-resistant infections
 - **Mobile Genetic Elements**
- Nosocomial (hospital acquired) infections
- Estimated (AU)\$2.5B per year in Australia
- **According to WHO, ABR is one of top three global health issues**



Gene Cassettes



Insertion Sequence



Transposition



Transposon



Integron

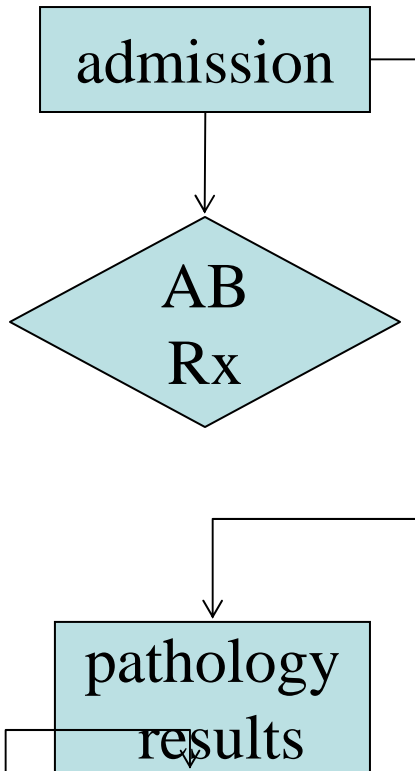


Composite
Transposon

The Challenge

- Most effective therapy for patient
- Minimal ecological impact
- Simple and quick pathology

Current Practice

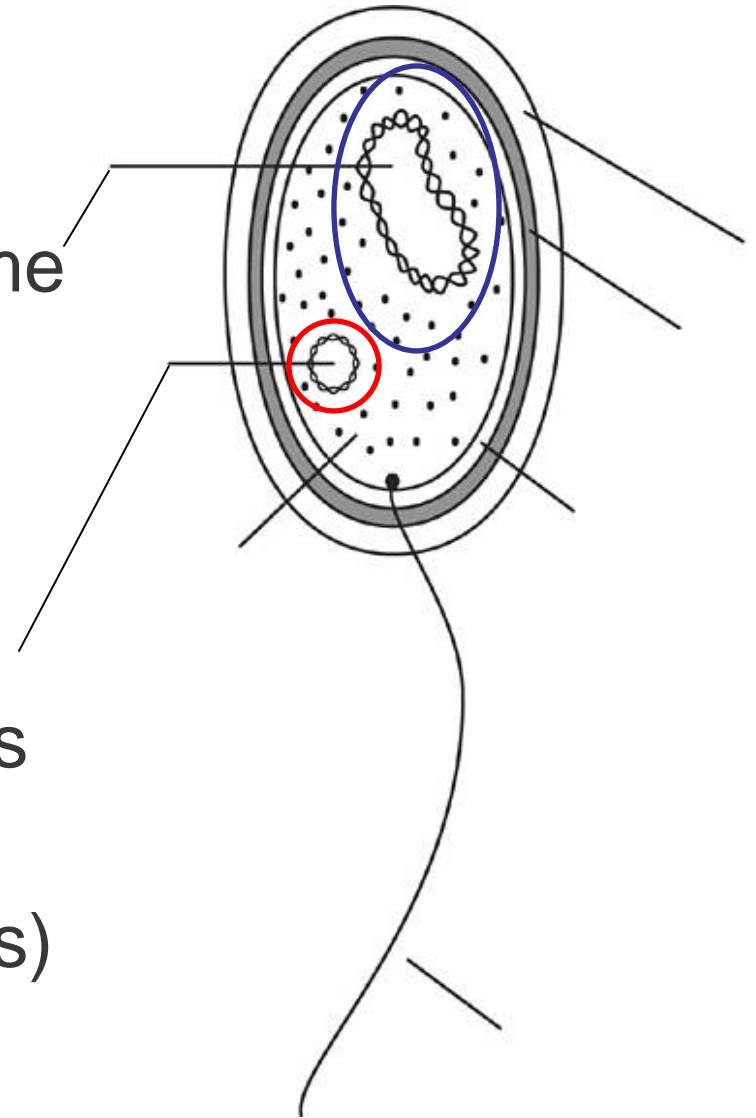


- “given patient symptoms, what is best practice?”
- Guidelines (pink book¹)
- Computed Guideline Support (Shahar et al.)
 - Not yet applied to AB
- Adaptive heuristic models based on any of: symptoms, patient records, pathology, risk
- Delayed pathology (2+ days)

¹ Centers for Disease Control and Prevention. *Epidemiology and Prevention of Vaccine-Preventable Diseases*. Atkinson W, Hamborsky J, McIntyre L, Wolfe S, eds. 10th ed. 2nd printing, Washington DC: Public Health Foundation, 2008.

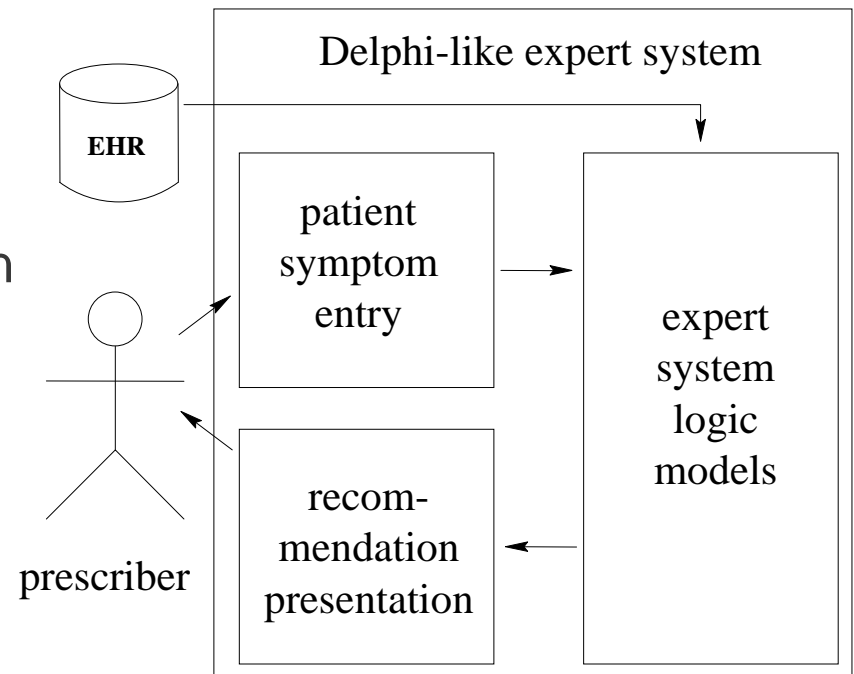
Current Pathology

- Measures the chromosome
 - Species
 - Strand
 - Morphology
 - Symptoms
- Should measure Plasmids
 - Multi-resistance regions
- Requires culturing (2 days)



Computational Support

- First developed 1976 (MYCIN)
 - shown some improvement in practice
- Modern systems:
 - Same model:
 - Clinician enters symptoms
 - Gets recommendations
 - Cost/benefit recommendation
 - Ecological impact
 - Risks to patient
 - Likely outcomes
 - Questionable adoption
 - No pathology

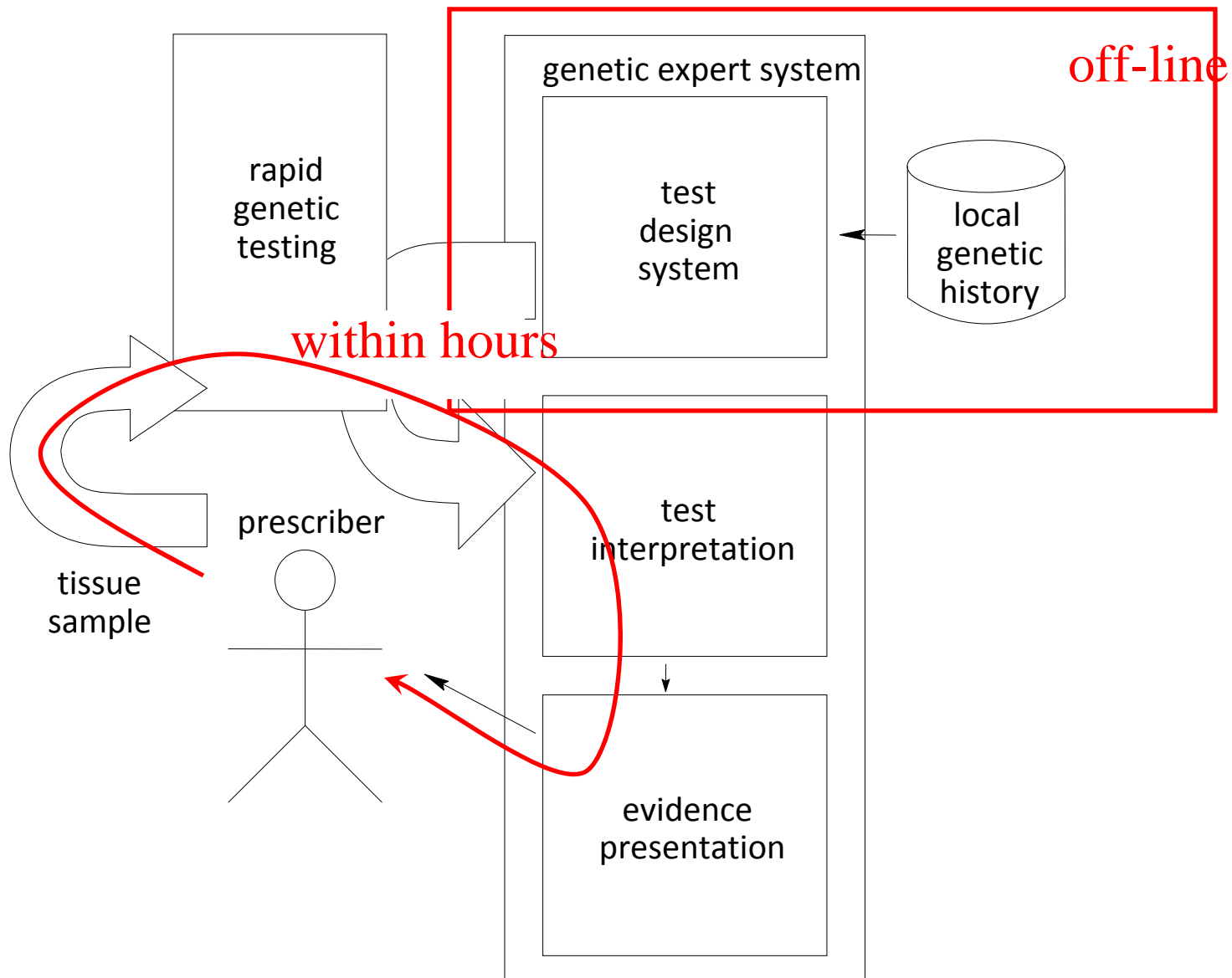


A new DSS Model for AB prescription

- Target the question:
 - “given a patient scenario and an environmental context, what is best action?”
- “Context” here means molecular data in the environment of the patient
 - Temporal
 - Monitored and adjusted
- Complements heuristics from existing systems

Approach

- Catalogue of molecular resistances
 - Mobile Genetic Elements
 - Species independent
 - Co-occurrences
- Specialised catalog per environment
 - Device, ward, hospital, city, country
 - Migration from other environments
 - Emergence from new sequences



Rapid Testing System

- A system that quickly checks for genes of interest without culturing
- Easily adjustable to new genes, tests
- Most likely PCR (2 under development)

Why PCR?

- PCR is a method for asking DNA yes/no questions for presence of sequence on interest
- Currently (likely always will be) cheaper than sequencing
- Cost per probe can be optimised further
 - [Hejlesen *et al.*, **DNA-Based Bayesian Inference and Analysis of Bacterial Antibiotic Resistance**, Scandinavian Health Informatics 2009]
- Extra probes can be used for monitoring



Offline set selection

- 2 structures
 1. 4 R genes: A, B, C, D
 2. 3 R genes: B, E, F
 3. 3 R genes: A, D, F
- $C \rightarrow B$
- $A \rightarrow C$
- $B \rightarrow C$
- $AB \rightarrow C$

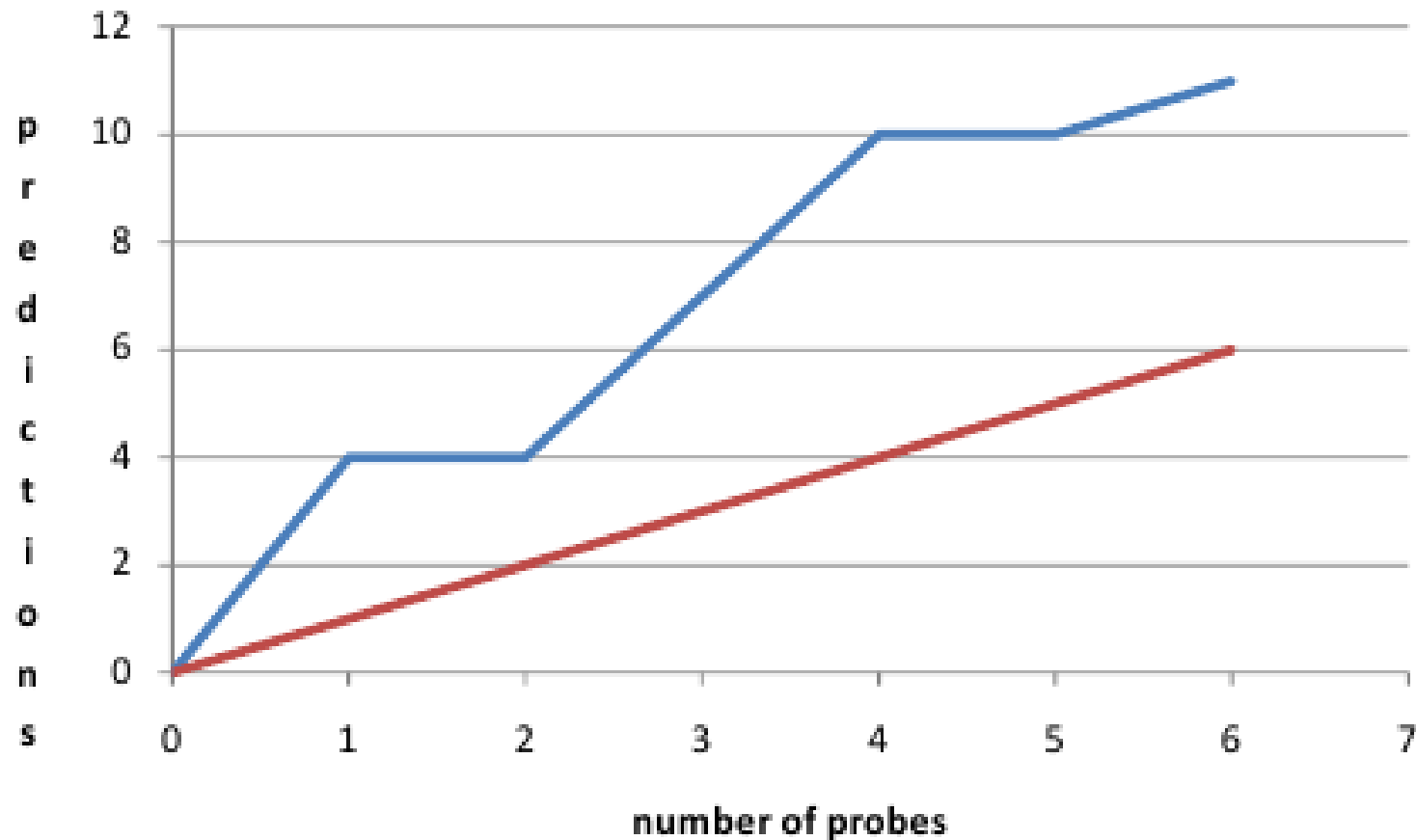
$$P = \frac{O(A,B)}{O(A)}$$

	A	B	C	D	E	F
A	2	1	1	2		1
B		2	1	1	1	1
C			1	1		
D				1		1
E					1	1
F						1



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Post-test results extrapolation



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Results Interpretation - Example

Offline selected probes (5/270):

aadB1a aacA4 dfrB17 blaVIM-1 dfrA1

Rapid test returns

x

x

x

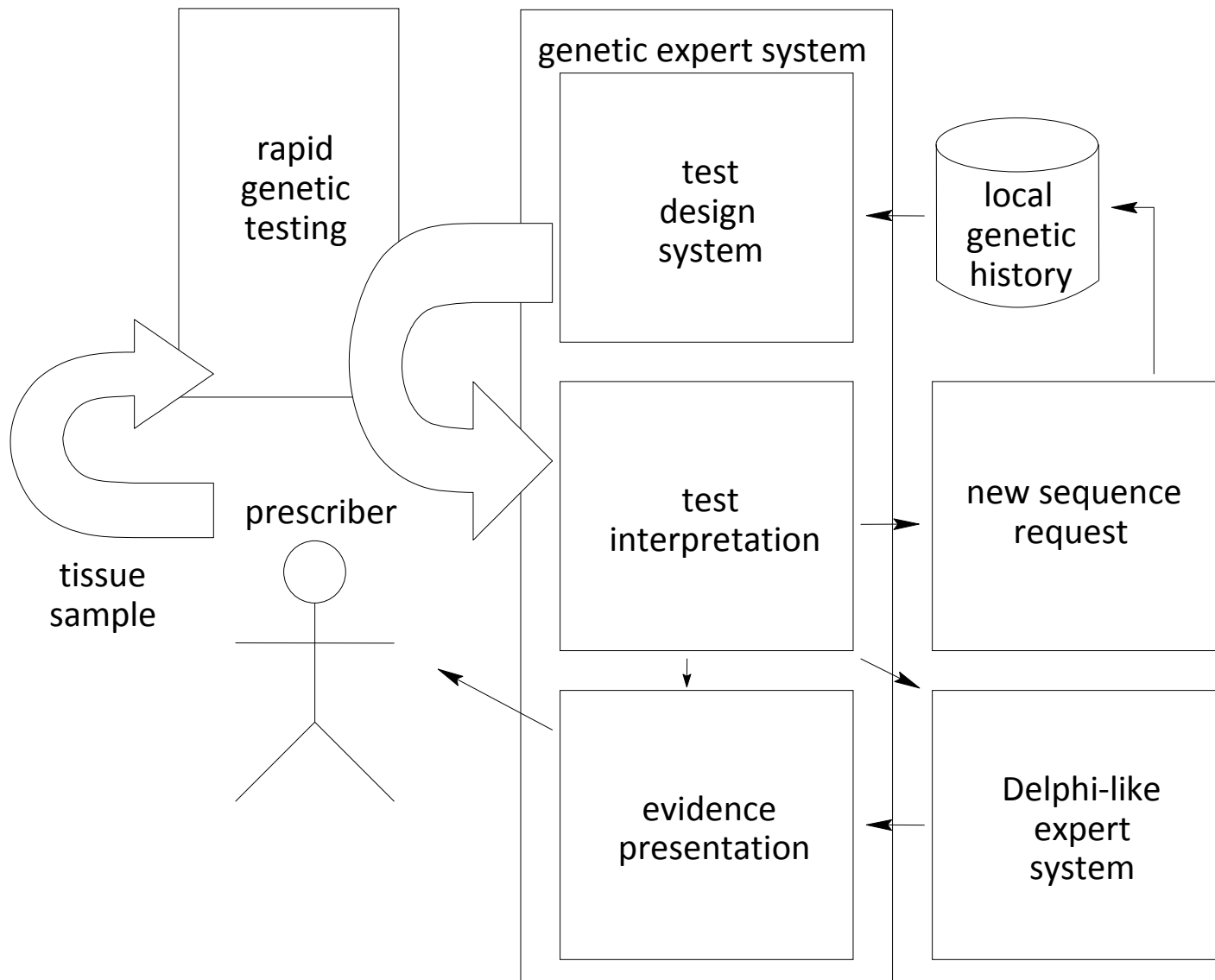
Interpretation adds

blaOXA-10

dfrB1

dfrB2

aadB





Thank You ...

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