



Session: Detecting antimicrobial resistance and treatment failure  
14.09.2015

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# Multiplex real-time PCR with High Resolution Melting analysis for detecting resistance mechanisms in *Neisseria gonorrhoeae*

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## Objectives/Methods

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Development of a **rapid diagnostic test** for the identification of *N. gonorrhoeae* and of potential **antimicrobial resistance** determinants directly from **clinical specimens**



**SybrGreen-based real-time PCR with High Resolution Melting** analysis of gDNA after direct extraction from eSwabs

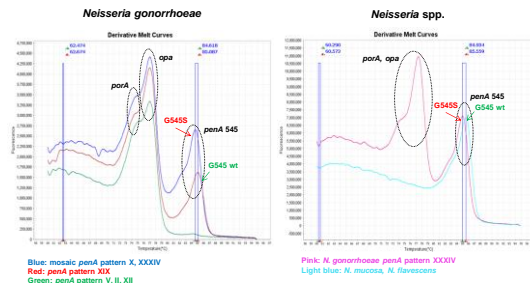
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## Results: Triplex *opa* + *porA* + *penA* 545

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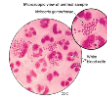
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## Background

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- > **Culture** ⇨ gold standard; antimicrobial susceptibility testing possible, but requires at least **2 days**
- > **NAATs** ⇨ sensitive and rapid, but **do not provide** information about **antimicrobial susceptibility**

Antibiotics	1998-2001		2010-2012	
	% non-S	MIC <sub>50</sub>	% non-S	MIC <sub>50</sub>
Penicillin	42.3	3	85.3	16
Spectinomycin	0.0	12	0.0	8
Ciprofloxacin	7.7	0.006	73.5	≥32
Cefixime	0.0	≤0.016	8.8	0.125
Ceftriaxone	0.0	0.004	0.0	0.047
Azithromycin	11.5	0.25	23.6	0.38

Endimiani et al., 2014

Antimicrobial resistance determinants

16S rRNA	C1192T
SS rRNA	Thr24Pro
GyrA	Ser91Phe
PBP2 ( <i>penA</i> )	Mosaic (i.e., Gly545Ser), Ala501Pro/Val7Thr
23S rRNA	A2059G - C2611T

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## Methods



**2 detection + 6 antimicrobial resistance targets**

Triplex	<i>opa</i> , <i>porA</i> , <i>penA</i> Gly545Ser
Duplex #1	GyrA Ser91Phe and 23S rRNA A2059G
Duplex #2	23S rRNA C2611T and <i>penA</i> 501
Duplex #3	16S rRNA C1192T and 5S rRNA Thr24Pro

Control strains used for method validation:

- > **REFERENCE STRAINS:**
  - Ceftriaxone-R (mosaic *penA* XXXIV, Ala501Pro)
  - Azithromycin-R (23S rRNA A2059; 23S rRNA C2611T)
  - Spectinomycin-R (16S rRNA C1192T; 5S rRNA)
  - Other WHO, ATCC strains
- > **FULLY CHARACTERIZED ISOLATES:**
  - GyrA Ser91Phe (n=22)
  - *penA* mosaics (XXXIV, n=7) and non-mosaics
- > **NON-GONOCOCCAL SPECIES** (n=10)

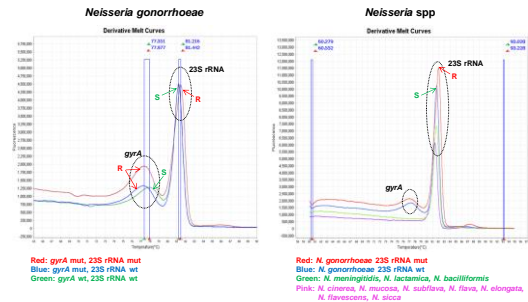
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## Results: Duplex #1 GyrA Ser91Phe + 23S rRNA A2059G

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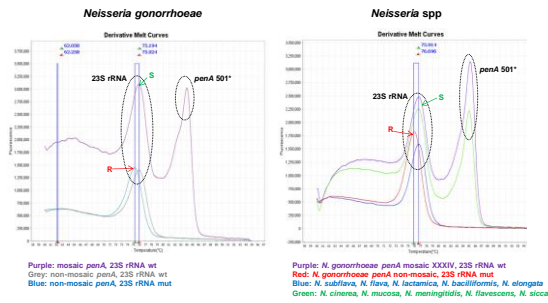


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## Results: Duplex #2

### 23S rRNA C2611T + penA 501

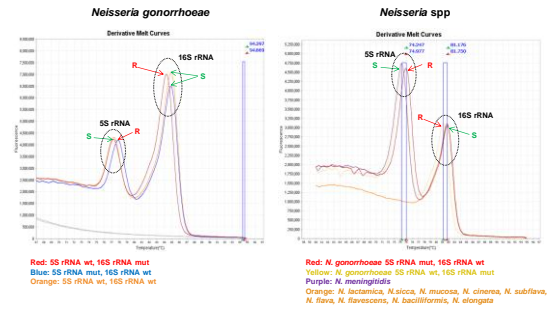


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## Results: Duplex #3

### 5S rRNA Thr24Pro + 16S rRNA C1192T



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## Results:



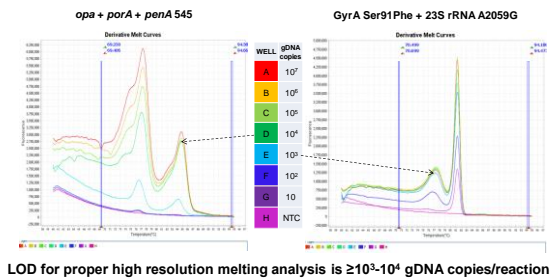
Control strains ( <i>Neisseria</i> spp., n=20)			Clinical isolates (n=99)		
Molecular target	sensitivity	specificity	Identification/AMR	sensitivity	specificity
opa	100%	100%	Detection of NG	100% (99/99)	NA
porA	100%	100%			
GyrA Ser91Phe	100%	100%	Ciprofloxacin	100% (20/20)	100% (79/79)
23S rRNA A2059G	100%	100%	Azithromycin	2.5%* (1/40)	100% (99/99)
23S rRNA C2611T	100%	100%			
Mosaic penA	100%	100%	Cephalosporins	NA	99%* (98/99)
16S rRNA C1192T	100%	100%	Spectinomycin	NA	100% (99/99)
5S rRNA Thr24Pro	100%	100%			

Gold standard: PCR/Seq      Gold standard: MALDI-TOF MS/MIC (Etest) (EUCAST 2015)

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## Results: Limit of detection (LOD)



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## Conclusions



### Advantages:

- Rapid (1.5 h) and cheap
- Multiplex 2 or 3 reactions/well
- Good sensitivity and specificity when testing main resistance targets in isolated cultures

### Disadvantages:

- Technical issues (i.e., G/C SNPs)
- Needs high limit of detection for proper high resolution melting analysis
- Cross-reaction with *Neisseria* spp.

Suitable for screening of isolated cultures but not for clinical specimen with low target gDNA (e.g., pharyngeal samples)

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