

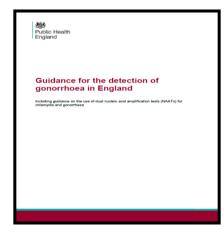
Prevalence of *por A* pseudogene deletion amongst *N. gonorrhoeae* isolates referred to GRASP

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Introduction

- PHE recommends confirmatory testing GC NAAT Positives >90%
- Can be problematic to confirm using commercial platforms
- Several In-house assays available: opa gene, 16s rRNA, porA pseudogene



- porA pseudogene conserved in NG & significant different in NM ideal sensitivity & specificity
- porA negative GC have been reported Australia, Sweden, Scotland & England
- **Aim:** Determine the prevalence of *porA* negative GC isolates in England & Wales using isolates referred to GRASP

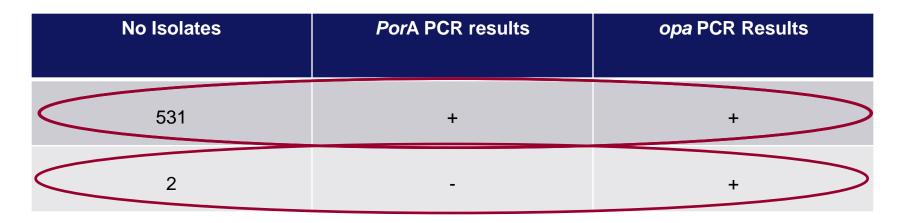




- GRASP sentinel surveillance study
- Regional representation of GUM clinic patients
- DNA lysates were prepared from 533 *N. gonorrhoeae* isolates 2011
 - 20 centres around England & Wales
- Tested using an in house RT-PCR for the *porA* and *opa* gene
- DNA sequencing of discrepant strains







99.6% - GC isolates porA positive

0.4% - GC isolates *porA* negative



DNA Sequencing Results

мс	ACTCTCCGGACTTTTCCGGTTTCAGCGGCAGCGTCCAATTCGTTCCGGCTCAAAACAGCA
GC	ATTCCCCCGGATTTTCCGGTTTCAGCGG <mark>CAGCATTCAATTTGTTCCGAGTC</mark> AAAACAGCA
	* ** ** * *****************************
МС	AGTCCGCCTATACGCCGGCTTATG-TGGATAAGGAGCAGGTGTCTCATGCGG
GC	AGTC <mark>CGCCTATACGCCTGCTACTTTCACGC</mark> TGGAAAGTAATCAGATGAAACCAGTTCCGG

- Primers and probes target gonococcal specific regions
- 2 porA PCR negative GC Isolates *N. mengitidies porA gene*
- False negative GC porA PCR results





- *por*A pseudogene is a popular target for confirmatory assays
- 0.4% (2) gonococcal isolates were identified as *porA* negative in England
- Both GC Isolates have incorporated MC porA gene into their genome
- Front line and confirmatory strategies remain a challenge for GC NAATs
- Neisseria are genetically fluid and competent all stages of their life cycle
- *porA* pseudogene is a nationally accepted target for a supplementary test
- Microbiologists must be vigilant to decreases in numbers of confirmatory tests



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