



Background

- Lopinavir co-formulated with ritonavir (LPV/r) has been an extensively used protease inhibitor (PI) against HIV
- Development of PI resistance reduces sequencing options
- Identifying predictors of PI resistance might better our ability to sequence therapy
- Many studies report few resistance mutations under lopinavir drug pressure (in virological) failure
 - can be difficult to identify LPV associated resistance pathways

Aims

- Evaluate patients in the UK failing LPV/r as their first PI
- Quantify the prevalence of PI resistance in this group
- Explore the patterns of PI mutations
- Identify possible factors contributing to the development of PI resistance

Methods

- *Pol* gene sequences retrieved from the UK HIV Drug Resistance Database
- Demographic and clinical data were obtained via linkage to UK CHIC
- Eligible patients were receiving LPV/r as their first PI
 - ART naïve
 - having previously received non-PI based regimen
- Virological failure defined at 6 months as:
 - >400 c/ml after previous suppression to <400 c/ml
 - >400 c/ml for the first 6 months





Pata from a large clinical cohort suggests prior (NNRTI based) ART failure does not compromise subsequent LPV/r response RAMs at positions 32 46 47 54 76 82 were associated with LPV/r failure 3056 patients included 811 (27%) failures 291 resistance tests 32 showed resistance (4% of failures but 11% of those failing with a test) No demographic factors found to be associated with risk of LPV/r failure



- We also looked at length of LPV exposure to detectable viraemia by looking at viral area under the curve (AUC)
- Increasing mutation risk in failures with high AUC suggests that maintaining LPV in a failing regimen may have significant genotypic resistance costs
 - although those with resistance had predicted sensitivity (Stanford) to other PIs – TPV (81%) – DRV (84%)

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