**Supporting information S3. STARD flowchart.**

Index test

n=110

Final diagnosis (genotyping)

* Wild-type=90
* Mutant-type=8
* Mixed=12

Reference standard (conventional sequencing) n=85

* Wild-type=65
* Mutant-type=8
* Mixed=12

No index test

n=0

No reference standard

n=0

No reference standard

n=0

No reference standard

n=25

Index test negative

n=0

Index test positive

n=110

Index test inconclusive

n=0

Eligible participants

n=110

Potentially eligible participants

n=110

Excluded

n=0