

Genotyping and phenotypic testing of RSV

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Northern France**



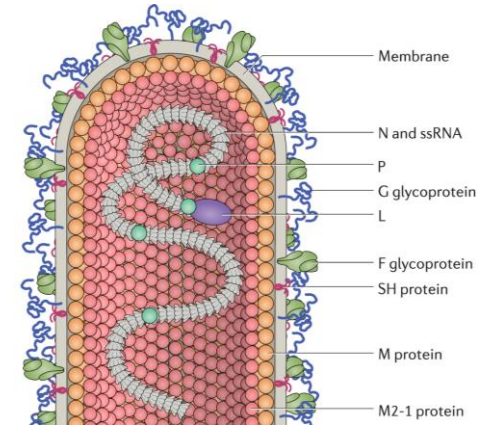
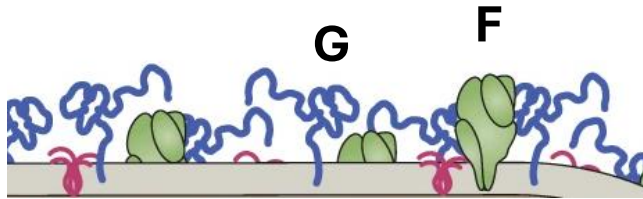
Objectives of virological surveillance ?

- **Description of RSV clades circulation**

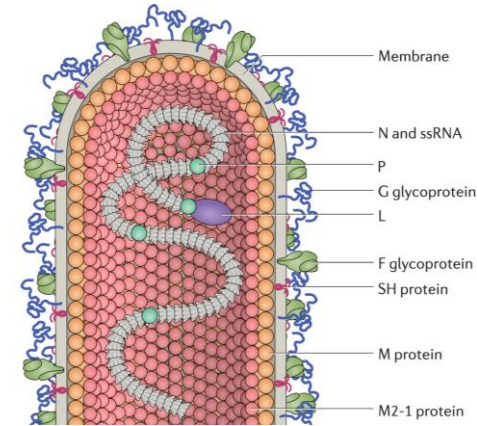
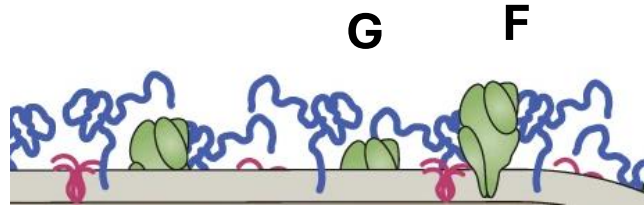
➡ Identification of clades of interest

- **Assessing and reporting susceptibility to**
 - **Monoclonal antibodies**
 - Vaccines
 - Antiviral drugs

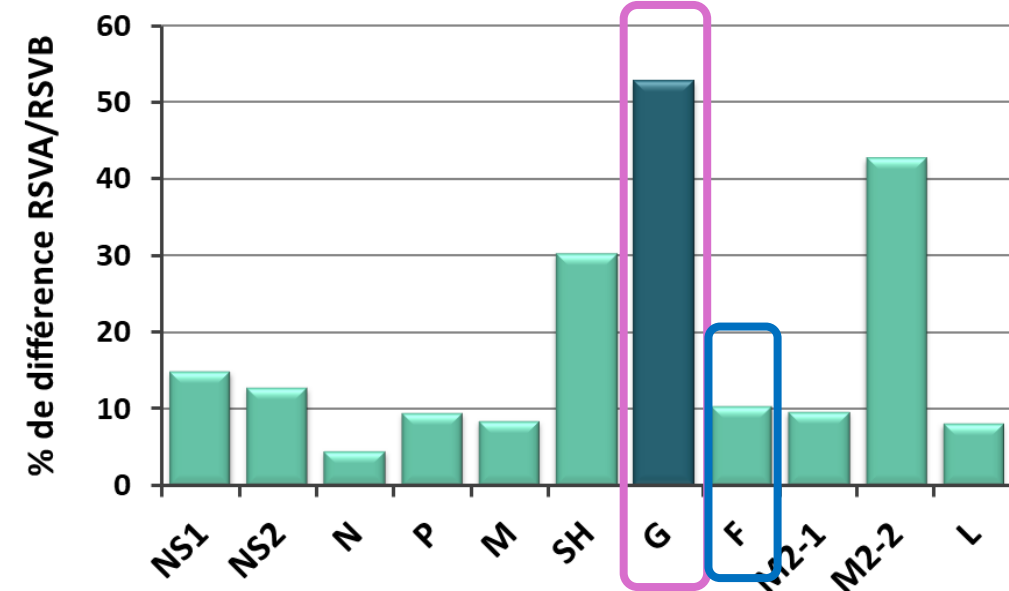
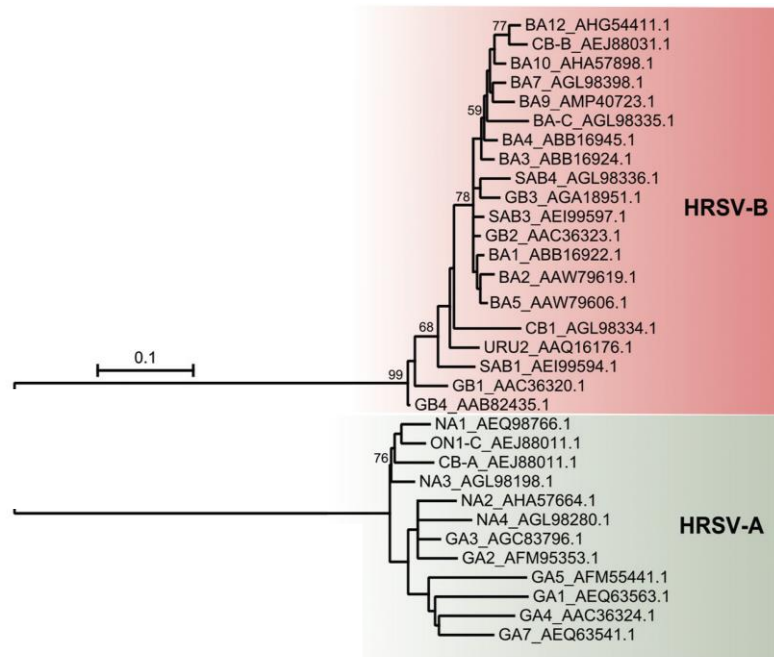
RSV has two major surface glycoproteins



RSV variability concentrated on G protein



2 groups : RSV-A and RSV-B



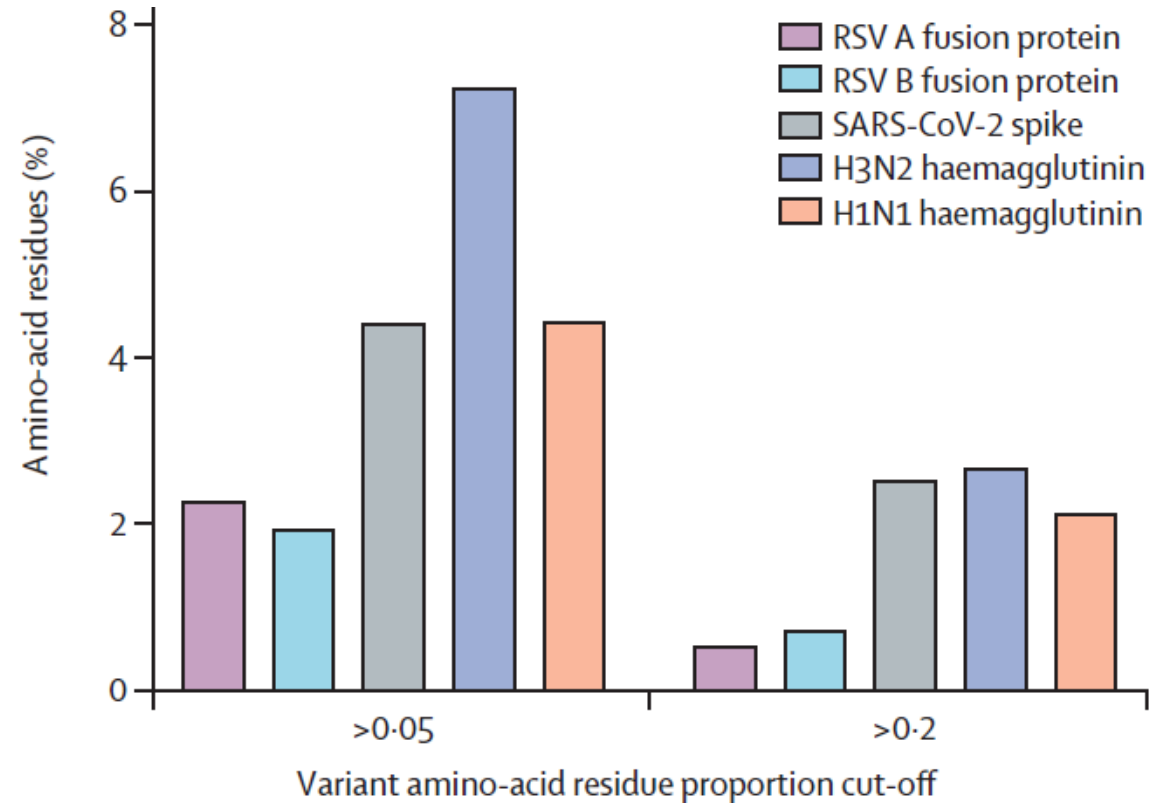
From data by Bose et al, 2015 (27 RSV-A and 19 RSV-B from virus origins over 10 years).

F protein is the main target for monoclonal Ab, vaccine (and antivirals)

F : highly conserved

structural constraints ?

limited selection pressure ?



Description of RSV clades: Recent definition of RSV lineages

Based on whole genome sequences from the first NS1 codon to the last L codon

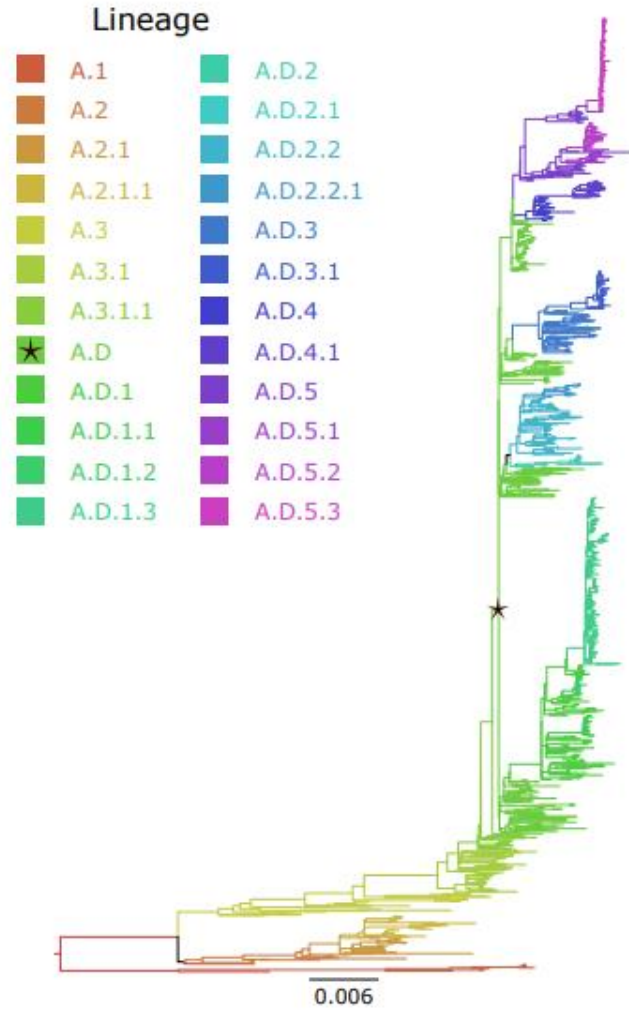
Minimal misclassification (1.2% error) was found in RSV-A and none in RSV-B when using only G sequence

New lineage definition: **>10 seq with more than 5 coding mutations**

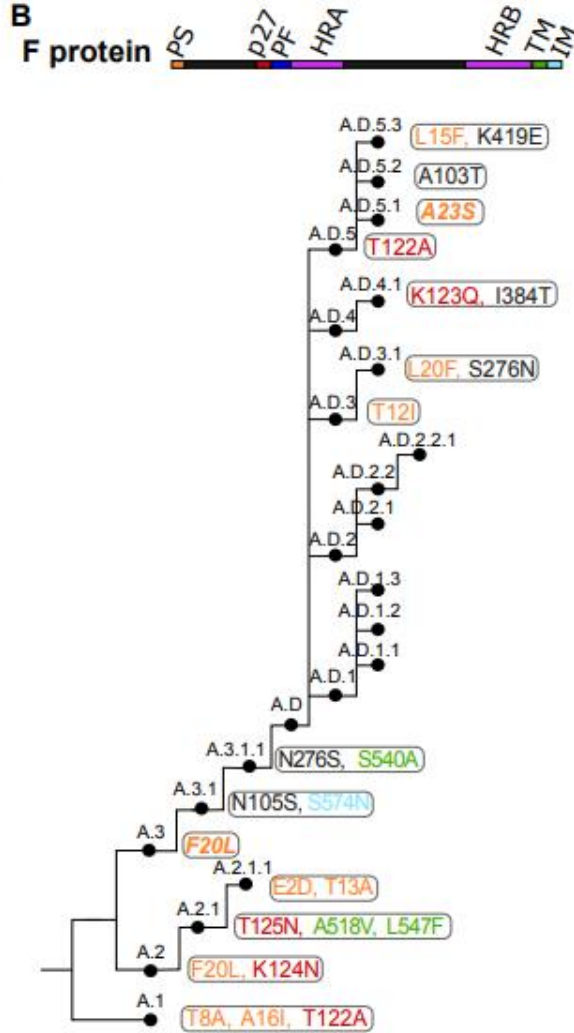
Recent definition of RSV lineages

ECDC NORMAL

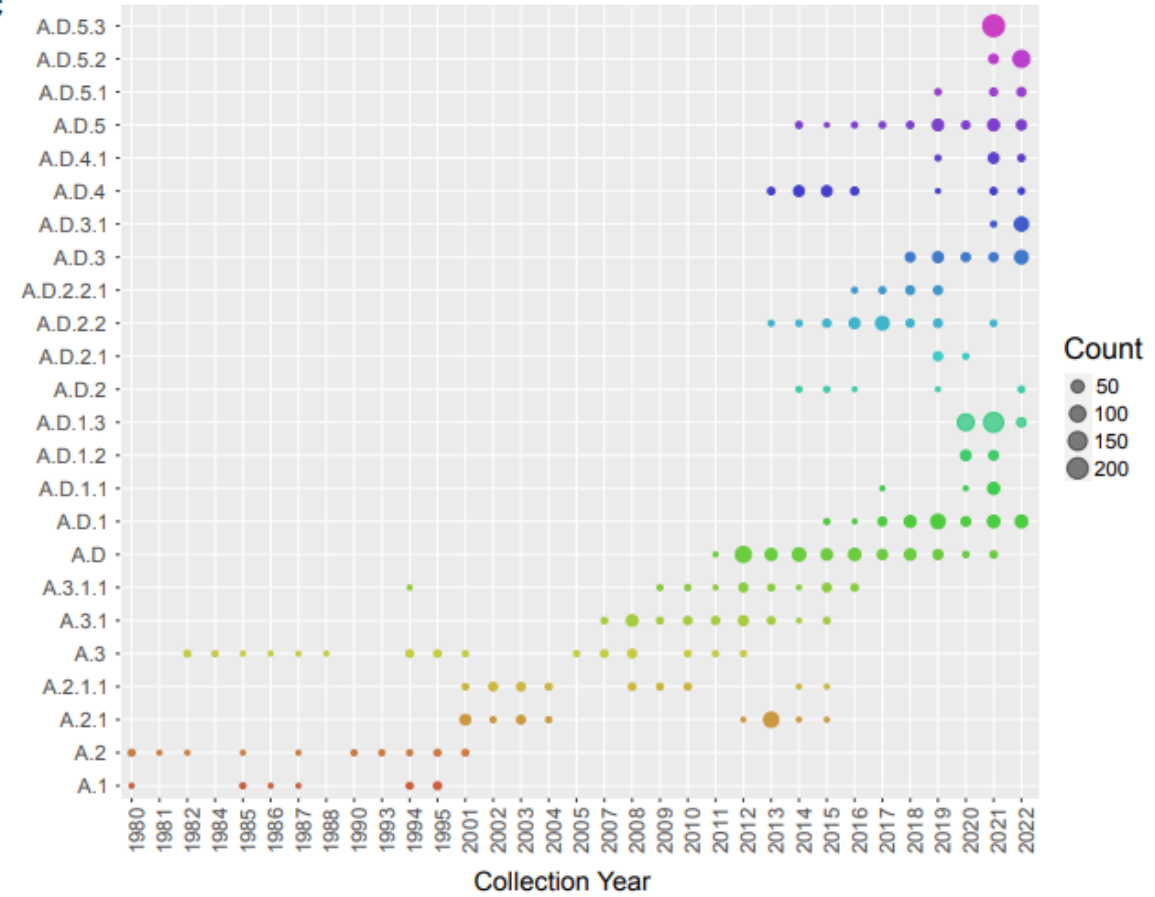
A



B



C



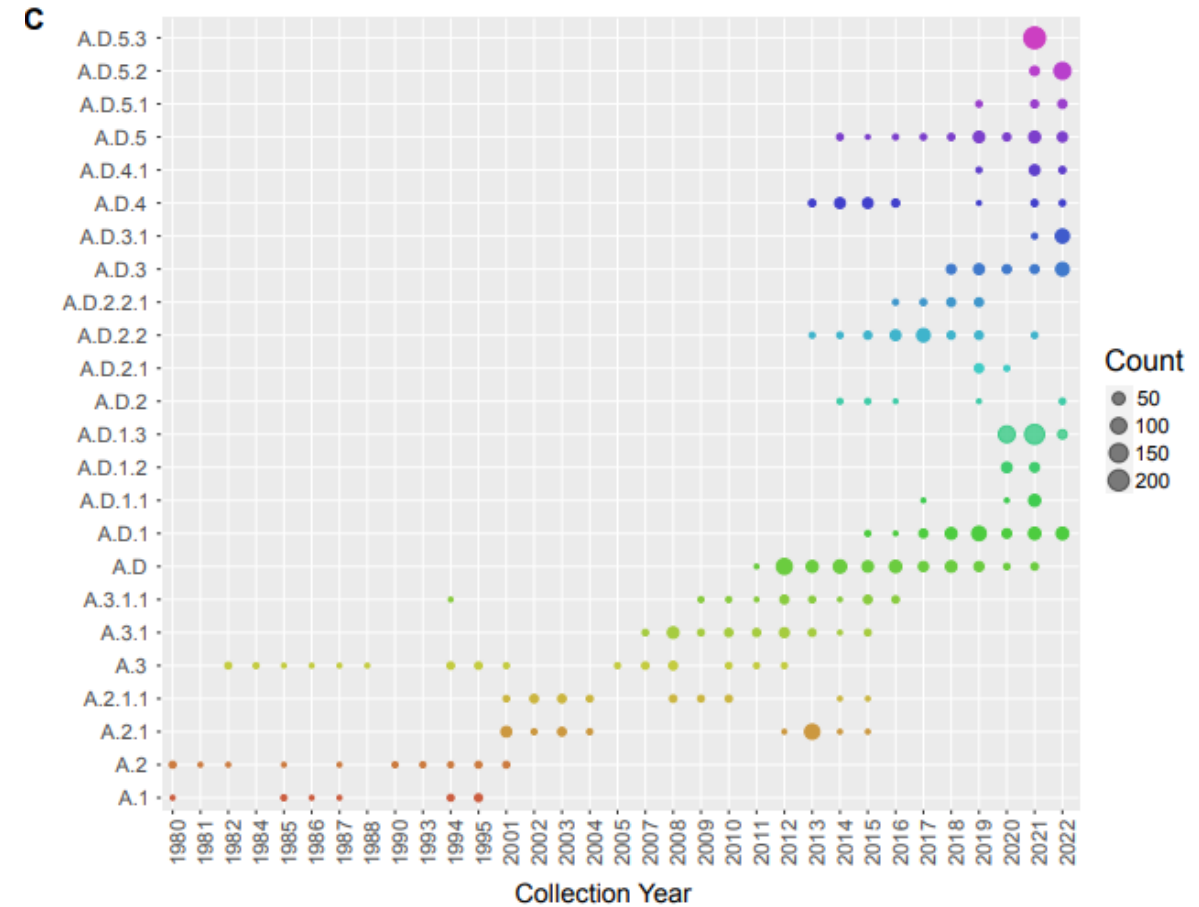
Recent definition of RSV lineages

ECDC NORMAL

Co-circulation of different lineages

Multiple introductions

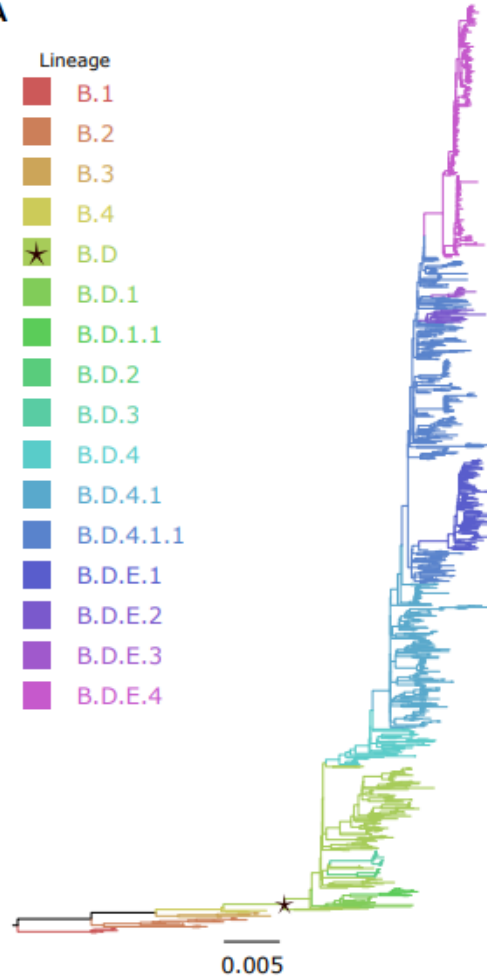
Persistence of some lineages



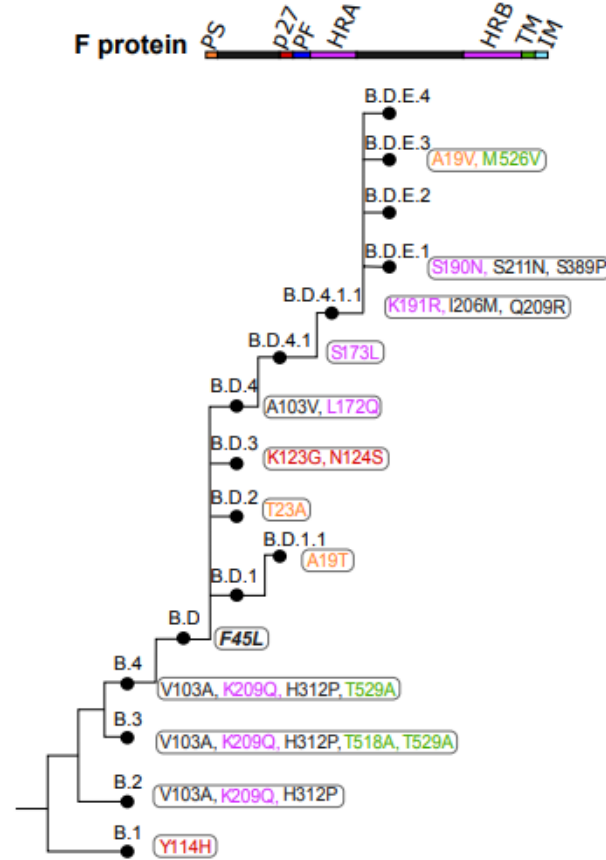
Recent definition of RSV lineages

ECDC NORMAL

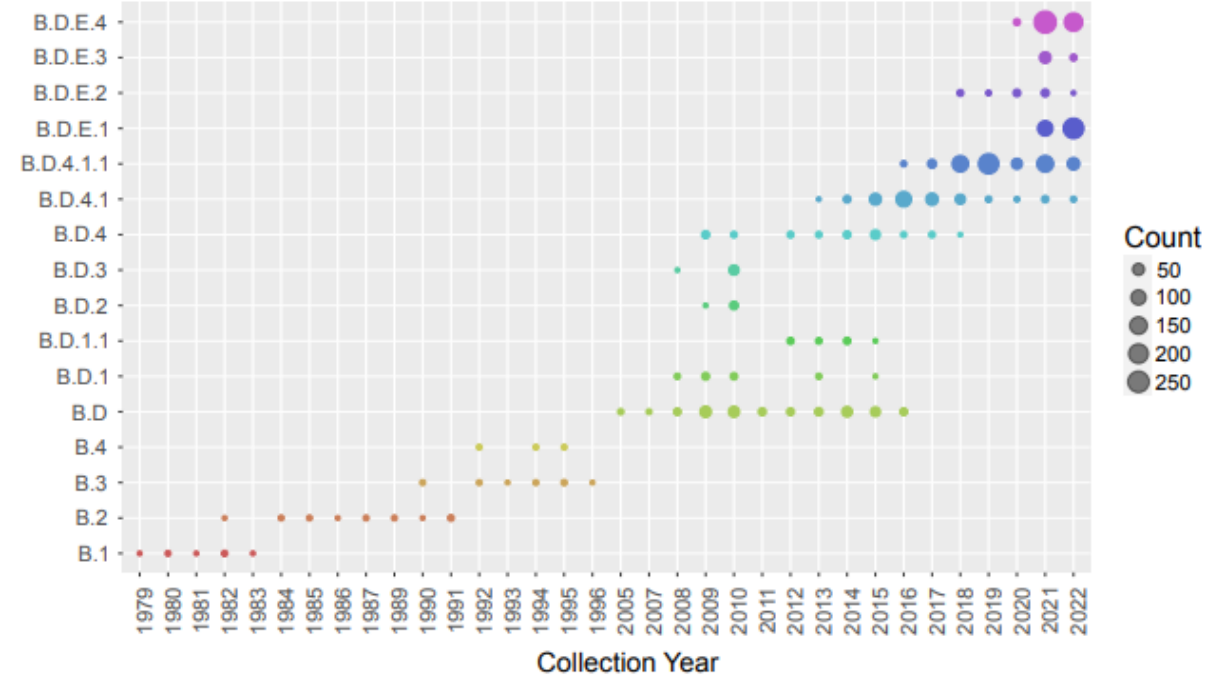
A



B



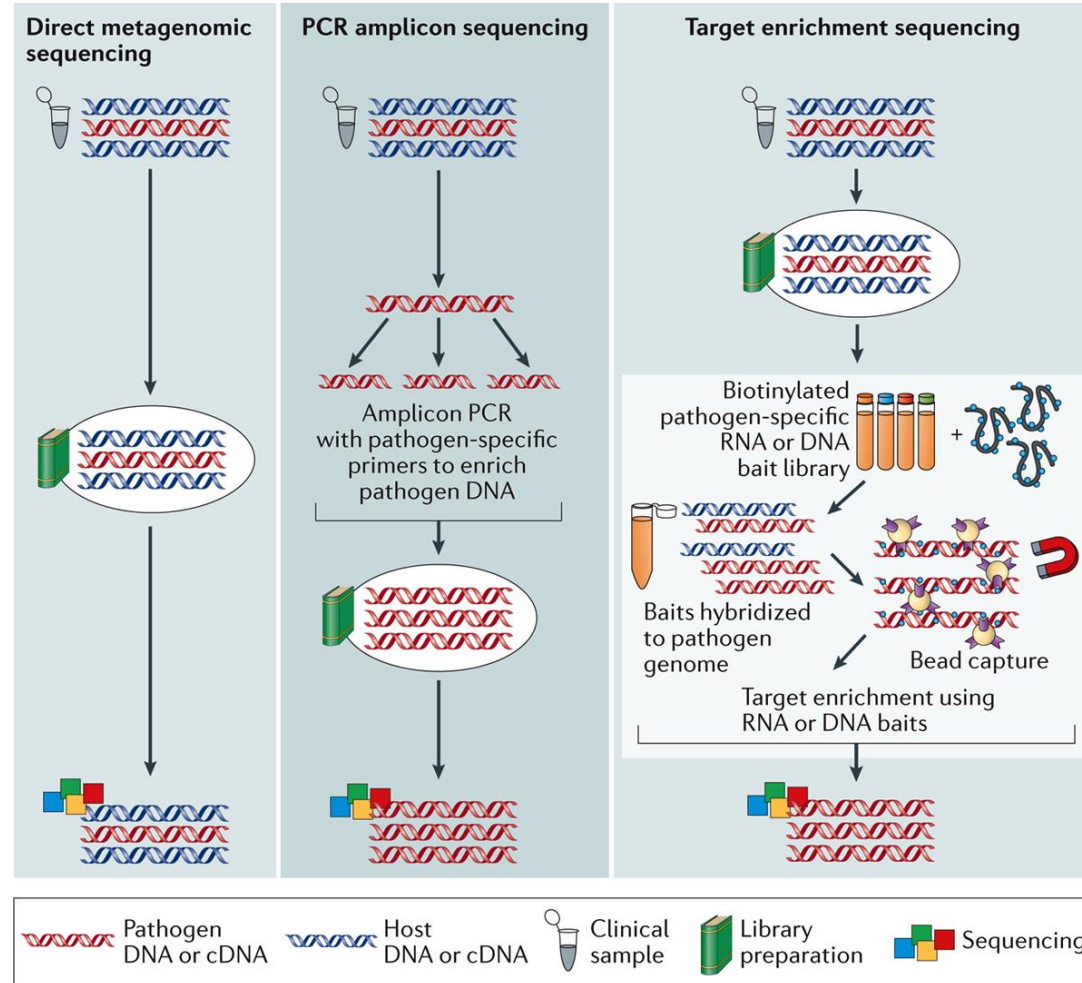
C



ECDC NORMAL Genotypic analysis of RSV

Sanger sequencing of G as a proxy for lineage assignment and of F to look for escape mutations

Reference method: whole genome sequencing



ECDC NORMAL

Genotypic analysis of RSV

Bruno Simon *et al*; Preliminary results of a comparison of methods

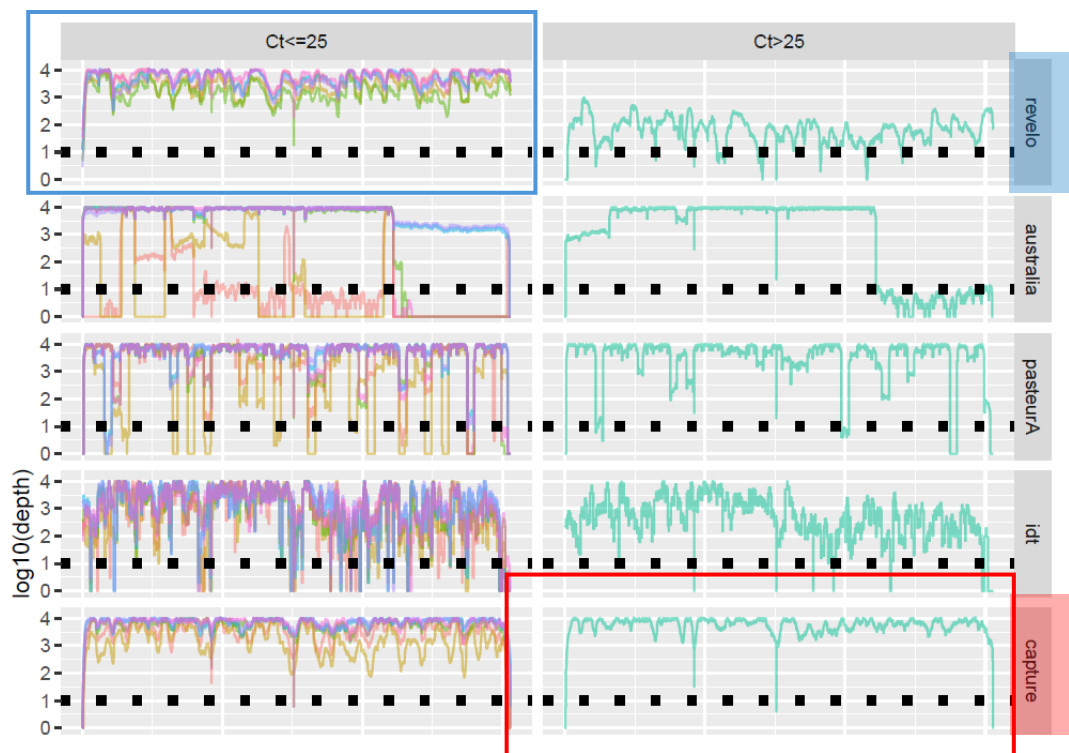


GENomique
EPIdémologique
des maladies
Infectieuses

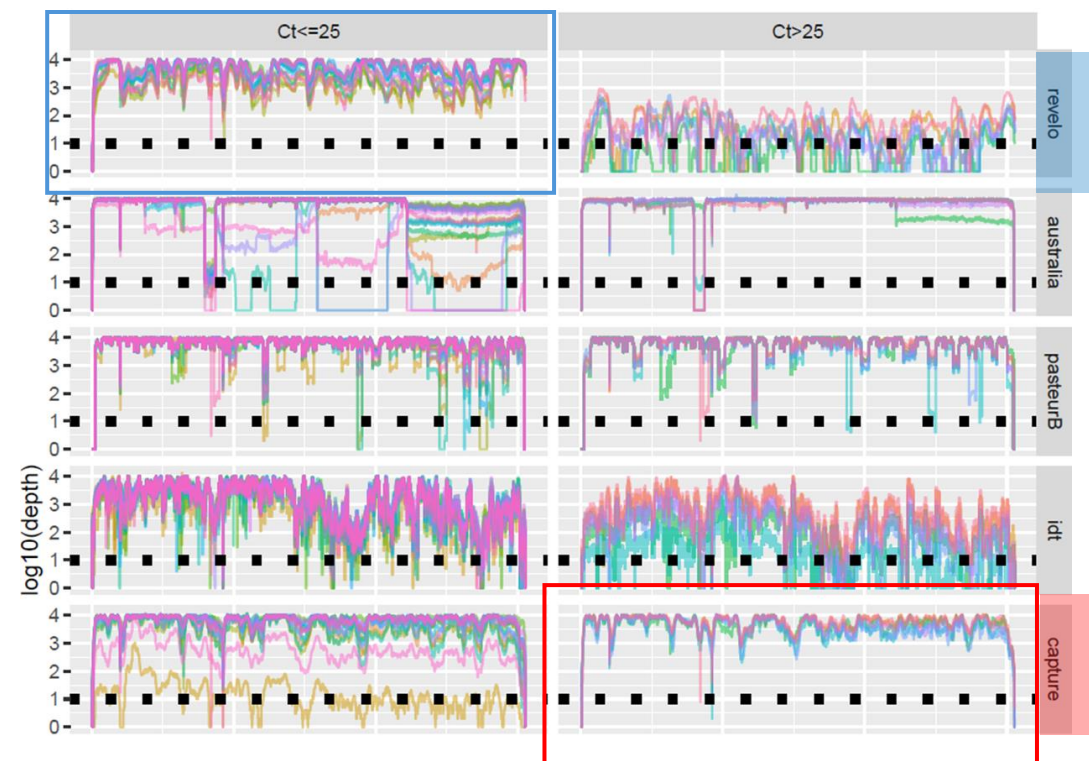


- **Metatranscriptomic: Revelo** RNA-Seq High Sensitivity (*Tecan Genomics*)
- **Amplicon: Australia** (*Dong et al, JCV, 2023*)
- **Amplicon: G5 GEVA, Pasteur** *Etienne Simon Lorière*
- **Amplicon: xGen Respiratory Virus Amplicon Panel (IDT)**
- **Capture** : Illumina RNA Prep with Enrichment Tagmentation – RVOP2 Panel

RSV-A n=24



RSV-B n=8



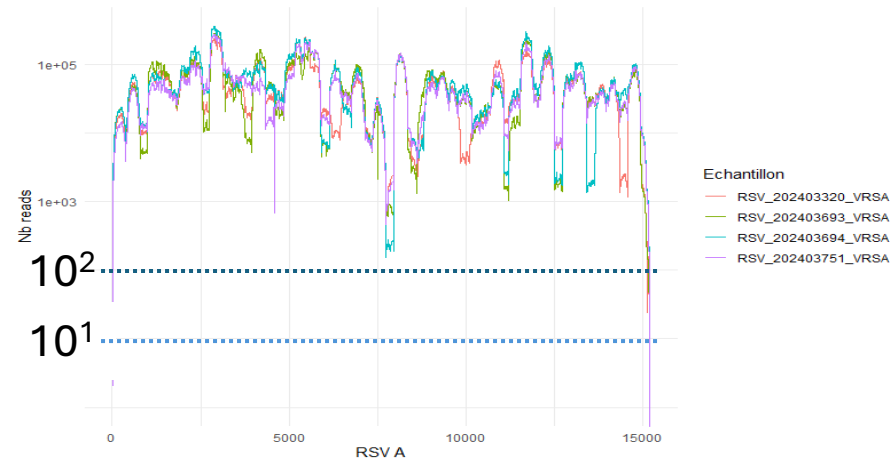
Amplicon methods

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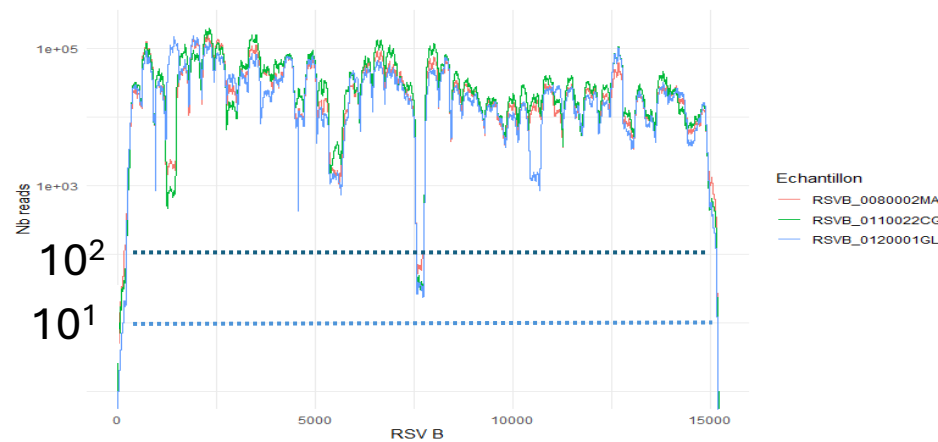
Update of Pasteur Protocol

Amplicons: 400 pb

RSV-A

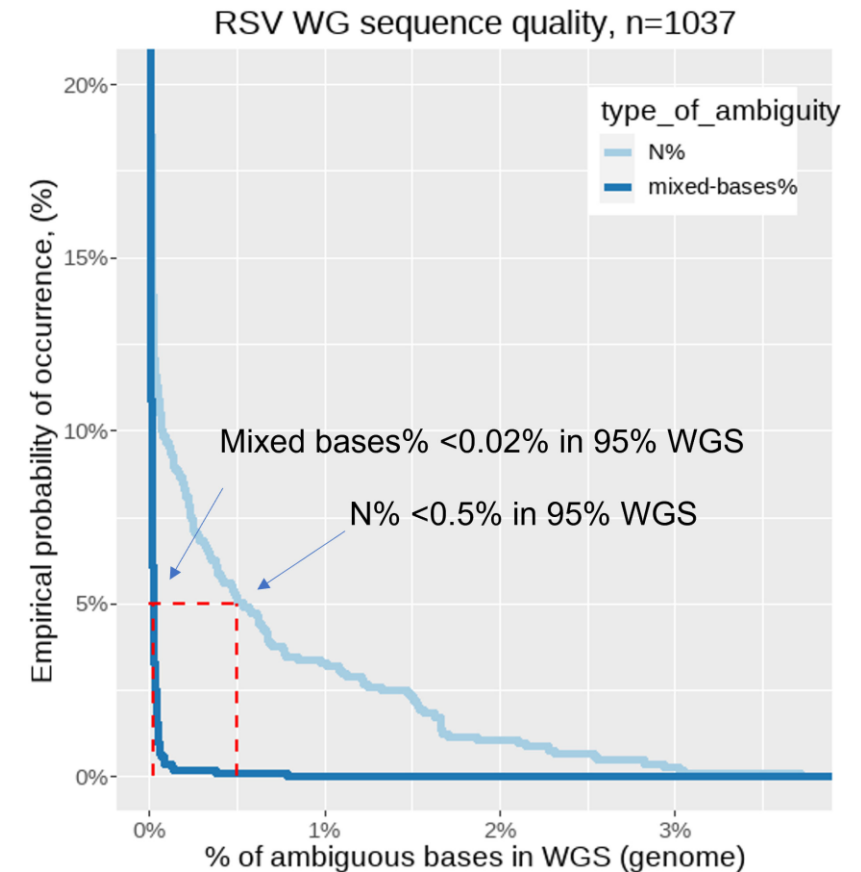


RSV-B



Talts et al, 2024 (UK Health Security Agency, London)

Amplicons: 750-1000 pb



ECDC NORMAL

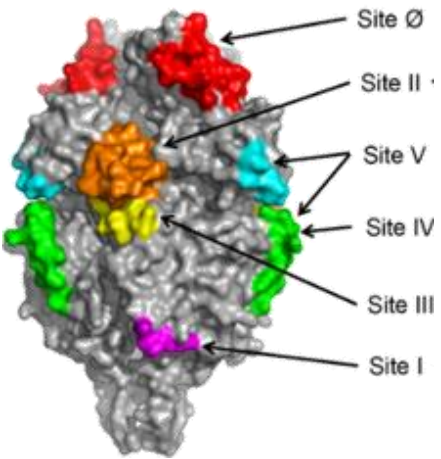
Escape mutations to nirsevimab

Description of mutations in the literature

Coming soon (this summer)

ANRS (French National Agency for Health Research)

Website with list of mutations



RSV-A genotypic drug resistance interpretation's algorithm			
October 2023 - Version n°1			
ANRS-MIE - Respiratory viruses group			
GENOTYPE INTERPRETATION: MONOCLONAL ANTIBODIES TO RSV F PROTEIN			
	Mutations associated with resistance	Mutations associated with "possible resistance"	Described polymorphisms in epitope Ø
Nirsevimab	<ul style="list-style-type: none">N67I + N208Y [4] <p><i>*Note that N67I or N208Y alone are not associated with resistance</i></p>	<ul style="list-style-type: none">K68E [1]	<p>N63T/S [3] I64V [1] K65R [1] E66K [3] K68N/R [1,4] N197K/H/D [1] I199M [1] L204I [1] I206T/I [1,3,4] V207I [1] Q210L [1]</p>

Escape mutations to nirsevimab

ECDC NORMAL

Description of mutations in the literature

Limits of genotypic analysis

Unknown mutations ?

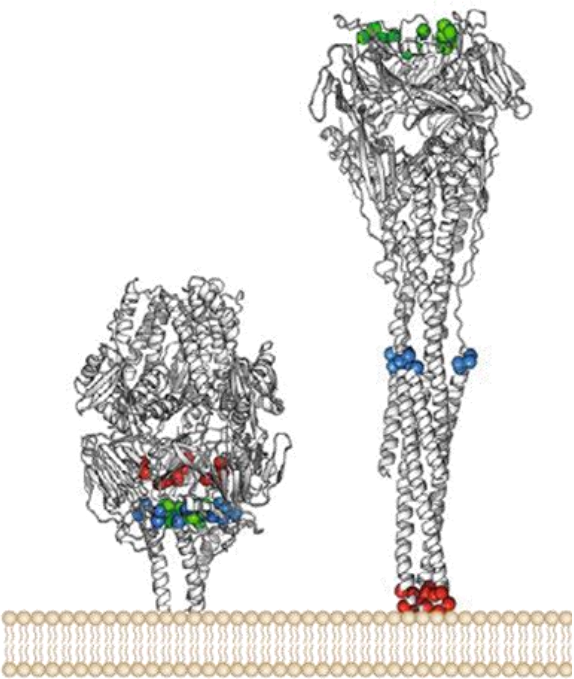
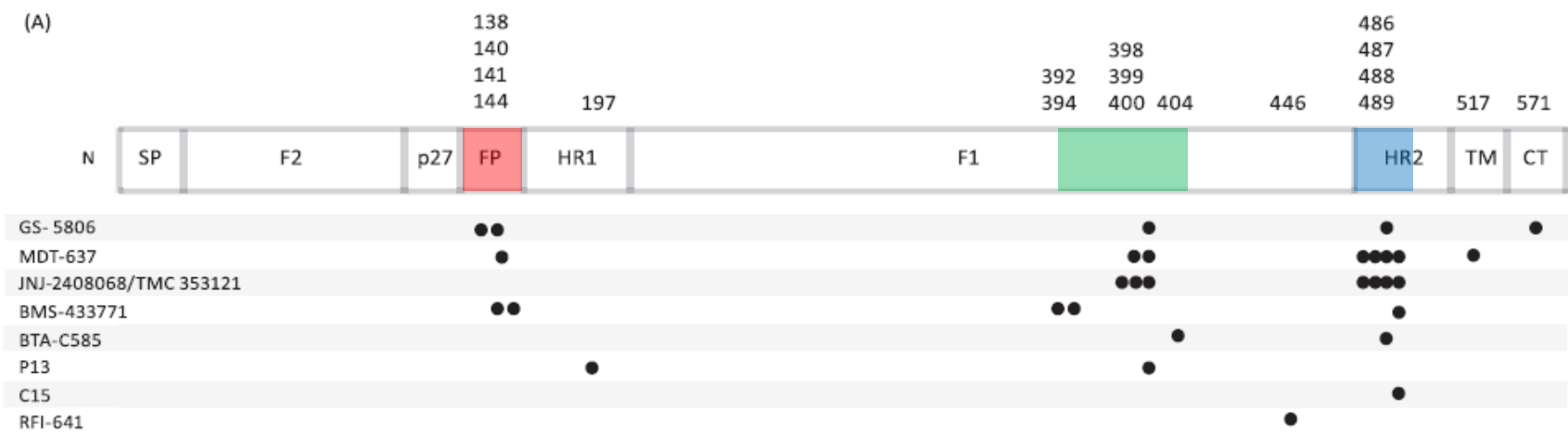
Escape mutations (might depend of context)

Ex: S201T mutation effect depends on 209 position

ECDC NORMAL

Escape mutations to antivirals

Known mutations on F for fusion inhibitors



Other inhibitors

Vaccine escape ?

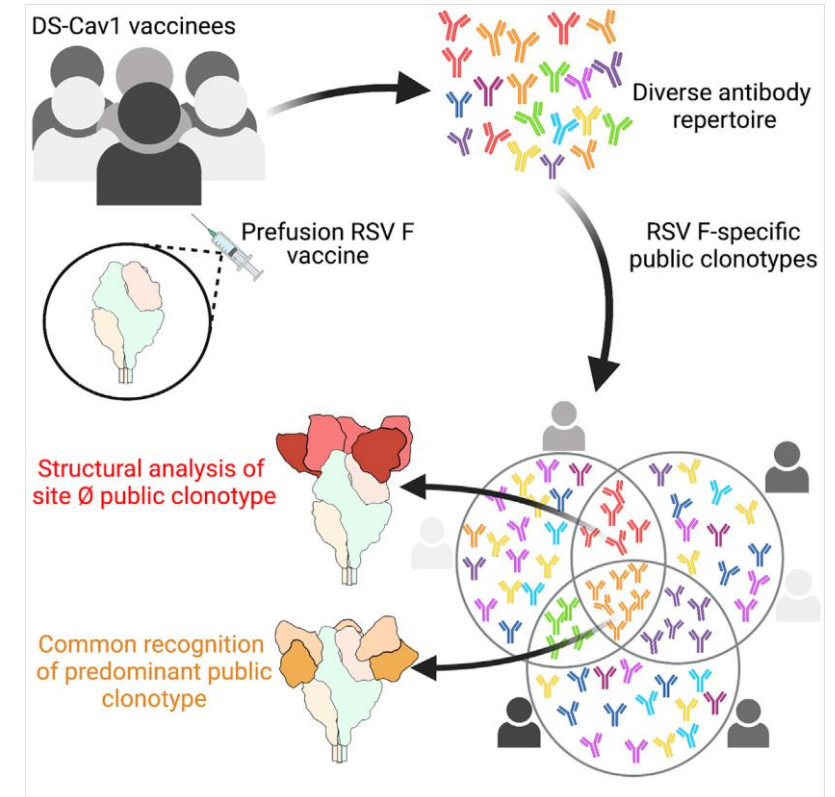
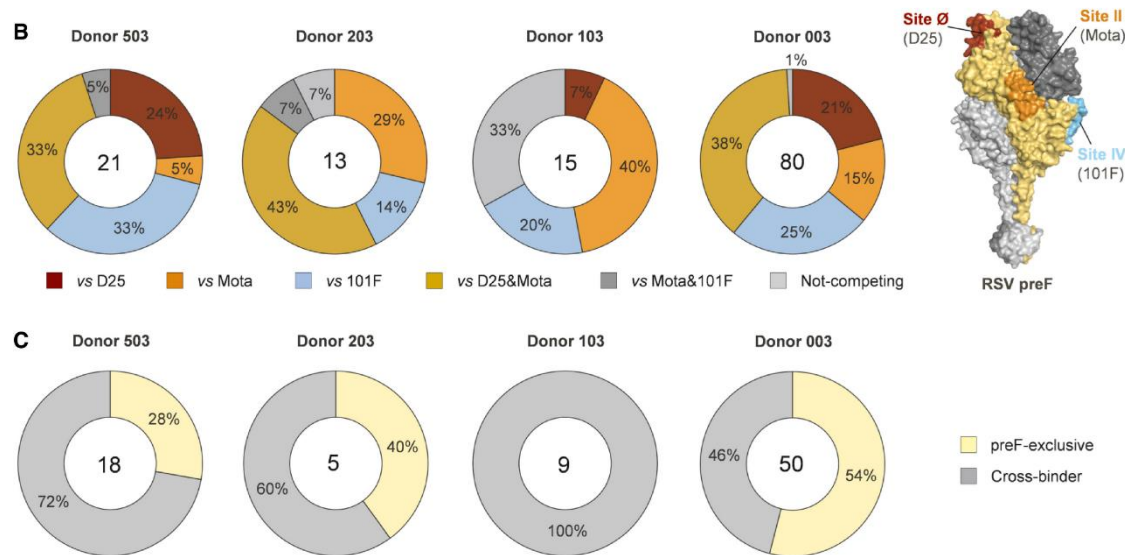
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Broad efficacy: GSK A vaccine only (A2 strain)

Response to F antigen (vaccine or natural infection):

Polyclonal

Efficient neutralisation of multiple RSV A and B variants, including those with escape mutations for a monoclonal, by sera from vaccinated individuals



Establish consistent and simple phenotypic protocols for characterisation of RSV

EDC NORMAL

Method of isolation: not fully agreed at this stage

2 major issues:

Difficult

Mutation during cell culture



Preservation conditions ?

Cells ?

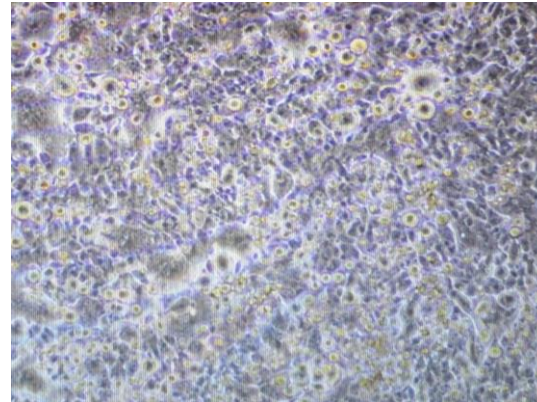
HEp-2

A549

Vero

MRC5

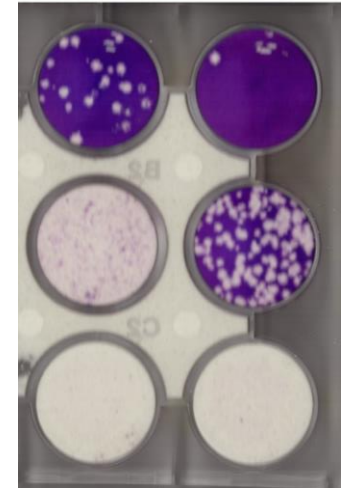
....



Detection of amplification

Cytopathic effect

Fluorescence, ELISA...



Quantification

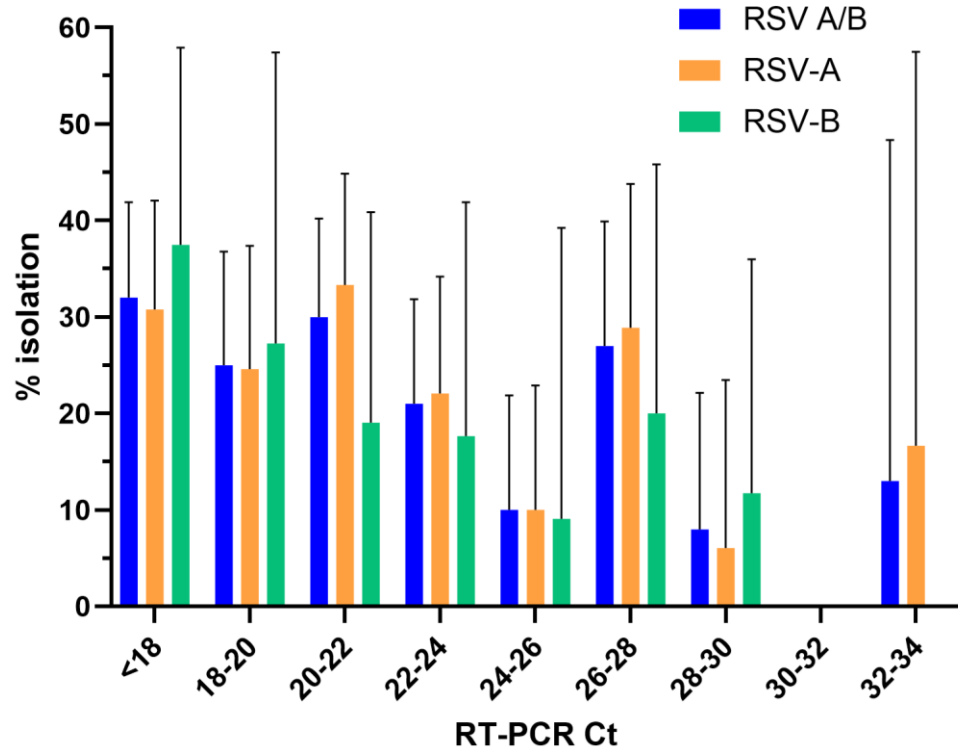
Plaque assay

Other (fluo, ELISA, RT-PCR..)

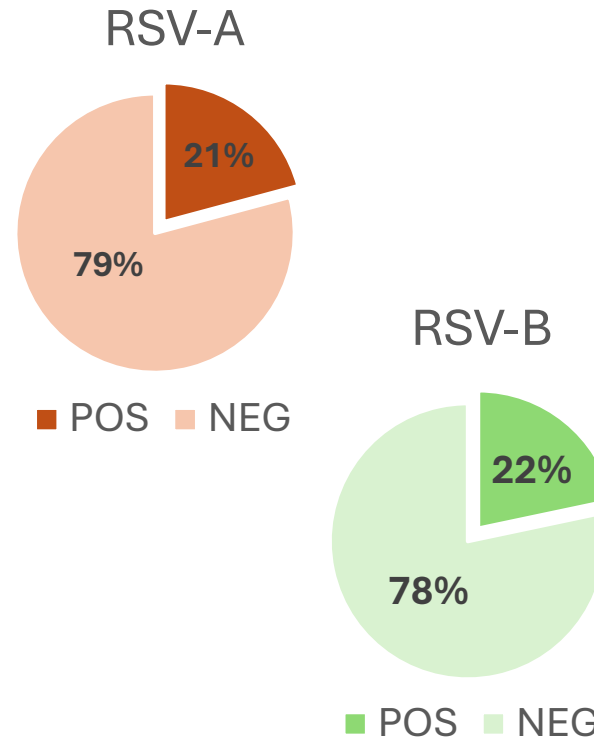
Factors affecting isolation efficiency

Isolation on HEp-2 cells monitoring cytopathic effect

Isolation effectiveness decreases with Ct

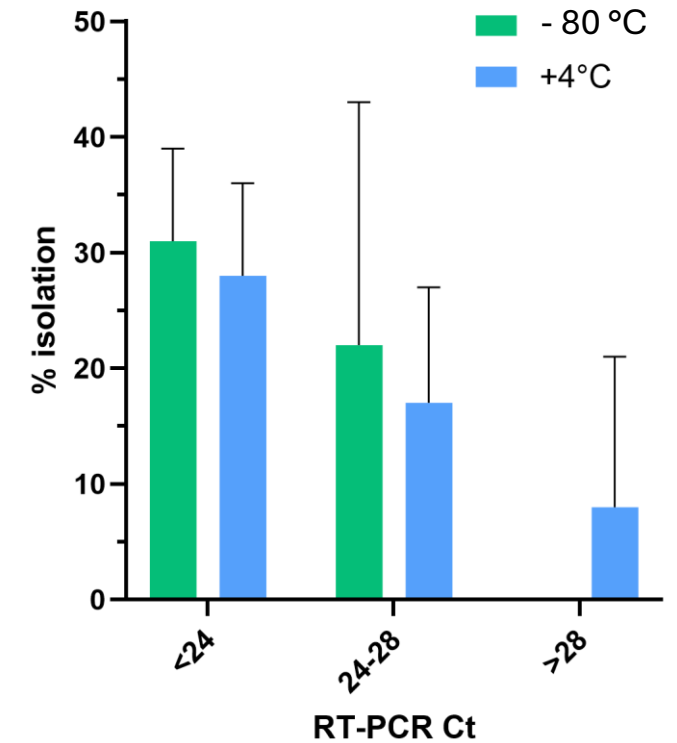


No difference between A/B



Limited effect of freezing ?

Bias for +4°C preservation

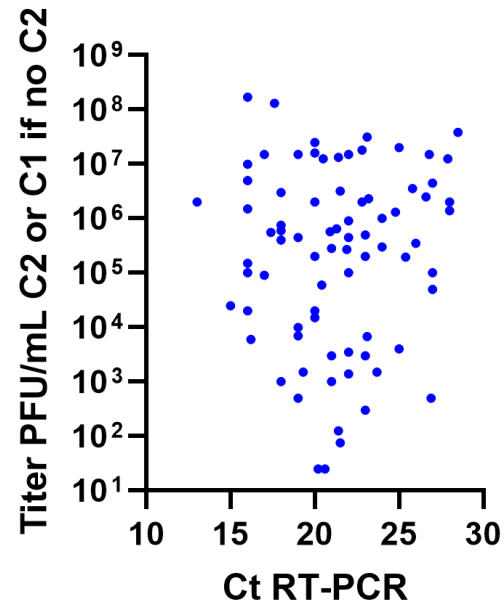
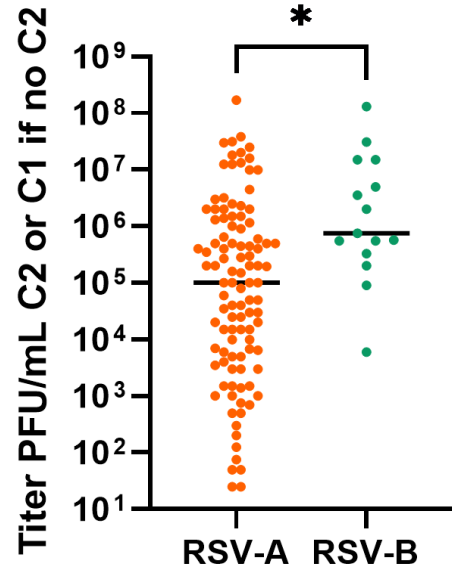


Titers and stability

ECDC NORMAL

No link between Titers and Ct

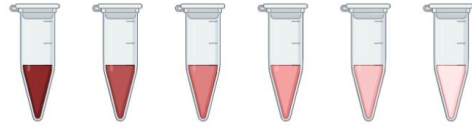
Stability assessment:
work still in progress....



Neutralisation Assay

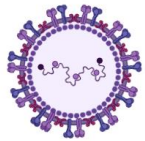
Principles

Serial dilutions of nirsevimab (or other)



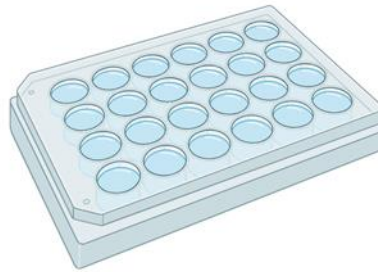
+

RSV



Incubation:
30-120min
RT or 37°C

Infection
Hep-2 or A549, Vero...



Incubation:
CO2 37°C
2-6 days

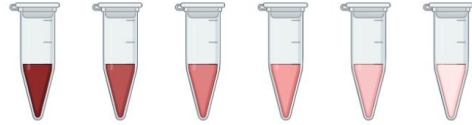
Revelation

TCID50 by ELISA or Fluo
FFU
PFU

Neutralisation Assay

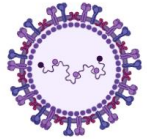
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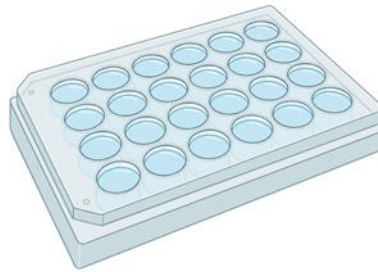
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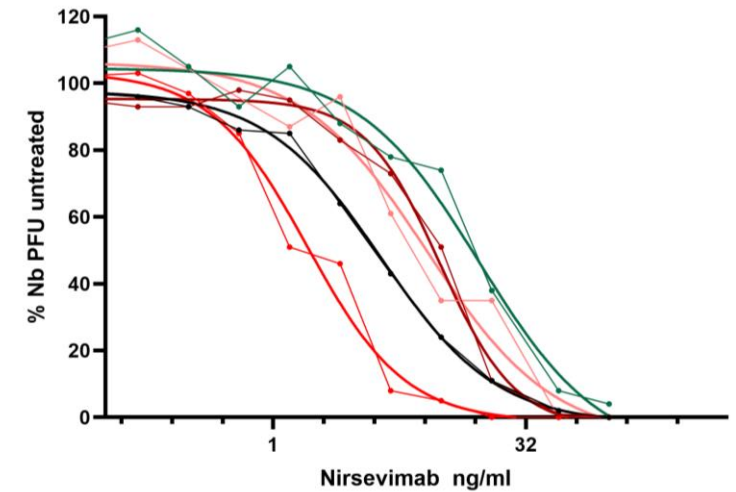
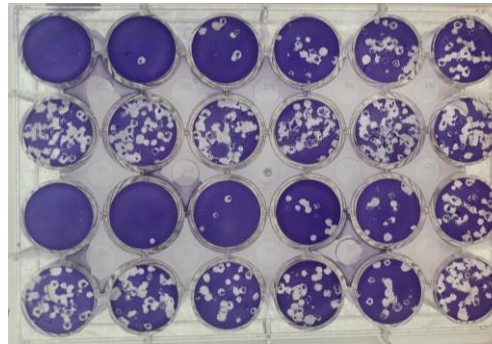
In our hands

Neutralisation assay on HEp-2 cells

Revelation of PFU after 6 days

Read out: PFU cumbersome

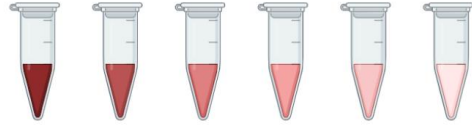
To set up: ELISA ? Fluorescence ?



Neutralisation Assay

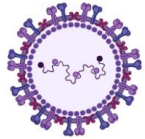
Principles

Serial dilutions of nirsevimab (or other)



+

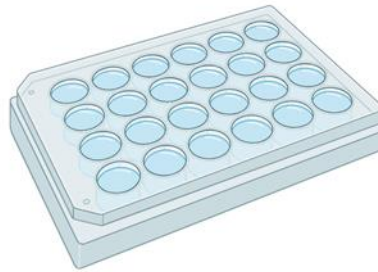
RSV



Incubation:
30-120min
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Infection

Hep-2 or A549, Vero...



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CO2 37°C
2-6 days

Revelation

TCID50 by ELISA or Fluo
FFU
PFU

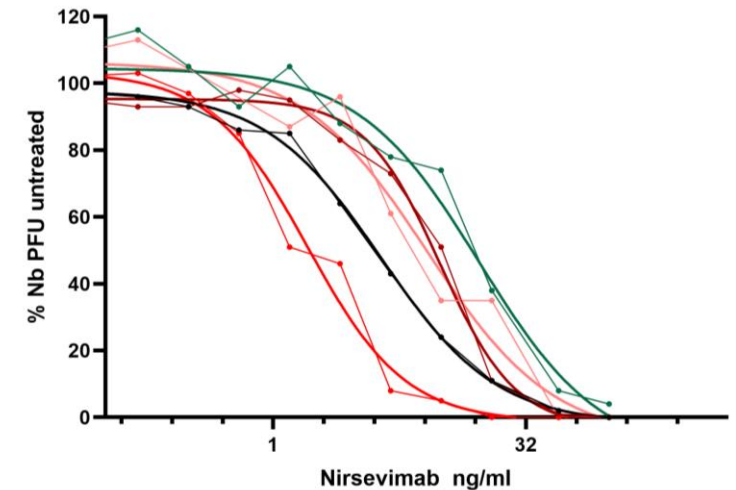
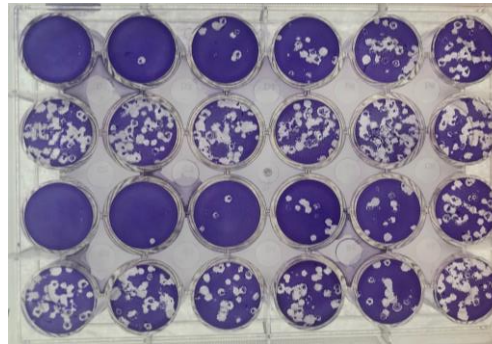
In our hands

Neutralisation assay on HEp-2 cells

Revelation after 6 days of PFU

Read out: PFU cumbersome

To set up: ELISA ? Fluorescence ?



Antigenic characterization : relevance for RSV ? Reagents ?

