

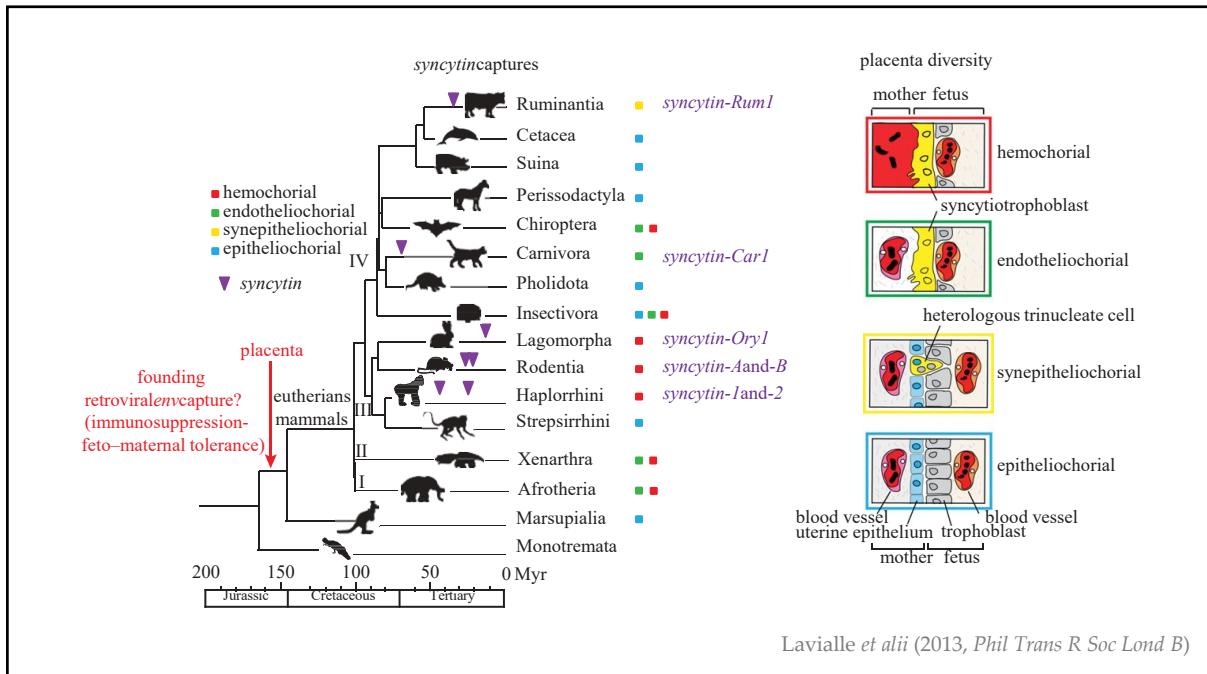
QUAND PASTEUR RENCONTRE DARWIN : L'ÉVOLUTION, LES VIRUS ET NOUS

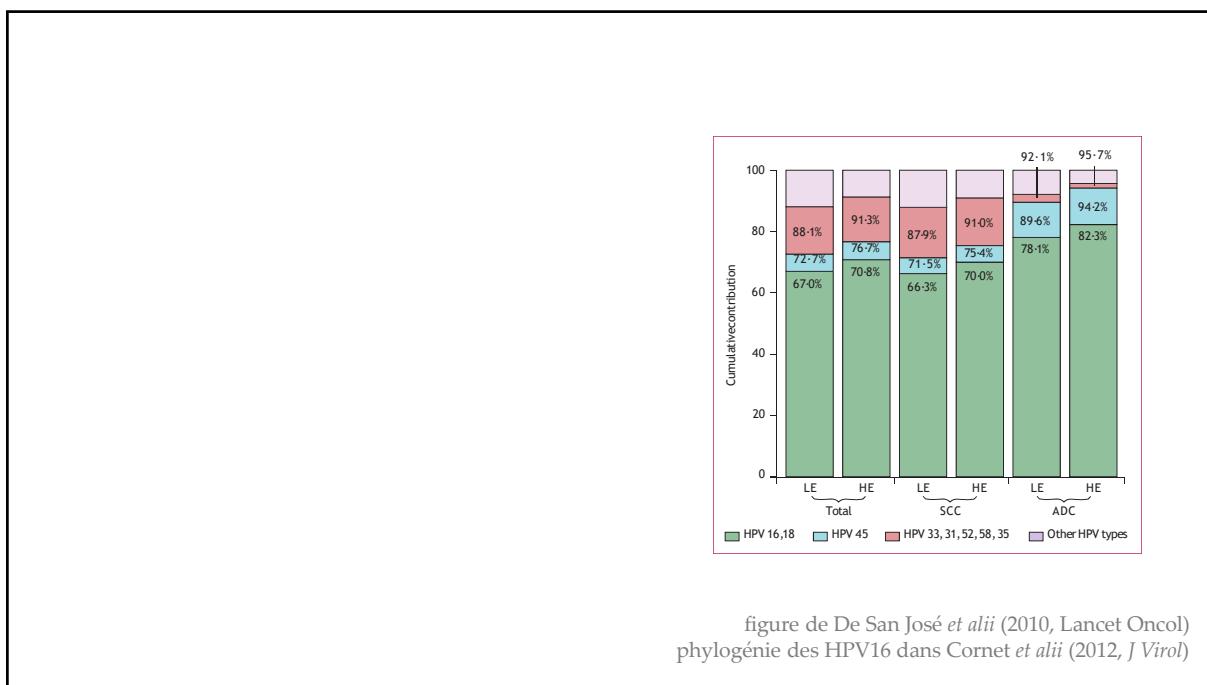
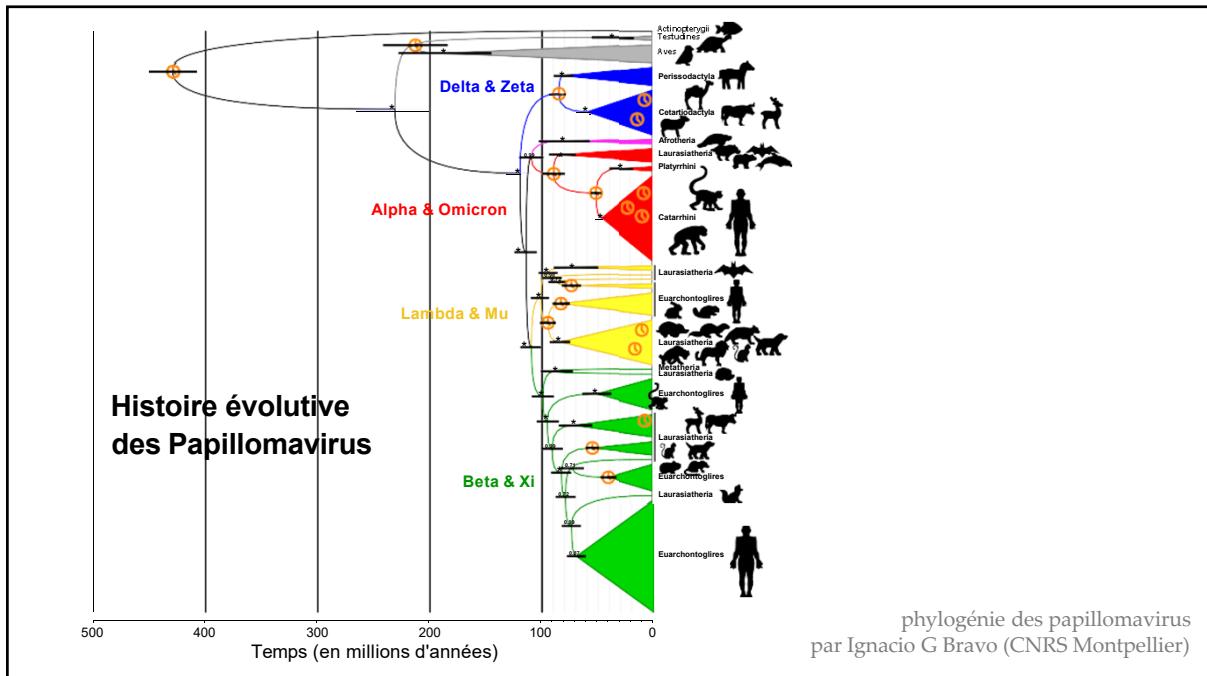
Samuel Alizon

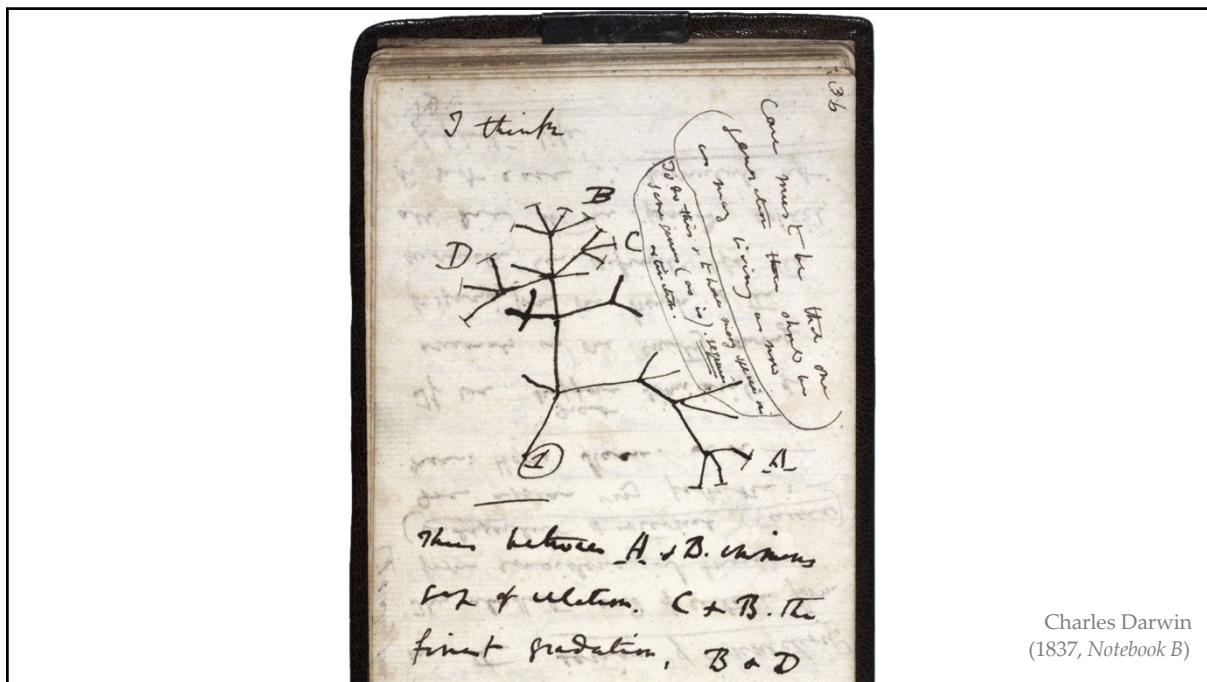
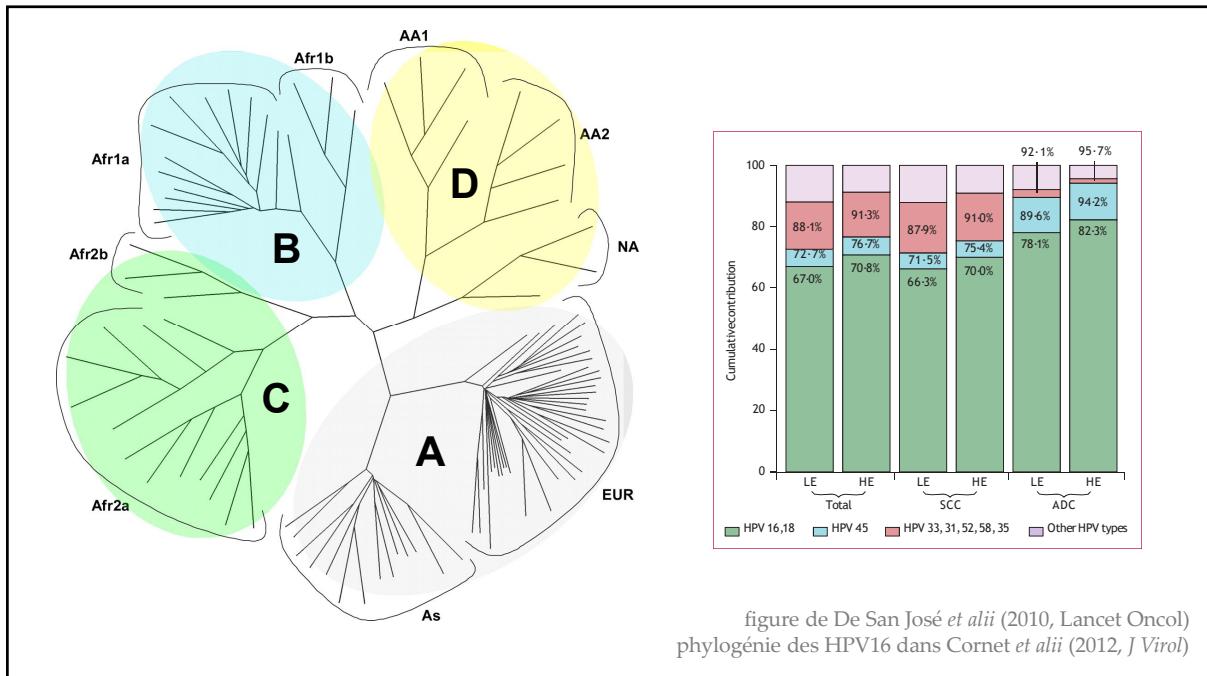
Directeur de recherche
Laboratoires MIVEGEC (Montpellier) et CIRB (Paris)
samuel.alizon@cnrs.fr
<https://alizon.ouvaton.org>

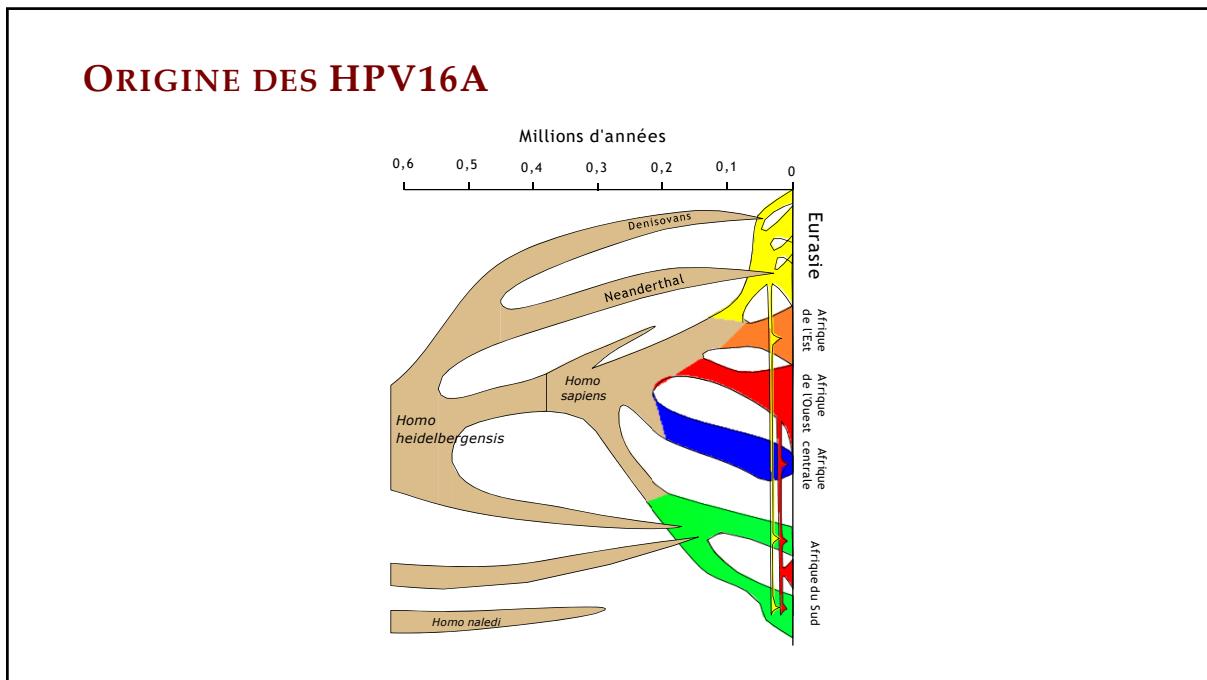
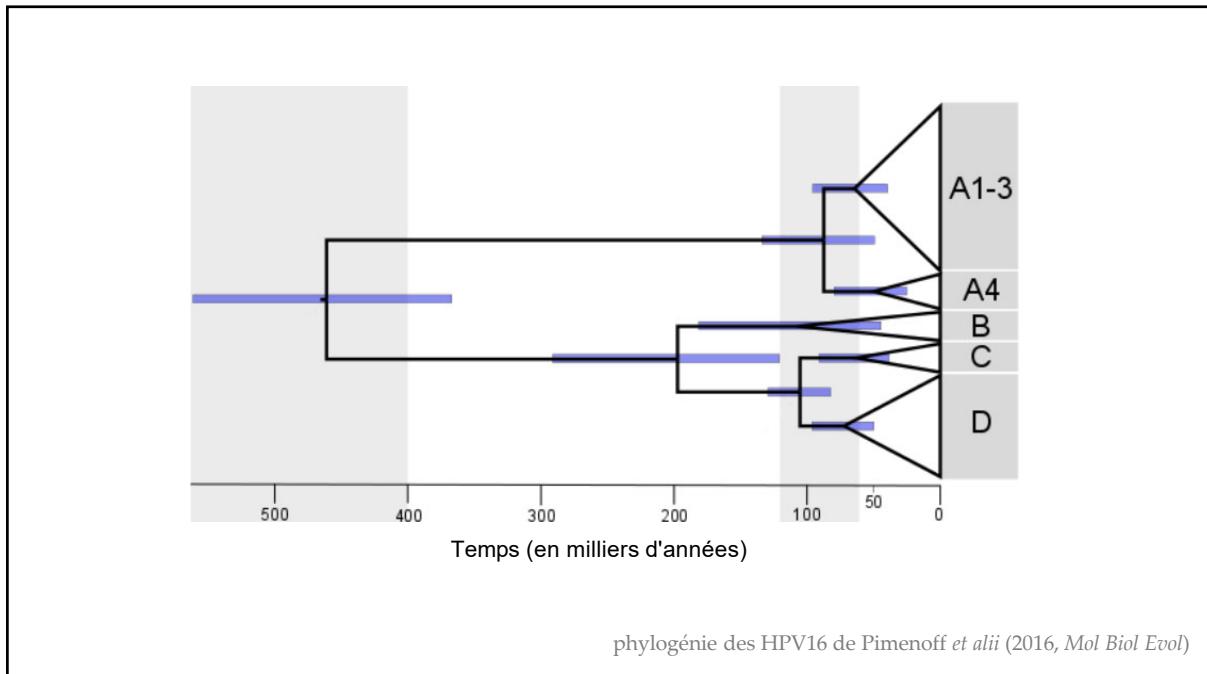
Soirée du résO InfectiO PACA-EST, Nice, 2 mars 2022



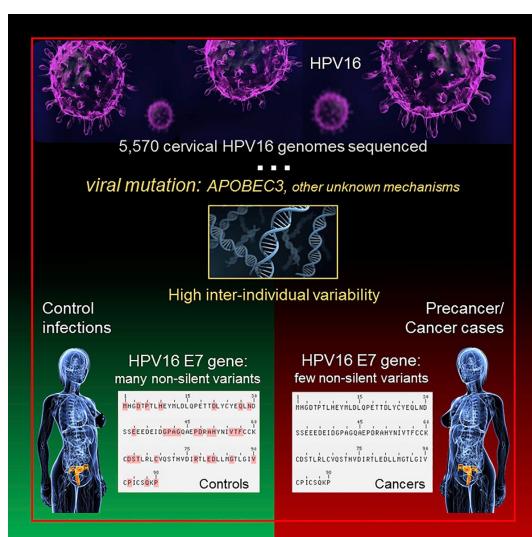
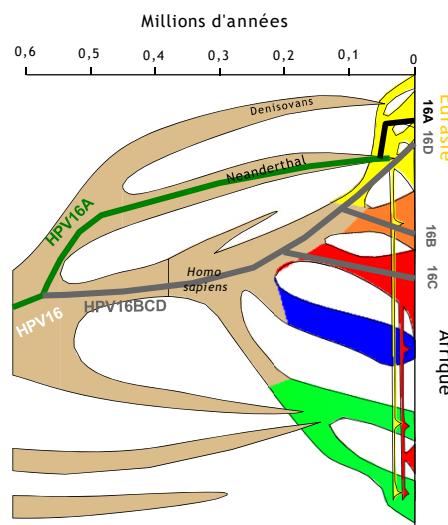






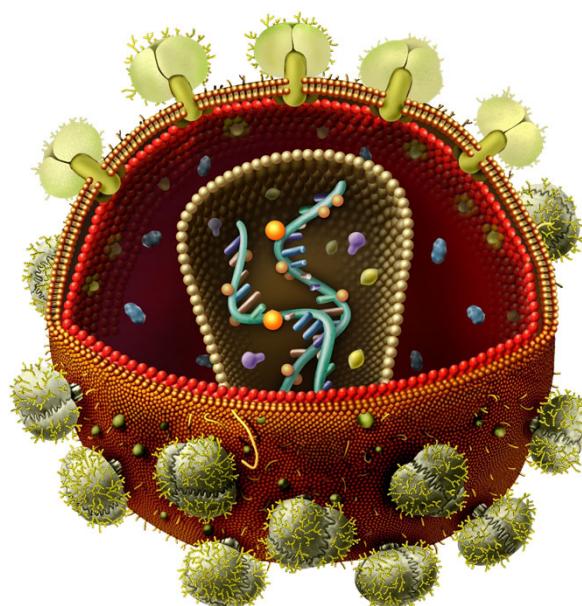


ORIGINE DES HPV16A



CO-ÉVOLUTION ENTRE *Homo sapiens* ET SES VIRUS

- Les virus façonnent notre génome (7 %)
- Les variants les plus virulents HPV sont apparus il y a environ 100.000 ans.
- Ces HPV16A seraient une IST transmise à *Homo sapiens* par *Homo neanderthalis*.
- La séquence du génome d'HPV semble associée à la virulence.



CENTERS FOR DISEASE CONTROL

MMWR

MORBIDITY AND MORTALITY WEEKLY REPORT

June 5, 1981 / Vol. 30 / No. 21

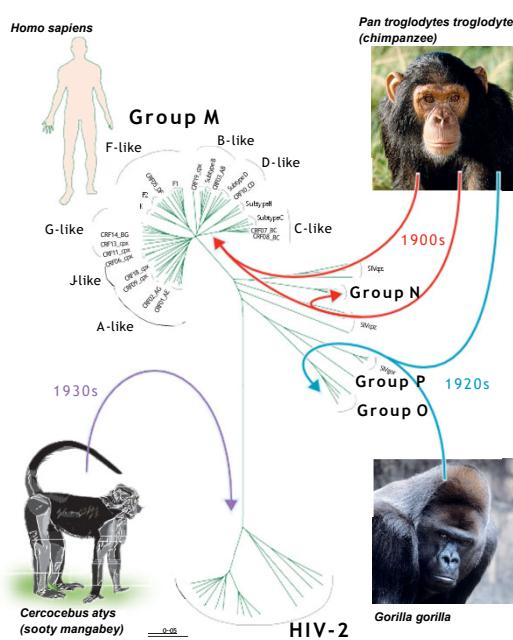
Epidemiologic Notes and Reports

- 249 Dengue Type 4 Infections in U.S. Travelers to the Caribbean
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Current Trends
 252 Measles — United States, First 20 Weeks
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International Notes
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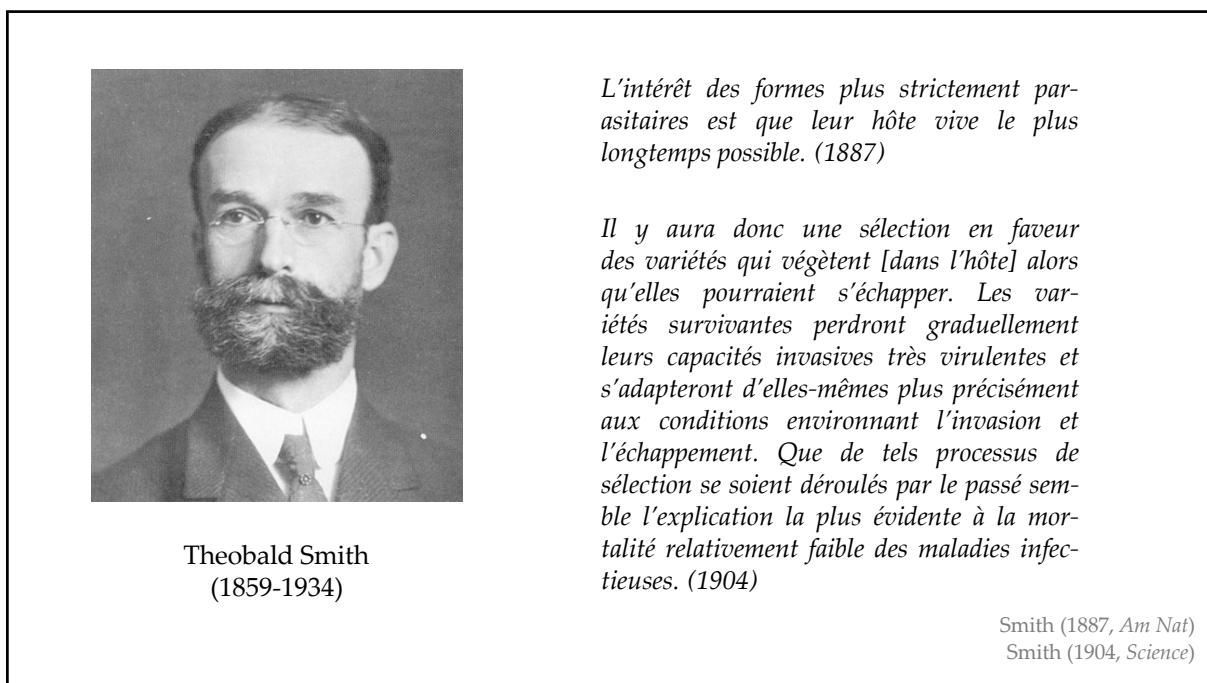
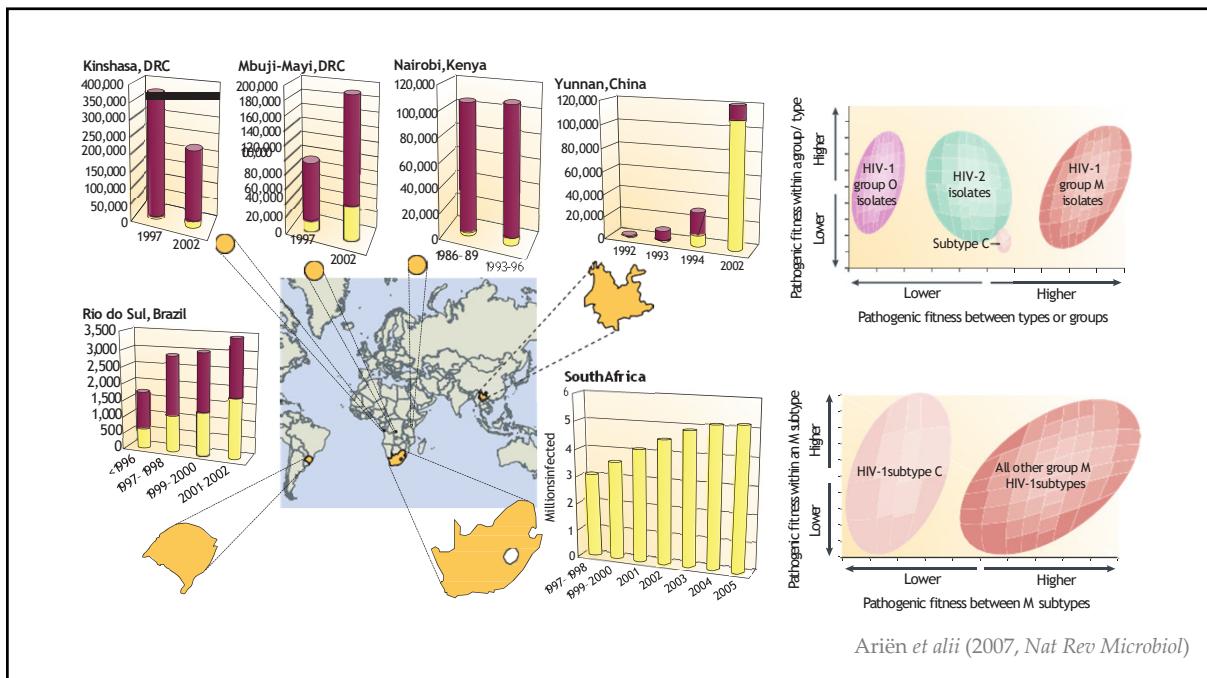
Pneumocystis Pneumonia — Los Angeles

In the period October 1980-May 1981, 5 young men, all active homosexuals, were treated for biopsy-confirmed *Pneumocystis carinii* pneumonia at 3 different hospitals in Los Angeles, California. Two of the patients died. All 5 patients had laboratory-confirmed previous or current cytomegalovirus (CMV) infection and candidal mucosal infection. Case reports of these patients follow.

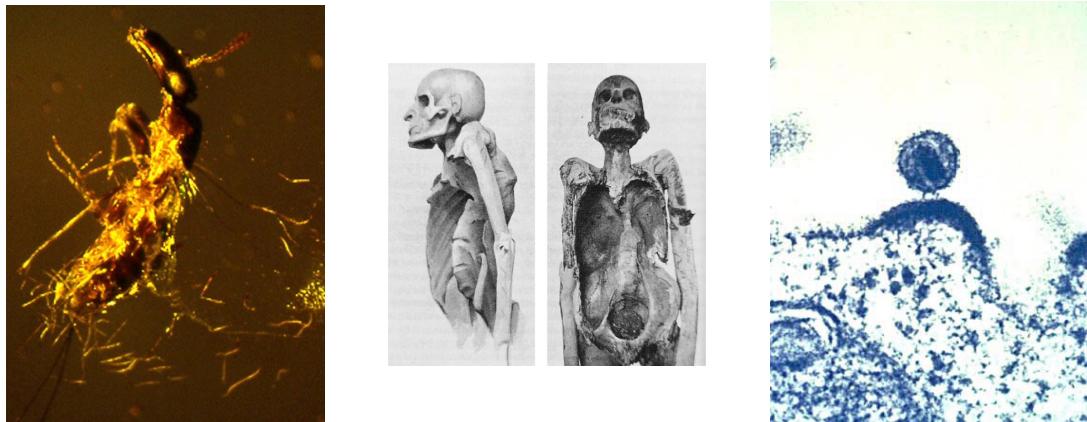
Patient 1: A previously healthy 33-year-old man developed *P. carinii* pneumonia and oral mucosal candidiasis in March 1981 after a 2-month history of fever associated with



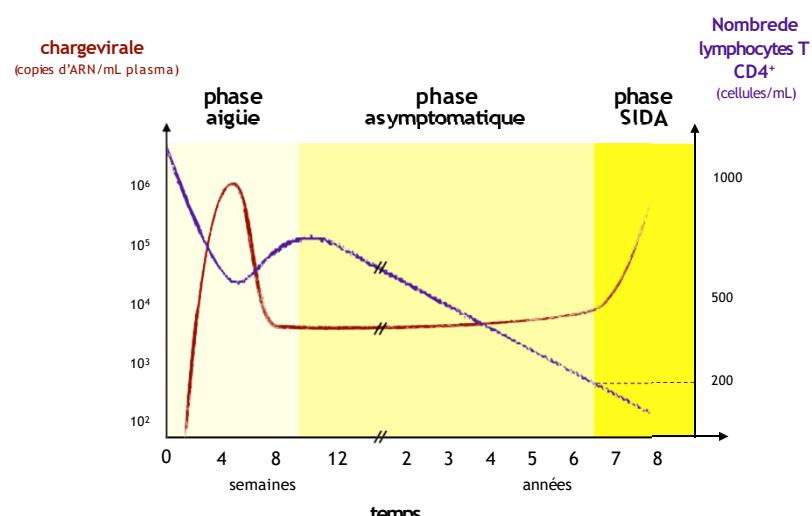
Tebit & Arts
(2011, *Lancet Inf Dis*)

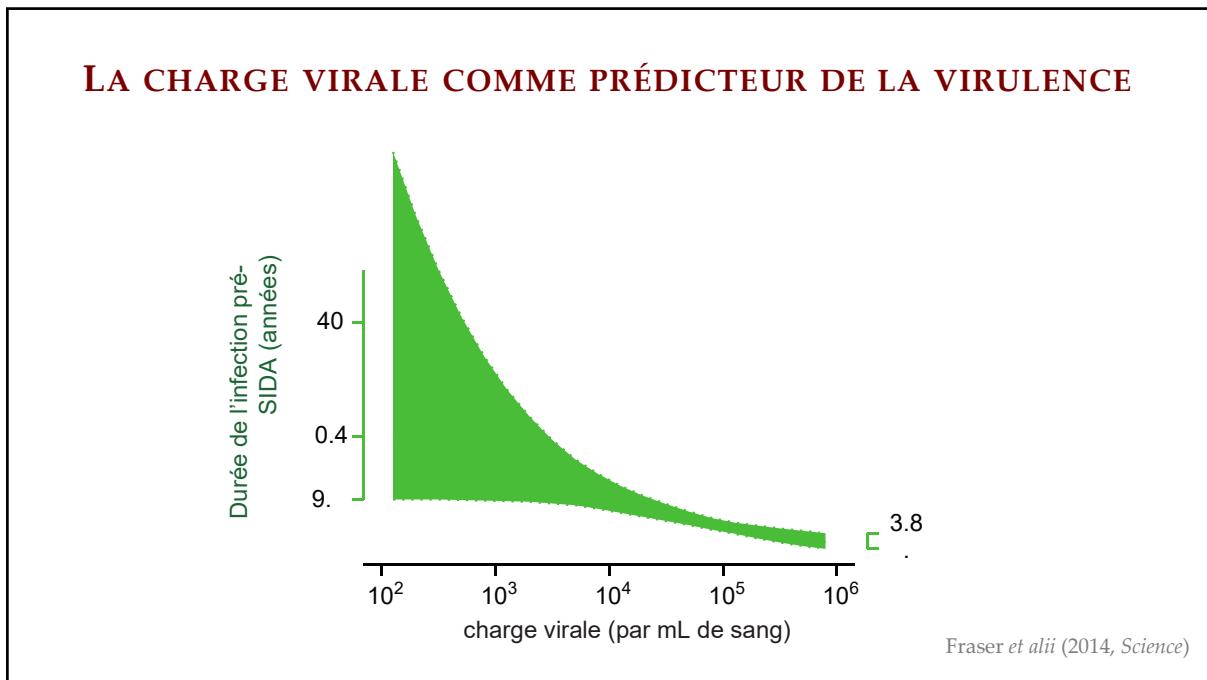
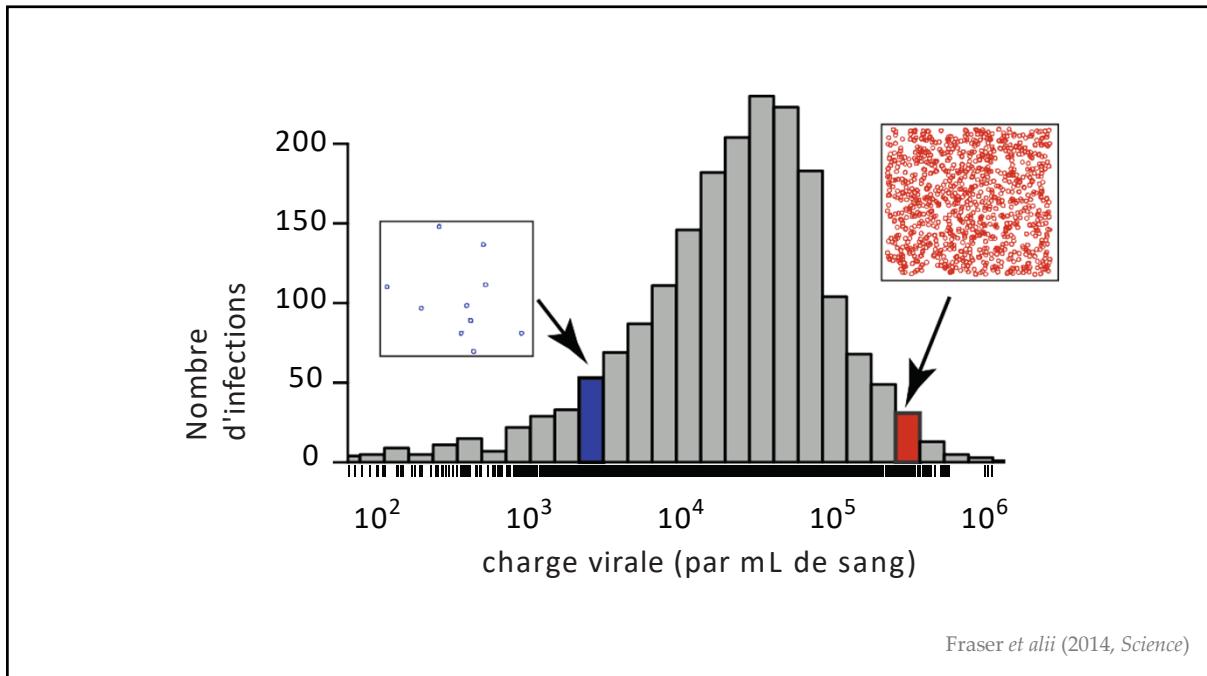


POURTANT LA VIRULENCE PERSISTE...

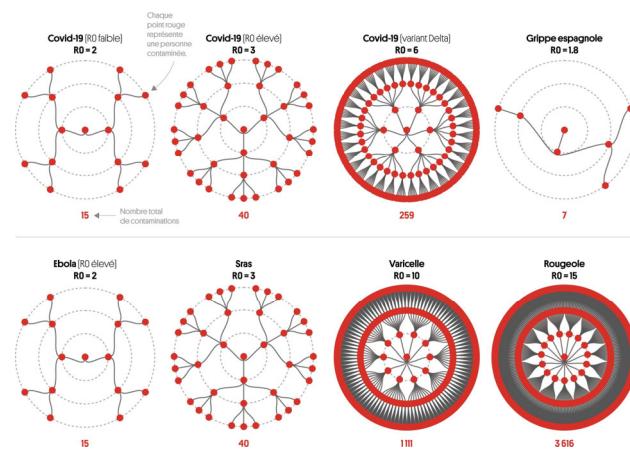


DÉROULEMENT D'UNE INFECTION PAR LE VIH



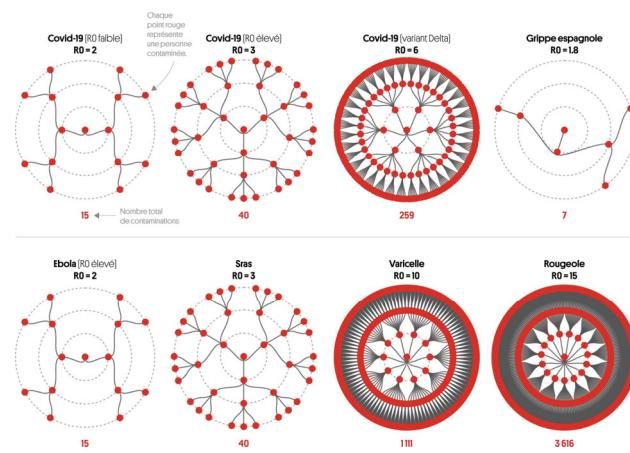


“VALEUR SÉLECTIVE” D’UNE MALADIE INFECTIEUSE

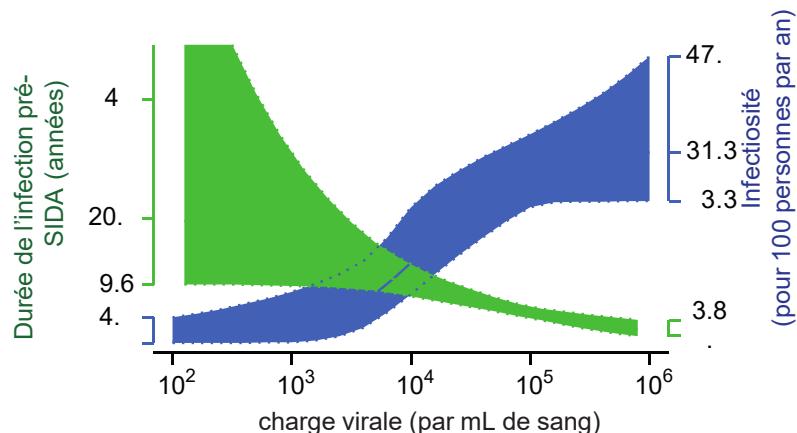


“VALEUR SÉLECTIVE” D’UNE MALADIE INFECTIEUSE

$$R_0 = \text{durée de l'infection} \times \text{contagiosité}$$

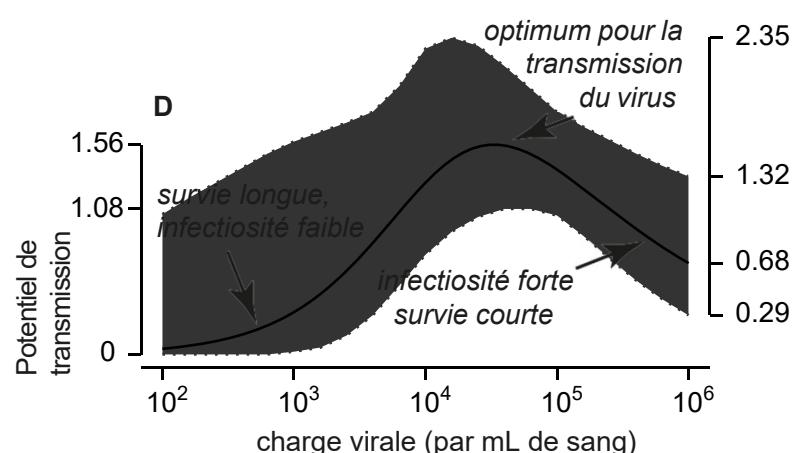


LA CONTAGIOSITÉ DÉPEND DE LA CHARGE VIRALE



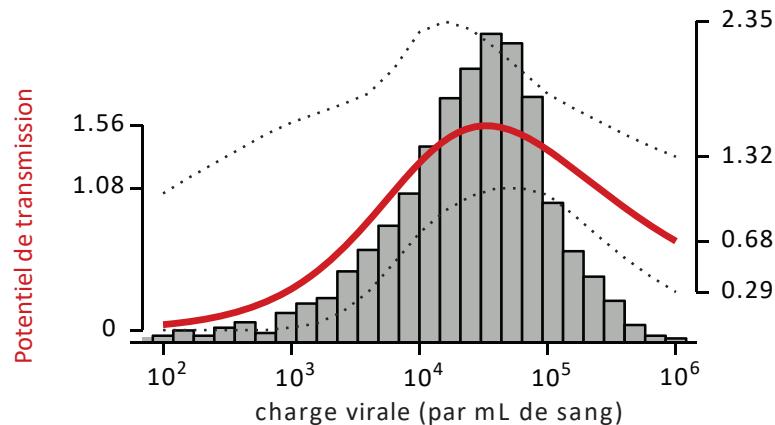
Fraser et alii (2014, Science)

COMPROMIS ÉVOLUTIF

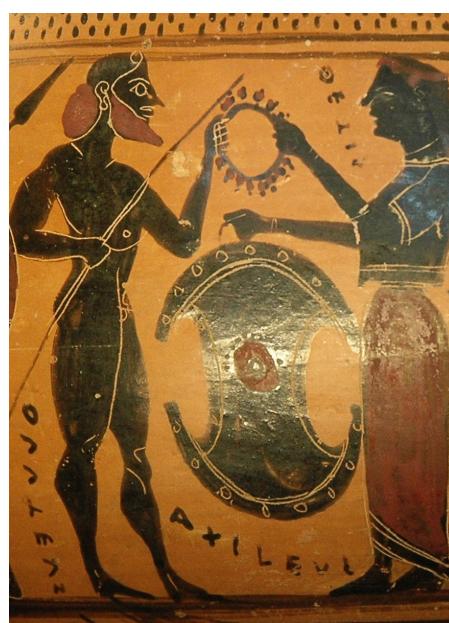


Fraser et alii (2014, Science)

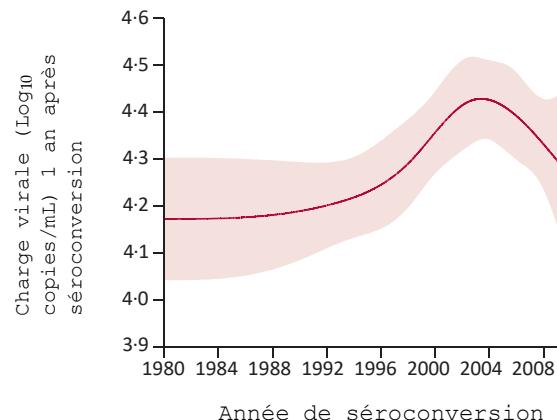
UN OPTIMUM ÉVOLUTIF ATTEINT ?



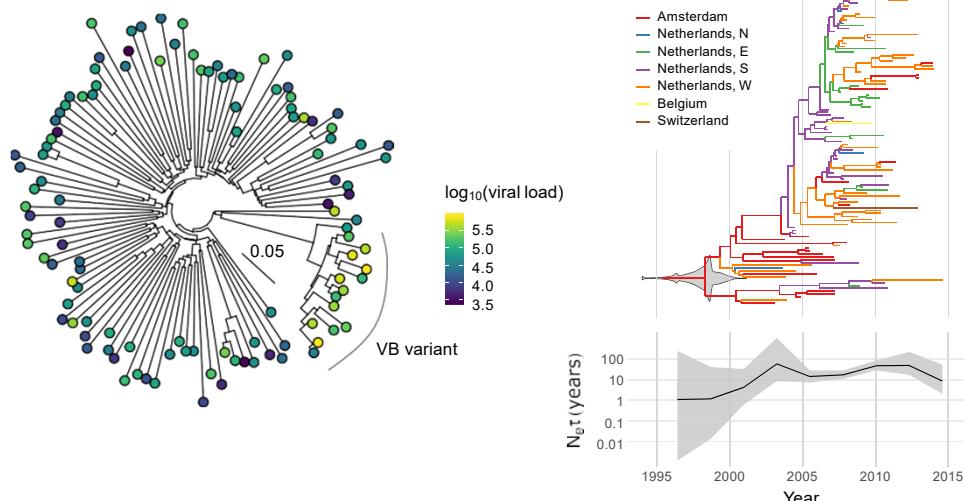
Fraser et alii (2014, Science)



LE VIH N'EST PAS DEVENU MOINS VIRULENT



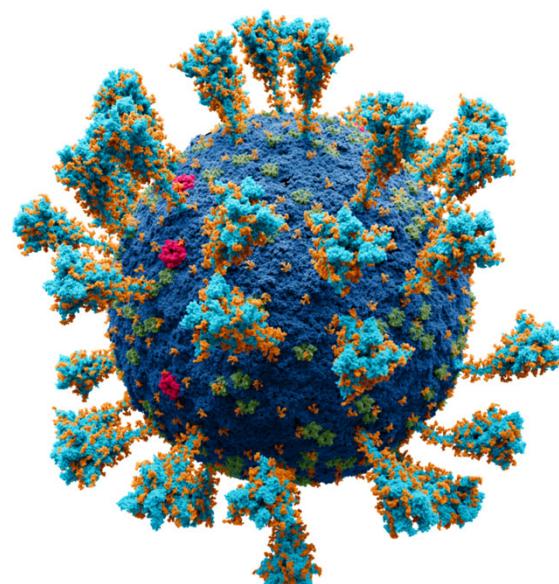
Pantazis *et alii* (2014, Lancet HIV)

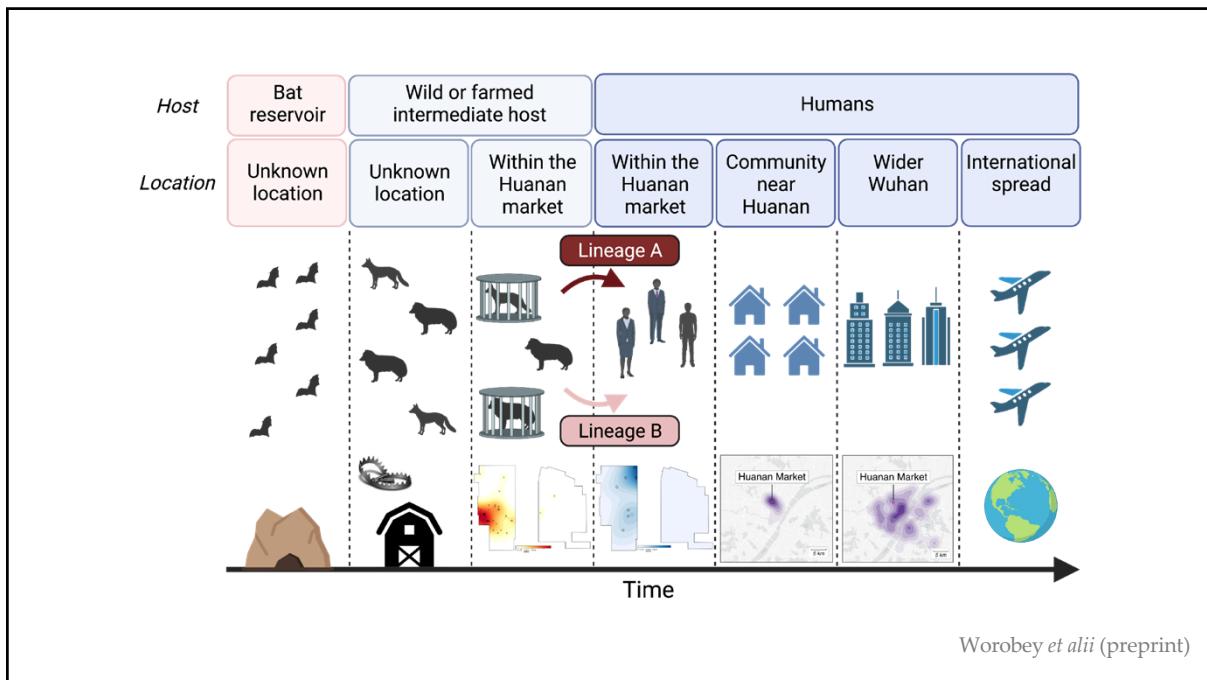
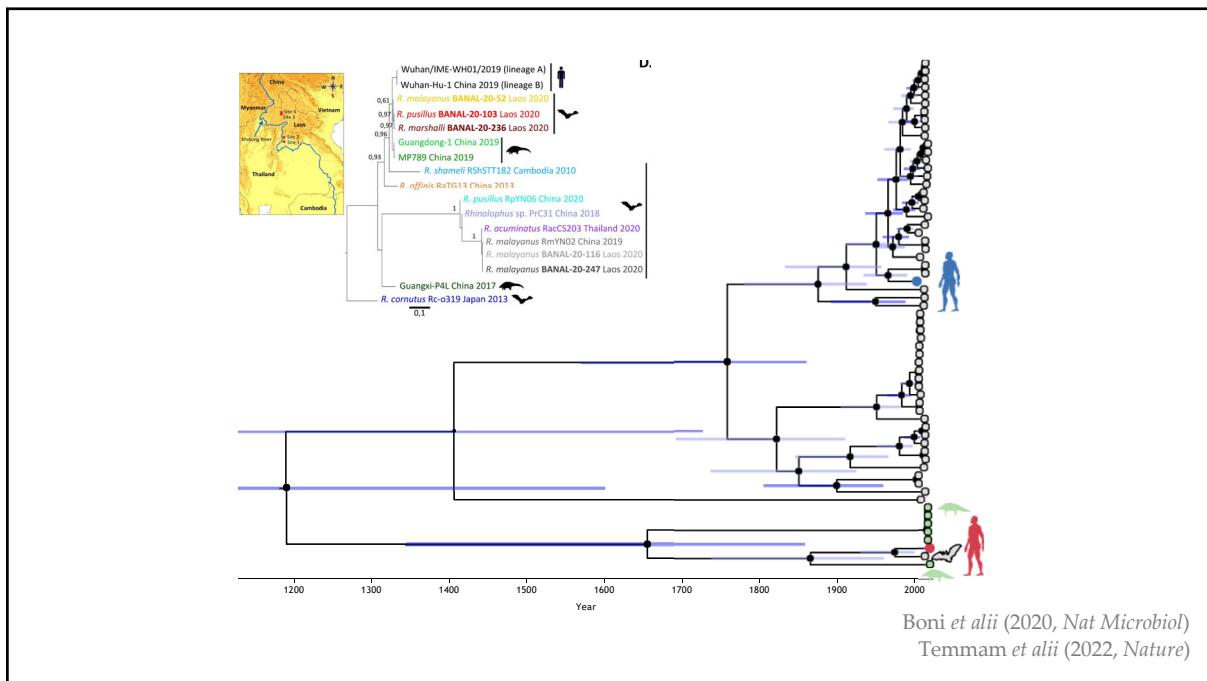


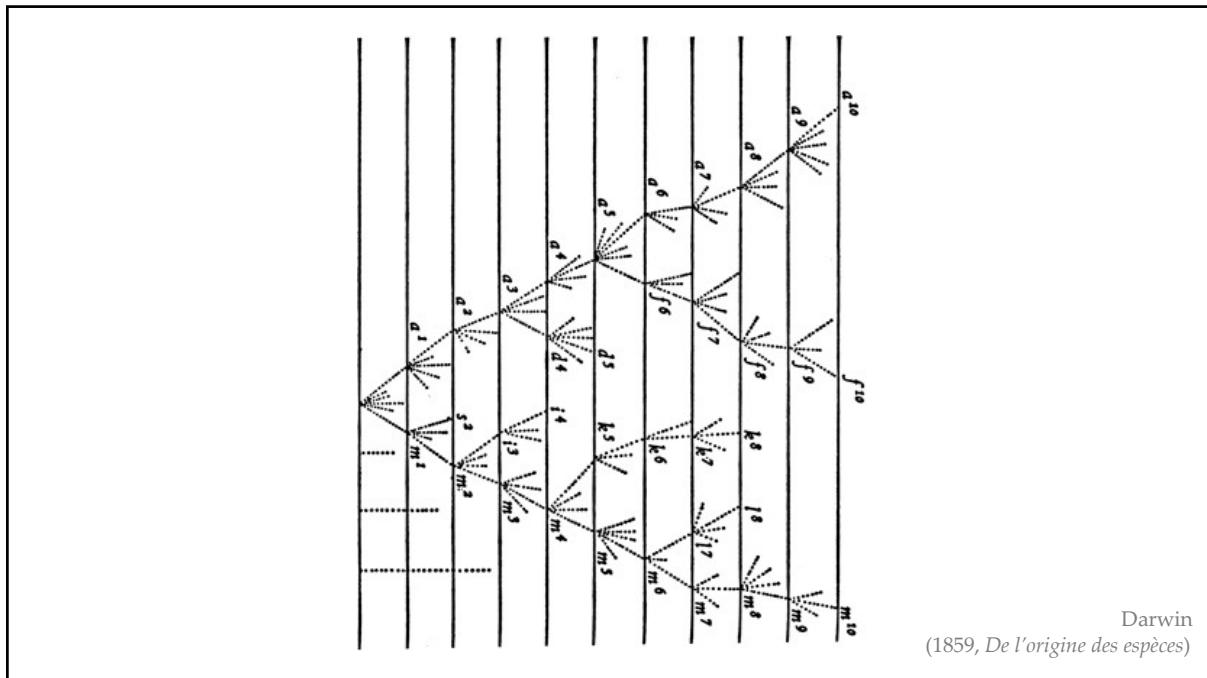
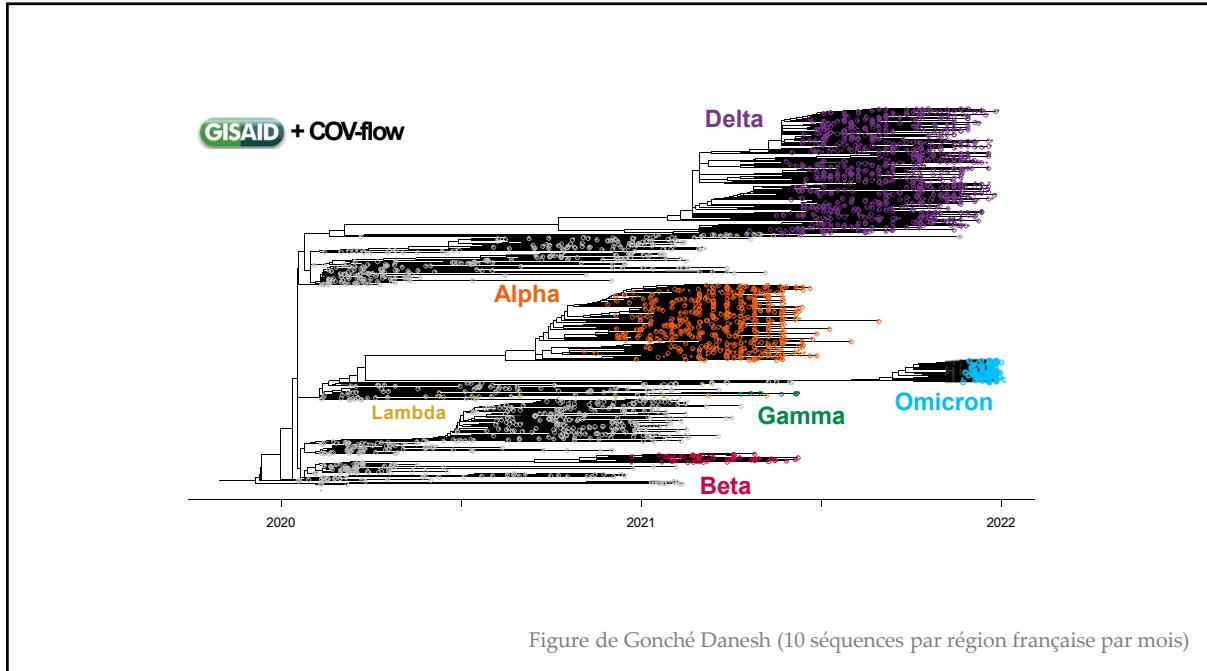
Wymant *et alii* (2022, Science)

ÉVOLUTION DE LA VIRULENCE D'UN VIRUS ÉMERGENT

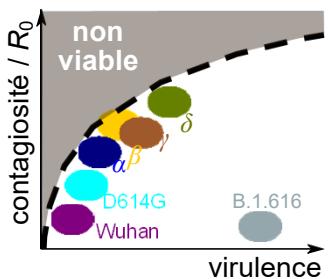
- Les VIH sont issus de plusieurs zoonoses il y a plus d'un siècle.
- Seul le groupe M, le plus virulent, est devenu pandémique.
- Il existe un compromis adaptatif entre contagiosité de l'une infection et virulence.
- La virulence du VIH sous-type B semble avoir augmenté depuis son émergence.







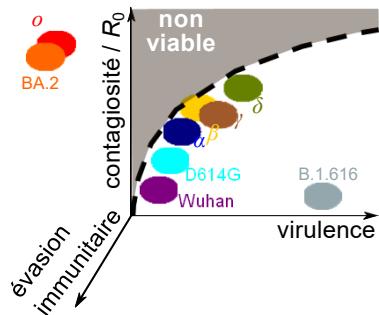
VARIANTS ET COMPROMIS ADAPTATIF



- corrélation entre contagiosité (R_0) et virulence (observé aussi pour le VIH)
- bonne immunité croisée (surtout vaccinale)

Alizon & Sofonea (2021, *J Evol Biol*)

VARIANTS ET ÉVASION IMMUNITAIRE



- Omicron contourne fortement l'immunité
- Deux "niches écologiques"?
- Quelle coexistence entre variants ?

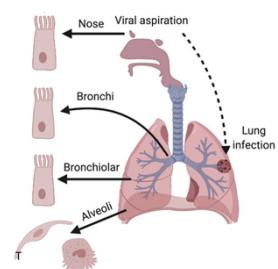


Figure issue de Hou *et alii* (2020, *Cell*)

SARS-CoV-2 : UN VIRUS EN ÉVOLUTION

- Le SARS-CoV-2 vient a priori d'un réservoir animal (chauve-souris).
- La pandémie résulterait d'au moins deux événements de transmission dans un marché de Wuhan.
- Les premiers variants ont été de plus en plus contagieux et virulents (en l'absence d'immunité).
- Les variants du type Omicron semblent explorer une nouvelle niche écologique et leur coexistence avec les autres est inconnue.



LA REINE ROUGE OU LA COURSE CO-ÉVOLUTIVE

