



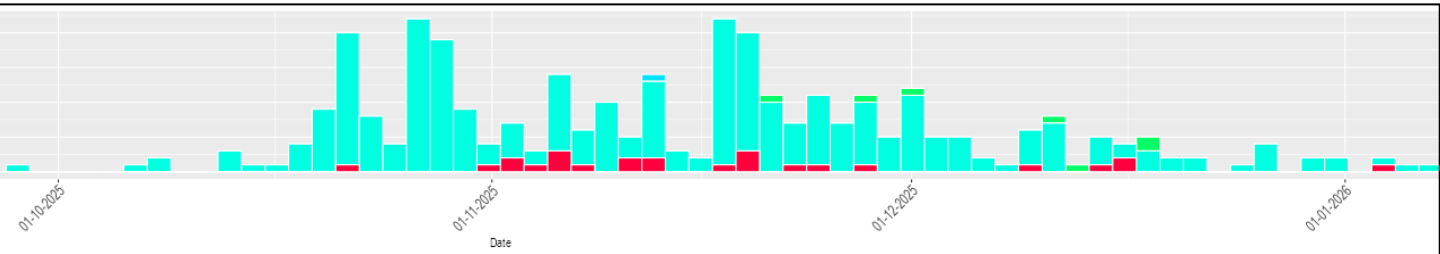
**anses**

# **ANALYSES PHYLOGÉNÉTIQUES DES VIRUS IAHP H5N1**

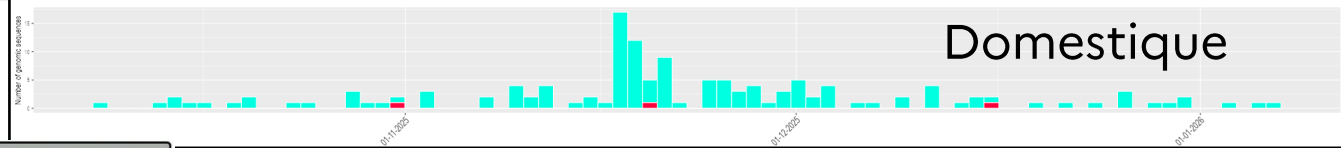
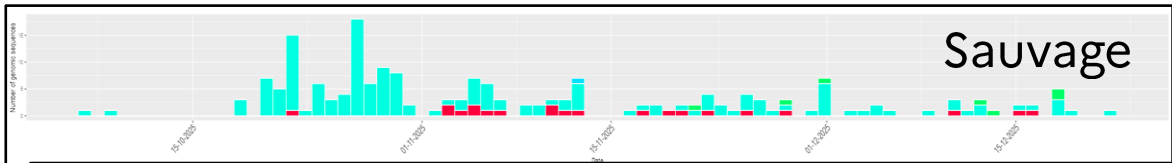
**FX BRIAND, LNR IA/MN -16/01/2026**

**CONNAÎTRE, ÉVALUER, PROTÉGER**

# Chronologie des foyers H5N1 HP séquencés depuis Oct 2025

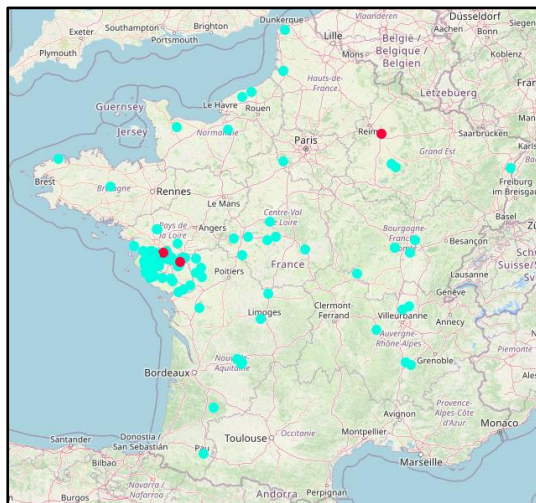


- Génotype FR20 (EA-2024-DI)
- Génotype FR19 (EA-2024-DT)
- Génotype FR21
- Non génotypable

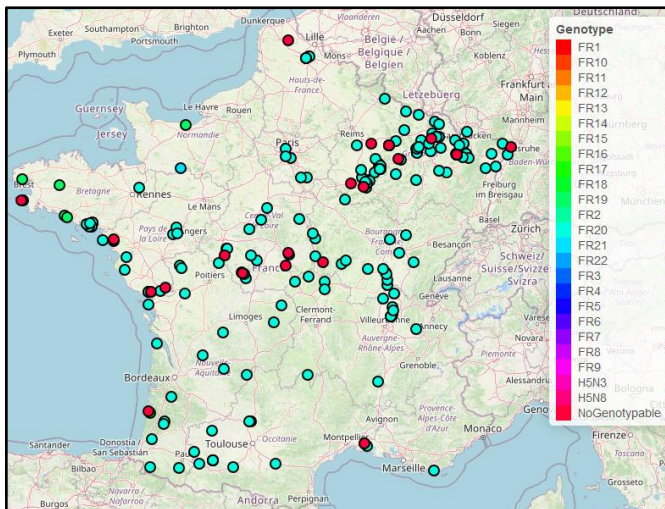


# Foyers IAHP H5 séquencés

## Foyers « élevage » séquencés



## Foyers « Sauvage » séquencés

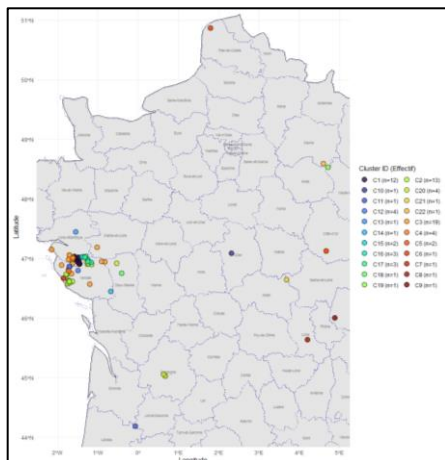


| Context  | FR19 | FR20 | FR21 | NoGenotypable | Total |
|----------|------|------|------|---------------|-------|
| backyard | 0    | 8    | 0    | 1             | 9     |
| captive  | 0    | 2    | 0    | 0             | 2     |
| poultry  | 0    | 79   | 0    | 1             | 80    |
| wild     | 1    | 123  | 1    | 13            | 138   |
| Total    | 1    | 212  | 1    | 15            | 229   |

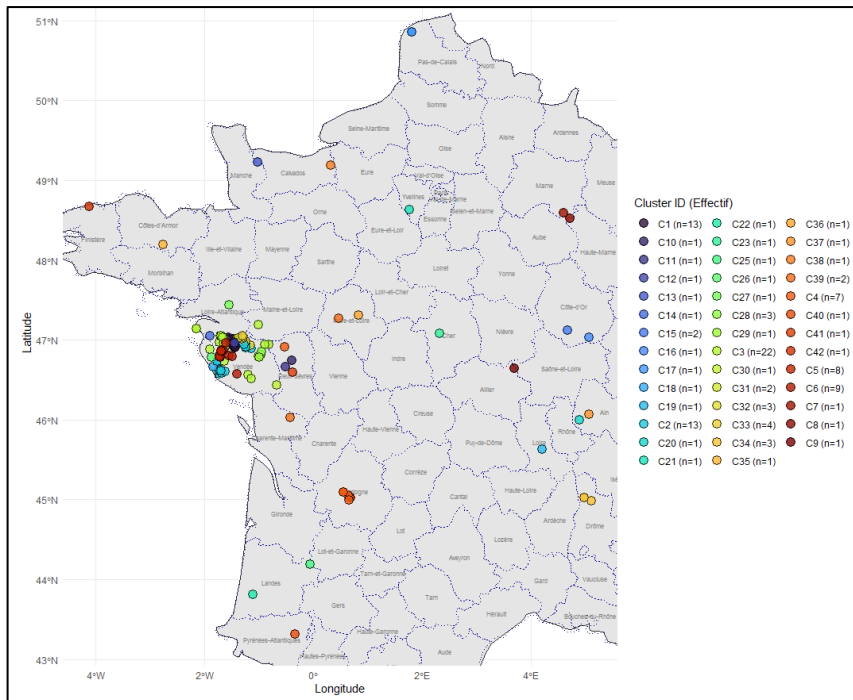


| Context  | FR19  | FR20     | FR21 | NoGenotypable | Total     |
|----------|-------|----------|------|---------------|-----------|
| backyard | 0     | 13(+5)   | 0    | 1             | 14(+5)    |
| captive  | 0     | 7(+5)    | 0    | 0             | 7(+5)     |
| poultry  | 0     | 116(+37) | 0    | 3(+2)         | 119(+39)  |
| wild     | 7     | 169(+46) | 1    | 21(+8)        | 198(+60)  |
| Total    | 7(+6) | 305(+93) | 1    | 25(+10)       | 338(+109) |

# Répartition géographique des clusters phylogénétiques



28/11/2025



15/01/2026

> 40 introductions primaires estimées en élevage +  
bassecours + captif (France)