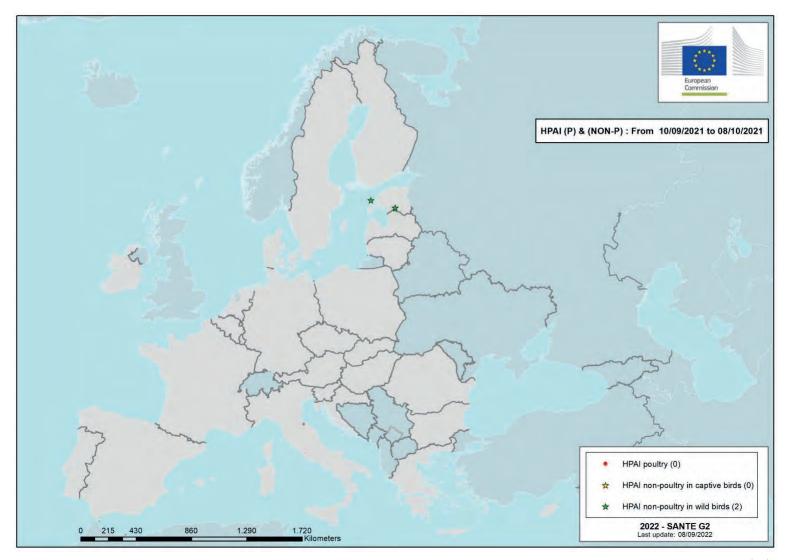
## EU situation Sept. 2021 – Sept. 2022





## HPAI viruses detected in EU during summer - information from EURL

Potential cause for the high number of HPAI H5N1 outbreaks in wild birds during summer:

 a large number of sites of sea bird species breeding in colony have been affected.

Based on the available genetic information, all the A(H5N1) viruses belong to three genotypes:

- two persistently circulating in Europe since the beginning of the 2021-2022 epidemic wave, and
- o one novel genotype, <u>mainly</u> identified <u>in gulls</u>, resulted from reassortment events with viruses of the gull-adapted A(H13) subtype. This new genotype has been identified in France, the Netherlands and Belgium.

