

The ENIGMA HPAI model – forecasting highly pathogenic avian influenza in Europe

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The ENIGMA HPAI model is an endemic/epidemic highly pathogenic avian influenza (HPAI) H5 forecasting model, incorporating within-country transmission, between-country transmission, short-distance transmission (from directly neighboring countries), and long-distance transmission. The model and its forecast publicly available at www.enigmahpai.org

The aim of the ENIGMA HPAI model is to enhance global preparedness for avian influenza, enabling countries in Europe to inform decision-makers and implement preventive measures based on the regularly updated data from WOAH-WAHIS.

The underlying model has been published in Scientific Reports: Kjær et al. (2023), Sci Rep 13, 15396 (2023).

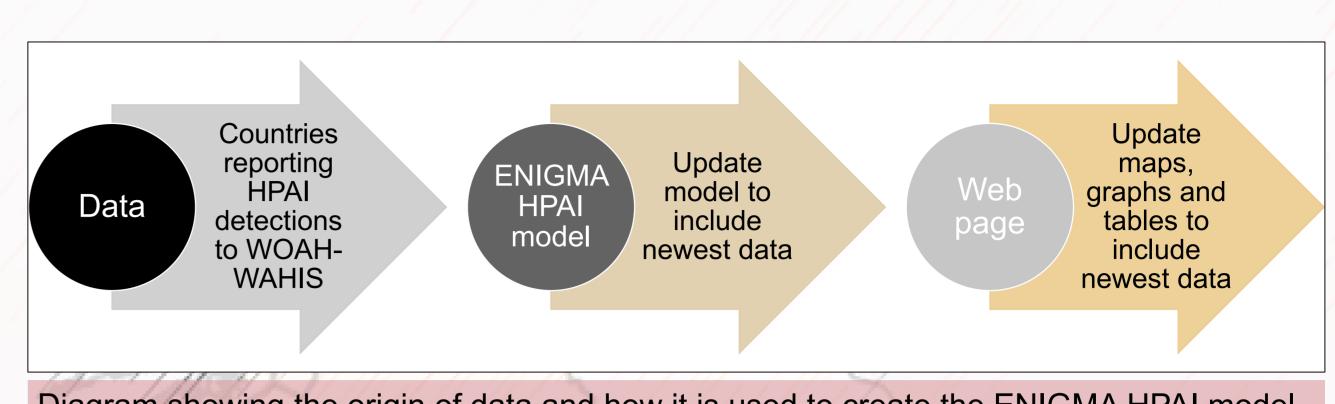
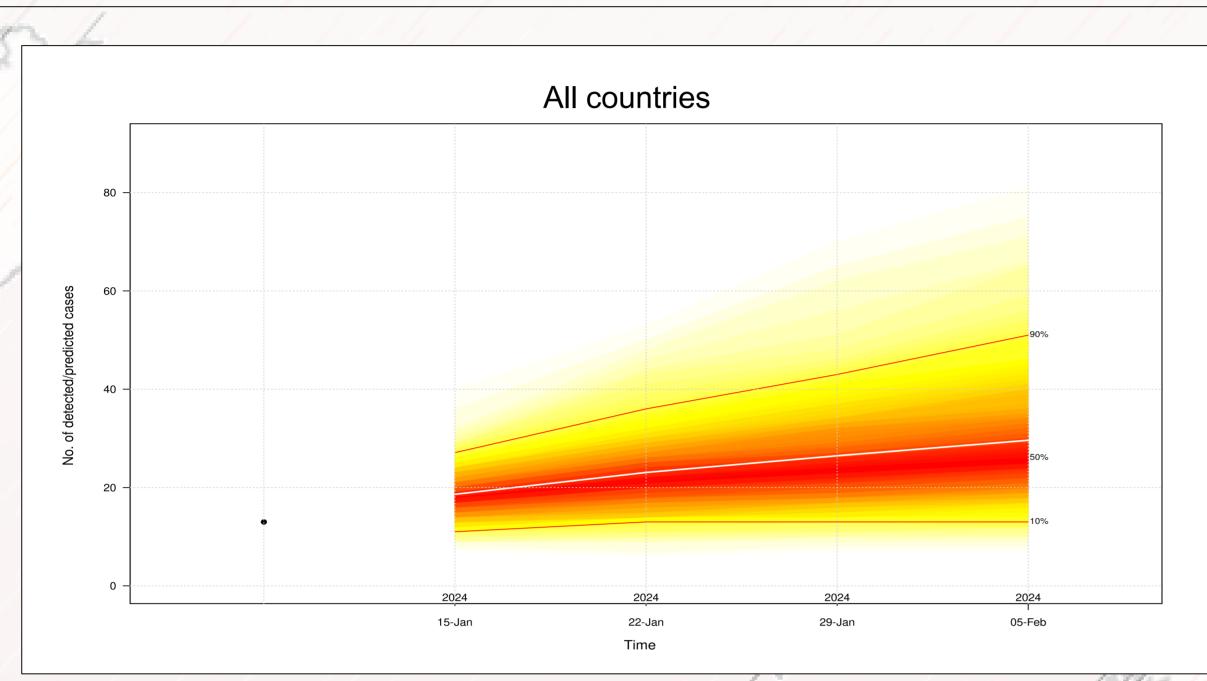
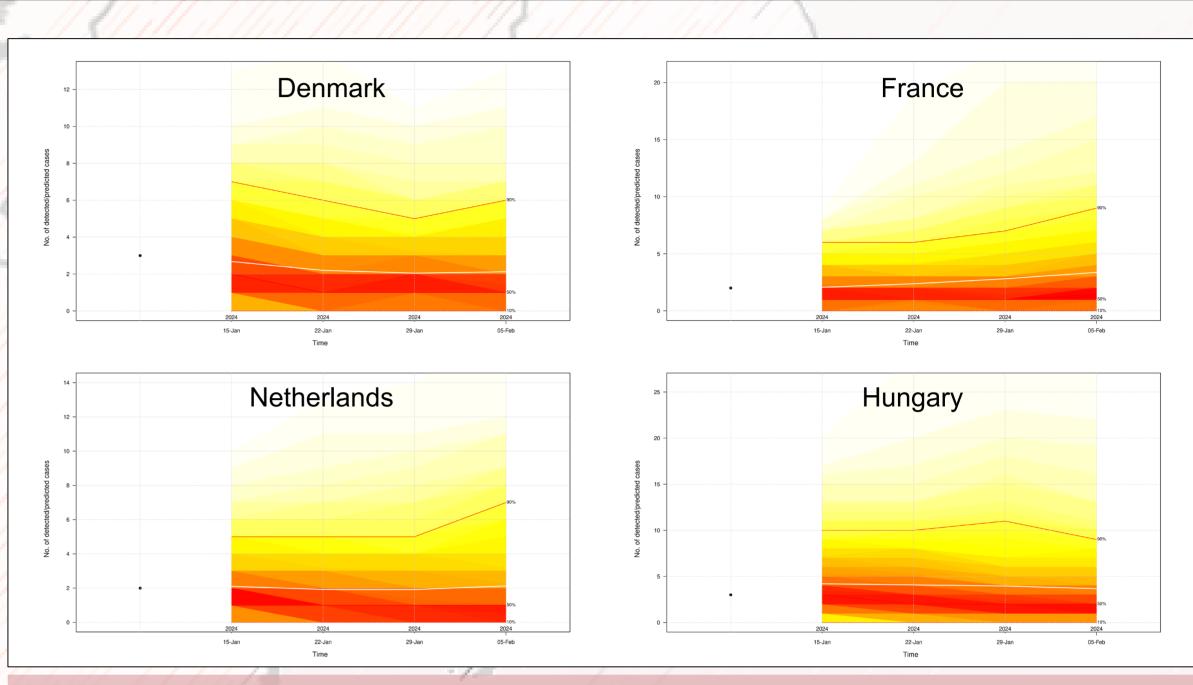


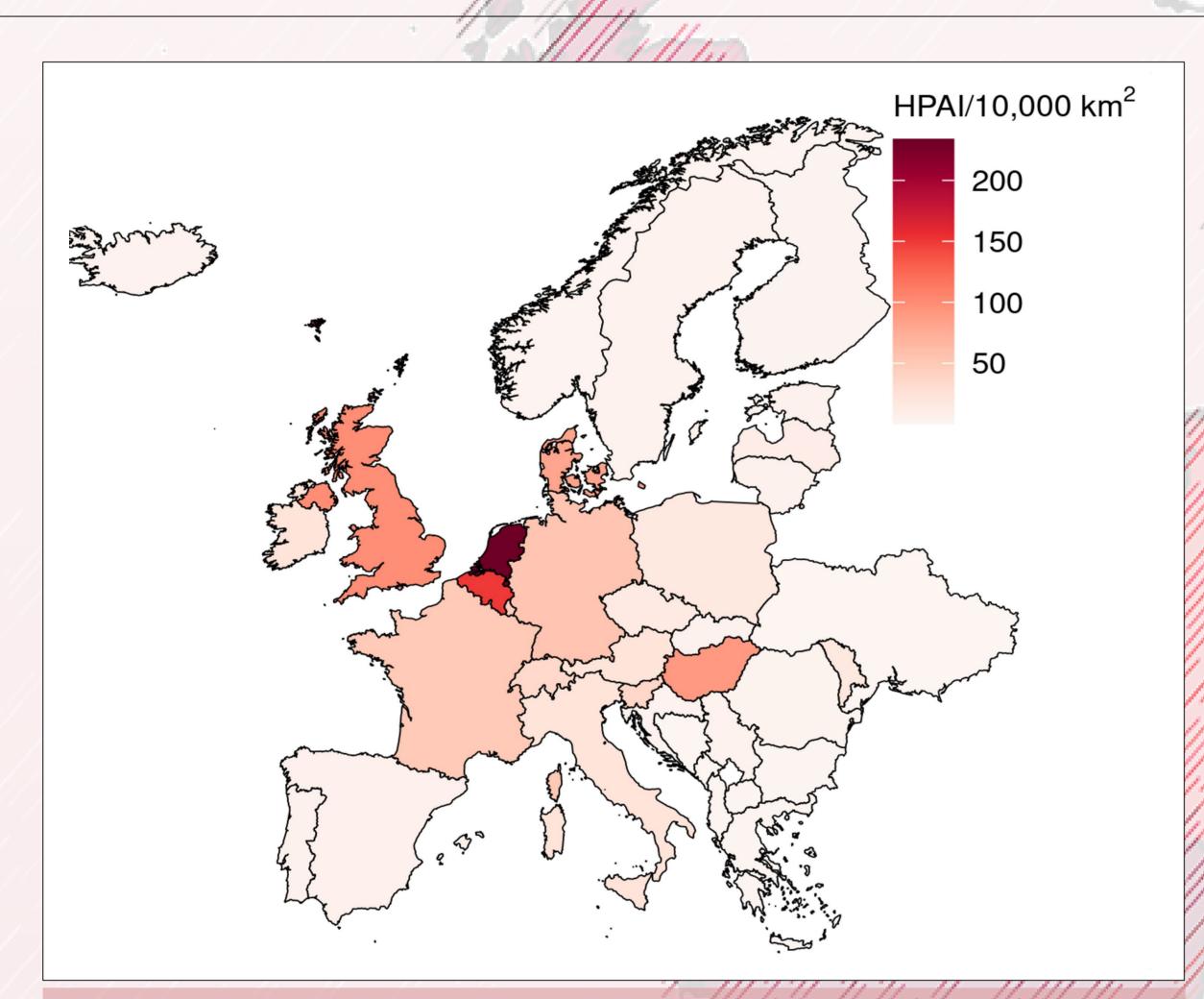
Diagram showing the origin of data and how it is used to create the ENIGMA HPAI model.



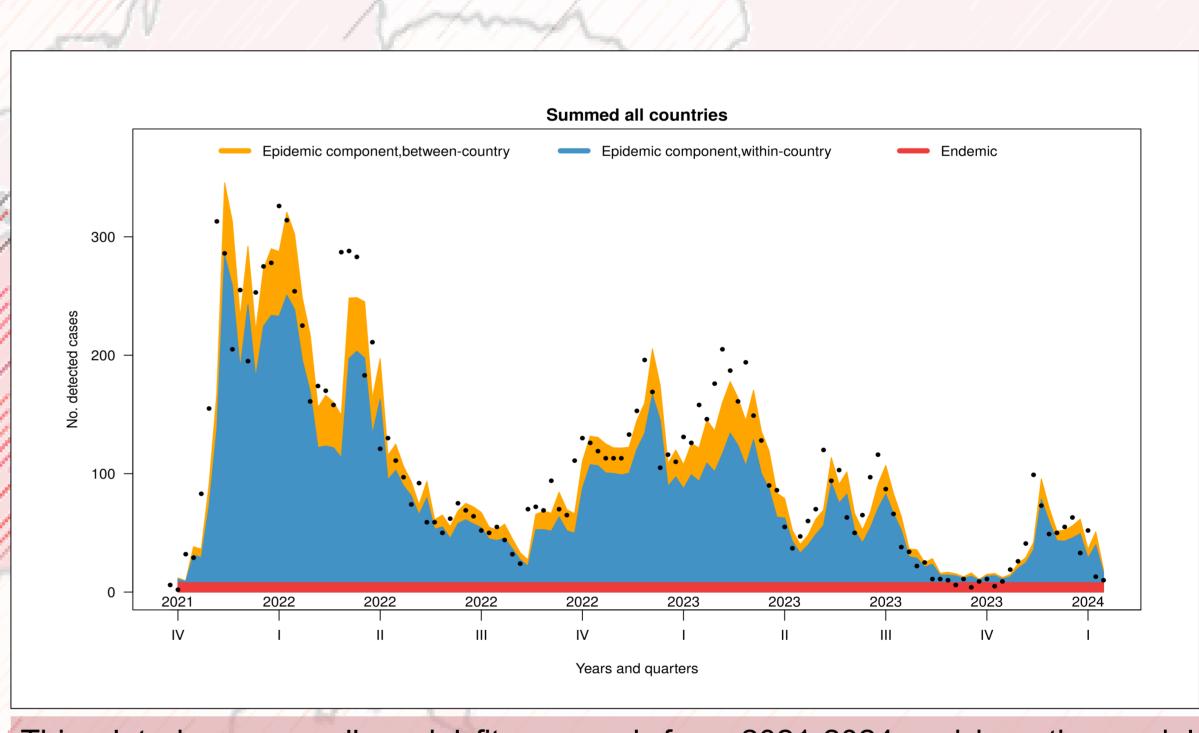
The ENIGMA HPAI model forecasts 4-week simulation-based HPAI forecast with 10%, 50%, and 90% quantiles based on 500 simulations. The mean is shown as white line, and the black dot signifies detections in the first week of 2024.



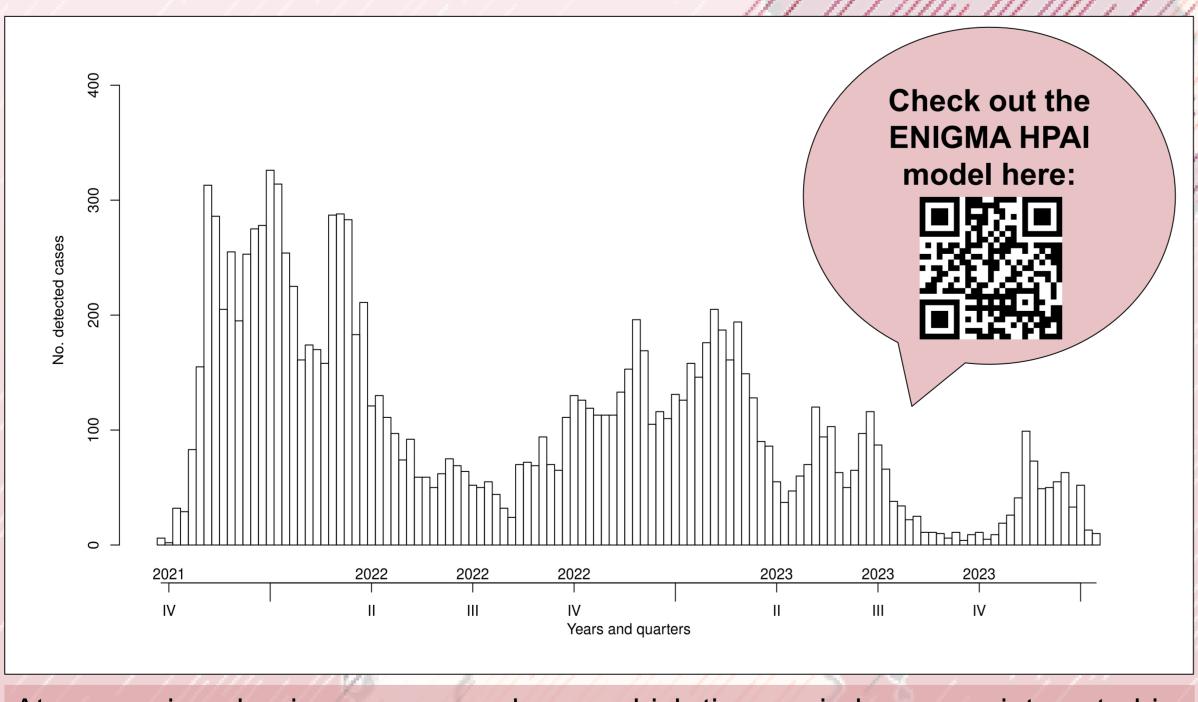
You can also find forecasts for individual countries, here for Denmark, France, The Netherlands, and Hungary.



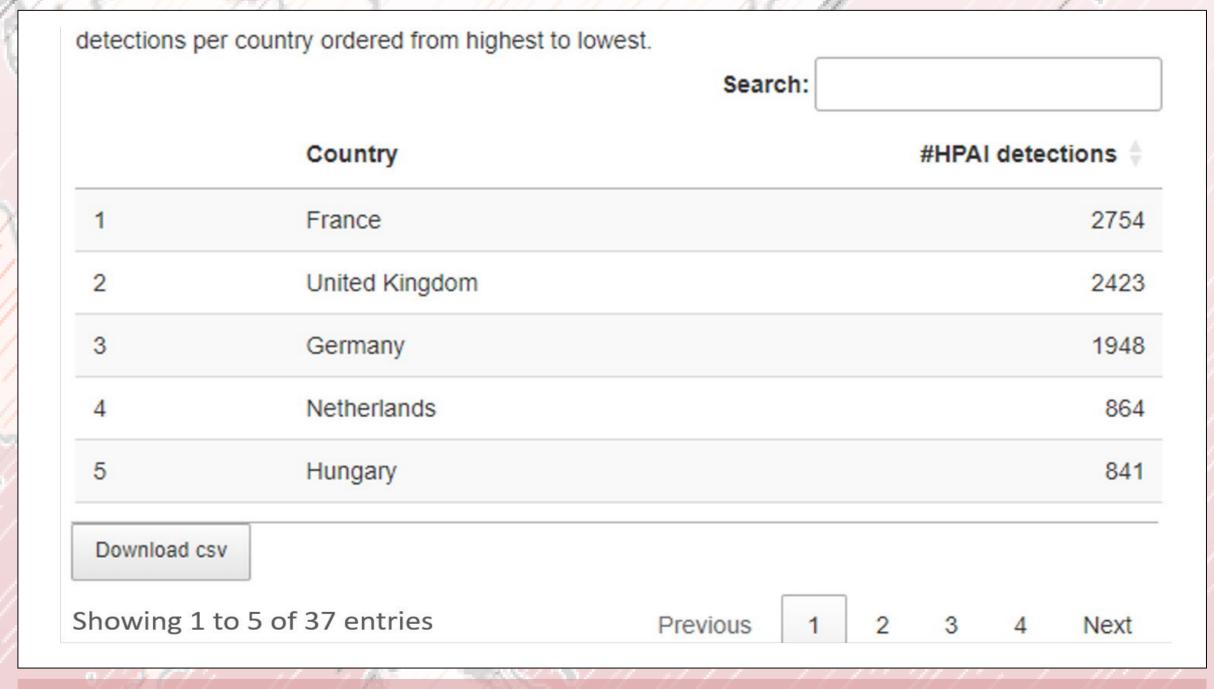
At www.enigmahpai.org you can also find maps of HPAI detections reported to WOAH-WAHIS within the selected time period.



This plot shows overall model fit per week from 2021-2024 and how the model divides predictions into endemic- and epidemic components.



At www.enigmahpai.org, you can choose which time period, you are interested in, and the model will make several plots based on the time period selection.



Tables of HPAI detections reported to WOAH-WAHIS within a selected time period can be downloaded as a csv-file.

All maps, graphs and tables on the home page can be downloaded for further use.

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