

# 1Landscape and spatial patterns of avian influenza virus in Danish wild birds, 2006-2020

2**Running title:** Spatial patterns of avian influenza in Danish wild birds

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16**Abstract** (max 300 words, now: 300)

17Avian influenza (AI) is a contagious disease of birds with zoonotic potential. AI virus (AIV) can  
18infect most bird species, but clinical signs and mortality vary. Assessing the distribution and factors  
19affecting AI incidence can direct targeted surveillance to areas at risk of disease outbreaks, or help  
20identify disease hotspots or areas with inadequate surveillance.

21           Using virus surveillance data from passive and active AIV wild bird surveillance,  
222006–2020, we investigated the association between a range of landscape factors and game bird  
23release and the presence of AIV. Furthermore, we assessed potential bias in the passive AIV

24surveillance data submitted by the public, via factors related to public accessibility. Lastly, we  
25tested the AIV data for possible hot- and cold spots within Denmark.

26           The passive surveillance data was biased regarding accessibility to areas (distance to  
27roads, cities and coast) compared to random locations within Denmark. We found significant effects  
28of variables related to coast, wetlands and cities for the passive and active AIV surveillance data  
29( $P < 0.01$ ), but found no significant effect of game bird release. We used these variables to predict  
30the risk of AIV presence throughout Denmark, and found high-risk areas concentrated along the  
31coast and fjords. For both passive and active surveillance data, low-risk clusters were mainly seen  
32in Jutland and northern Zealand, whereas high-risk clusters were found in Jutland, Zealand, Funen  
33and the southern Isles such as Lolland and Falster.

34           Our results suggest that landscape affects AIV presence, as coastal areas and wetlands  
35attract waterfowl and migrating birds and therefore might increase the potential for AIV  
36transmission. These findings have enabled us to create risk maps of AIV incidence in wild birds and  
37pinpoint high-risk clusters within Denmark. This will aid targeted surveillance efforts within  
38Denmark and potentially aid in planning the location of future poultry farms.

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40**Keywords:** Avian influenza, AIV surveillance, wild birds, high-risk clusters, landscape, spatial  
41patterns

42

## 431 **Introduction**

44Avian influenza (AI) is a contagious disease of birds with zoonotic potential. It is caused by  
45Influenza A viruses (AIV), and can be classified as low pathogenic (LPAI) and high pathogenic  
46(HPAI) subtypes based on their pathogenic phenotype. Only AIV of subtype H5 and H7 are known  
47in the HPAI form. LPAI is a persistent problem worldwide. LPAI is spread between most species of

48birds, and LPAI subtypes H5 and H7 have the potential to mutate into HPAI, which can cause great  
49economic loss and animal welfare problems when farmed birds are infected (Rao et al., 2009;  
50Monne et al., 2014). Furthermore, some AIV subtypes have zoonotic potential with high case-  
51fatality for humans (Lai et al., 2016), thus, it is crucial to monitor and prevent the spread of AIV in  
52both wild and farmed birds. Control measures to prevent the spread of AI include transport  
53restrictions between areas at risk, contact tracing, hygiene measures and culling exposed animals  
54(Stegeman et al., 2004).

55

56Several countries have implemented surveillance programs to monitor the distribution of AI and  
57evaluate the spatio-temporal risk, both for wild and farmed birds (Buscaglia et al., 2007; Hesterberg  
58et al., 2009; Bevins et al., 2014; Machalaba et al., 2015). Data obtained from these surveillance  
59programs can aid in developing statistical spatio-temporal models to identify high-risk areas and  
60critical time periods, which can optimise surveillance for AI. Prediction models for AI have, to a  
61large extent, focused on landscape use, which can indicate the density of specific birds with higher  
62risk of transmitting AIV. Studies have also been able to identify continental hotspots for subtypes of  
63AIV (Bevins et al., 2014), showing that it is possible to identify risk factors on a large geographical  
64scale. Denmark has dense wild birding areas that intersect with many bird migration routes,  
65including routes coming to and from Europe (Bregnballe et al., 1997), Africa (Tøttrup et al., 2018)  
66and Siberia (Dick et al., 1987). Therefore, there is a high potential for AIV incursions from other  
67regions. In particular, migratory birds from Siberia seem to be a risk factor, as Siberia has  
68previously been identified as a major hub for AIV spread (Li et al., 2014; Lai et al., 2016).  
69Additionally, a large number of game birds are released every year for hunting in Denmark  
70(Kanstrup et al., 2009; Gamborg et al., 2016). Some of these game birds originate from other

71countries (The Danish Hunting Association, 2020; Ministry of Environment and Food of Denmark,  
722020), increasing the potential of introducing AIV into Denmark.

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74Since 2002, the Danish authorities have carried out surveillance for AIV in wild birds. We obtained  
75data from this surveillance system generated between 2006 and 2020 and explored potential  
76patterns of AIV occurrence and spatial risk factors in Denmark. The aim of the study was to identify  
77areas with high or low occurrence of AIV and possible factors associated with these occurrences, in  
78order to optimize future surveillance for AIV. We furthermore aimed to assess bias in the Danish  
79passive AIV surveillance data submitted by the public by assessing variables related to human  
80accessibility. Potentially, our results can be applied to future planning efforts; for example, high risk  
81areas should be excluded when planning the location of future poultry farms.

82

## 832 **Materials and methods**

### 842.1 **Passive and active AIV surveillance data**

85We obtained virus detection data from both passive (2006–2020) and active (2007–2019) wild bird  
86AIV surveillance. Passive surveillance data were from the EU mandatory passive surveillance  
87programme in Denmark, where dead and diseased wild birds are tested for AIV and particularly H5/  
88H7 subtypes, whereas the active surveillance data are based on samples from healthy birds, captured  
89for sampling or ringing, submitted by hunters, or from bird dropping samples. Some of the  
90observations in the active AIV surveillance data were pooled samples, whereas others were from  
91individual birds, which had to be taken into account when analysing the data (see section 2.4).

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93All data were manually checked for entry errors and plotted in ArcMap 10.6.1 (Environmental  
94Systems Research Institute, 2017) to check for any errors in the coordinates (such as coordinates not

being located within Denmark). The passive location data all had UTM coordinates for where the birds were found. As the birds from the passive AIV surveillance data were found by the public, we suspected it to be biased due to varying detection probabilities as well as human accessibility to wildlife areas. To assess this, we compared various accessibility variables of the passive AIV surveillance location data to random locations within Denmark (See section 2.3 and 2.4). The active surveillance data only had precise UTM-coordinates from 2007–2010. From 2011–2019 the active surveillance data only registered the postal code of where the sample was collected. To create one single dataset for the active surveillance data, we converted the 2007–2010 coordinates to postal codes instead, and conducted all our analyses on active surveillance data at the postal code level. We also created a single wild bird AIV surveillance dataset by combining the passive and active AIV surveillance data, leaving us with three datasets to conduct our analyses on – the passive AIV surveillance data, the active AIV surveillance data, and a combined wild bird AIV surveillance dataset. When combining the active and passive AIV surveillance data, we converted all the passive surveillance data coordinates to postal codes, producing a combined dataset based on postal codes alone. We refer to this combined dataset as the wild bird AIV surveillance data throughout this paper.

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## 2.2 Data on game birds

We obtained data from 2018–2019 on game birds bred and released for hunting from the Danish Environmental Protection Agency. This data had addresses only and no coordinates, thus we used ArcGIS World Geocoding Service (Environmental Systems Research Institute, 2017) to transform all addresses to UTM coordinates. In some cases, only a postal code was reported for the release site, and not a complete address. In those cases, we used the centroid coordinates of the total area of that particular postal code. These centroid coordinates were obtained from a shape file of all Danish

119postal codes and their areas (Danish Map Supply; Kortforsyningen, 2020). There was no  
120information on the origin of the released birds in the data. To test if game bird releases affected AIV  
121presence/absence in the passive and active AIV surveillance data, we extracted observations from  
1222018–2020 from the surveillance data. We included the year 2020, as we allowed for game bird  
123release to have occurred up to 8 months prior to an observation in the surveillance data. For each  
124observation in the passive AIV surveillance data, we then calculated the nearest game bird release  
125within the last 8 months prior to the observation and identified the species released and the number  
126of birds released. For the active and wild bird AIV surveillance data, we calculated the number of  
127game bird releases and the total number of birds released up to 8 months prior to the observation  
128within the same postal code as the observation.

129

### 1302.3 Landscape variables

131We obtained Corine land cover data as a 100 m<sup>2</sup> resolution raster consisting of 100x100m pixels  
132(European Environment Agency, 2018). For each observation in the passive AIV surveillance data,  
133we extracted the land cover types for the UTM coordinates using the raster package (Hijmans,  
1342019) in R 3.5.2 (R Development Core Team, 2018). We furthermore calculated distance to coast  
135and distance to wetlands for the passive surveillance data in ArcMap 10.6.1 (Environmental  
136Systems Research Institute, 2017). Distance to wetlands was calculated by selecting only Corine  
137land cover types defined as wetlands (inland marshes, peat bogs, salt marches, salines, intertidal  
138flats). We then calculated the closest distance from our observations to a wetland pixel centroid. To  
139calculate distance to coast line, we obtained a shape file of the Danish coast line (Danish Map  
140Supply; Kortforsyningen, 2020) and added a 1 km buffer. We then calculated the closest distance  
141from our observations to this buffered coastline. To assess the effect of accessibility on passive AIV  
142surveillance locations, we furthermore calculated distance to roads and distance to cities as well as

143population density at each location. To calculate distance to roads, we obtained a shape file of all  
144roads in Denmark (Danish Map Supply; Kortforsyningen, 2020) and calculated the closest distance  
145to a road for each location. Population density at a location was extracted from the Gridded  
146Population of the World dataset (raster with 1 km<sup>2</sup> resolution; Socioeconomic Data and  
147Applications Center, NASA, 2015). We also used this raster data to calculate distance to nearest  
148city, defining a city to be a raster grid cell with  $\geq 200$  inhabitants/km<sup>2</sup>. Distance to nearest city pixel  
149centroid was then calculated for each location in the passive AIV surveillance data. All distance  
150calculations were done in ArcMap 10.6.1 (Environmental Systems Research Institute, 2017).

151  
152As the active and wild bird AIV surveillance data were at the postal code level, instead of distances,  
153we calculated the area of wetlands, coast and city within a postal code. We chose the area of city as  
154a measure of whether the area within a postal code was mostly rural with a low density of people or  
155if it was more densely inhabited. Area of wetland and coast were calculated using the 100 m<sup>2</sup>  
156resolution Corine land cover data (European Environment Agency, 2018), whereas area of city was  
157calculated using the Gridded Population of the World dataset (raster with 1 km<sup>2</sup> resolution;  
158Socioeconomic Data and Applications Center, NASA, 2015). As with the passive AIV surveillance  
159data calculations, a city was defined as having  $> 200$  inhabitants/km<sup>2</sup>. These calculations were done  
160in R 3.5.2 (R Development Core Team, 2018), using the raster package (Hijmans, 2019).

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## 1622.4 Bias in the passive AIV surveillance data

163To assess any potential bias in data submitted by the public, we compared our passive AIV  
164surveillance data locations to random locations within Denmark in regards to accessibility. We  
165created random locations and extracted distance to coast, distance to roads, distance to cities, and  
166population density for each of these locations, using the same methods as in section 2.3.

167

## 1682.4 Statistical analysis

169To test for bias in the passive AIV surveillance data, we compared accessibility variables from these  
170locations to the random generated locations using a Kolmogorov-Smirnov test in R 3.5.2 (R  
171Development Core Team, 2018)

172

173We used mixed generalized linear models (GLMs) in the lme4 package (Bates et al., 2015) in R  
1743.5.2 (R Development Core Team, 2018) to test for associations between landscape and game bird  
175variables and passive, active and wild bird AIV surveillance data. For the passive AIV surveillance  
176data, we used year and month of the observations as random effects, since we knew that  
177observations varied over the months and years. We could not estimate prevalence due to the nature  
178of the data, and our focus was on whether AIV was present at a location or not. Thus, if multiple  
179birds from the same location were observed on the exact same date (meaning they were probably  
180found together), we aggregated these multiple observations into a single observation with presence  
181of AIV if any of the observations were AIV positive (see section 3.1 regarding the differentiation of  
182subtypes in the data). Exact locations very rarely reoccurred on separate dates (see Section 3.1), and  
183thus location was excluded as a random variable. For the active and wild bird AIV surveillance  
184data, we also used year and month as random variables. These data were based on postal codes and  
185the same postal codes did reoccur between months and years, thus postal code was also used as a  
186random effect. In the active AIV surveillance data, an observation could be anything from a single  
187bird, to a pooled sample of multiple birds. To avoid any errors or misrepresentations arising from  
188this – and as we were only interested in whether AIV had been confirmed within a postal code in a  
189given month – we summarized observations from the same month and postal code into one  
190observation. If any of the multiple observations within the same month and postal code were AIV



191positive, the summarized observation was classified as positive (see Section 3.1). This procedure  
192was also used on the wild bird AIV surveillance data.

193  
194Effect of game bird release was analysed for the years 2018–2020, and we included the year and  
195month of the observations as random effects. As above, we aggregated multiple observations from  
196the same location or postal code on the exact same date (passive AIV surveillance) or from the  
197same month and year (active and wild bird AIV surveillance) into one single presence/absence  
198observation. For the active and wild bird AIV surveillance data, we then calculated the number of  
199releases and the total number of birds released up to 8 months prior to the summarized data for that  
200month and postal code. For active and wild bird AIV surveillance data, we also included postal code  
201as a random effect. We only used the GLM with variables pertaining to game birds, as we wanted to  
202investigate any possible association.

203  
204When needed, for all GLMs, we used backwards stepwise elimination by removing the variable  
205with highest P-value, and re-running the mixed GLM. We also performed an ANOVA between the  
206original and the reduced model to check whether reduction in the residual sum of squares (SS) was  
207statistically significant or not, and compared AIC-values between models. Lastly, we checked the  
208final models for spatial autocorrelation by plotting the residuals.

209  
210If the landscape variables were found to be associated with AIV presence, we wanted to use these  
211variables and the GLM models to predict the probability of AIV presence throughout Denmark. To  
212measure predictive power of our GLM models, we reran the models using a leave-one-out cross  
213validation (LOOCV) scheme. This method fits the model as many times as there are observations  
214and each time, withholds one location. We then used the model to predict the withheld location. By

215withholding all locations, one-by-one, we achieved a measure of predictive power – i.e. how well  
216we could predict the AIV status of each location based on the other locations. As the models could  
217not predict using unknown factor levels in the LOOCV (for example unique postal code or unique  
218Corine land cover), we had to exclude observations whose factor level only appeared once in the  
219dataset. We did this because when leaving out an observation with a unique factor level in the  
220LOOCV, the model based on the remaining factor levels does not recognize the one left out, and  
221thus cannot predict using this factor level. We also investigated the predictive power by estimating  
222accuracy, sensitivity, and specificity to assess the validity of using the model to predict unknown  
223locations.

224

225For the passive AIV surveillance models, we wanted to predict a map of Denmark in a 1 km<sup>2</sup>  
226resolution. To do so, we created three 1 km<sup>2</sup> raster maps that each covered the entire area of  
227Denmark. We obtained Corine land cover data in a 1 km<sup>2</sup> raster resolution (European Environment  
228Agency, 2018), and removed land cover types not observed in the location data, as we would not  
229predict to unobserved land covers. For the other two rasters, for each raster pixel centroid within the  
230rasters, we calculated the distance to coast or to wetlands, and thus created two rasters that for each  
2311 km<sup>2</sup> in Denmark depicted the distance to coast and the distance to wetlands, respectively. We used  
232Corine land cover (1 km<sup>2</sup>, European Environment Agency, 2018) to calculate the distances to coast  
233and wetlands. For the active and wild bird AIV surveillance data, we created data of the area of  
234coast, wetlands and city for each postal code in Denmark (based on Corine land cover 100 m<sup>2</sup>  
235resolution raster, thus the units are in 100 m<sup>2</sup>). All calculations were done in R 3.5.2 (R  
236Development Core Team, 2018).

237

## 2382.5 Cluster analysis

To identify potential clusters of AIV within Denmark, we used the program SatScan and the package `rsatscan` (Kleinman, 2015) in R 3.5.2 (R Development Core Team, 2018). For passive, active and wild bird AIV surveillance data, we performed spatial scan analyses for summarized years and for separate years with an elliptical scanning window, using the Bernoulli probability model and a maximum spatial window size of less than or equal to 50% of the total population at risk. This form of analysis identifies significant spatial clusters where there is a higher (hotspots) or lower (cold spots) number of positive cases within the scanning window than expected based on the Bernoulli probability of the entire study area. SatScan then reports the ODE, which is the ratio of observed number of positive cases within a cluster to the expected number. Interpretation of an ODE of 1 means that there is no difference from the expected number of cases. We used the Gini coefficient (Han et al., 2016) for cluster selection, as it measures the heterogeneity of the cluster collection, aiding us in which clusters to report (multiple smaller clusters versus large joint clusters). All analyses focused on presence or absence of AIV at a specific site or postal code – not the number of cases reported.

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### 2543. Results

#### 2553.1 Passive and active AIV surveillance

Positive data on AIV found in wild birds were sparse, and thus we did not differentiate the positive data by AIV subtype, but only registered whether AIV was detected at a location or not. The different AIV subtypes found are summarised in Table 1. For the same reason, we did not differentiate the data by bird species. The passive AIV surveillance dataset consisted of 2,089 observation entries, with 1,601 unique site locations (Figure 1). Of these 2,089 entries, 189 were AIV positive (Table 1). When summarizing same-date and same-location observations for the mixed GLMs, 208 of the 1,601 unique sites had multiple entries ranging from 2–55 birds. The

summarized dataset used in the GLM contained 1,614 observations, as 11 locations had multiple entries on different dates within the same year (9 locations with 2 dates, and 2 locations with 3 dates, Figure 2). Of the 1,614 observations, 144 were AIV positive. We found significant differences for all accessibility variables when the 1,601 unique locations were compared to 1,601 random locations (all  $P < 0.0001$ , Figure 3), but as positive and negative AIV locations were equally biased, we proceeded with our analyses described in Section 3.3 and 3.4.

The active AIV surveillance dataset consisted of 8,912 observations within 234 unique postal codes (Figure 4). There were 1,066 observations in this dataset that tested positive for AIV (the AIV subtypes are summarised in Table 1). Summarizing over month, year and postal codes for the GLMs produced 873 observations, of which 319 were AIV positive (Figure 2). Combining wild bird AIV surveillance data resulted in 11,001 observations within 480 unique postal codes, and 1,255 AIV positive observations (Figure 5). Summarizing this dataset over month, year and postal code produced 1,977 observations, of which 426 were AIV positive (Figure 2).

The number of observations in both passive and active AIV surveillance differed over the years (Figure 6) and over the months (Figure 7). For passive surveillance, most observations were from January to April with a small peak in November and most of the positive observations were in March and November. In the active AIV surveillance data, most observations were from September to December, which were also the months with the most positive observations. Several different bird species in the surveillance data tested positive for AIV, most often duck species, swans and raptors (Figure 8).

285

### 3.2 Data on game birds

287A total of 2,268 game bird releases were recorded from 2018–2019 at 1,179 unique  
288locations. The total number of birds released was 1,558,302; of these 92.7% were pheasants  
289(*Phasianus colchicus*), 6.6% were mallards (*Anas platyrhynchos*), and 0.7% were grey partridges  
290(*Perdix perdix*).

291

### 2923.3 Landscape and AIV incidence

293For the passive AIV surveillance data, distance to coast and distance to wetlands were significant  
294( $P < 0.01$ , odds ratio (OR) = 0.9994 and 0.9992, respectively), whereas land cover at the location was  
295not. However, we kept the land cover variable in the model, since a comparison of the full and  
296reduced model showed significant differences in the residual SS ( $P < 0.0001$ ) and removing land  
297cover increased the AIC and reduced the  $R^2$  (Table 2, Figure 2). The OR indicates that for every  
298meter increase in the distance from the coast, the likelihood of AIV presence decreases by 0.06%.  
299This decrease in likelihood was 0.08% for wetlands (Table 2). Accounting for both fixed and  
300random variables, the  $R^2$  for the full model was 0.86. For the active surveillance data, only city  
301proved to be significant ( $P < 0.01$ , OR = 0.9822, Table 2), with the OR indicating that for every  
302increase in the area of city (in units of 100 m<sup>2</sup>), the likelihood of AIV decreased by 1.78%. We  
303chose the final model to include area of city and area of coast as variables, as this model was not  
304significantly different from the full model (no significant differences in the residual SS,  $P < 0.05$ ,  
305same  $R^2$  and a reduction in AIC, Table 2, Figure 2).  $R^2$  was 0.52 for the final model. In the wild bird  
306AIV surveillance data, we found that the area of coast ( $P < 0.01$ , OR=1.0008) and the area of city  
307( $P < 0.01$ , OR=0.9887) were significant. We used the reduced model without the wetlands variable,  
308because a comparison of the full and reduced models showed no significant differences in the  
309residual SS ( $P > 0.05$ , Table 2, Figure 2) and we observed a smaller-AIC value and no change in the  
310 $R^2$ . The OR for area of coast indicates that for every unit the area of coast increases (here unit is 100

311m<sup>2</sup>), the likelihood of AIV presence increases 0.08%. For area of city, the OR indicates that for  
312every increase in a unit area of city (unit is 100 m<sup>2</sup>), the likelihood of AIV presence decreases by  
3131.13%. R<sup>2</sup> was 0.43 when both fixed and random variables were included. Detailed results for all  
314mixed GLMs are shown in Table 2, and an overview of the data used and the final GLMs are shown  
315in Figure 2. Residual plots of all final models indicated that the active AIV surveillance model had  
316spatial autocorrelation in the residuals (Supplementary Figure S1), which was further confirmed  
317with Moran's I (I = 0.05, z = 5.70, P < 0.0001). Spatial autocorrelation of the residuals (Moran's I: I  
318= 0.04, z = 4.71, P < 0.0001) was still present when postal code centroid coordinates were included  
319as independent variables in the model, therefore coordinates were excluded from the final model.  
320However, a spline (cross-) correlogram of the final model residuals showed that the spatial  
321autocorrelation was generally weak, with a weak negative autocorrelation (correlation coefficients <  
322-0.20) at distances of 300 km (Supplementary Figure S2). Although the spatial autocorrelation was  
323weak, these results indicate that we did not account for all of the spatial variation within the data.  
324

325We ran the LOOCV for the passive AIV surveillance data on 1,612 out of the 1,614 observations in  
326the summarized dataset, as two land cover types were only found once in the dataset. The LOOCV  
327produced an accuracy of 0.91 when using the default threshold value of 0.5 for classification  
328(probability of AIV presence above 0.5 is classified as a presence, whereas anything below or equal  
329to 0.5 is classified as an absence). However, this accuracy equalled the proportion of AIV negative  
330observations in the data, meaning that the model was not better than predicting all observations to  
331be AIV negative. Hence, the model sensitivity was 0 and the specificity was 1, meaning that none of  
332the positive observations were classified as positive. We could change the threshold to obtain a  
333higher sensitivity (which would then lower the specificity), but we were not able to obtain an

accuracy higher than the proportion of absences (0.91). Thus, predictions for this model should be viewed with caution.

336

For the active AIV surveillance model, the LOOCV was performed on 801 observations, as 72 of the observations in the summarized dataset ( $n = 873$ ) had postal codes only appearing once. With a threshold of 0.5, the active model had an accuracy of 0.77, a sensitivity of 0.62 and a specificity of 0.85. As the proportion of absences (majority class) was 0.65, this model performed better than if all observations were predicted to be absences. We performed LOOCV on 1,836 out of the 1,977 observations in the summarized wild bird AIV surveillance dataset, as 141 postal codes only appeared once in the summarized dataset. With a default threshold value of 0.5, the accuracy was 0.82, with a sensitivity of 0.32 and a specificity of 0.96. Here the proportion of absences was 0.78, thus the model was more informative than a model only predicting absences.

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For the full passive AIV surveillance model, we used the Corine 1 km<sup>2</sup> land cover data and the coast- and wetlands-distance rasters to predict the probability of AIV throughout Denmark. We set the random effects to zero to predict over all years and all months. We found high-risk areas along the coast and around the fjords (Figure 9A). We also used the active AIV surveillance model to predict the probability of AIV presence based on postal code level area of coast and area of city. We chose the active AIV surveillance model over the wild bird surveillance model, as the sensitivity was higher, thus predicted positive postal codes were more likely to be correctly classified in this model than in the wild bird surveillance model. Again, we set the random variables to zero to predict over all the years, months and postal codes. Here we also found the highest probabilities of AIV presence in postal codes with coastline or along fjords (Figure 9B).

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For the game bird release data, we found no significant association with distance to bird release site, bird species released or number of birds released for the passive AIV surveillance data (Table 2). We also did not find any significant association with number of releases and total number of birds released for the active and wild bird AIV surveillance data (Table 2).

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### 3.4 Cluster analysis

The SatScan analysis detected several significant clusters both for the passive, active and wild bird AIV surveillance data. For all the AIV surveillance data, hotspots were mostly found in the southern parts of Denmark, comprising southern Zealand, Lolland/Falster and Funen, whereas cold spots were found in northern Zealand and Jutland (see summarized results in Figures 1, 4 and 5). For the individual years, not all years had detectable clusters or the amount of data were insufficient to perform cluster analysis (See Supplementary Figures S3, S4 and S5, and Table S1). The summarized presence/absence data used for the cluster analyses are shown in Supplementary Table S1.

372

## 3.4. Discussion

We analysed 11,001 observations from the Danish AIV surveillance program collected from 2006 to 2020 and found associations between landscape variables and AIV incidence. We furthermore detected spatial hot and cold spots of AIV incidence within Denmark. We found differences in AIV incidence across months of the year and between years. A higher number of positive samples in the active AIV surveillance were found from September to November, however, these were also the months where most observations occurred. Many of the observations in the active AIV surveillance data originated from hunted birds, thus the higher number of observations from September to December was expected as this time period coincides with the hunting season of many Danish bird



382species (The Danish Hunting Association, 2020). For the passive AIV surveillance data, most  
383positive samples were found between March and November. This coincides with the timing of bird  
384migration, when migratory birds are in transit through Denmark (DOF BirdLife, 2020). We found  
385the largest number of observations in 2006. This coincides with the first outbreak of HPAI in wild  
386Danish birds (Bragstad et al., 2007), an outbreak that occurred in several European countries and  
387caused the EU to fund compulsory active wild bird surveillance in all member countries  
388(Hesterberg et al., 2009; European Commission, 2020). This compulsory active surveillance lasted  
389until 2011, after which the Danish authorities continued active AIV surveillance in wild birds, albeit  
390at a smaller scale (Hjulsager et al., 2018). The increase and decrease in the intensity of surveillance  
391measures can be seen in the increasing number of observations in the active AIV surveillance data  
392from 2007–2010, and the subsequent decrease in the number of observations from 2011–2020. The  
393most sampled species were buzzard, swans and mallards, and the distribution of sampled species  
394shows large variation, reflecting public interest and accessibility to the bird habitats. Therefore, it  
395was not possible to quantify the risk of AIV for the different species.

396

397We found that the passive AIV surveillance data were biased regarding the geographical location of  
398sample sites. The majority of recorded locations were within 35 km of a larger city and within 500  
399m of roads. Public access to Danish beaches might also explain numerous records close to the coast,  
400suggesting that accessibility to wildlife areas biases Danish passive surveillance data. However,  
401passive surveillance is not easy to control as it depends on the willingness and efforts of the general  
402public. Implementation of information campaigns can be of great assistance to reinforce sampling in  
403areas with sparse information or hotspots, and would be a valuable contribution to the ongoing  
404surveillance program.

405

406 For the passive AIV surveillance data, we found that distance to coast and distance to wetlands had  
407 a significant effect on presence of AIV. For the active AIV surveillance data, we furthermore found  
408 an effect of the area of coast and the area of city. Other studies have found effects of landscape  
409 variables and anthropogenic factors on AI incidence in both wild and domestic birds. In Thailand,  
410 Paul et al. (2010) found a positive effect of free grazing ducks, high rice-cropping intensity areas,  
411 densely populated areas, short distances to a highway junction, and short distances to large cities on  
412 AIV incidence in poultry. Gilbert et al. (2008) identified duck abundance, human population  
413 density, and rice cropping intensity as risk factors in South East Asia. In Romania, Ward et al.  
414 (2008; 2009) found associations between distance to migratory waterfowl sites, distance to major  
415 roads and distance to rivers or streams and HPAI outbreaks. Using a machine learning (ML)  
416 approach, Belkhiria et al. (2018) found spatial risk areas for AIV in wild birds in California, where  
417 land cover and distance to coast were some of the most important predictors in their model. The  
418 poor performance of our passive AIV surveillance model and the relatively low sensitivity of our  
419 active and wild bird AIV surveillance models, indicate that other factors not considered in this study  
420 might be important for predicting AIV incidence. We did attempt using ML methods on the  
421 summarised data and included environmental MODIS variables. Unfortunately, this did not  
422 improve the models, and thus the simple GLM's were chosen to make prediction maps of AIV (See  
423 Supplementary File for the ML description and results). Migratory birds have long been suspected  
424 of spreading AIV between regions (Sullivan et al., 2018; van der Kolk, 2019) and adding data on  
425 bird migration to our models could potentially be of value. However, no fine-scale data are  
426 available on bird migration routes within Denmark that would enable us to distinguish between  
427 individual locations within the same region. We found no significant association between game bird  
428 releases and both the passive and active surveillance. This could mean that the current legislation  
429 with testing and quarantine for imported game birds is effective to prevent spread of AI into wild

430birds. Our results could also be explained by a lack of data from other years, as we could only  
431perform our analyses for the years 2018–2020.

432

433Our cluster analyses identified several hot and cold spots for AIV presence within Denmark. We  
434generally found hotspots in the southern parts of Denmark, whereas cold spots were found in  
435northern Zealand and Jutland. The southern parts of Denmark lay on the main migration routes of  
436duck and geese, mainly coming from north-eastern Russia and Siberia, whereas the northern and  
437western parts of Denmark lay on migration routes of birds coming mostly from Fennoscandia and  
438north-western Russia (Bregnballe et al., 2003). The Wadden Sea along the south western coast of  
439Denmark is a well-known stop-over for migratory birds on their way south or north (Lotze, 2005).  
440Thus, we could expect a cluster of positive samples here. However, we found no hotspots in the  
441western part of the country. This could be due to biased sampling, as only few people venture into  
442the Wadden Sea region, and dead birds are quickly washed away. It could however also be due to  
443the origin of migrating birds, as migrating birds in the southern parts of Denmark could have  
444travelled from Siberia, which is known to be a hot spot for transmission of AIV (Li et al., 2014; Lai  
445et al., 2016). Thus, land areas within the migration routes coming from Siberia might pose an  
446increased risk of AIV incidence, whereas land areas within other migration routes might not be as  
447exposed to infected birds.

448

449Our predictive maps of AIV in Denmark predicted high-risk areas located around the coast and  
450fjords in Denmark. This suggests that any risk-based surveillance should be concentrated in these  
451areas, particularly high-risk areas that are not extensively covered in the present Danish AIV  
452surveillance, such as the coast and Fjords in northern Jutland. The cluster analysis found hotspots in  
453the southern parts of Denmark, areas that our predictive maps also highlight as being high-risk.

454 These areas should also be included in risk-based surveillance. Knowing which parts of Denmark  
455 constitute high-risk areas for potential AIV transmission might aid in the planning of future poultry  
456 farms. Organic- or free-ranging poultry farms, where the farmed birds can come into contact with  
457 wild birds are of particular concern and any location of such farms in high-risk areas should be  
458 avoided. It is important to note that although we did not divide any of our analyses into AIV  
459 subtypes, the majority of subtypes for the passive AIV surveillance data belonged to the HPAI  
460 types, whereas the majority of subtypes in the active AIV surveillance data belonged to the LPAI  
461 types (Table 1). Thus, our separate passive and active AIV data models can approximately be  
462 interpreted as predicting the risk of HPAI and LPAI occurrence respectively.

463

464 At the beginning of November 2020, a HPAI positive peregrine falcon (*Falco peregrinus*) was  
465 found dead near Sakskøbing on the island of Lolland in the southern parts of Denmark, an  
466 observation not included in our datasets (Ministry of Environment and Food of Denmark, 2020) .  
467 The area where the falcon was found coincides with high-risk areas predicted by both our passive  
468 and active AIV surveillance models. Furthermore, mid-November 2020, there was an outbreak of  
469 HPAI in a poultry farm east of Randers in Jutland (Ministry of Environment and Food of Denmark,  
470 2020), also an observation not included in our datasets. This particular poultry farm kept all their  
471 animals indoors, with little risk of contact to wild birds. Thus, the occurrence of HPAI is puzzling  
472 and as of now, there is no knowledge of how HPAI was transmitted to the farm. This particular area  
473 coincides with predicted low-risk areas in both the passive and active AIV surveillance model,  
474 which further emphasizes this surprising outbreak. Thus, even though our models have deficiencies  
475 regarding predictive power, we were still able to predict and in these cases validate possible areas  
476 where AIV are likely or unlikely to occur.

477

478The results of our study highlight some of the deficiencies in the current Danish AIV surveillance  
479program. The active AIV surveillance program is mostly used by the authorities to study LPAI virus  
480epidemiology whereas the aim of passive AIV surveillance program is the early detection of HPAI  
481viruses. Despite the different objectives of the programs, more knowledge on the epidemiology and  
482transmission of both LPAI and HPAI demands thorough coverage of Denmark in order to be able to  
483determine variables important for transmission. Both our passive and active AIV surveillance  
484models predicted high probabilities of AIV occurrence in the north-western parts of Denmark; an  
485area that is one of the least covered areas in the active surveillance program. More knowledge on  
486AIV presence in these areas is needed, and our findings may elicit implementation of more  
487thorough surveillance in these north-western parts of Denmark. The sparse data on AIV occurrence  
488in Denmark and the variation over the years, makes generalising over our results difficult.  
489Moreover, we were not able to conduct analyses of individual subtypes of AIV. More  
490comprehensive studies and analysis demand more consistent sampling and a stratified sampling  
491scheme for the future surveillance of AIV.

492

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497

#### 498**Conflict of interest**

499The authors declare no conflict of interest.

500

#### 501**Data availability**

502Data is subject to confidentiality and is not freely available.

503

#### 504**Ethics statement**

505The authors confirm that the ethical policies of the journal, as noted on the journal's author

506guidelines page, have been adhered to. No ethical approval was required as this article does not use

507original research data, but data obtained through the Danish authorities.

508

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616

617**Tables**

618**Table 1.** The amount of observations in the Passive and active AIV surveillance data divided into  
619AIV subtypes. In some cases, only H5/H7 was screened for in a test positive for Influenza A virus,  
620thus no further subtyping was performed (“not H5/H7”).

Data	Totals	AIV subtype	# observations
Passive AIV surveillance data		H3 N2	1
		H5	24
		H5 N1	22
		H5 N6	43
		H5 N8	81
		H7	1
		not H5/H7	17
	Total AIV positive		189
	Total AIV negative		1900
	Total observations		2089
Active AIV surveillance data		both H5 and H7	3
		H1 N1	1
		H1 N2	1
		H3 N2	1
		H3 N8	3
		H5	177
		H5 N2	1

	H6 N1	1
	H6 N2	4
	H7	9
	H7 N1	2
	H11 N9	1
	H12 N5	2
	not H5/H7	860
		8912
Total AIV positive		1066
Total AIV negative		7980
Total observations		8912

Table 2. Mixed logistic GLM results for passive, active and wild bird AIV surveillance data. The Corine land cover variable is not shown for the full passive model, as this factor variable had over 20 classes, none of which were significant. The ANOVA P-values are from comparing the reduced model to the full model. The R<sup>2</sup>-values depicted are Nakagawa and Schielzeth's R<sup>2</sup> for mixed models from the MuMIn package (Barton, 2009) in R 3.5.2 (R Development Core Team, 2018). These values show the R<sup>2</sup> for fixed variables only as well as the R<sup>2</sup> for fixed and random variables combined. Abbreviations are explained in the footnote.

Data	Fixed variables	z-value	P-value	Random variables, variance/stddev	ANOVA, P-value	OR	R <sup>2</sup> fixed only/all	AIC
Passive AIV	Corine LC			Month: 0.0055/0.074			0.79/0.86	820.3
	DistToCoast,	-3.31	< 0.001	Year: 1.81/1.35		0.9994		
	DistToWetlands	-2.48	<0.05			0.9992		
	DistToCoast,	-3.98	P<0.0001	Month: 0.03/0.18	< 0.0001	0.9999	0.065/0.40	842.2
	DistToWetlands	-2.78	P<0.01	Year: 1.85/1.36		0.9999		
Active AIV	Coast	1.50	0.13	Month: 2.00/1.42		1.0007	0.033/0.52	985.7
	Wetlands	1.07	0.29	Year: 2.12E-10/1.42		1.0002		
	City	-2.30	P<0.01	PC: 1.79/1.34		0.9823		

	Coast	1.70	0.089	Month: 1.55/1.24	0.29	1.0008	0.028/0.52	984.8
	City	-2.70	P<0.01	Year: 0.00/0.00		0.9822		
				PC: 1.82/1.35				
Wild birds AIV	Coast	2.54	<0.05	Month: 1.05/1.02		1.0008	0.020/0.43	1702.7
	Wetlands	0.18	0.86	Year: 0.26/0.51		1.0000		
	City	-2.69	<0.01	PC: 1.01/1.01		0.9887		
	Coast	2.62	<0.01	Month: 1.05/1.02	0.85	1.0008	0.020/0.43	1700.7
	City	-2.69	<0.01	Year: 0.26/0.51		0.9887		
				PC 1.02/1.01				
Game birds vs. passive AIV	Pheasant	-0.05	0.96	Month: 3.37E-10/2.52E-5		0.9487	0.036/0.19	86.6
	Mallard	0.70	0.48	Year: 0.62/0.79		2.3342		
	NearestRL	-0.35	0.73			1.0000		
	NumBirds	0.28	0.78			1.0001		
	NearestRL	-0.18	0.86	Month: 0.00/0.00	0.47	1.0000	0.006/0.16	84.1
	NumBirds	0.29	0.78	Year: 0.60/0.78		1.0001		

	NumBirds	0.35	0.73	Month: 6.99E-10/2.64E-5	0.65	1.0001	0.003/0.16	82.2
				Year: 0.62/0.79				
Game birds vs.	TotBirds	-1.27	0.20	Month: 1.07/1.04		0.9999	0.026/0.31	139.9
active AIV	NumRL	1.06	0.29	Year: 0.00/0.00		1.0641		
				PC: 0.28/0.53				
	TotBirds	-0.77	0.44	Month: 1.10/1.05	0.32	1.0000	0.096/0.29	138.9
				Year: 0.00/0.00				
				PC: 0.21/0.46				
Game birds vs.	TotBirds	-1.66	0.10	Month: 0.51/0.71		0.9999	0.016/0.25	339.5
wild bird AIV	NumRL	1.58	0.11	Year: 0.14/0.37		1.0743		
				PC: 0.40/0.64				
	TotBirds	-0.68	0.50	Month: 0.55/0.74	0.18	0.1000	0.003/0.25	339.3
				Year: 0.13/0.36				
				PC: 0.39/0.63				
	NumRL	0.10	0.92	Month: 0.53/0.73	0.13	1.0025	5.21E-5/0.24	339.8

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Year: 0.13/0.36

PC: 0.38/0.62

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627Abbreviations: LC = land cover, DistToCoast = distance to coast in meters, DistToWetlands = distance to wetlands in meters, Coast = area of coast within postal

628codes (in units of 100 m<sup>2</sup>), Wetlands = area of wetland within postal code (in units of 100 m<sup>2</sup>), City = area of city within postal code (in units of 100 m<sup>2</sup>), PC = postal

629code, OR = odds ratio, stdev = standard deviation, NearestRL = distance to nearest release site, and NumBirds = the number of birds released there. TotBirds = total

630amount of birds released within the postal code (up to 8 months prior to an observations) and NumRL = number of releases within that postal code.



## 631 Figures

632 **Figure 1.** Passive AIV surveillance data and estimated clusters for the combined years 2006-2020.

633 Clusters were analysed using SatScan on presence/absence of AIV and only significant clusters  
634 with the maximum Gini coefficient are depicted. Satscan calculates ODE, which is the observed  
635 AIV cases divided by expected AIV cases based on the Bernoulli probability of the entire study  
636 area.

637

638 **Figure 2.** Overview of the total amount of data, the data used to run GLMs and the final GLM  
639 models for the passive, active and wild bird AIV surveillance data. PC = postal code, Corine LC =  
640 Corine land cover, DistToCoast = distance to coast in meters, DistToWetlands = distance to  
641 wetlands in meters, Coast = area of coast within postal codes (in units of 100 m<sup>2</sup>), City = area of  
642 city within postal code (in units of 100 m<sup>2</sup>).

643

644 **Figure 3.** Density plots of locations recorded through passive AI surveillance in Denmark, 2006-  
645 2020 (red) and random locations in Denmark (blue) in relation to population density, distance to  
646 nearest city ( $\geq 200$  inhabitants/km<sup>2</sup>), distance to coast and distance to nearest road. All x-axes have  
647 been truncated to omit low density observations. As the kernel density calculations replace each  
648 observation by a small probability density, negative values around observation zeroes will occur.

649

650 **Figure 4.** Active AIV surveillance data and estimated clusters for the combined years 2007-2019.

651 Clusters were analysed using SatScan on presence/absence of AIV and only significant clusters  
652 with the maximum Gini coefficient are depicted. Satscan calculates ODE, which is the observed  
653 AIV cases divided by expected AIV cases based on the Bernoulli probability of the entire study  
654 area.

**Figure 5.** Wild bird AIV surveillance data and estimated clusters for the combined years 2006-2020. Clusters were analysed using SatScan on presence/absence of AIV and only significant clusters with the maximum Gini coefficient are depicted. Satscan calculates ODE, which is the observed AIV cases divided by expected AIV cases based on the Bernoulli probability of the entire study area.

**Figure 6.** Yearly number of observations and AIV diagnosis results from the Danish A) passive AIV surveillance program (2006-2020), B) active AIV surveillance program (2007-2019).

**Figure 7.** Monthly number of observations and AIV diagnosis results from the Danish A) passive AIV surveillance program (2006-2020), B) active AIV surveillance program (2007-2019).

**Figure 8.** Recorded bird species and AIV diagnosis results from the Danish A) passive AIV surveillance program (2006-2020), B) active AIV surveillance program (2007-2019). Only species with at least one positive AIV diagnosis are depicted.

**Figure 9.** Predicted probabilities of AIV presence, based on the A) the passive AIV surveillance data model with variables land cover, distance to coast and distance to wetlands, and B) the active AIV surveillance data with variables area of coast and area of city.