**Outbreak summary:** Total outbreaks = 1 (Submitted)

**Species**

<table>
<thead>
<tr>
<th>Species</th>
<th>Susceptible</th>
<th>Cases</th>
<th>Deaths</th>
<th>Destroyed</th>
<th>Slaughtered</th>
</tr>
</thead>
<tbody>
<tr>
<td>Birds</td>
<td>21170</td>
<td></td>
<td></td>
<td></td>
<td>0</td>
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</tbody>
</table>

**Epidemiology**

**Epidemiological comments**

On 24 February, the National Reference Laboratory confirmed avian influenza virus H5N8 by RT-PCR and, on the next day, by sequencing partial HA and NA sequence fragments. The amino acid sequence at the hemagglutinin cleavage site is consistent with highly pathogenic avian influenza (HPAI). Preliminary results suggest that the sequences are very closely related to the Asian and also to the four European H5N8 HPAI virus sequences reported so far. Whole genome sequencing is in progress.

**Source of the outbreak(s) or origin of infection**

- Unknown or inconclusive

**Measures applied**

- control of wildlife reservoirs
- stamping out
- quarantine
- movement control inside the country
- screening
- zoning
- disinfection of infected premises/establishment(s)

**Animals treated**

Vaccination Prohibited

No

**Diagnostic test results**

<table>
<thead>
<tr>
<th>Laboratory Type</th>
<th>Name of Laboratory</th>
<th>Species</th>
<th>Test Type</th>
<th>Date Results Provided</th>
<th>Result</th>
</tr>
</thead>
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<tr>
<th>Laboratory Type</th>
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<th>Species</th>
<th>Test Type</th>
<th>Date Results Provided</th>
<th>Result</th>
</tr>
</thead>
<tbody>
<tr>
<td>National laboratory</td>
<td>Veterinary Diagnostic Directorate of National Food Chain Safety Office (National Reference Laboratory)</td>
<td>Birds</td>
<td>reverse transcription - polymerase chain reaction (RT-PCR)</td>
<td>24/02/2015</td>
<td>Positive</td>
</tr>
</tbody>
</table>

**Future Reporting**

The event is continuing. Weekly follow-up reports will be submitted.