

Salmonella in Animals 2018

Table 1: *Salmonella* in breeding and commercial poultry flocks

Age/Stage	Tested	Positive (%)	Serotype
Census Sampling¹			
Broilers – before slaughter (<i>Gallus gallus</i>)	4,158	32 (0.8)	<i>S. 6,7:e,h:-</i> (n=5) <i>S. Braenderup</i> (n=9) <i>S. Brandenburg</i> (n=3) <i>S. Infantis</i> (n=1) <i>S. Kentucky</i> (n=6) <i>S. Llandoff</i> (n=1) <i>S. Mbandaka</i> (n=2) <i>S. Nottingham</i> (n=3) <i>S. Orion</i> (n=1) <i>S. Tennessee</i> (n=1)
Broilers - grandparent breeding flocks (<i>Gallus gallus</i>)	8	0	
Broilers – laying hens (<i>Gallus gallus</i>)	339	7 (2.1)	<i>S. Llandoff</i> (n=7)
Broilers – parent breeding flocks (<i>Gallus gallus</i>)	182	1 (0.5)	<i>S. Derby</i> (n=1)
Broilers – parent breeding flocks for egg production (<i>Gallus gallus</i>)	53	0	
Turkeys – fattening	432	51 (11.8)	<i>S. Derby</i> (n=41) <i>S. Senftenberg</i> (n=8) <i>S. Typhimurium</i> (n=2)
Turkey – parent breeding flock	6	0	
Turkey – rearing	4	1 (25)	<i>S. Kentucky</i> (n=1)
Total	5,182	92 (1.8)	<i>S. 6,7:e,h:-</i> (n=5) <i>S. Braenderup</i> (n=9) <i>S. Brandenburg</i> (n=3) <i>S. Derby</i> (n=42) <i>S. Infantis</i> (n=1) <i>S. Kentucky</i> (n=7) <i>S. Llandoff</i> (n=8) <i>S. Mbandaka</i> (n=2) <i>S. Nottingham</i> (n=3) <i>S. Orion</i> (n=1) <i>S. Senftenberg</i> (n=8) <i>S. Tennessee</i> (n=1) <i>S. Typhimurium</i> (n=2)
Objective Sampling²			
Broilers – before slaughter (<i>Gallus gallus</i>)	68	2 (2.9)	<i>S. Braenderup</i> (n=1) <i>S. Kentucky</i> (n=1)
Turkeys – fattening	29	8 (27.6)	<i>S. Derby</i> (n=6) <i>S. Senftenberg</i> (n=2)

¹ When the totality of a population, on which the data are reported, is controlled.

² Strategy based on the selection of a random sample from a population on which the data are reported

Total	97	10 (10.3)	<i>S. Braenderup</i> (n=1) <i>S. Kentucky</i> (n=1) <i>S. Derby</i> (n=6) <i>S. Senftenberg</i> (n=2)
Overall total	5,279	102 (1.9)	<i>S. 6,7:e,h:-</i> (n=5) <i>S. Braenderup</i> (n=10) <i>S. Brandenburg</i> (n=3) <i>S. Derby</i> (n=48) <i>S. Infantis</i> (n=1) <i>S. Kentucky</i> (n=8) <i>S. Llandoff</i> (n=8) <i>S. Mbandaka</i> (n=2) <i>S. Nottingham</i> (n=3) <i>S. Orion</i> (n=1) <i>S. Senftenberg</i> (n=10) <i>S. Tennessee</i> (n=1) <i>S. Typhimurium</i> (n=2)

Source: Department of Agriculture, Food and the Marine (DAFM)

Table 2: *Salmonella* spp. in cattle

Animal	Tested	Positive	Serotype
Suspect Sampling			
Cattle – (Bovine unspecified)	1,618	20 (1.2)	<i>S. Dublin</i> (n=19) <i>S. Montevideo</i> (n=1)
Cattle – (Bovine foetus/stillbirth)	1,914	83 (4.3)	<i>Salmonella</i> spp. unspecified (n=1) <i>S. Dublin</i> (n=72) <i>S. Indiana</i> (n=1) <i>S. Typhimurium</i> (n=9)
Overall total	3,532	103 (2.9)	<i>Salmonella</i> spp. unspecified (n=1) <i>S. Dublin</i> (n=91) <i>S. Indiana</i> (n=1) <i>S. Montevideo</i> (n=1) <i>S. Typhimurium</i> (n=9)

Source: DAFM

Table 3: *Salmonella* spp. in animal feed materials

Type of feed material	Tested	Positive	Serotype
Objective Sampling			
<i>Feed material for land animals</i>			
Cattle	61	0	
Horses	6	0	
Pigs	9	0	
Poultry	14	0	
Sheep	12	0	
Dairy cattle	1	0	
<i>Feed material of marine animal origin</i>			
Fish meal	9	0	
Fish oil	5	0	
<i>Feed material of cereal grain origin</i>			
Barley	18	0	
Maize	71	0	
Other cereal grain	21	0	
Wheat	23	0	
<i>Feed material of oil seed or fruit origin</i>			
Other oil seeds	5	0	
Palm kernel	10	0	
Rape seed	69	1	<i>S. Typhimurium</i> (n=1)
Soya	50	0	
Sunflower seed	16	0	
Other feed material			
Forages and roughages	8	0	
Legume seeds and similar products	1	0	
Other plants	11	0	
Other seeds and fruits	3	0	
Tubers, roots and similar products	24	0	
Overall total	447	1 (0.2)	<i>S. Typhimurium</i> (n=1)

Source: DAFM

References:

Department of Agriculture, Food and the Marine (DAFM)