Infectious bursal disease in Ethiopian village chickens



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selected



Poultry production in Ethiopia and Infectious bursal disease virus (IBDV)

- The majority of Ethiopian poultry are indigenous birds kept in small backyard flocks, belonging to rural smallholders, and are particularly important to landless in society, and also to women.
- Disease is reported by smallholders to be a major constraint to production, but most outbreaks are attributed to Newcastle disease without any investigation of the pathogens responsible.
- IBDV was first reported in Ethiopia in 2002 in a commercial flock, but the very virulent (vvIBDV) strain type has since been identified in all production systems. This strain is associated with high mortality in birds between 3 and 6 weeks of age.
- · Only birds over 3 weeks and less than around 10 weeks of age will have clinical disease $(diarrhoea, dehydration, echymotic\ haemorrhages).\ Adult\ birds\ will\ seroconvert\ without$ showing clinical signs. Chicks under 2 weeks do not normally develop disease, but become immunosuppressed, as the developing B-cells are depleted by the virus, impairing the bird's ability to produce antibodies.

Modelling risk factors

- Multi-level intercept only models were fitted to assess the variation in antibody levels (as measured by the S:P ratio) at each level of the dataset, and the amount of variation between
- Intraclass clustering coefficients indicated ~66% of variation was contributed by individual birds. and ~ 13% by the farm. Market shed and region did not contribute significantly, however, the ELISA plate was an important source of variation. Household and plate were therefore fitted in the final multilevel mixed model as random effects, and village and season were included as
- Other household and bird level variables were screened for inclusion in the final model, and retained where there was evidence that they significantly (p<0.05) improved model fit (Table 1.)
- Due to the non-normality of the residuals, bootstrap estimates of the standard errors (SE) were estimated. The simulated SE's were lower for the effects of Salmonella titre and outbreak history and only slightly increased for the village effect without altering the significance, providing evidence that our estimates are generally robust, despite non-normality. However, the bootstrapped model suggested that Season B was not significantly different to the other seasons (p=0.09).
- Bird-level residuals were examined for spatial clustering, by comparing them with simulated sets of residuals with no spatial correlation. The variogram shown below suggests that, in adjacent villages H1A and H1B, in Season D, there was more similarity between birds found close to each other, as we might expect with an infectious agent. However, spatial clustering of residuals was not evident for all villages in all seasons.

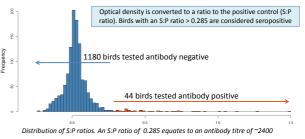
Table 1: Fixed effects	Coefficient	SE	P	o
Intercept	0.060	0.026	0.022	
Continuous variables				9 -
Salmonella s:p ratio	0.012	0.003	0.000	8 6
Categorical variables				6 6
Season A	Reference			so 0 0
Season B	0.040	0.018	0.028	
Season C	0.031	0.027	0.242	
Season D	0.033	0.027	0.230	
Village H1A	Reference			8 0 0
Village H1B	-0.054	0.027	0.045	0.00 0.02 0.04 0.06 0.08 0.10
Village H2A	-0.064	0.027	0.020	Maria and with talance limits of the anciduals
Village H2B	-0.070	0.026	0.006	Variogram with tolerance limits of the residuals
Village J1A	-0.094	0.030	0.002	(from the model in Table 1) for villages H1A and H1B in Season D, showing evidence of spatial
Village J1B	-0.055	0.029	0.052	autocorrelation over short distances (p=0.02)
Village J2A	-0.116	0.028	0.000	dutocorrelation over short distances (p=0.02)
Village J2B	-0.056	0.029	0.051	Season A
No outbreak in last 12 months	Reference			• Season B
Outbreak in chicks	0.014	0.024	0.556	Season C
Outbreak in growers	0.059	0.026	0.022	· Season D
Outbreak in chicks and growers	0.097	0.043	0.025	
Outbreak in adults	0.011	0.012	0.337	
Outbreak in chicks and adults	0.001	0.025	0.978	
Outbreak in growers and adults	0.040	0.028	0.152	H1A)
Outbreak in all age groups	-0.003	0.021	0.879	
This work is part of a larger colla looking at the infectious disease genetic and socio-economic aspe keeping in Ethiopia. Further information can be found www.ch4d.wordpress.com	epidemiology, ects of poultry	1	H1B	

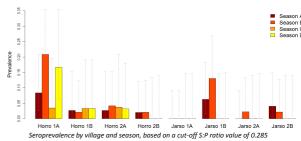
Sampling Methods Data was clustered at 4 levels 2 Regions (Horro & Jarso) Both in highland areas, but with different social demographics 4 Market sheds (groups of villages dependent on a single market) 8 Villages Each village 640 Households H1A was visited 4 May 2011 25 25 25 25 25 25 25 20 200 times, and Oct 2011 25 25 25 25 25 24 25 25 199 different May 2012 15 15 14 16 15 15 15 15 120 households Oct 2012 16 15 were randomly

2 birds of over 6 months of age were randomly selected from each household flock



- · Management data for the household were collected in a questionnaire
- Blood was collected into an anticoagulant and transported to the laboratory, where samples were tested for antibodies to IBD using a FlockScreen Antibody ELISA (x-OvO, Inverkeithing,UK).





- Infectious bursal disease is or has been circulating in at least seven out of the eight villages in our study.
- The association of seropositivity in a household flock with recent deaths in growers, but not chicks or adults fits with the biology of the disease, and is consistent with IBDV contributing to mortality in this population.
- All seropositive birds had moderate to high antibody levels to Salmonella. which makes it unlikely that they were infected as chicks. It is probable that all immunosuppressed birds have died by the age of 6 months.
- The differences between and possibly within villages highlights the need for control strategies to be tailored to the local area, with regard to regional differences in the socio-economic importance of poultry, rather than blanket measures applied.







Residuals (from model in Table 1) for villages H1A and H1B









