

Machine learning models for diagnostics and risk analysis

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- Goal:
 - Augment the results of the Single Intradermal Comparative Cervical Tuberculin (SICCT) test with surrounding epidemiological risk factors:
 - to improve herd-level test sensitivity,
 - thus improve early detection and reduce onward transmission.
- Outcomes:
 - Integration of a large number of risk factors, including:
 - Farm characteristics, movements, testing,
vet practice, tuberculin batch, . . .
 - Herd-level prediction of bTB breakdowns using a machine learning model.
 - Risk factor breakdown using feature importance
 - and deeper analysis.

- Initial idea from:

Article | [Open Access](#) | [Published: 26 January 2021](#)

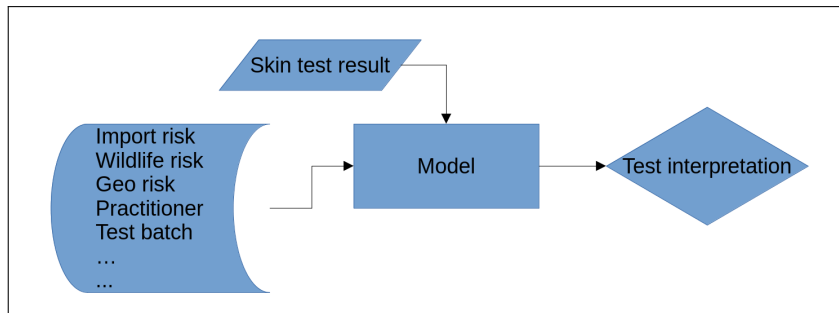
Using machine learning improves predictions of herd-level bovine tuberculosis breakdowns in Great Britain

[K. Stański](#) , [S. Lycett](#), [T. Porphyre](#) & [B. M. de C. Bronsvoot](#)

[Scientific Reports](#) **11**, Article number: 2208 (2021) | [Cite this article](#)

- Improved by:
 - Fewer (streamlined) risk factors, more historical data
 - Histogram-based Gradient Boosted Tree model:
 - Tuned and cross-validated model performance
 - Simplifying the treatment of missing data
 - Including new data from UK Farmcare (Vet practice and tuberculin batch).

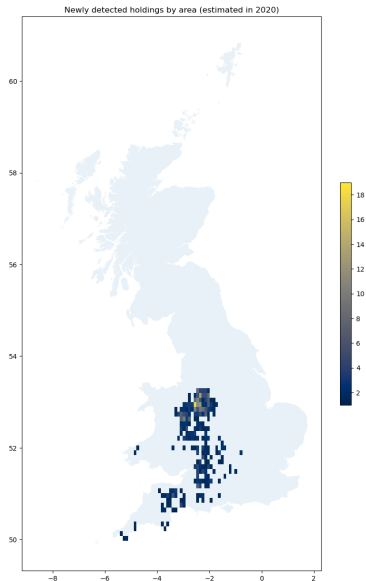
Model and Data



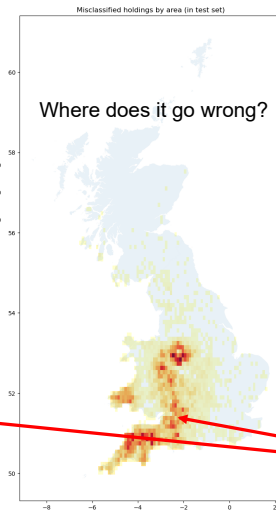
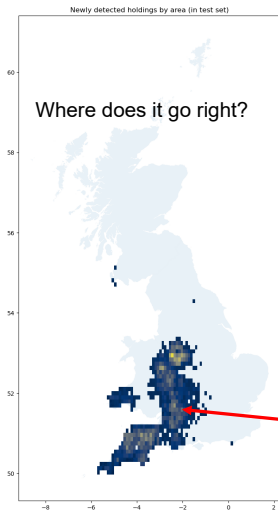
- Data extracted from APHA bTB surveillance database (SAM) and the Cattle Tracing System (CTS) database.
 - Every recorded SICCT test event between January 2012 and September 2021 (1.3m records)
 - Metadata on herd, location, movements, ...
- Data from UKFarmcare on vet and tuberculin batch for some tests:
 - 400 vet practices, covering 120k tests
 - 650 tuberculin batches, covering 57k tests

Results

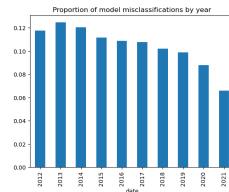
- Model goal is to predict a new breakdown within 90 days of testing.
- Balance of sensitivity vs. specificity can be tuned in the model.
- Here we take the same herd-level specificity as the skin test alone, maximising sensitivity.
- **Increase in herd-level sensitivity 12%.**
- Over one year (2020) we find **around 400** negative herd-level tests, that went on to have breakdowns, but were **identified as positive by the model.**



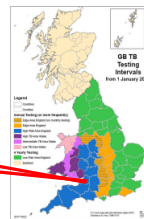
Results – Spatiotemporal



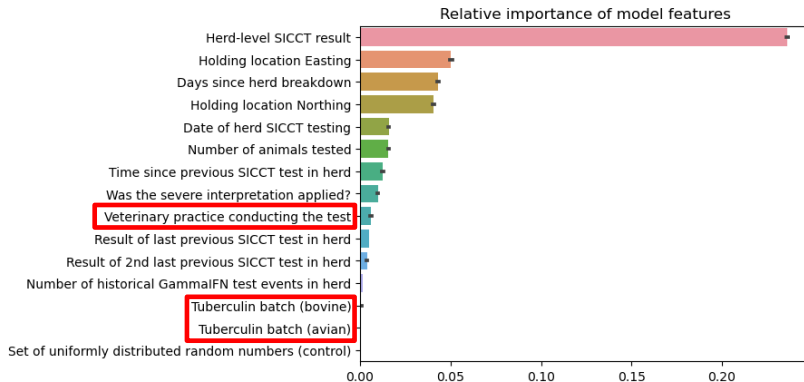
When does it go wrong?



HRA



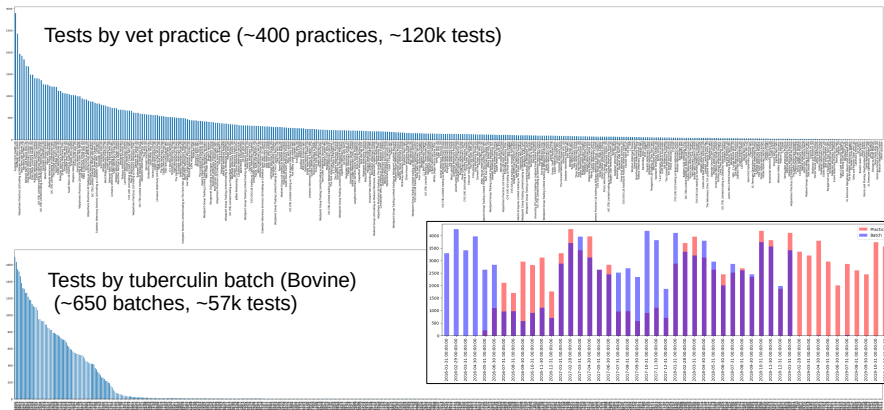
Analysis – Risk factor importance



- Risk factors ranked by permutation-based importance testing.
- Bar length shows accuracy reduction when factor removed from model.

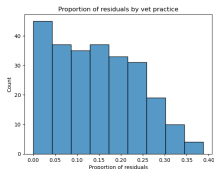
Vet data – Coverage

- Can we further explain risk factor importance of vet data?
- Data coverage may be reducing effectiveness:

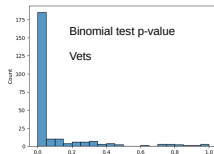


Vet data – Vet/tuberculin accuracy

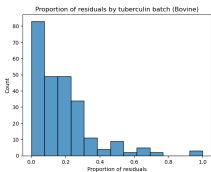
• What can we say about vet / tuberculin performance?



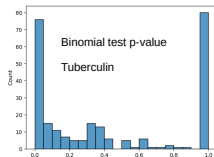
Fairly wide
distribution of
vet performance



74% perform
outside expected
range



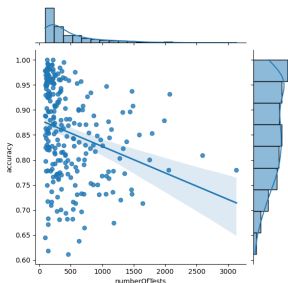
Longer tailed
distribution of
tuberculin performance
(small number very
bad)



30% perform
outside expected
range

Vet data – Performance by practice type

• Which practices perform worse?



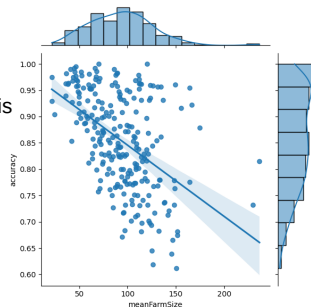
Size of vet practice by number of tests conducted has a weak correlation

$$r = -.27$$

Mean size of herd dealt with by the vet is reasonably well correlated

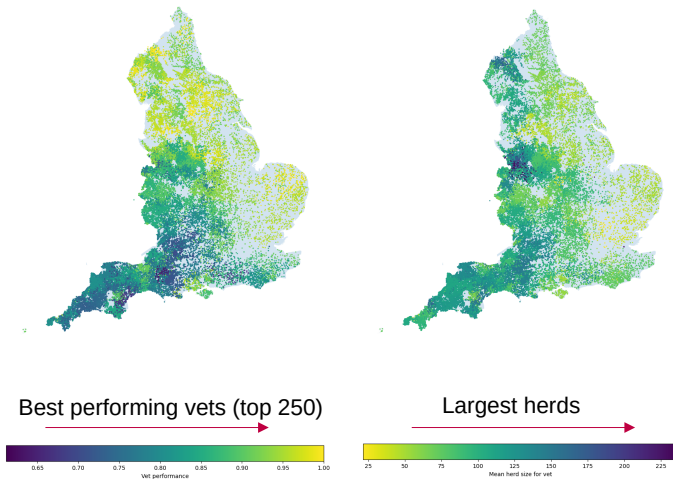
$$r = -.48$$

Vets that deal with larger herds perform worse



Vet data – Performance by practice type

- Geographically?

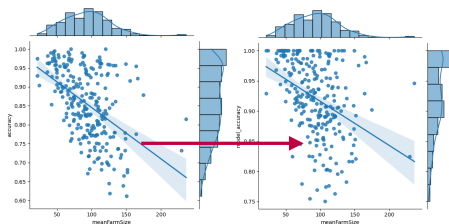
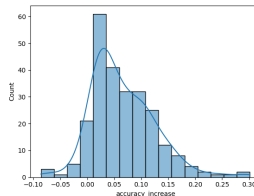


What is the point of all this?

How does the model help?

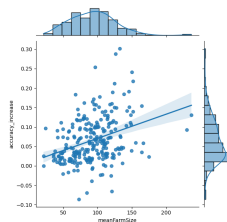
Model augmented testing improves accuracy for most vets.

Some up to 30%:



Model reduces accuracy correlation with mean farm size from $r = -.48$ to $r = -.38$

It increases performance with larger herds:



Where next?

- UK Farmcare to conduct further analysis on vet results
- Possible extension to use skin test measurements instead of binary results
- How could this be used in the field?
 - App interface to model for vets to use at testing?
 - A guide to re-testing high-risk herds?
- Other situations: BVD, Johne's.