

Pooling data for international evaluations for feed intake and efficiency

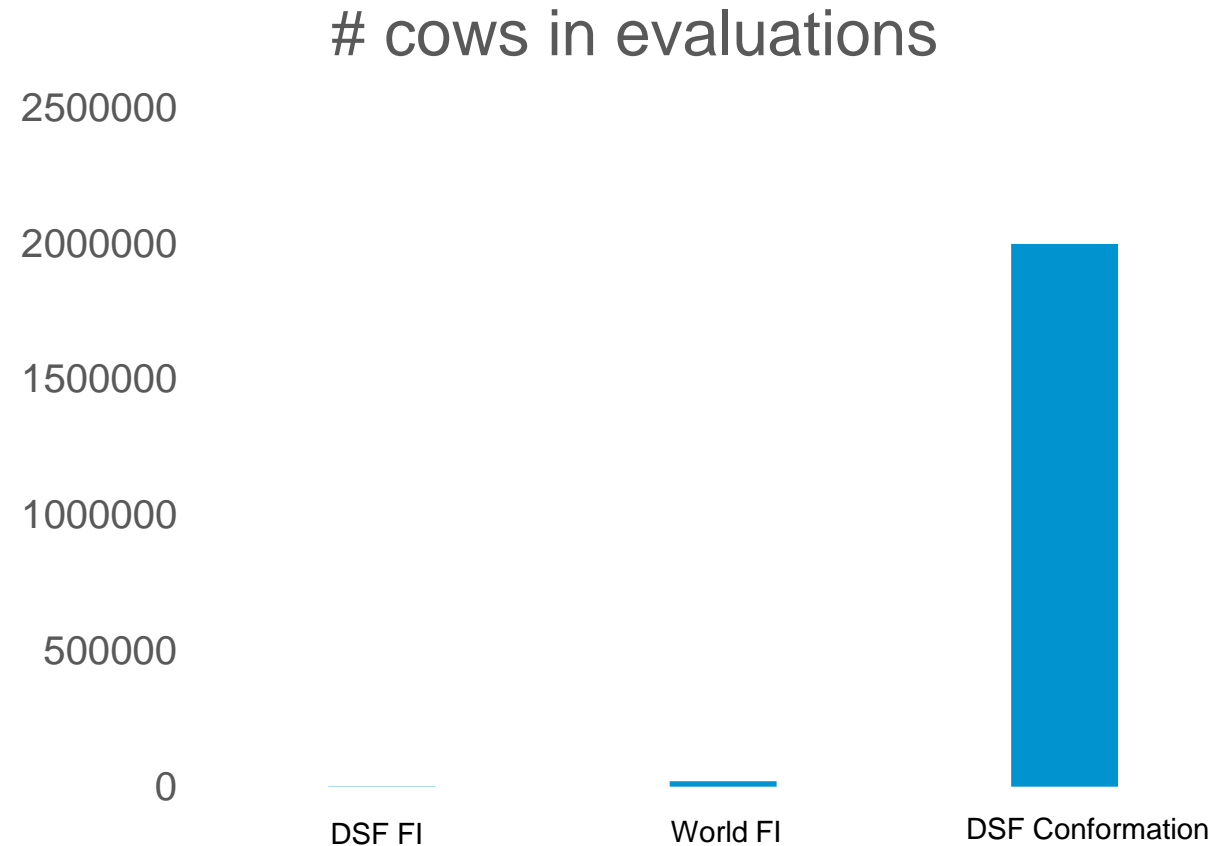
Presented by Jan Lassen



The genomics revolution



”With genomic selection we will be able to select for scarcely measured traits such as efficiency.”





Current status

- 🕒 VERY limited data all over the world
- 🕒 Almost all Holstein
- 🕒 Very complex biology
- 🕒 Huge interest



Challenges

- Genetic links between countries
- $G * E$ is probably substantial
- Registrations protocols are different
- Many phenotypes are old



Data ownership

- 📍 Very different from country to country
- 📍 Often owned by University or specific researcher
- 📍 Very costly registrations
- 📍 A challenge for proper collaboration

RFI models

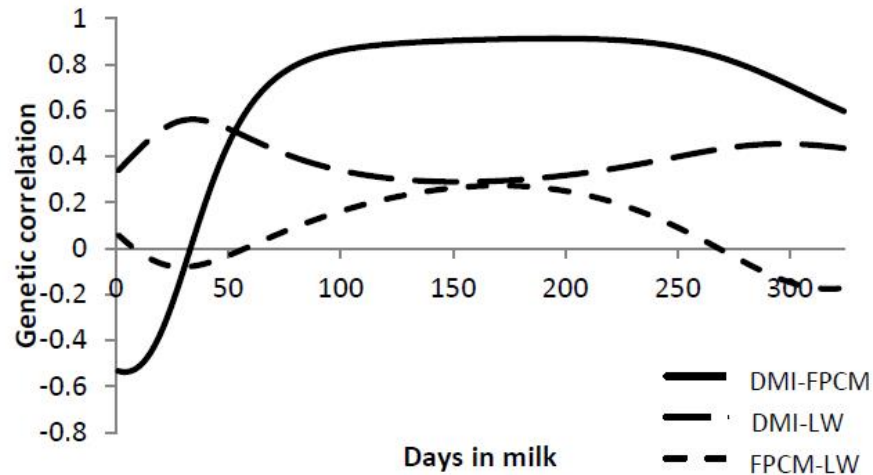
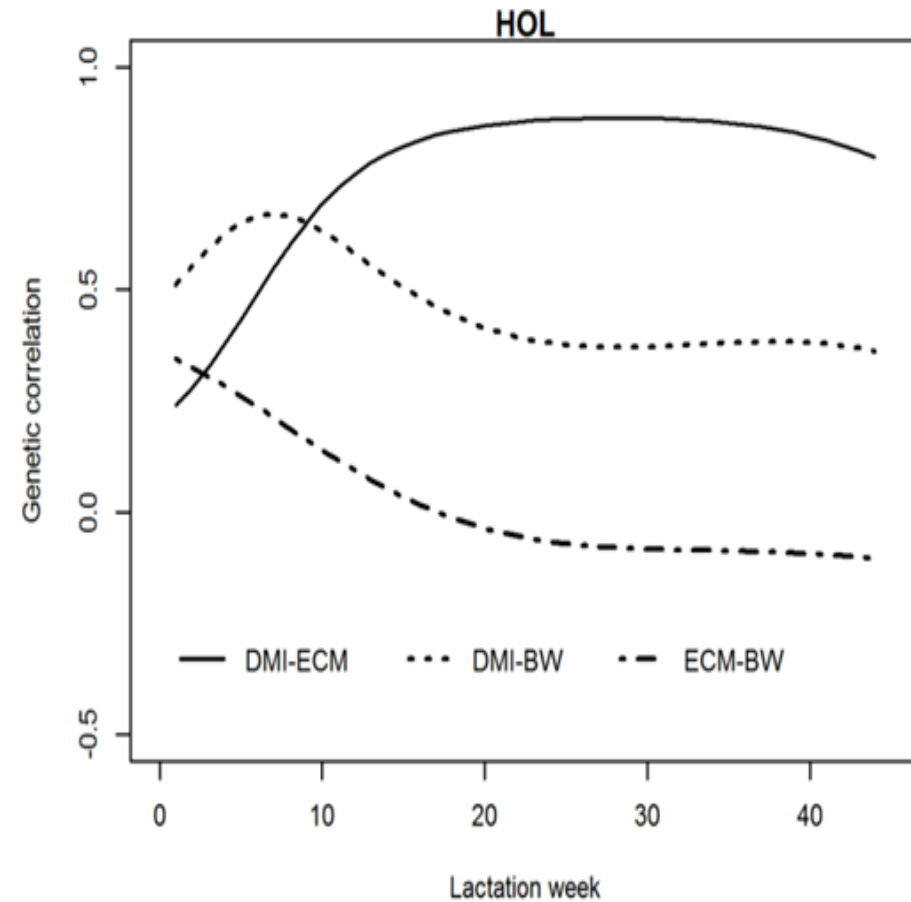


Figure 2.6 Pairwise genetic correlations when two traits are measured on the same day from 1 to 324 days in milk (DIM) between 1. dry matter intake and fat and protein corrected milk (DMI-FPCM, SE of median=0.06, of 3rd quartile=0.09), 2. dry matter intake and live weight (DMI-LW, SE of median=0.11, of 3rd quartile=0.10), and 3. fat and protein corrected milk and live weight (FPCM-LW, SE of median=0.12, of 3rd quartile=0.13).

Manzanilla Pech et al., 2016 JDS



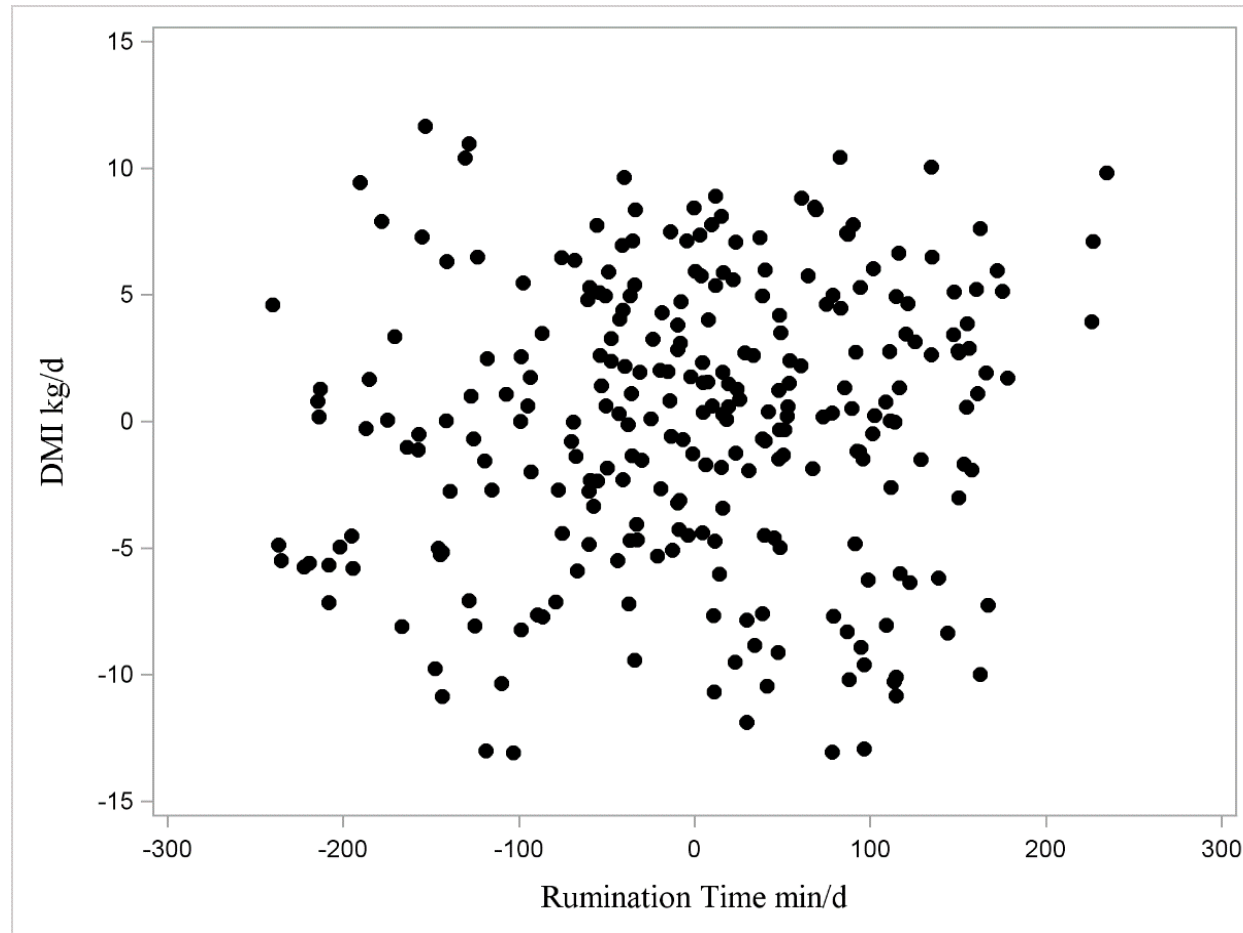
Li et al., 2018 JDS



Indicators

- 🕒 Double counting
- 🕒 No documentation without direct measures
- 🕒 Should be easier and cheaper to measure than the direct trait

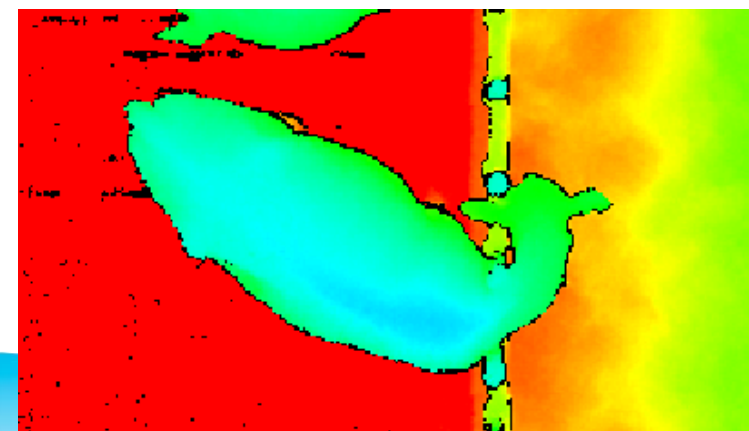
Rumination vs DMI



Efficiency



- 🔍 More data is needed
- 🔍 Not sure indicators will help
- 🔍 Biology is hardly adequately described
- 🔍 How to quantify progress
- 🔍 Direct measures



Cattle Feed Intake CFIT

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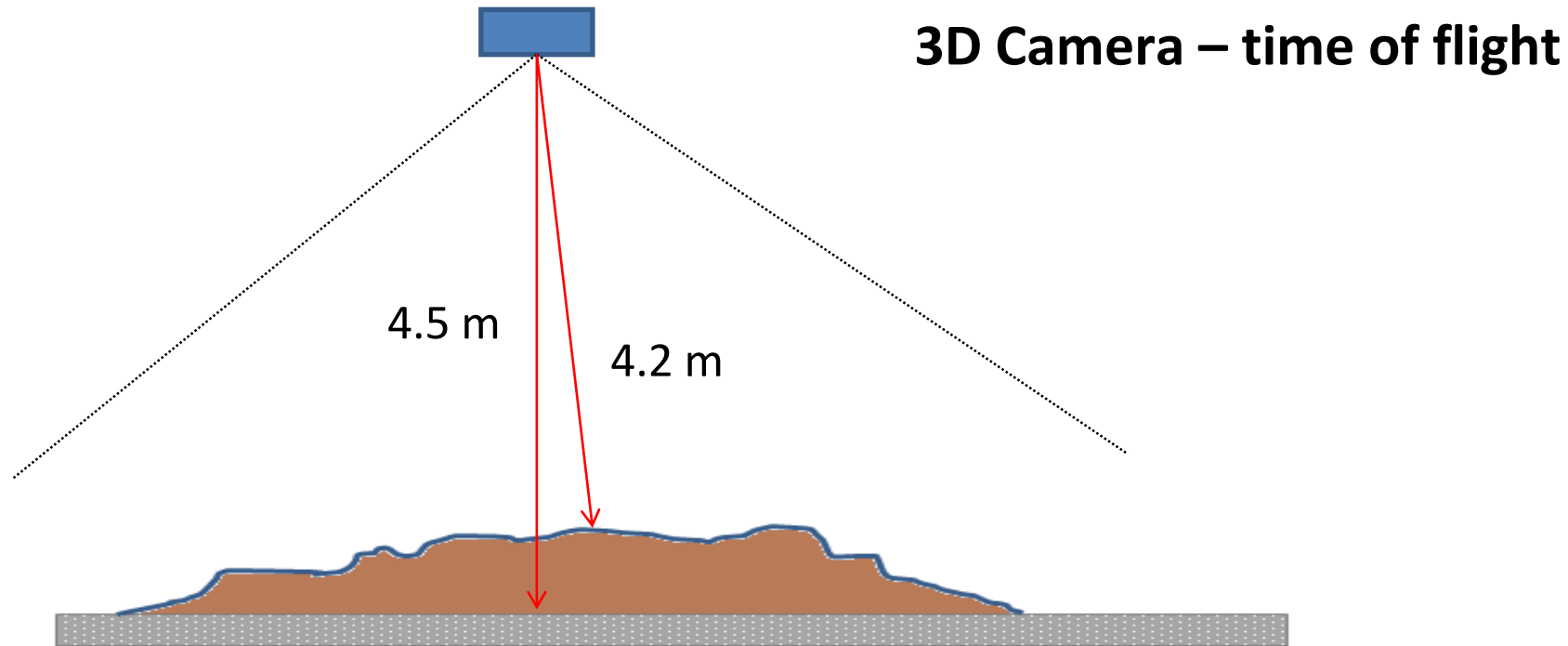


Aim and purpose

- To develop a 3D camera system that can measure feed intake at individual cow level at each visit
- May not:
 - Disturb daily routines on farm
 - Disturb cow behaviour
- Should be same system as for identification



System setup

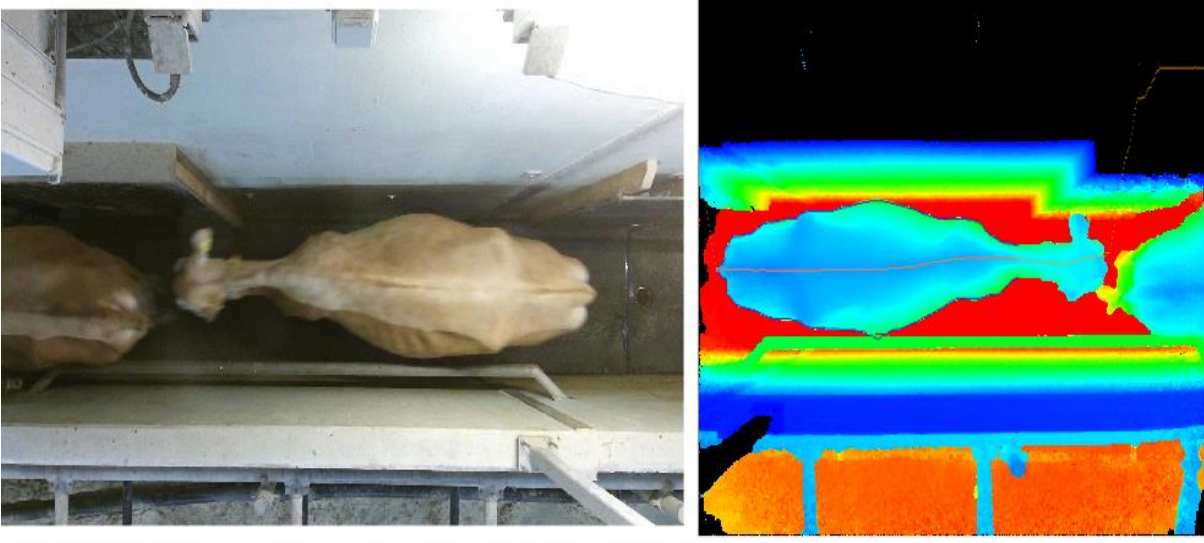


Zero calibration of floor at each feeding

System setup



Reference unit



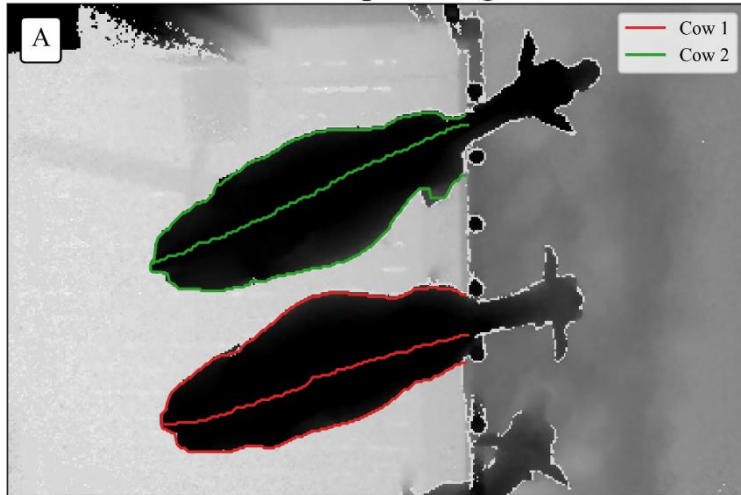
Prediction unit



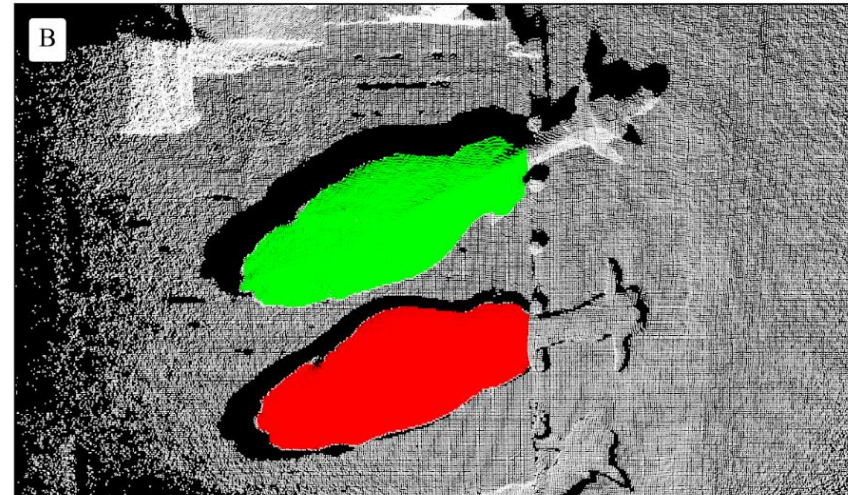
Cow detection and 3D correction



Raw Depth Image



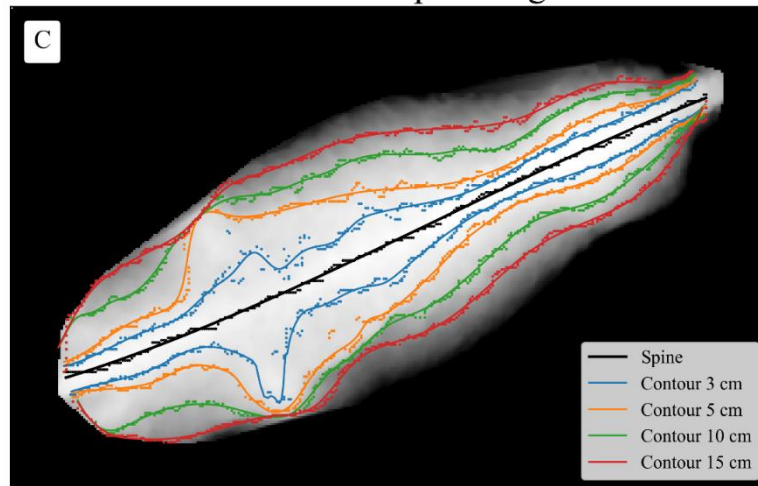
Corrected Point Cloud



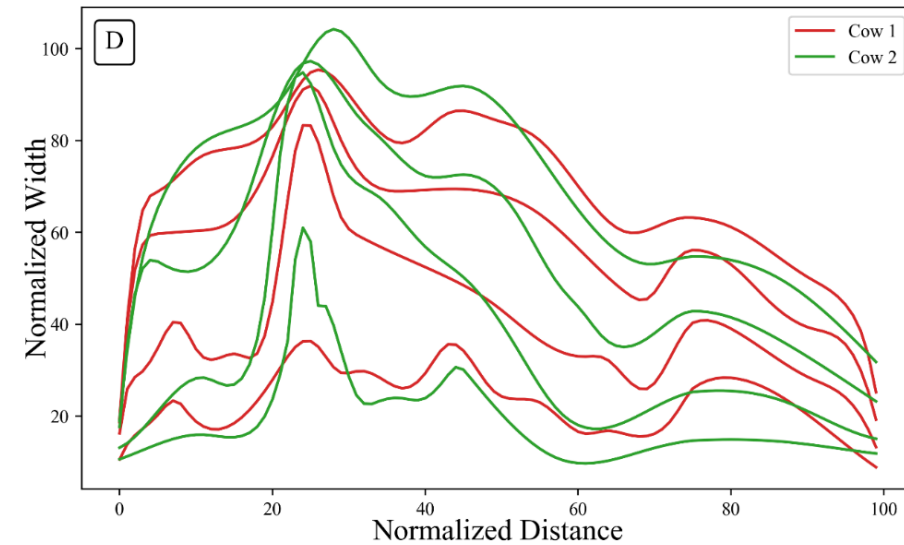
Feature extraction



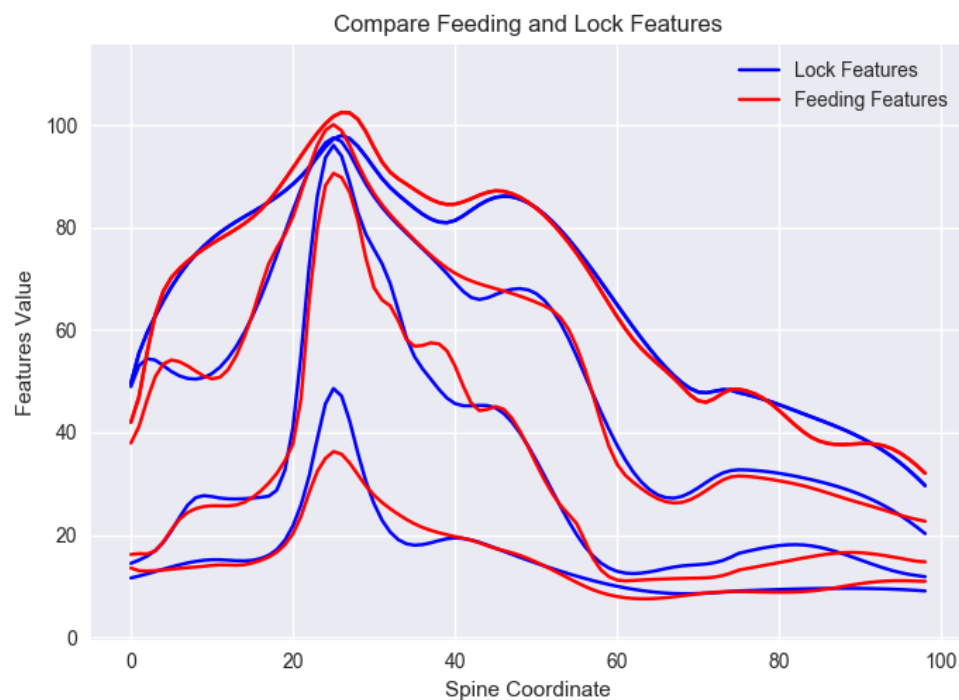
Corrected Depth Image



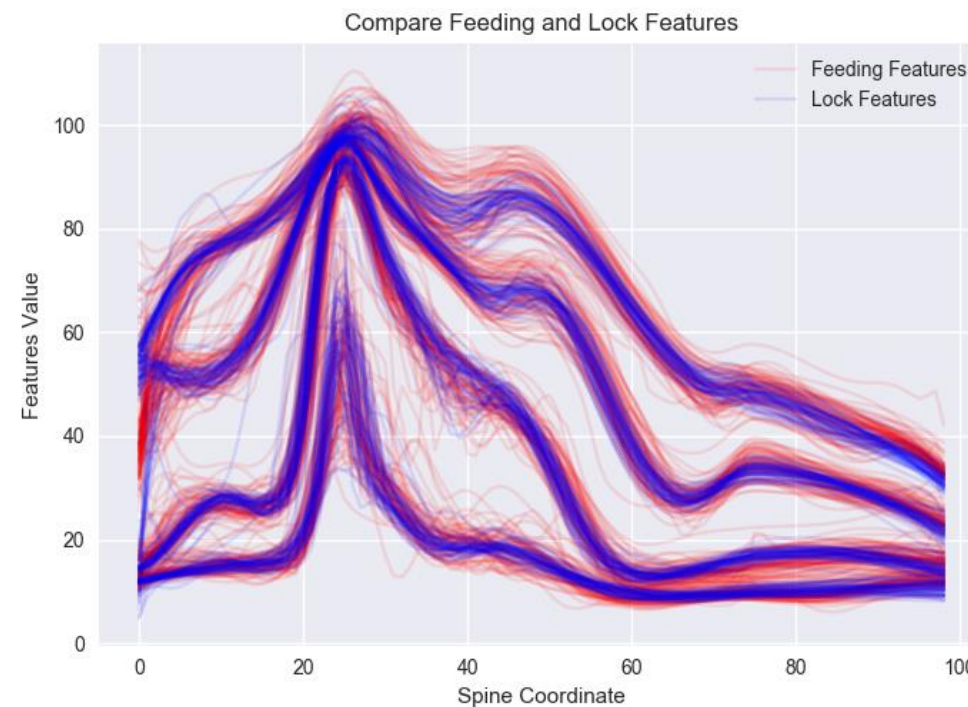
Contour Features



Match of reference and prediction



Match of a cow

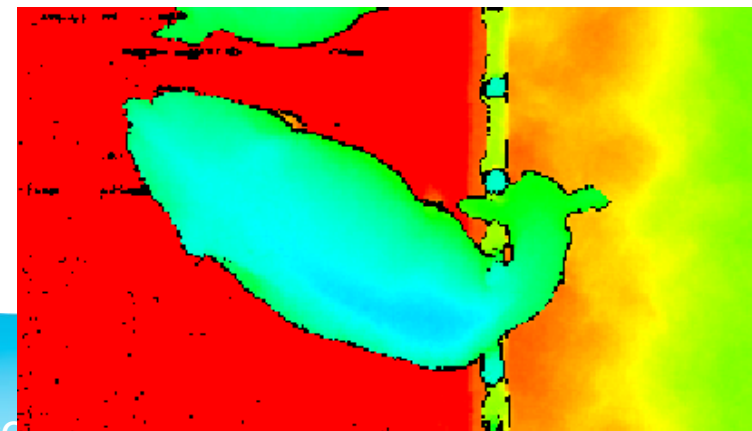


Variation for one cow - more observations

Validation of prediction accuracy



- 97 labelled Jersey cows
- 18 to 50 images from each cow in the reference unit
- 6357 manually labelled cow visits at the feeding table over 5 days

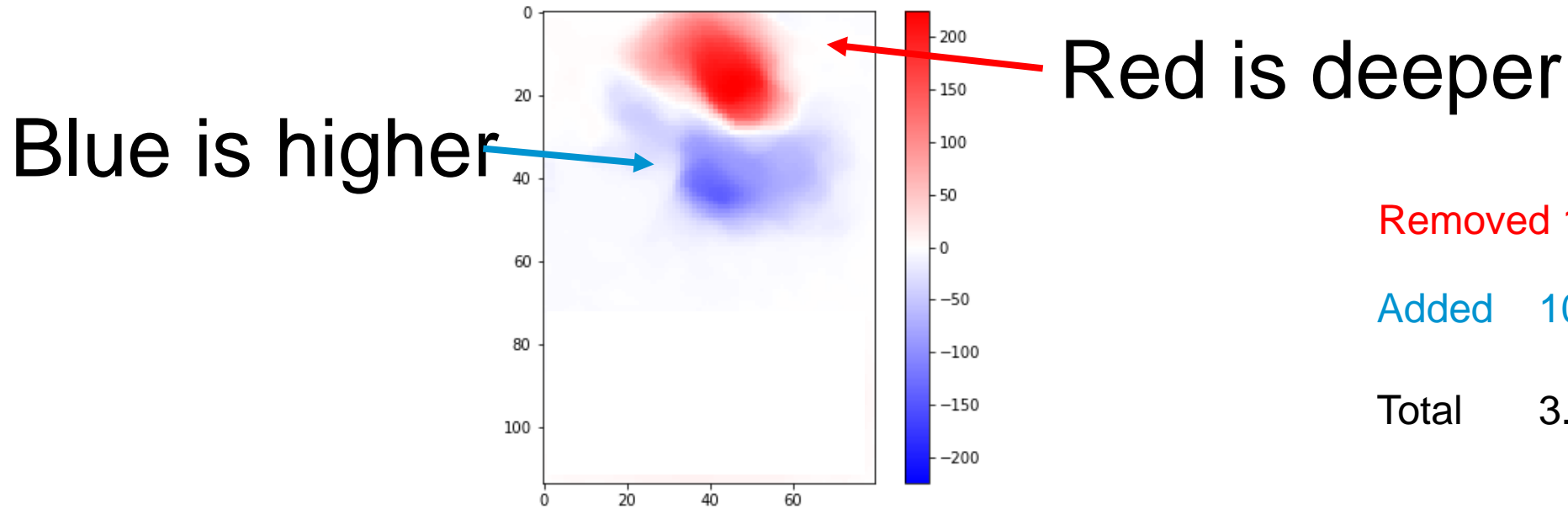


Results of validation



Sample	Count	Fraction
Correctly predicted cow-id	6022	95 %
Wrongly predicted cow-id	335	5 %

Example of feed intake from a visit

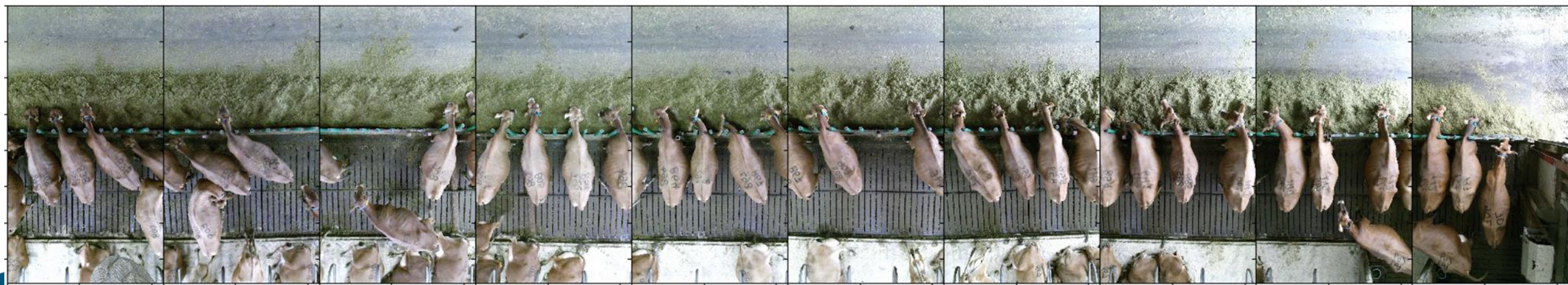


Total is difference between red and blue

Data

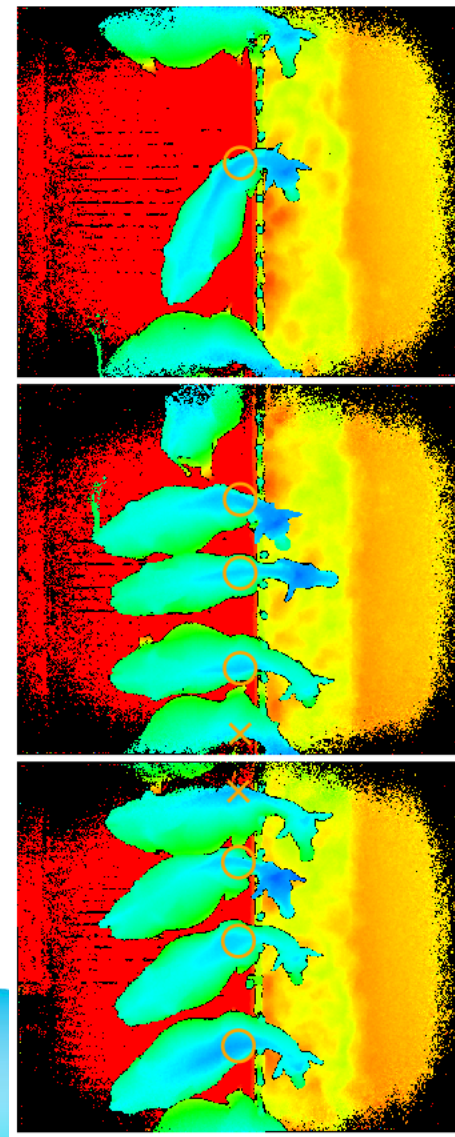


- 📍 340 Jersey cows measured for 14 weeks in 2 herds
- 📍 Weekly phenotype
- 📍 Converted to kilo of feed from density



Model

FI = μ + wil (β_1 dim) fixed reg
+ β_2 dim fixed reg
+ herd class
+ parity class
+ animal random
+ res random



Results



 Repeatability = 0.62

 Corr EKM, FI = 0.50





Next steps

- All systems up and running on 4 farms
 - + 1000 cows (350 JER, 450 HOL, 250 RDC)
- Data presentation for farmers
- Improvement of identification algorithm
- Improvement of feed intake algorithm

Sum up



- 🔍 Pooling of data and collaboration is needed to make evaluations for feed intake
- 🔍 Identification of cows based on the geometry of their back and patterns can be performed using 3D camera technology
- 🔍 Feed intake can be measured using 3D camera technology (CFIT)
- 🔍 Repeatability of 0.62 between weeks (gDMI 0.53)