

Calculation of dry matter intake and energy balance based on MIR spectral data at European level

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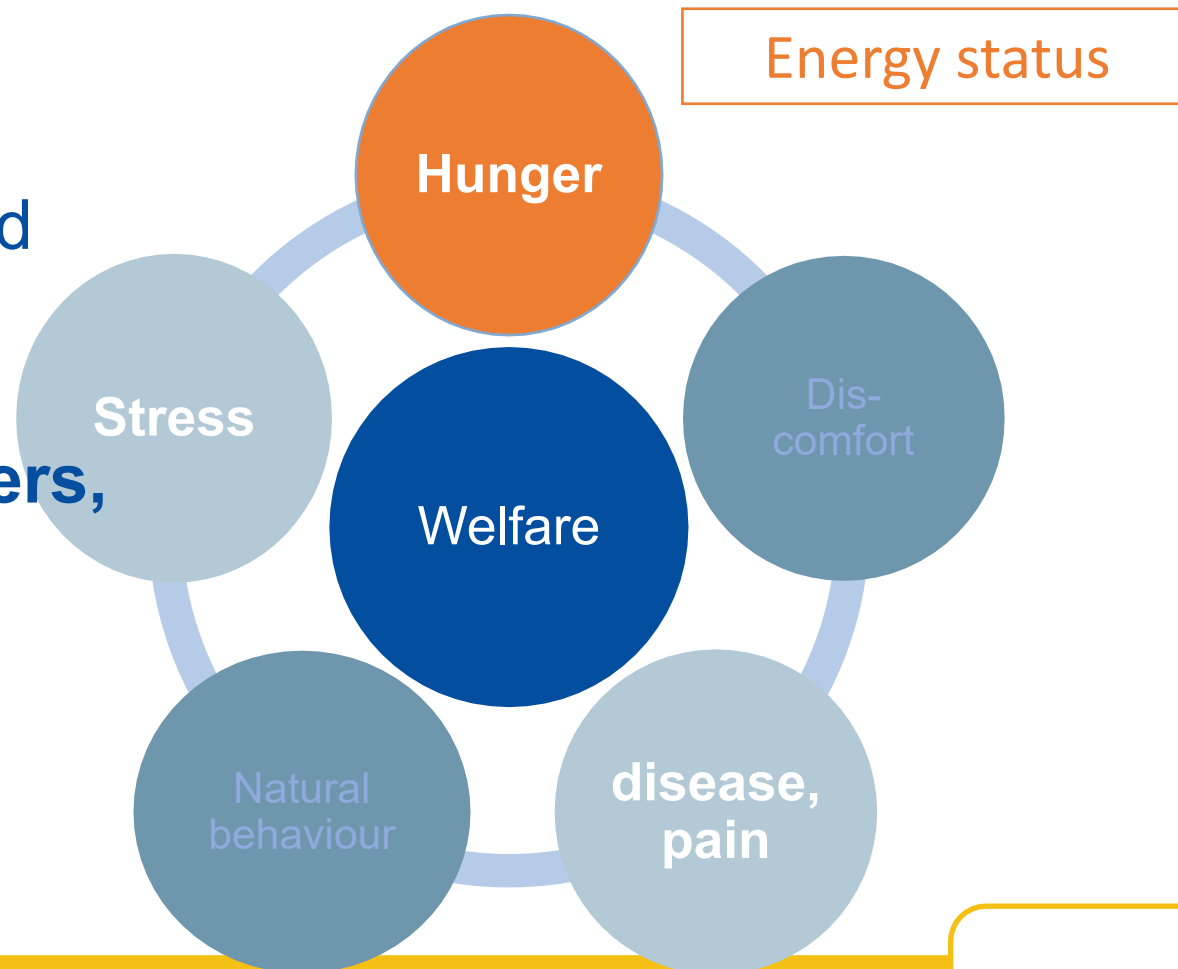
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EAAP, 31st August, Lyon

HappyMoo – objective

Use milk mid-infrared spectral analysis to **predict** welfare-related traits (molecules, phenotypes)

→ **Monitoring tools for the farmers, vets and extension workers**



Impact of Negative Energy Balance

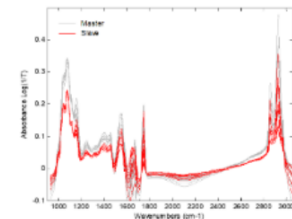
- Failure to cover energy needs can lead to metabolic disorders, stress of the immune system, breeding problems and reduced milk production



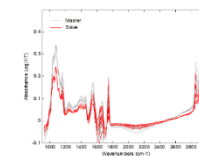
Important to have **fast, recurrent and cheap** detection of negative energy balance



Use milk mid-infrared spectral analysis to **predict** negative energy balance traits



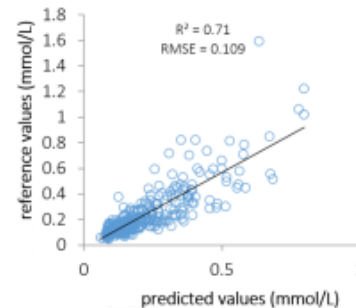
HappyMoo Energy balance models – how?



Reference values

Standardised spectra

Models



Prediction of:
Energy Balance (EB_NEL)
Dry Matter Intake (DMI)
Feed Efficiency (FE_ECM)

Reference values to calculate Energy Balance status



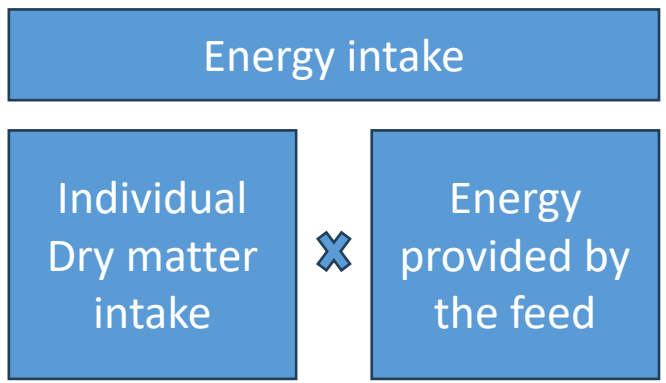

 Data collected in experimental farms in France (Trinottières) and Germany


QUALITAS[®]


LKV
 Baden-Württemberg




 GfE (2001)



Energy requirements

- For Maintenance $(0.293 * (\text{BodyWeight}^{0.75}))$
- For lactation $(0.38 * \text{Fat} + 0.21 * \text{Protein} + 0.95 + (0.1)) * \text{MilkYield}$
- For Pregnancy $(0.044 * \exp(0.0162 * \text{GestationStage})) / 0.29$

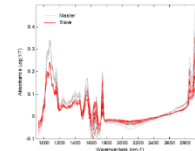
HappyMoo Energy balance models – how?

Energy Balance (EB - NEL)

Dry Matter Intake (DMI)

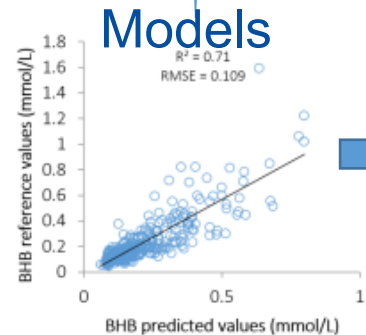
Feed Efficiency (FE_ECM)

(as a function of milk yield corrected with fat, and protein content and the DMI)



Reference values

Standardised spectra



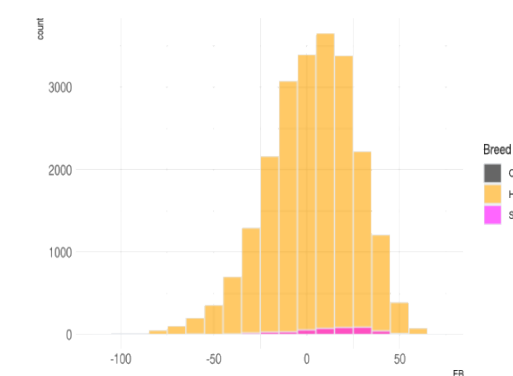
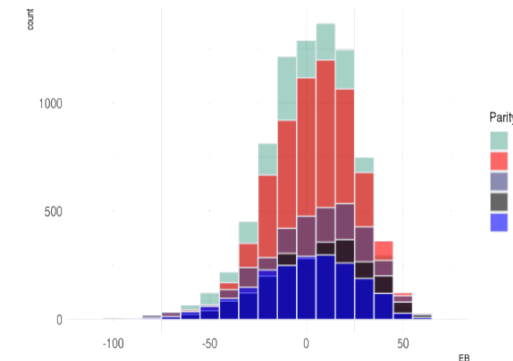
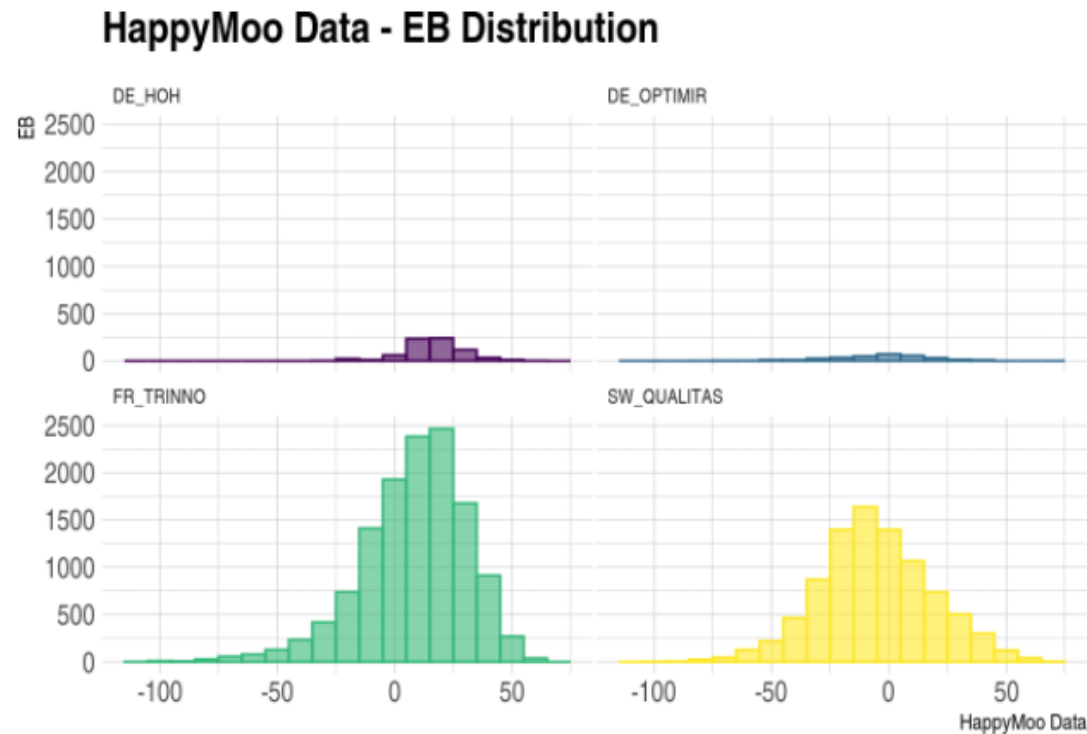
Prediction of:
Energy Balance (EB_NEL)
Dry Matter Intake (DMI)
Feed Efficiency (FE_ECM)

Descriptive statistics – Energy balance, Dry matter intake, Feed efficiency

Parameters	French	Swiss	German – OptiMIR	German – LKVBW	HappyMoo
Number of rows	15579	10917	708	2093	22813
Mean EB	7.20	-6.87	-4.68	18.32	2.88
SD EB	27.51	29.22	25.05	14.50	23.96
Mean DMI	22.35	21.18	21.98	----	22.11
SD DMI	3.86	3.98	3.65	----	3.85
Mean FE	1.50	1.49	1.68	----	1.51
SD FE	0.37	0.43	0.35	----	0.37

Outliers	
Fat Diff – 0.5 g/dl	2911
GH > 8	787
DIM ≥ 5	127

Descriptive statistics – EB parameter



Energy balance modelling

Daten	Calibration			Validation		
	N	Mean	SD	N	Mean	SD
EB	15971	2.96	23.96	6842	2.68	23.94

Material

Animal Related Data:

Parity: 1-5, >5=5
Breed: OBM, OBD, OBF
Milking Moment: AM/PM, AM, PM
Milk Yield Classes:
<25, 25-30, 30-35, 35-40, >40
Days in Milking (DIM)

Milk Related Data:

MIR-Spectral Data

MIR-Spectral Data:

Spectra + DIM Polynomial correction

Method

PLS - Partial least square regression: OSCORES, SIM

PCR – Principal component regression

CPPLS - canonical correlation analysis

GLMNET Regression

1st Validation - **Spectra Selection**

- 70% calibration model
- 30% validation model

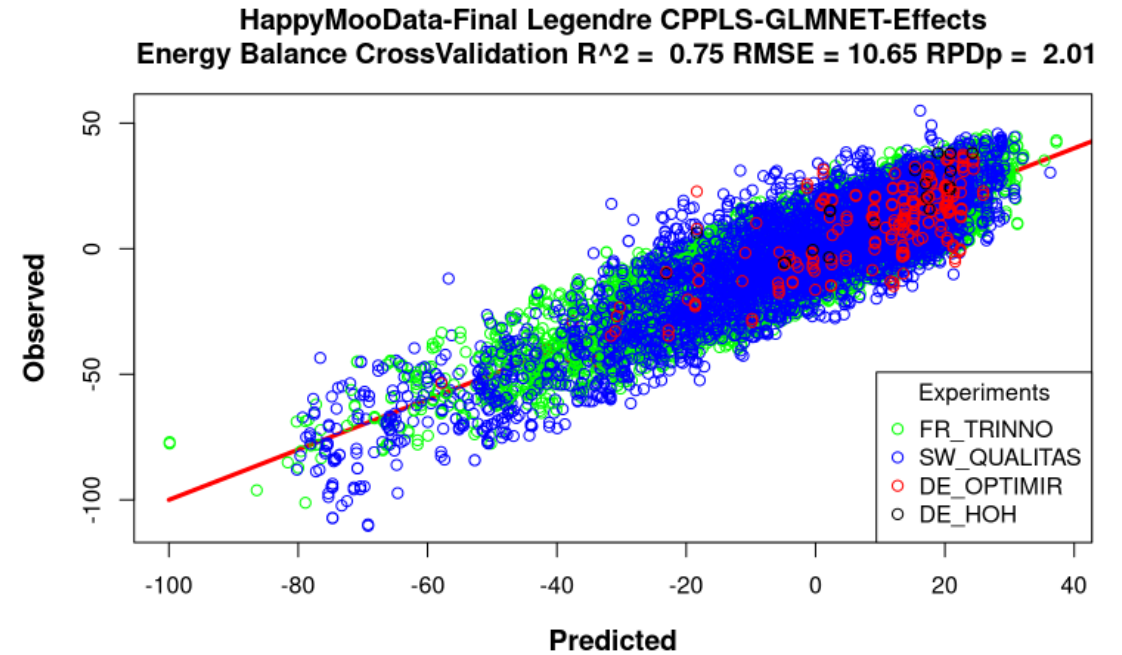
2nd Validation - **Animal Selection**

- 70% calibration model
- 30% validation model

Prediction of Energy balance EB_NEL

HappyMooData - Statistical Parameters Energy Balance equations calibration and validation

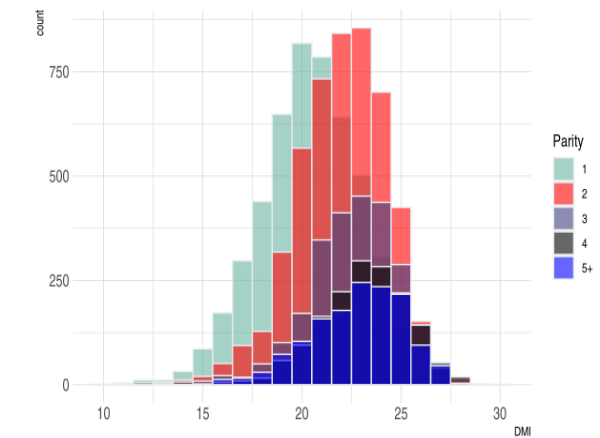
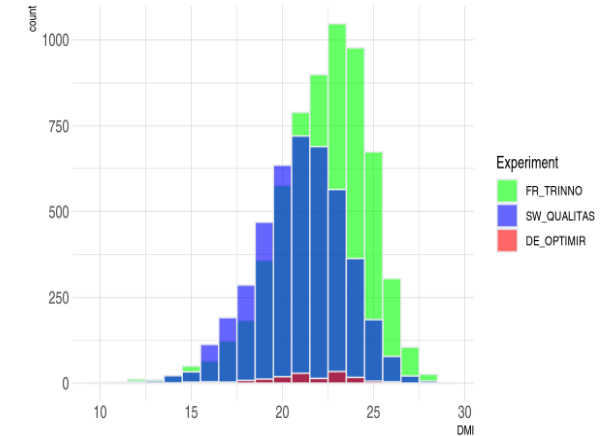
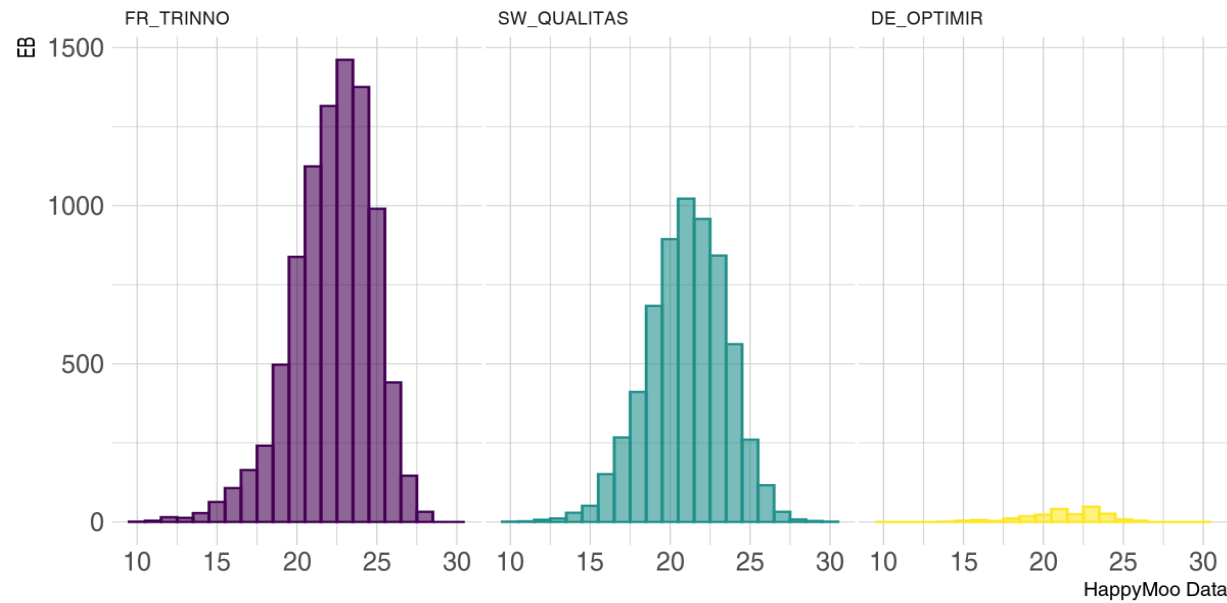
	R ² c	RMSEC	R ² p	RMSEP	RPDp
PLS_SpectraSelection	0.68	12.02	0.62	13.86	1.62
PLS_AnimalSelection	0.69	11.5	0.64	14.17	1.67
CPPLS_SpectraSelection	0.68	12.02	0.62	13.86	1.62
CPPLS_AnimalSelection	0.69	11.5	0.64	14.17	1.67
GLMNET_SpectraSelection	0.69	11.84	0.63	13.59	1.65
GLMNET_AnimalSelection	0.69	11.34	0.66	13.9	1.7
EB_GLMNET_Spectra_P_M_B	0.7	11.73	0.66	13.13	1.71
EB_GLMNET_Animal_P_M_B	0.7	11.25	0.69	13.33	1.77
LegGLMNET_SpectraSelection	0.7	11.62	0.66	13.12	1.71
LegGLMNET_AnimalSelection	0.7	11.17	0.67	13.63	1.73
Leg_CPPLS_GLMNET_Spectra	0.76	10.56	0.7	12.41	1.81
Leg_CPPLS_GLMNET_AnimalSelection	0.76	10.06	0.68	13.27	1.78



	n	Mean	SD	SEC	R2	SECV	R2cv	RPDcv	Use
EB NEL [MJ /d]	16993	0.32	21.41	10.6	0.75	10.65	0.75	2.01	0

Descriptive statistics – DMI parameter

HappyMoo Data - Dry Matter Intake Distribution

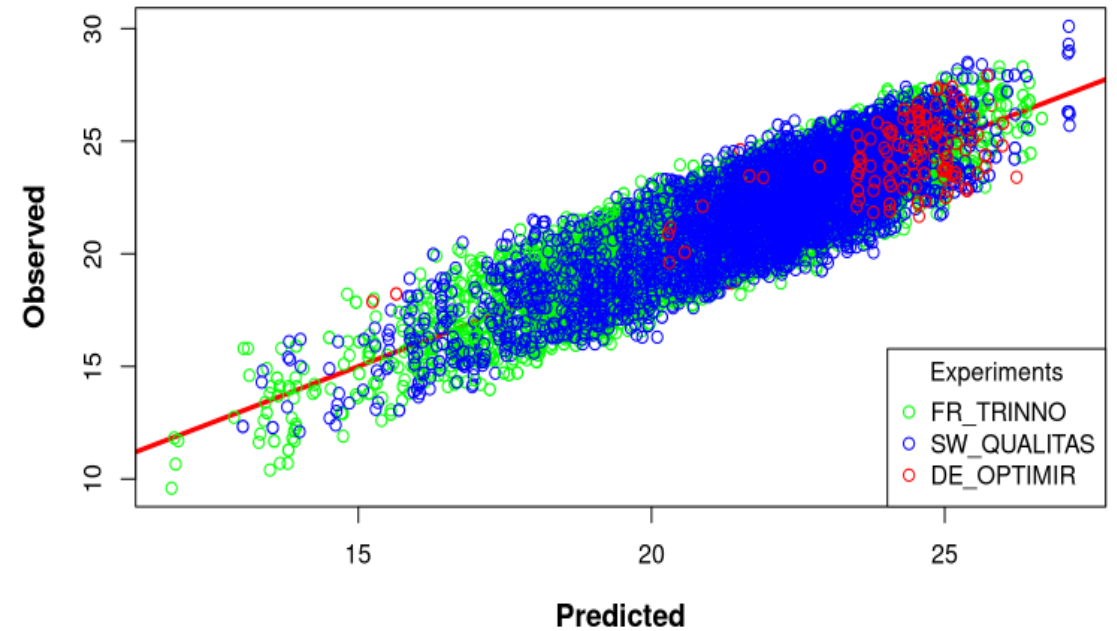


Prediction of dry matter intake (DMI)

HappyMooData - Statistical Parameters Dry Matter Intake equations calibration and validation

	R ² c	RMSEC	R ² p	RMSEP	RPDp
PLS_SpectraSelection	0.66	1.46	0.62	1.56	1.61
PLS_AnimalSelection	0.66	1.45	0.62	1.6	1.62
CPPLS_SpectraSelection	0.66	1.46	0.62	1.56	1.61
CPPLS_AnimalSelection	0.66	1.45	0.62	1.6	1.62
GLMNET_SpectraSelection	0.67	1.45	0.62	1.56	1.62
GLMNET_AnimalSelection	0.66	1.44	0.62	1.6	1.63
EB_GLMNET_Spectra_P_M_B	0.69	1.41	0.65	1.5	1.69
EB_GLMNET_Animal_P_M_B	0.68	1.41	0.66	1.51	1.72
LegGLMNET_SpectraSelection	0.67	1.45	0.62	1.55	1.63
LegGLMNET_AnimalSelection	0.67	1.43	0.63	1.59	1.64
Leg_CPPLS_GLMNET_Spectra	0.75	1.23	0.62	1.54	1.63
Leg_CPPLS_GLMNET_AnimalSelection	0.7	1.36	0.62	1.6	1.62

HappyMooData-Final Legendre CPPLS-GLMNET-Effects
Dry Matter Intake CrossValidation R² = 0.7 RMSE = 1.38 RPDp = 1.83



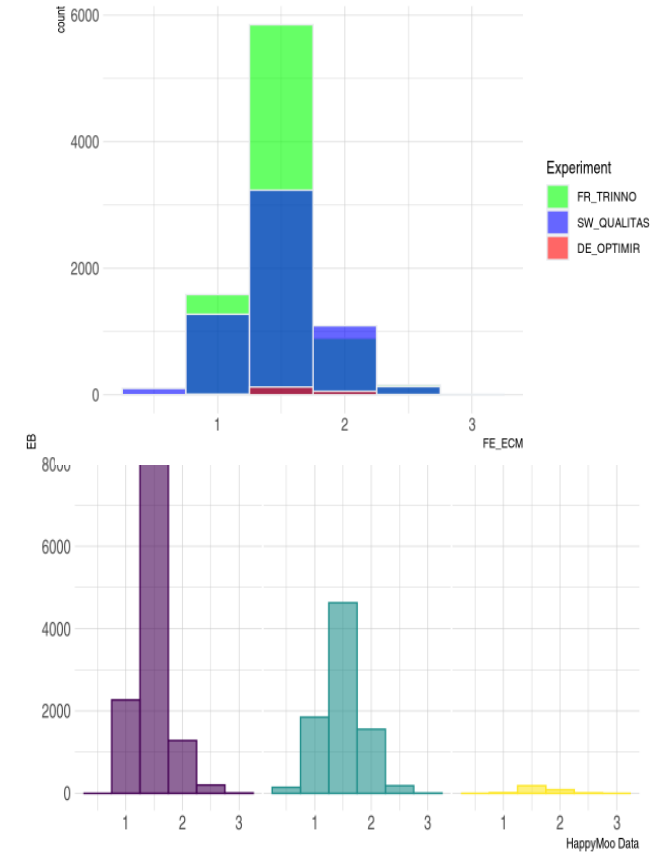
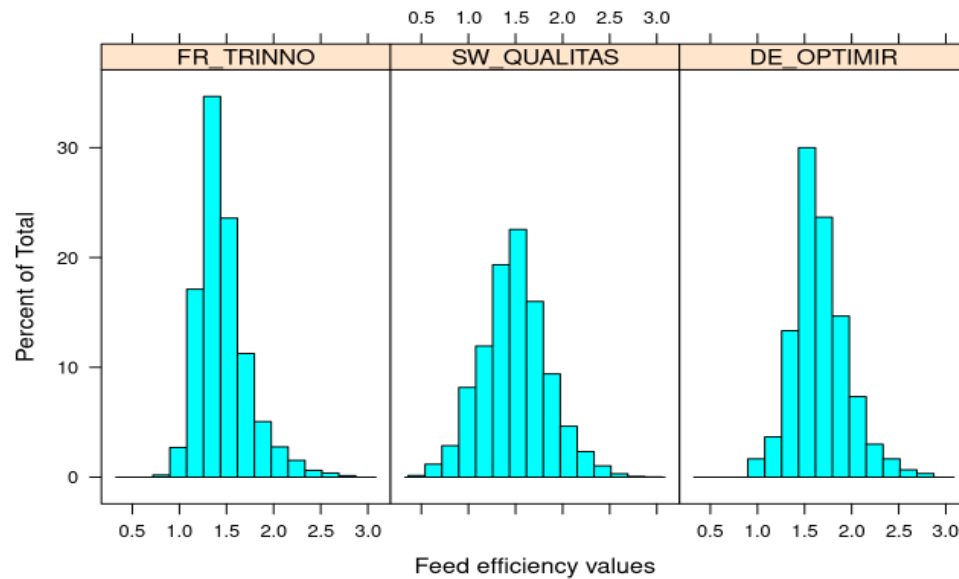
	n	Mean	SD	SEC	R2	SECv	R2cv	RPDcv	Use
DMI [kg]	14804	21.77	2.52	1.36	0.71	1.38	0.7	1.83	0

Descriptive statistics – FE parameter

Definition of Feed Efficiency

$$FE = ECM / DMI$$

$$ECM = MY24H * ((1.05 + 0.38 * FC + 0.21 * PC24H) / 3.28)$$

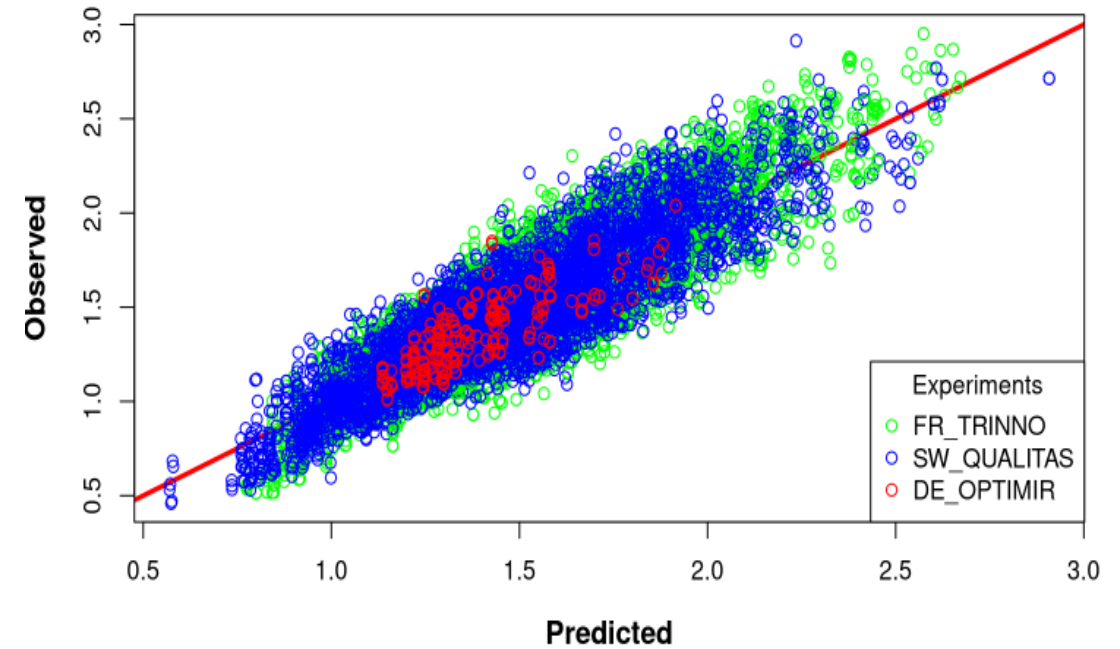


Prediction feed efficiency (FE_ECM)

HappyMooData - Statistical Parameters Feed efficiency equations calibration and validation

	R ² c	RMSEC	R ² p	RMSEP	RPDp
PLS_SpectraSelection	0.66	0.18	0.66	0.18	1.72
PLS_AnimalSelection	0.65	0.18	0.67	0.18	1.74
CPPLS_SpectraSelection	0.66	0.18	0.66	0.18	1.72
CPPLS_AnimalSelection	0.65	0.18	0.67	0.18	1.74
GLMNET_SpectraSelection	0.67	0.18	0.67	0.18	1.74
GLMNET_AnimalSelection	0.66	0.18	0.68	0.18	1.77
EB_GLMNET_Spectra_P_M_B	0.71	0.17	0.7	0.17	1.84
EB_GLMNET_Animal_P_M_B	0.66	0.18	0.68	0.18	1.77
LegGLMNET_SpectraSelection	0.68	0.17	0.67	0.17	1.75
LegGLMNET_AnimalSelection	0.68	0.17	0.67	0.18	1.72
Leg_CPPLS_GLMNET_Spectra	0.72	0.16	0.71	0.16	1.87
Leg_CPPLS_GLMNET_AnimalSelection	0.71	0.16	0.7	0.17	1.79

HappyMooData-Final Legendre CPPLS-GLMNET-Effects
 Feed efficiency CrossValidation R² = 0.76 RMSE = 0.15 RPDp = 2.03



	n	Mean	SD	SEC	R2	SECV	R2cv	RPDcv	Use
FE [ECM /DM kg]	20747	1.48	0.31	0.15	0.76	0.15	0.76	2.03	0

Conclusion

Energy balance

Best model with CPPLS, GLMNET based on Legendre polynom and fix effects: breed, parity and milk yield classes.
RPD of 2 → a good prediction ability

Dry matter intake

DMI is interesting to know if cow has enough to eat.
RPD of 1.83 → model need some improvement

Feed efficiency

Feed efficiency is more linked to the performance of the cow to convert feed.
RPD of 2 → good prediction ability

Thank you for your attention!

