



APPROXIMATING GREENHOUSE GAS EMISSIONS FOR A FARM NETWORK USING READILY AVAILABLE DATA

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1 Introduction

IP-SUISSE

- Swiss farming association IP-SUISSE (20,000 Members (out of 50,000 Swiss Farmers; 40%))
- 10,000 Members produce for the **IP-SUISSE Label**
- Label producers must satisfy **additional standards**, e.g. ones that promote biodiversity
- If the **additional standards** are met, farmers receive **higher prices** for their products

1 Introduction

Current project (IP-Suisse/ Agroscope)

- ✓ Extension of Label standards with climate-change mitigation measures (e.g., solar panels, heat recovery, biogas, covering of slurry pit, etc.)
- ✓ Aim of this collaboration: 10% reduction in GHG emissions for IP-SUISSE Label producers in 2022 compared to 2016

⇒ Requires estimate of total GHG emissions of all IP-SUISSE Label farms in 2016.

Problem: Full LCA based on SALCA/SimaPro too time-consuming

Solution: Apply simple method to approximate GHG emissions at farm level (using readily available input data at farm level)

2 Materials and Methods

Data

- (1) **Computed** global warming potential (GWP) for **33 IP-Suisse pilot farms** (covering typical production systems in Switzerland) using SALCA/ SimaPro
- (2) **Farm Structure Survey**: Land area/ Livestock numbers
→ **Preselection** of relevant variables taking into account the main drivers for methane CH₄, fossil carbon dioxide CO₂, nitrous oxides N₂O.

Land	Livestock
Utilised agricultural area (UAA)	Total livestock [LU]
Open arable land (OA)	Total cattle [LU]
Permanent grassland	Pigs [LU]
Biodiversity-promotion areas	Poultry [LU]



2 Materials and Methods

After eliminating insignificant variables by stepwise model selection using AIC, the following approach was discovered:

Model: Quadratic polynomial regression (POLMOD)

$$y = a + \beta_1 * x_1 + \beta_2 * (x_1)^2 + \beta_3 * x_2 + \beta_4 * (x_2)^2$$

Explaining variables

x_1 = Livestock density (TLD) [LU/ha UAA]

x_2 = Proportion of open arable land (POA) []

Target (dependent) variable

y: GHG emissions per UAA [t CO₂eq /ha]

3 Results

Estimation of coefficients for polynomial regression (POLMOD)

$$y = 10.56 - 2.73 * TLD + 1.48 * (TLD)^2 - 23.61 * POA + 17.91 * (POA)^2$$

y = GHG emissions per UAA [t CO₂eq /ha]

TLD = Livestock density (TLD) [LU/ha UAA]

POA = Proportion of open arable land (POA) []

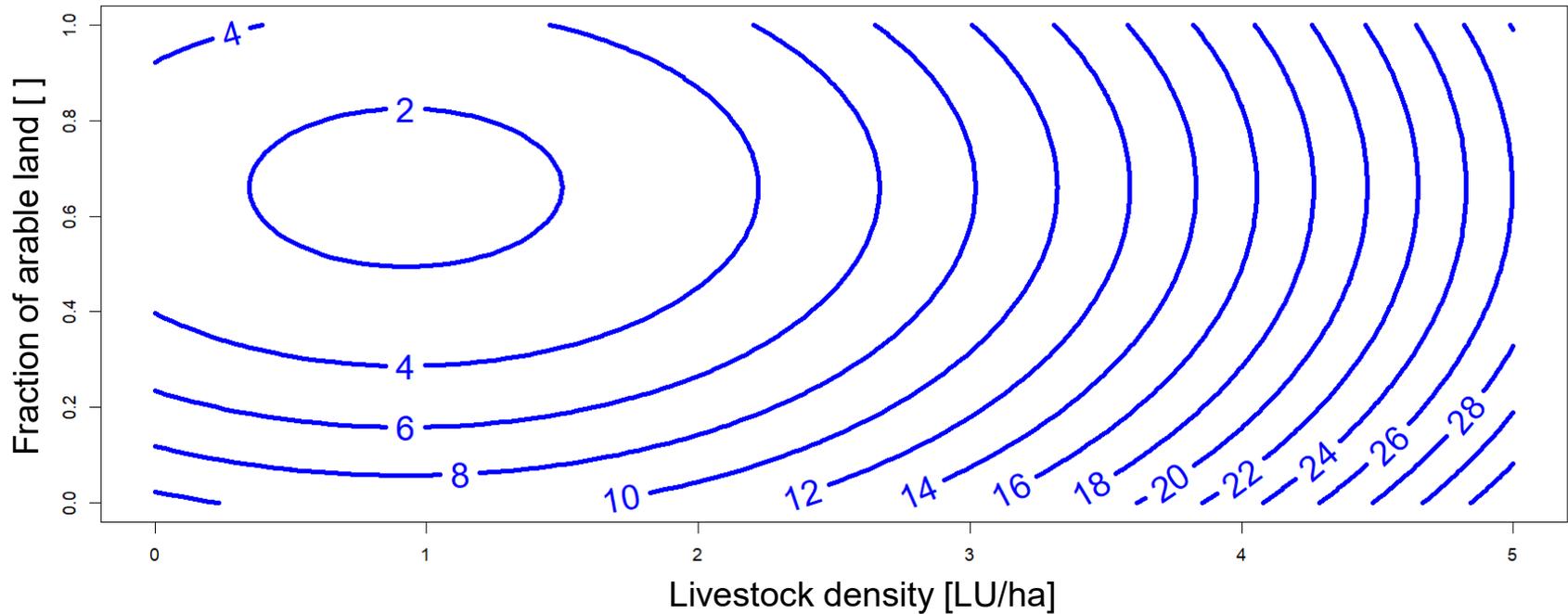
The following comments are worth noting:

- (i) Small sample → avoid overfitting → use limited number of explaining variables
- (ii) Model assumptions are well satisfied (e.g. residuals are randomly distributed)
- (iii) Application of robust methods did not improve the results



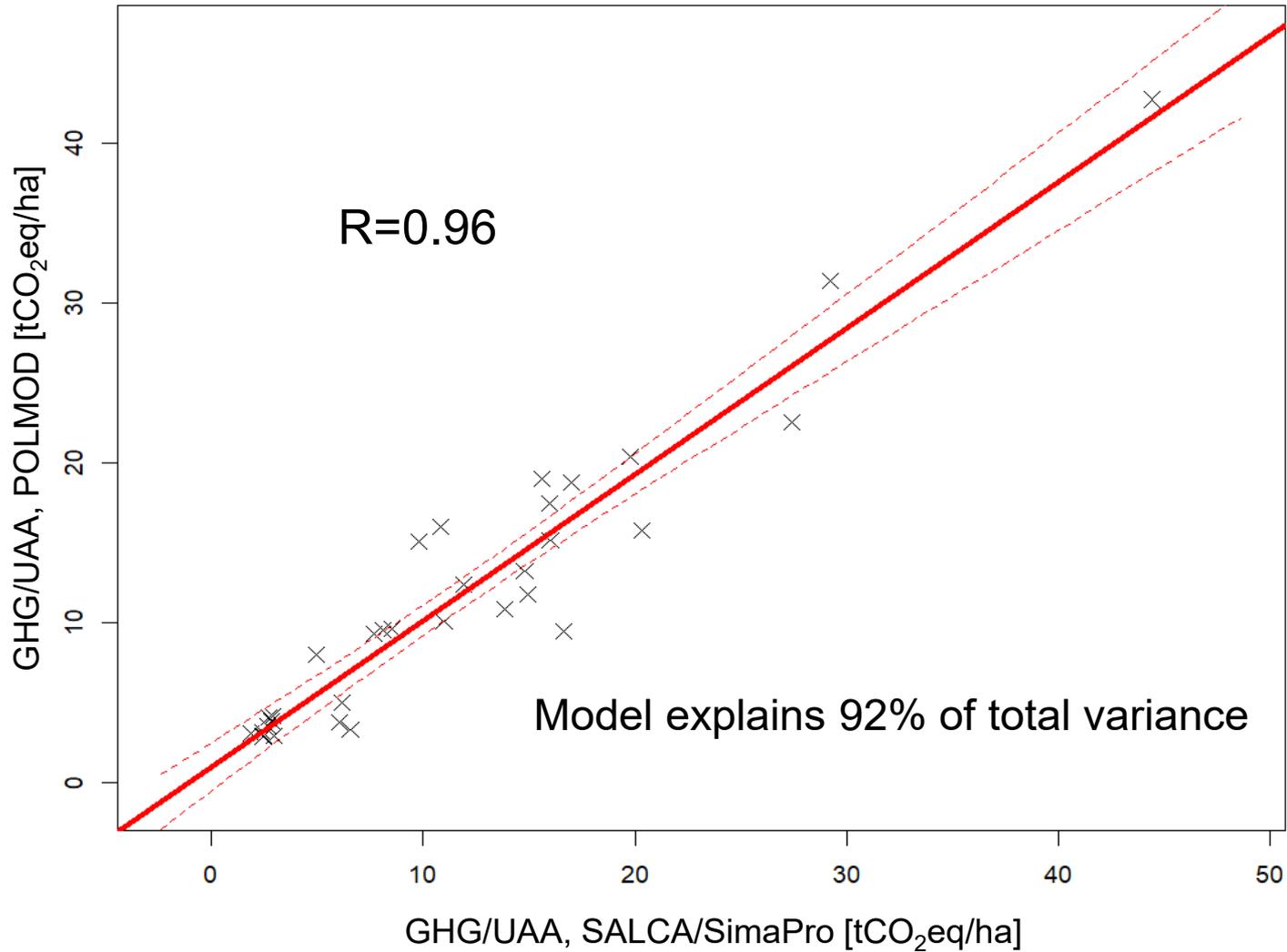
3 Results

Estimated GHG Emissions [t CO₂eq] per ha
Method: POLMOD





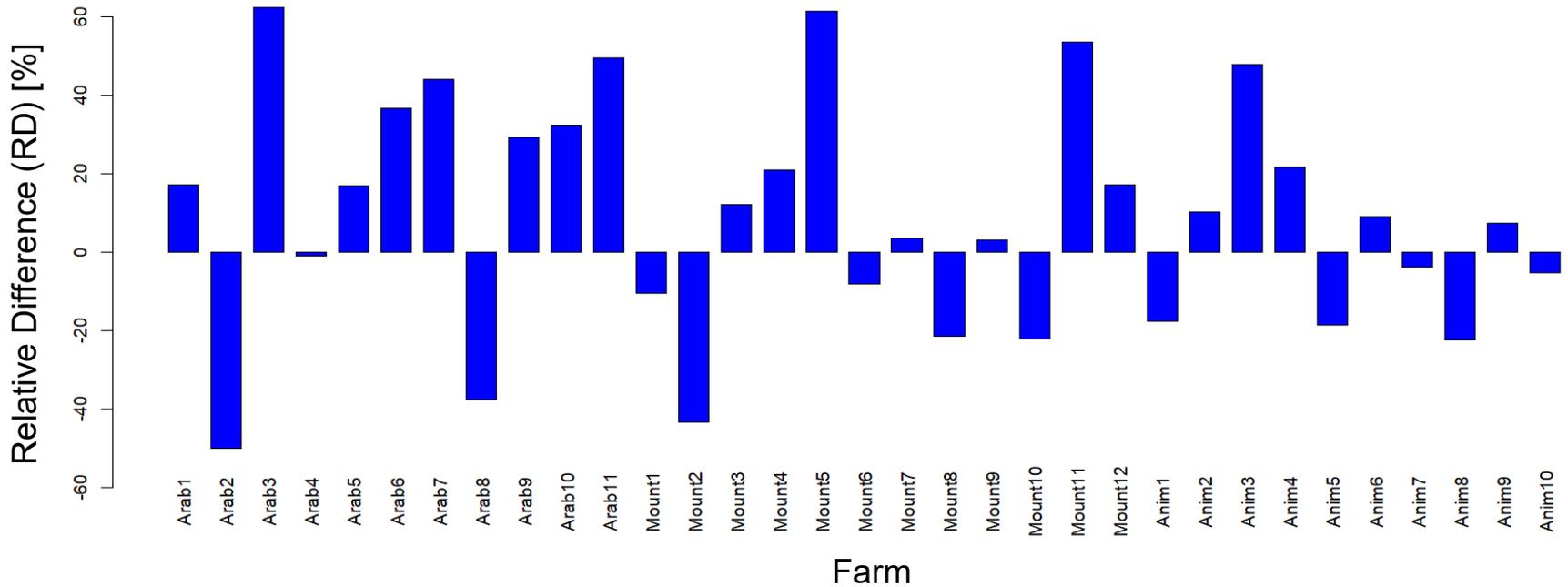
3 Results





3 Results

Relative GHG difference: POLMOD minus SALCA



Mean = 9% , std. dev. = 30%

16 out of 33 farms $RD < 20\%$, 4 out of 33 farms $RD > 50\%$

3 Results

Extrapolation to Swiss agricultural sector (2016)

Application of POLMOD to all Swiss farms

=> total GHG emissions from the Swiss agricultural sector

GHG Emissions CH = 6.93 +/- 1.24 million t CO₂eq

Plausibility check/ Verification by independent source
(Bretscher et al., 2020)

GHG Emission CH = ~ 7.5 million t CO₂eq



Good agreement



4 Discussion/ Conclusions

- POLMOD allows the computation of **GHG emissions** at farm level based on **two readily available variables**
- The POLMOD method is well suited to estimating GHG emissions at farm level (**92% of the variance** can be explained)
- POLMOD allows **extrapolation** to all IP-Suisse Label farms or to the entire Swiss agricultural sector
- **Percentage deviations** from SALCA/Simapro computed GHG estimates **may exceed 50%** for certain farms
- **Small size** of sample may be critical
- More **independent evaluations** (based on larger samples) are required



Thank you very much for your attention

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