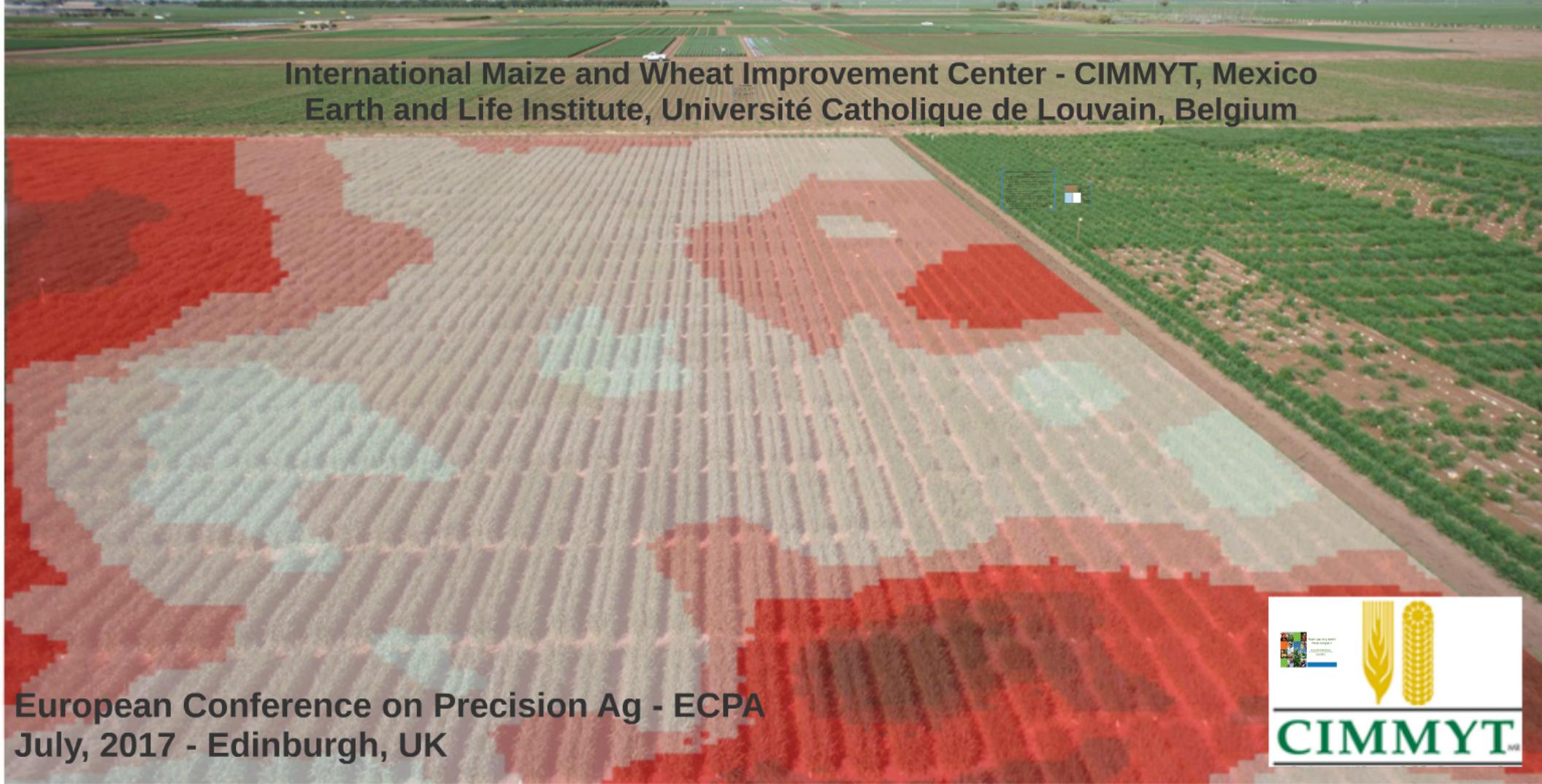


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... and many others

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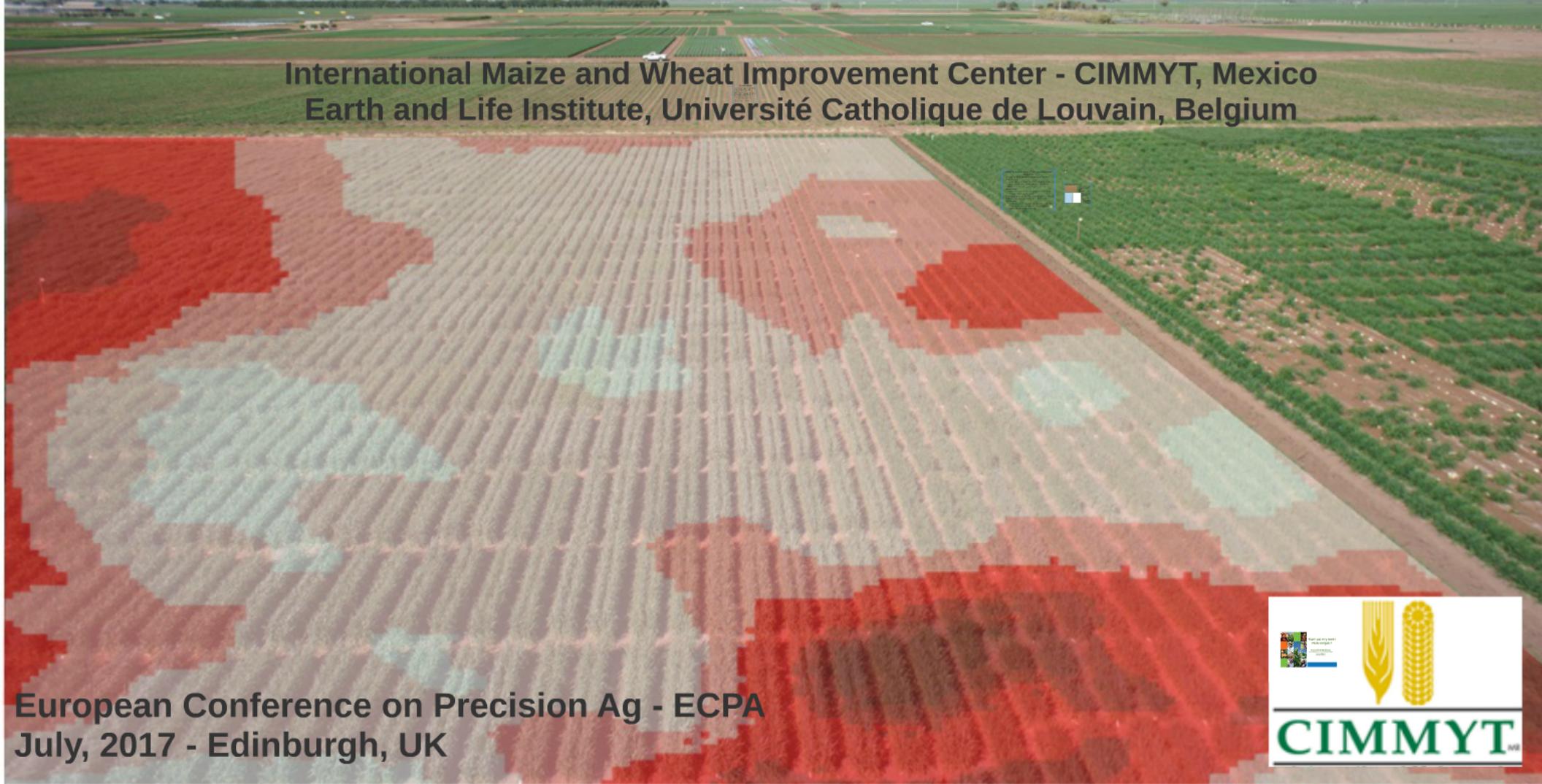


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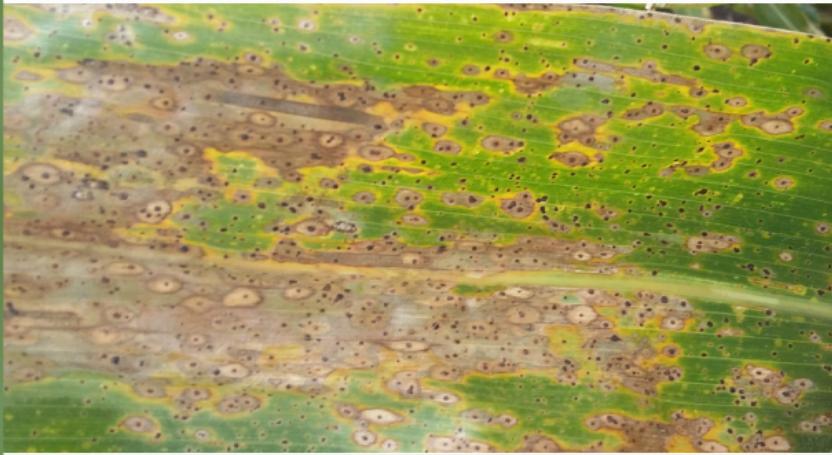
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# Maize Tar Spot Complex (TSC) importance and motivation

- One of the most important diseases of maize in tropical areas
  - grain loss may reach 50-75% - depending on susceptibility and environmental conditions
- Selection for TSC resistance starts at early generations
  - more advanced lines are screened in multiple locations for several years.
- Phenotyping maize for TSC resistance is mostly conducted by visual observations by breeders/pathologists
  - A 1 to 5 scale is used for the disease evaluation
  - Depends on personal experience and results may vary due subjectiveness

# TSC development...



- dark spots on the leaf surface (both lower and upper leaves)
- usually 2-3 weeks before flowering
- chlorotic circles due to the development of the second parasite
  - chlorotic circles take over the entire leaves, becoming necrotic - affecting the photosynthetic activity - reducing the grain yields.
    - High-throughput phenotyping through UAV may facilitate TSC resistance evaluation

# CIMMYT'S Experimental Station - Agua Fria, Mexico - April, 2016



Square lattice design - 25 genotypes in two main blocks

714 - fungicide

715 - without fungicide

three randomized replications.

plot - four rows, 4.5m x 0.75m



fixed effect

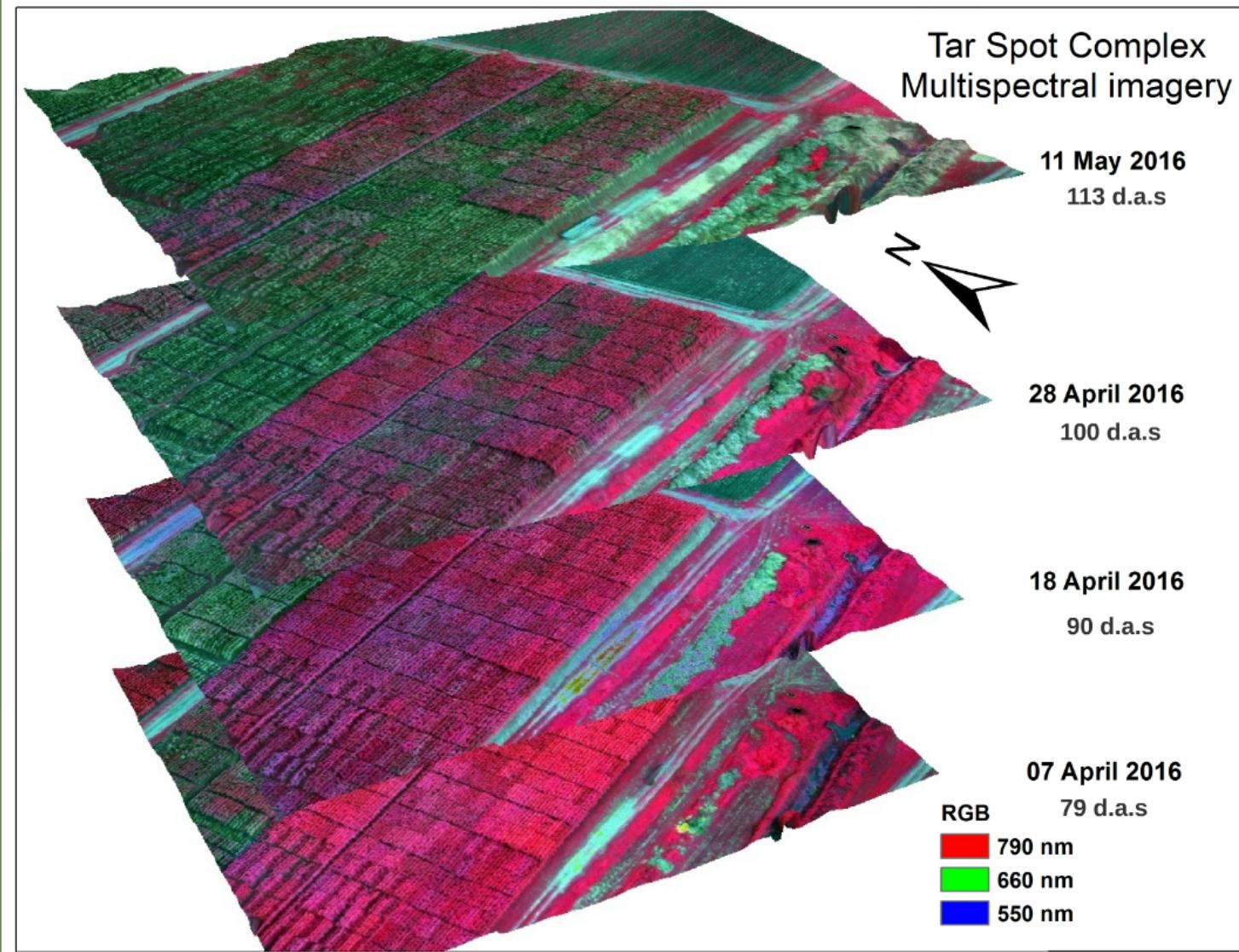


random effect

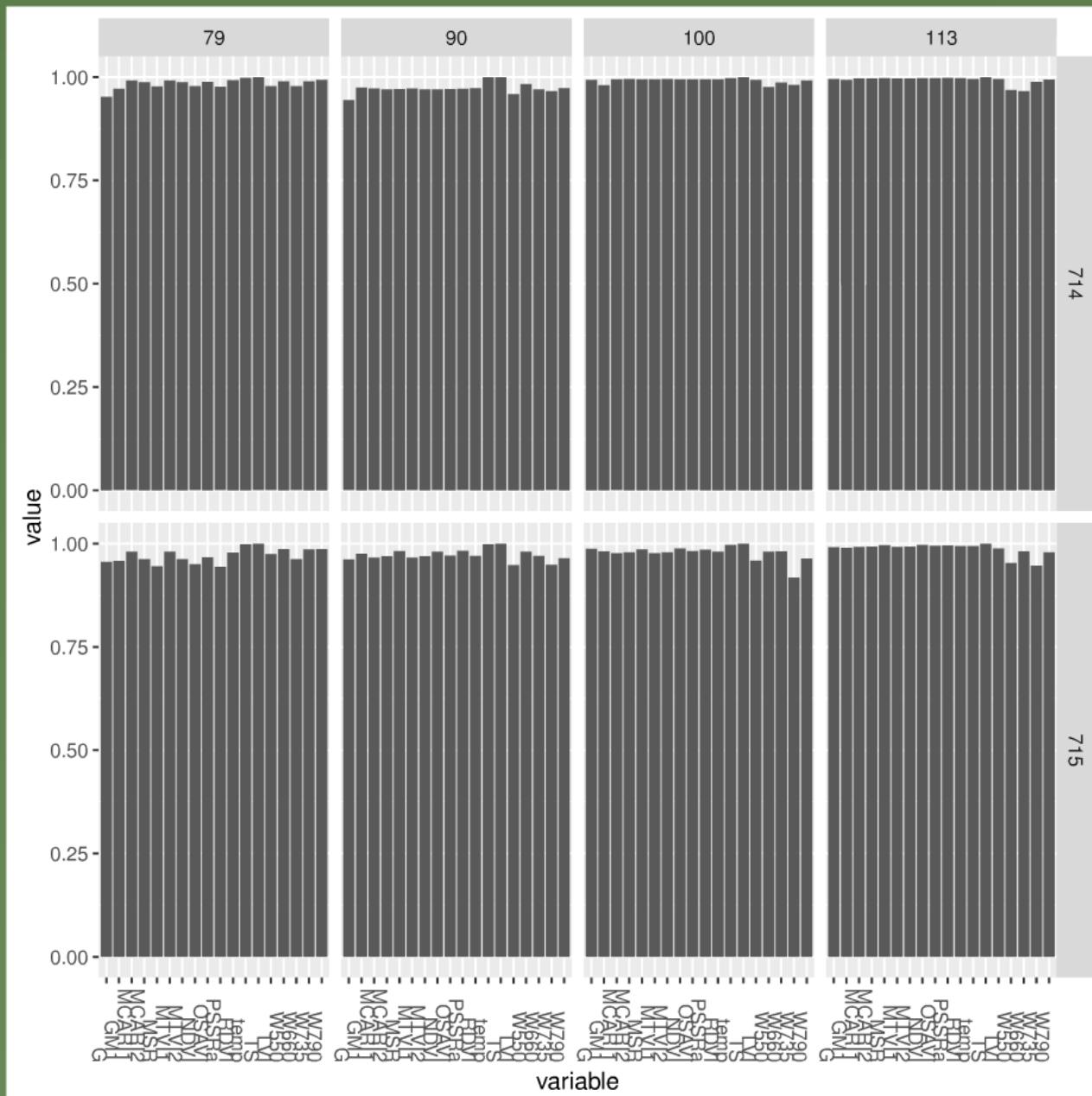
RML model



- MultiSpectral
  - (550, 660, 735, 790 nm) - 6 cm/px
- Thermal images – 12 cm/px



# Results



**Table 1 – Phenotypic correlation between yield, image data and visual scoring (*n* = 75)**

|                | 714    |       |        |       | 715    |         |        |        |
|----------------|--------|-------|--------|-------|--------|---------|--------|--------|
|                | 79     | 90    | 100    | 113   | 79     | 90      | 100    | 113    |
| <b>NDVI</b>    | 0.31*  | 0.31* | 0.47*  | 0.54* | 0.42*  | 0.51*   | 0.63*  | 0.57*  |
| <b>RDVI</b>    | 0.28*  | 0.36* | 0.51*  | 0.55* | 0.55*  | 0.59*   | 0.67*  | 0.59*  |
| <b>OSAVI</b>   | 0.31*  | 0.36* | 0.50*  | 0.55* | 0.55*  | 0.58*   | 0.66*  | 0.59*  |
| <b>MSR</b>     | 0.30*  | 0.31* | 0.49*  | 0.56* | 0.42*  | 0.52*   | 0.68*  | 0.58*  |
| <b>MCARI1</b>  | 0.22** | 0.35* | 0.47*  | 0.53* | 0.51*  | 0.60*   | 0.65*  | 0.58*  |
| <b>MCARI2</b>  | 0.29*  | 0.35* | 0.48*  | 0.54* | 0.57*  | 0.60*   | 0.66*  | 0.58*  |
| <b>TVI</b>     | -0.09  | 0.27* | 0.25** | 0.42* | 0.15   | 0.58*   | 0.57*  | 0.53*  |
| <b>GM1</b>     | 0.11   | 0.36* | 0.39*  | 0.48* | 0.23** | 0.49*   | 0.64*  | 0.60*  |
| <b>PSSRa</b>   | 0.29*  | 0.31* | 0.50*  | 0.58* | 0.42*  | 0.53*   | 0.72*  | 0.61*  |
| <b>G</b>       | -0.02  | 0.05  | 0.12   | 0.34* | 0.12   | 0.38*   | 0.52*  | 0.51*  |
| <b>Thermal</b> | -0.12  | 0.35* | -0.36* | 0.19  | -0.13  | 0.01    | -0.46* | 0.00   |
| <b>TS</b>      | 0.08   | -0.01 | -0.05  | -0.07 | -0.30* | -0.24** | -0.39* | -0.43* |

**Table 2 – Genetic correlation between yield, image data and visual scoring (*n* = 25)**

|                | 714      |         |         |         | 715    |         |        |        |
|----------------|----------|---------|---------|---------|--------|---------|--------|--------|
|                | 79       | 90      | 100     | 113     | 79     | 90      | 100    | 113    |
| <b>NDVI</b>    | 0.29     | 0.21    | 0.40**  | 0.45**  | 0.09   | 0.26    | 0.58*  | 0.54*  |
| <b>RDVI</b>    | 0.56*    | 0.38*** | 0.44**  | 0.45**  | 0.53*  | 0.48*   | 0.60*  | 0.55*  |
| <b>OSAVI</b>   | 0.50*    | 0.34*** | 0.43**  | 0.45**  | 0.52*  | 0.45**  | 0.60*  | 0.55*  |
| <b>MSR</b>     | 0.26     | 0.22    | 0.39*** | 0.46**  | 0.09   | 0.30    | 0.62*  | 0.54*  |
| <b>MCARI1</b>  | 0.56*    | 0.46**  | 0.43**  | 0.43**  | 0.51*  | 0.52*   | 0.60*  | 0.54*  |
| <b>MCARI2</b>  | 0.55*    | 0.39**  | 0.43**  | 0.43**  | 0.57*  | 0.51*   | 0.61*  | 0.55*  |
| <b>TVI</b>     | 0.36***  | 0.44**  | 0.36*** | 0.39**  | 0.42** | 0.54*   | 0.57*  | 0.54*  |
| <b>GM1</b>     | 0.00     | 0.19    | 0.23    | 0.38*** | -0.25  | 0.13    | 0.52*  | 0.53*  |
| <b>PSSRa</b>   | 0.24     | 0.22    | 0.38*** | 0.46**  | 0.09   | 0.32    | 0.65*  | 0.55*  |
| <b>G</b>       | 0.24     | 0.24    | 0.32    | 0.37*** | 0.26   | 0.38*** | 0.60*  | 0.54*  |
| <b>Thermal</b> | -0.36*** | 0.56*   | -0.50*  | -0.27   | -0.17  | -0.1    | -0.57* | -0.33* |
| <b>TS</b>      | -0.19    | -0.1    | -0.11   | -0.14   | -0.61* | -0.51*  | -0.62* | -0.60* |

Where: NDVI, RDVI, OSAVI, MSR, MCARI1 and MCARI2 – structural indices; TVI, GM1, PSSRa - chlorophyll indices; G – RGB ratio; thermal – canopy temperature; TS – visual disease scoring; 714 – fungicide treatment; 715 – Non-fungicide treatment. 79, 90, 100 and 113 days after sowing. \* Coefficients of correlation statistically significant at 1% probability; \*\* coefficients of correlation statistically significant at 5% probability; \*\*\* coefficients of correlation statistically significant at 10% probability.

# Conclusions

- VIs (structural and chlorophyll) proved to be a promising tool for the estimation of yield losses caused by TSC and offering new opportunities for high throughput phenotyping for resistance of maize to this highly important foliar disease;
- High  $r^2$  ( $<0.8$ ) between visual scores and wavelengths
- 2017 maize growing cycle was carried out to account for possible environmental variability and to ensure the repeatability of the methodology.
  - Room for improvements = different approaches for image data extraction
    - multivariate analysis are the next steps - squeeze data!
- Challenges
  - Early detection.. before eyes can see it!
  - Upscale it!



**Thank you very much !  
Muito obrigado !**

***Francelino Rodrigues***  
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