

# Reconstructing the root system development of spring barley using minirhizotron data

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## Background

Increasing computer power favors the use of **detailed mesoscopic models** to predict root water uptake at field scale. These models explicitly consider the **3-D root architecture development** of a plant. However, a lack of high-quality data to **calibrate and validate** these models remains, especially for plants in undisturbed, layered soils. Combination of root architecture simulations and minirhizotron measurements at several depths may offer a way out...

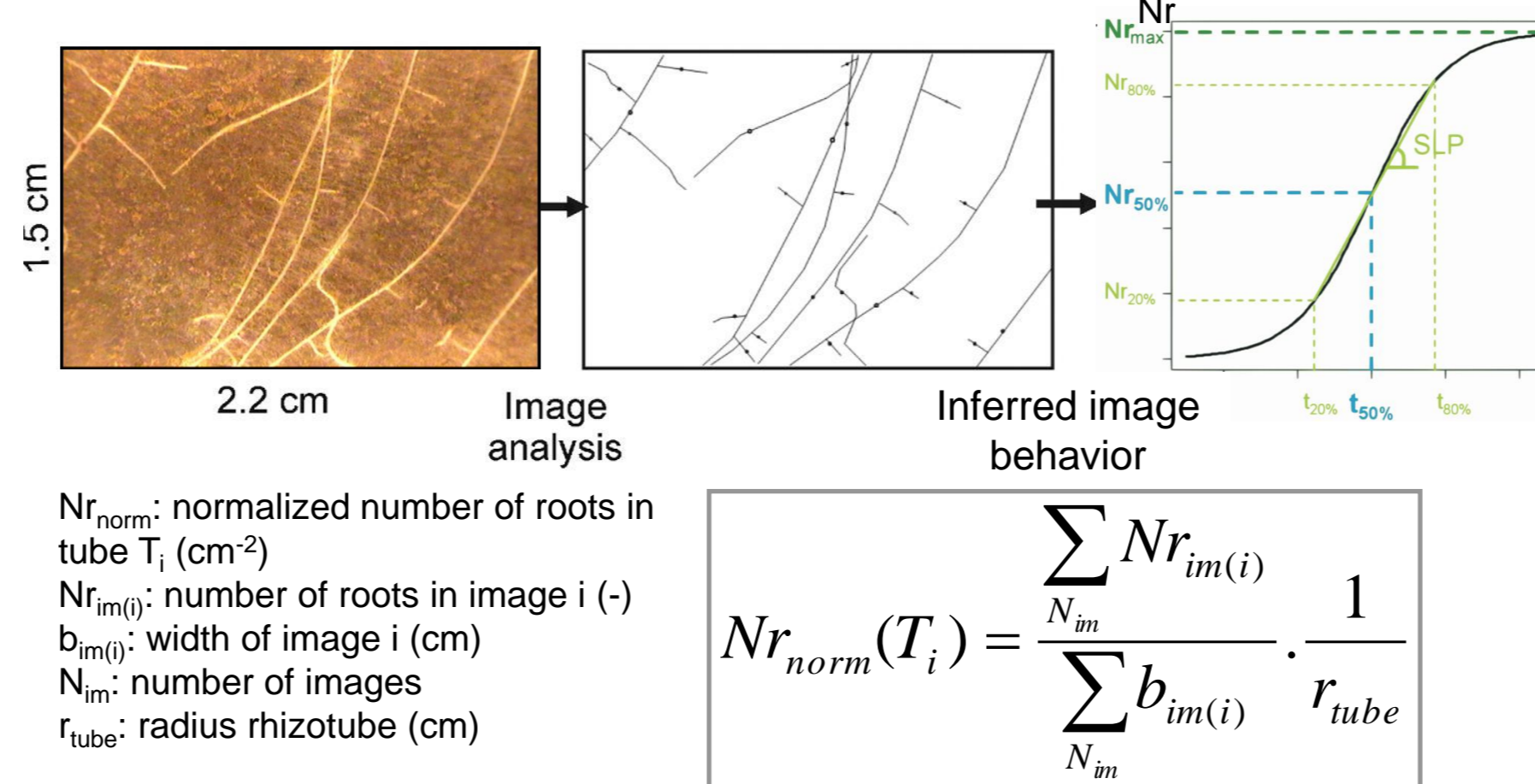
Objectives:

- explore the value of the **information** that can be extracted from minirhizotron images in horizontally installed rhizotubes at four depths;
- assess the **sensitivity** of predictions of variables that can be observed in minirhizotrons by the RS model RootTyp to changes of its model parameters;
- estimate **RS model parameters** and investigate their information content concerning the root architecture

## Methodology

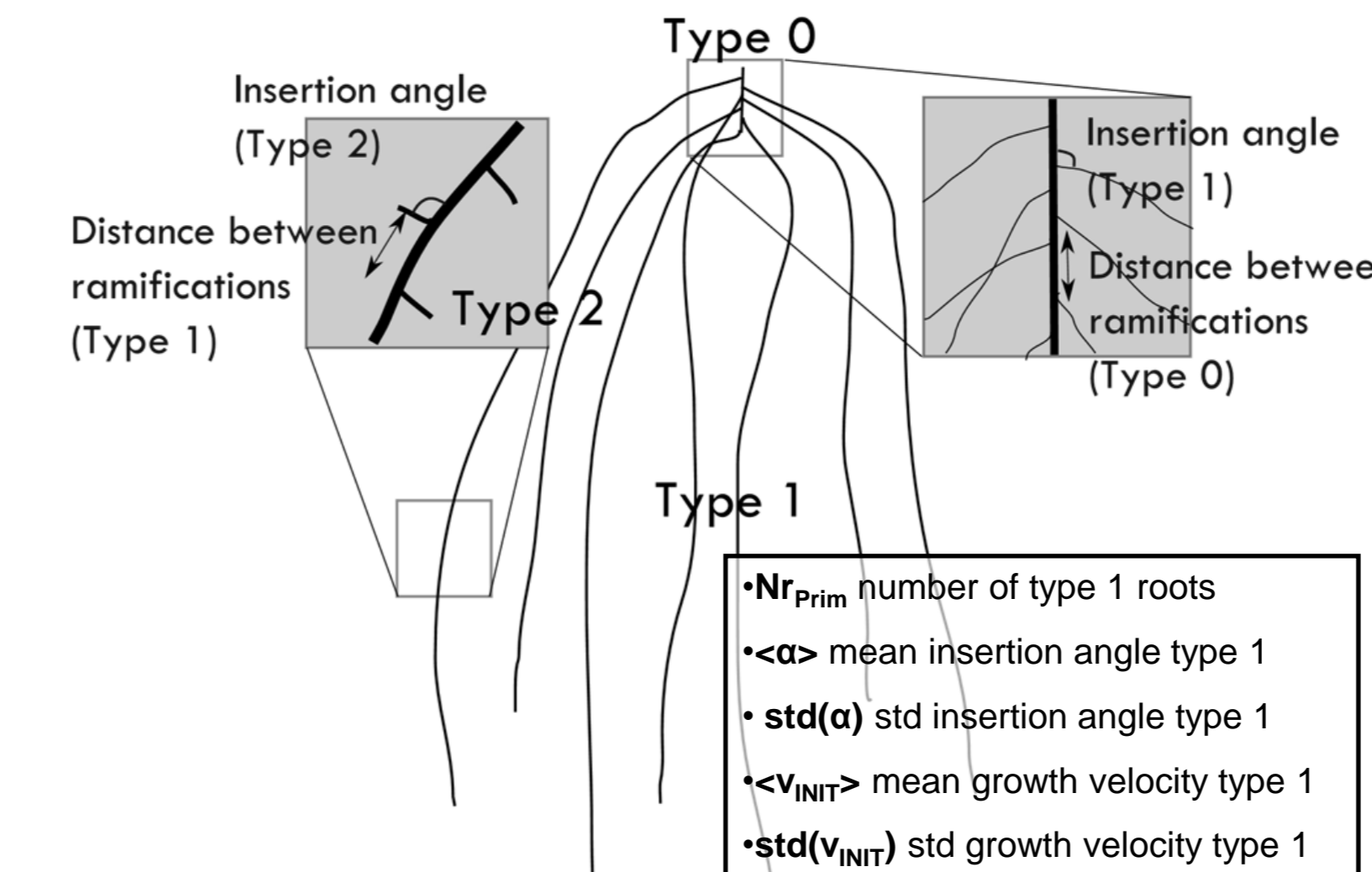
### Minirhizotron measurements

- two undisturbed soil monoliths S1 & S2  
h=150 cm,  $\phi=116$  cm
- orthic Luvisol:  $A_p$  (0-40 cm),  $B_t$  (41-70 cm),  $B_{v1}$  (71-100 cm) and  $B_{v2}$  (>100 cm)
- 4 rhizotubes T1-T4 per monolith:  
-22.5, -47.5, -72.5 and -122.5 cm depth
- 5x2 images per tube



- For each rhizotube: characteristics of root breakthrough curve (rBTC)  $\rightarrow$   $Nr_{Max}$ ,  $t_{50\%}$ , SLP

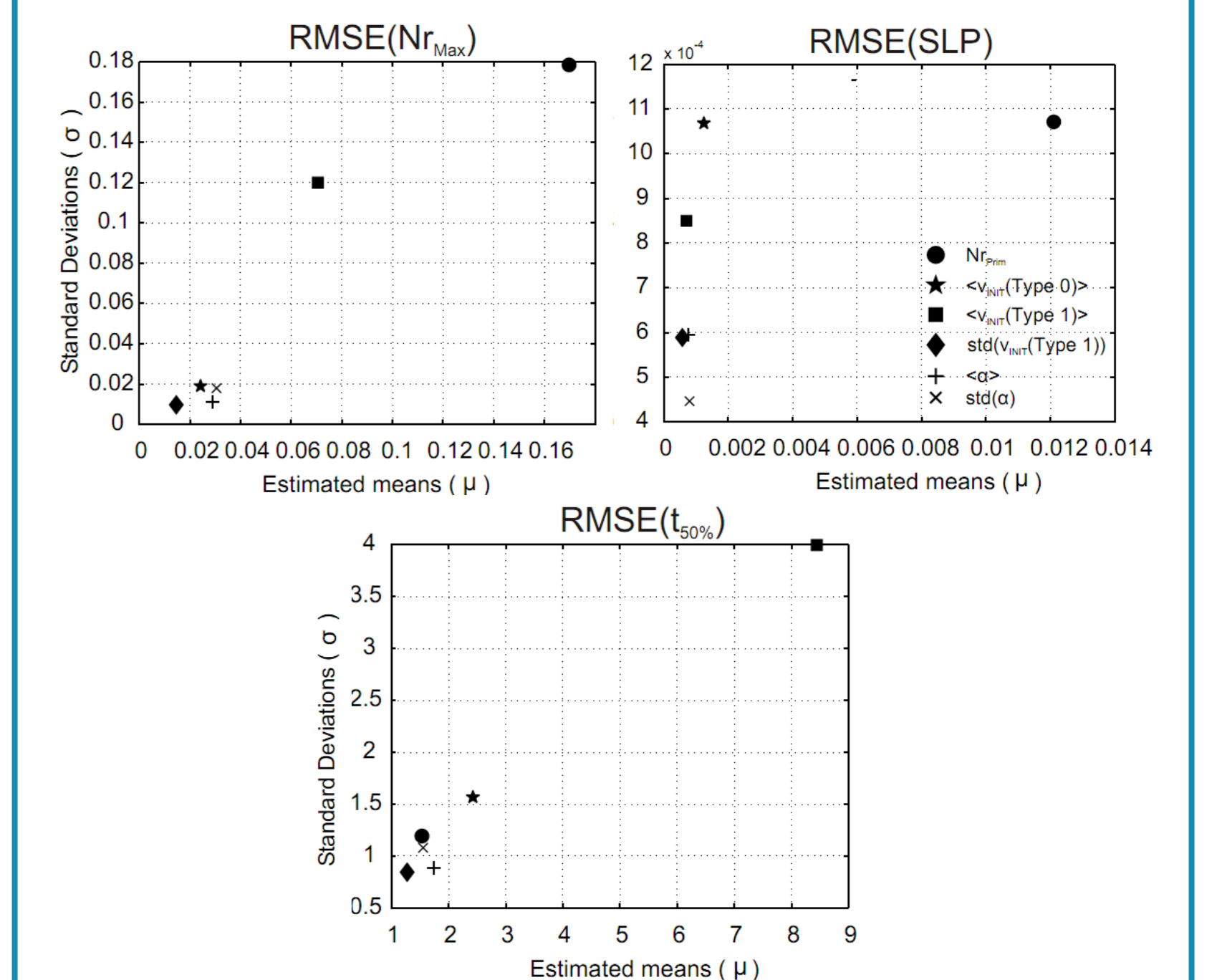
### Root architecture simulations



- Simple model:** 5 'basic' parameters  
 $Nr_{Prim}$ ,  $\langle \alpha \rangle$ ,  $std(\alpha)$ ,  $\langle v_{INIT} \rangle$ ,  $std(v_{INIT})$
- Simulations:** 200 plants, 7 rows 24 cm apart, 2 cm between plants/row ( $\approx$  real lysimeter)
- Model optimization:** comparison of minirhizotron data and simulations using rBTC characteristics  $Nr_{Max}$ ,  $t_{50\%}$ , SLP and overall course of the rBTC

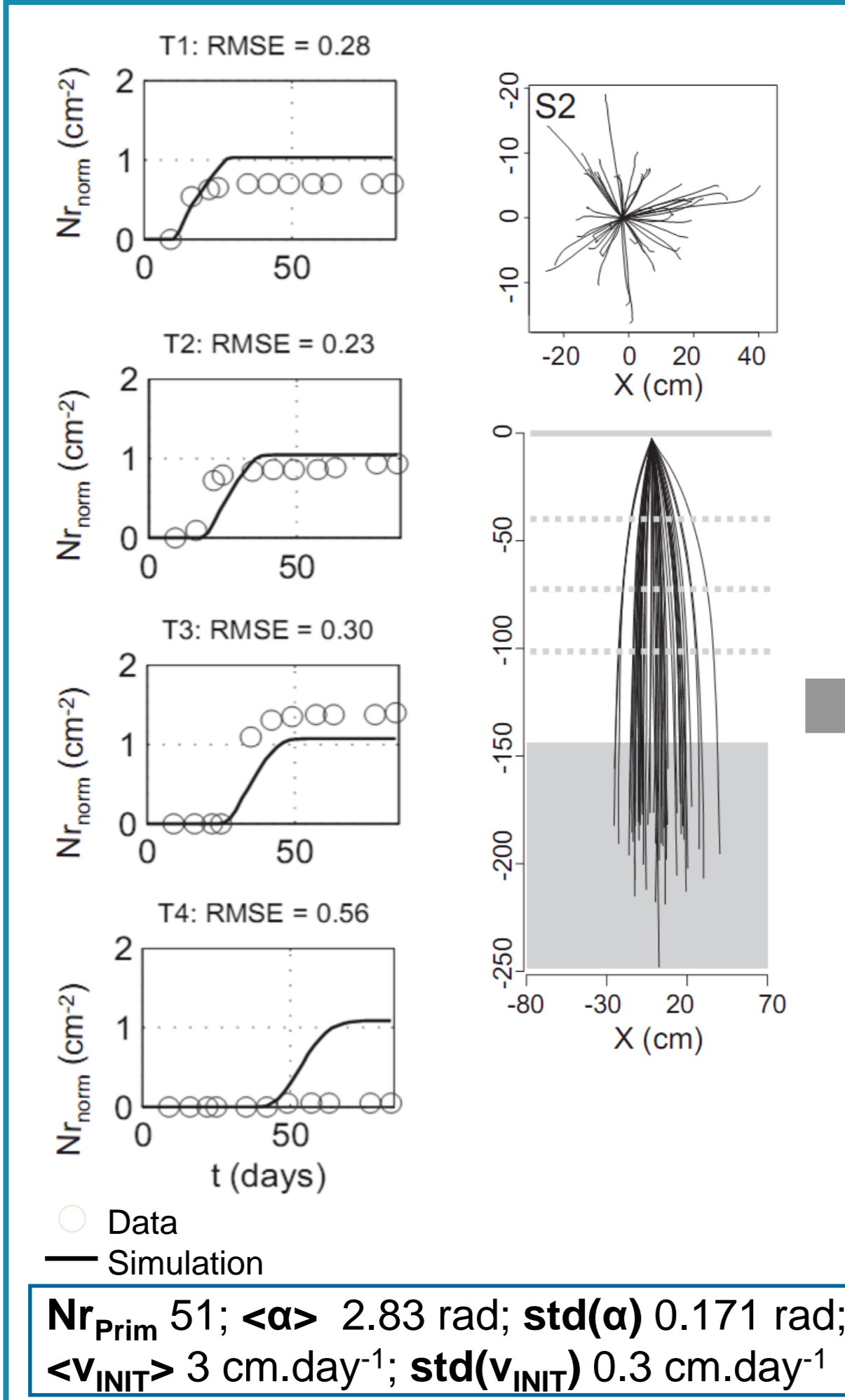
### Sensitivity analysis

- the Morris One-at-a-time test
- effect of  $\langle v_{INIT}(\text{Type } 0) \rangle$ ,  $\langle v_{INIT}(\text{Type } 1) \rangle$ ,  $std(v_{INIT}(\text{Type } 1))$ ,  $\langle \alpha \rangle$ ,  $std(\alpha)$  and  $Nr_{Prim}$
- 70 model evaluations



## Results

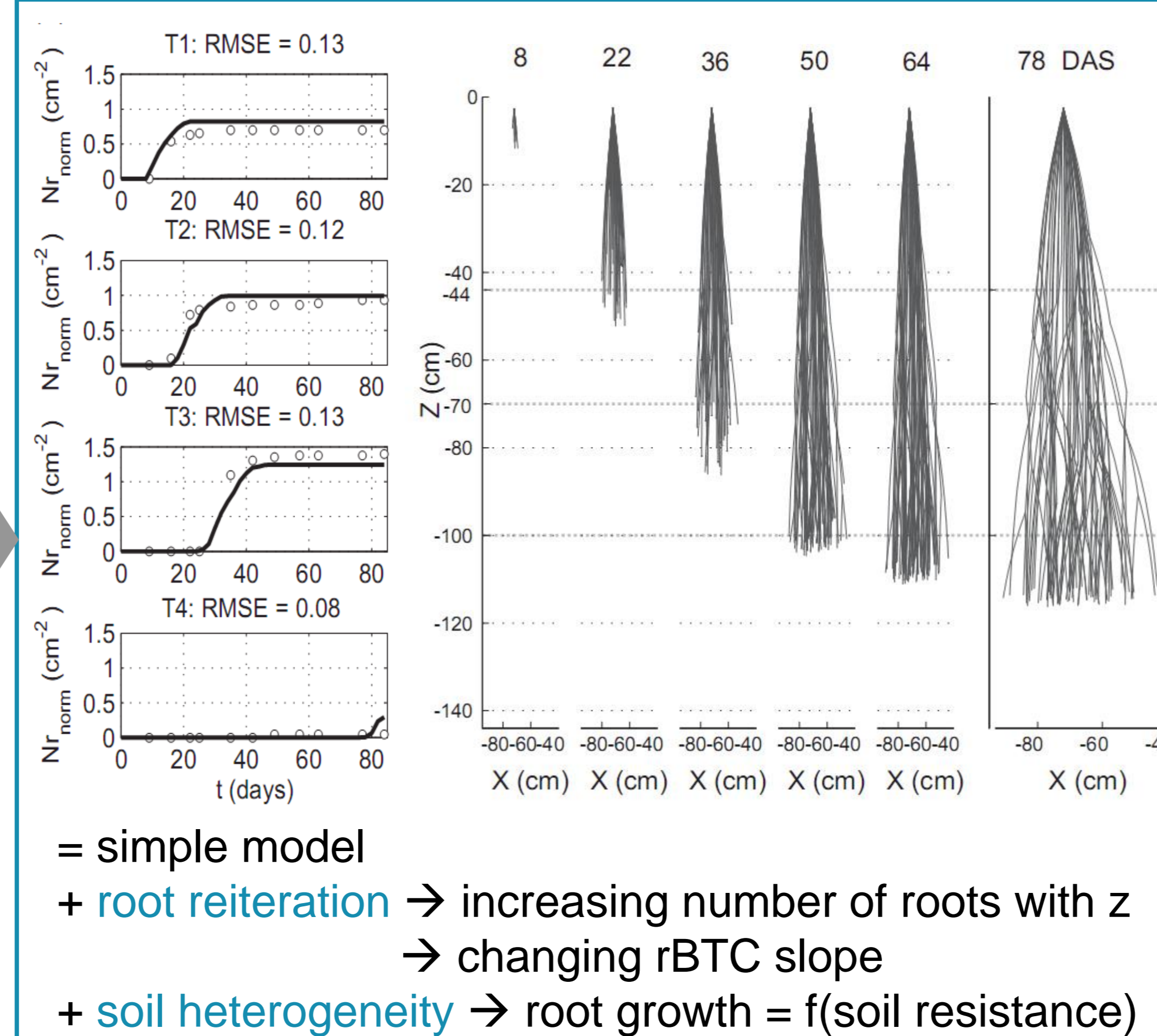
### Best simple model S2



### Observations

- Highest number of roots is **not** observed at the tube nearest to the soil surface  $\rightarrow$  new thick and long roots originating deeper in the soil profile?
- Slope of the root breakthrough curves changes with depth  $\rightarrow$  idem 1?
- Effective growth velocity of the roots changes with depth  $\rightarrow$  varying soil resistance may cause roots to change growth velocity or tortuosity of their paths within certain horizons?

### Complex model S2



## Conclusion

Minirhizotrons are the only way to retrieve **dynamic information on root growth and root architecture (RA)** in an undisturbed soil environment at the lysimeter or field scale without destroying parts of the root system. RA model parameters greatly influencing the root breakthrough curve at a certain depth are the **root growth velocity of the primary roots and the number of primary roots emitted**. The deviation between the model results and the observations indicated that other processes which were not considered in the simple simulations, such as **reiteration and varying soil resistance**, played an important role in the root development during this experiment. However, the minirhizotron technique does not provide enough information to restrain a RA model with reiteration and soil layering in a satisfying way without **being combined with additional information**.

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