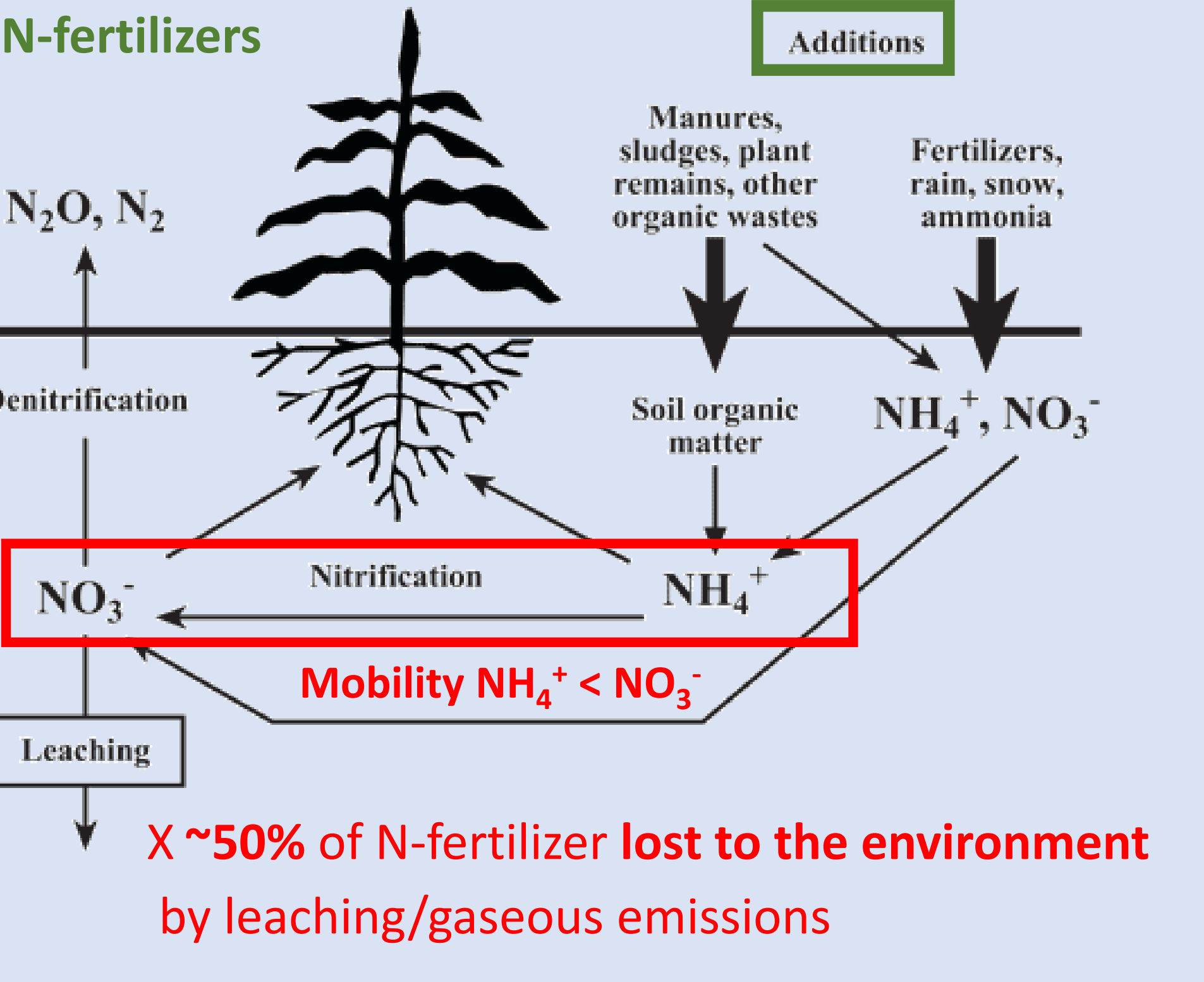


Overview: Biological Nitrification Inhibitors

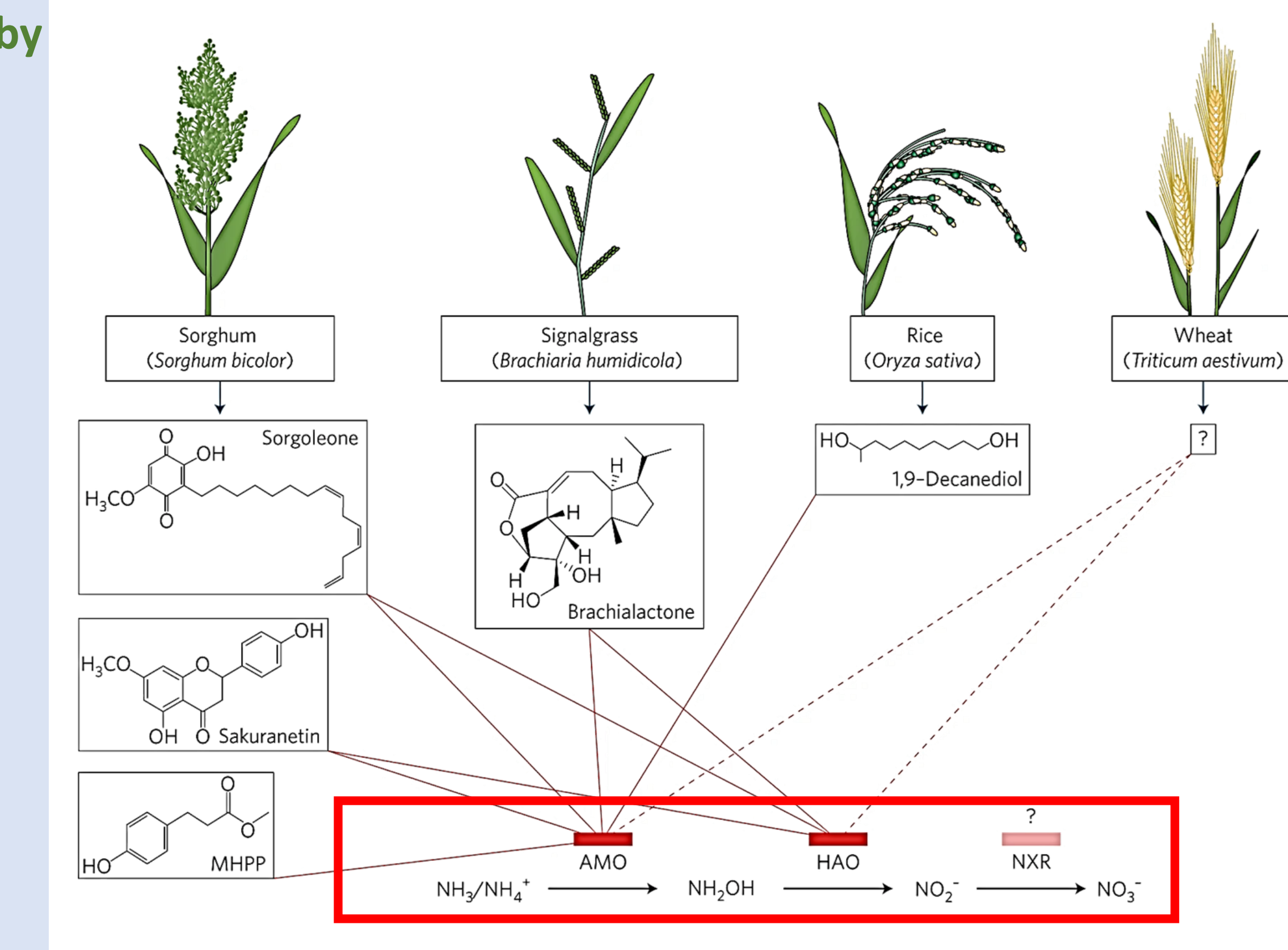
✓ 1/2 half the food produced in the world supported by N-fertilizers



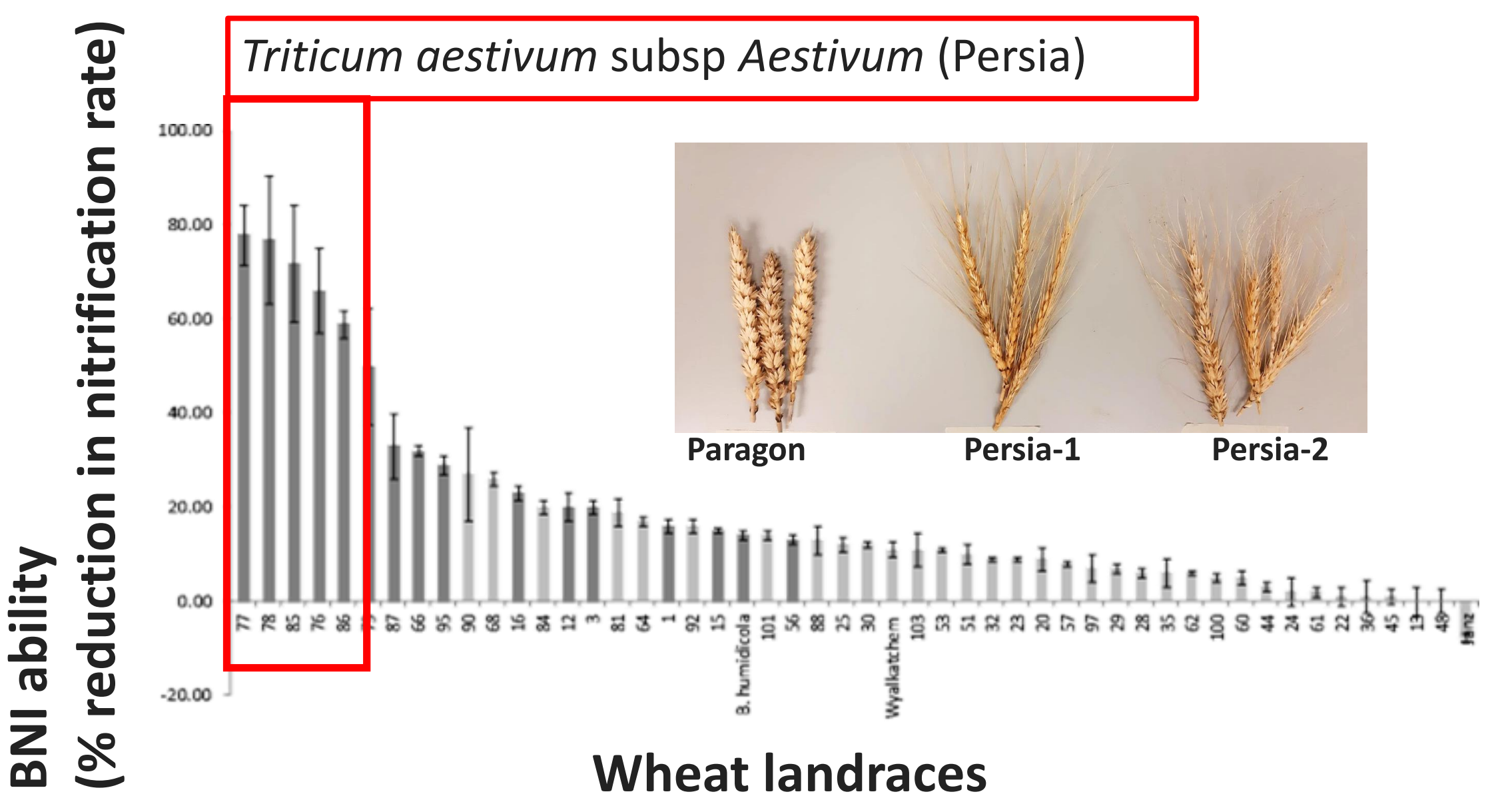
Biological nitrification inhibitors (BNIs) in root exudates reduce nitrification rates [1]:

- increasing N retention time in the root zone – mobility NH₄⁺ << NO₃⁻;
- decreasing fertilizer need and the associated environmental impact;
- providing an alternative to synthetic nitrification inhibitors.

Potential for breeding this trait into modern, elite wheat cultivars.



Historical wheat landraces with higher BNI capacity than equivalent modern wheat cultivars [2]:



Objectives:

- To develop a set of robust, complementary methodologies for rhizosphere phenotyping
- To quantify the variation across wheat landraces in the impact on rhizosphere nitrification rates;
- To link rhizosphere nitrification rates with trait loci in wheat landraces.

References:

1. Coskun *et al.* 2017 Nat. Plants 3: 17074
2. O'Sullivan *et al.* 2016 Plant Soil 404:61-74
3. Winfield *et al.* 2018 Plant Biotechnol. J. 16:165–175
4. <https://www.seedstor.ac.uk/search-browsecollections.php>
5. Cai *et al.* 2016 Anal. Chem. 89: 1178-1184
6. Kautsar *et al.* 2017 Nucleic Acids Res. 3;45:W55-W63

Phase I: Optimization of methodologies to measure nitrification rate

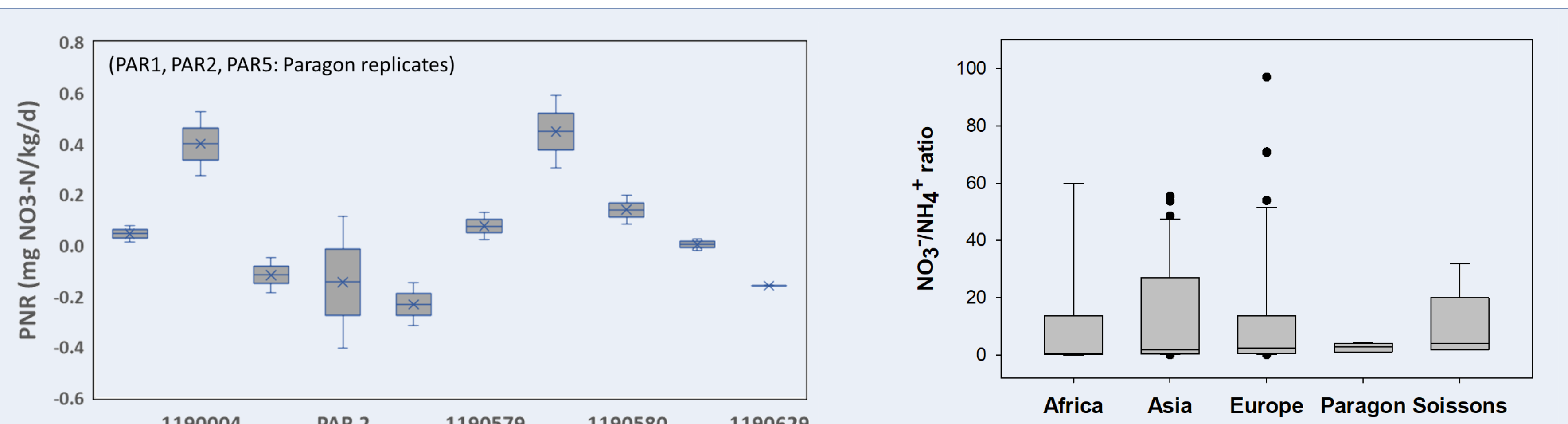
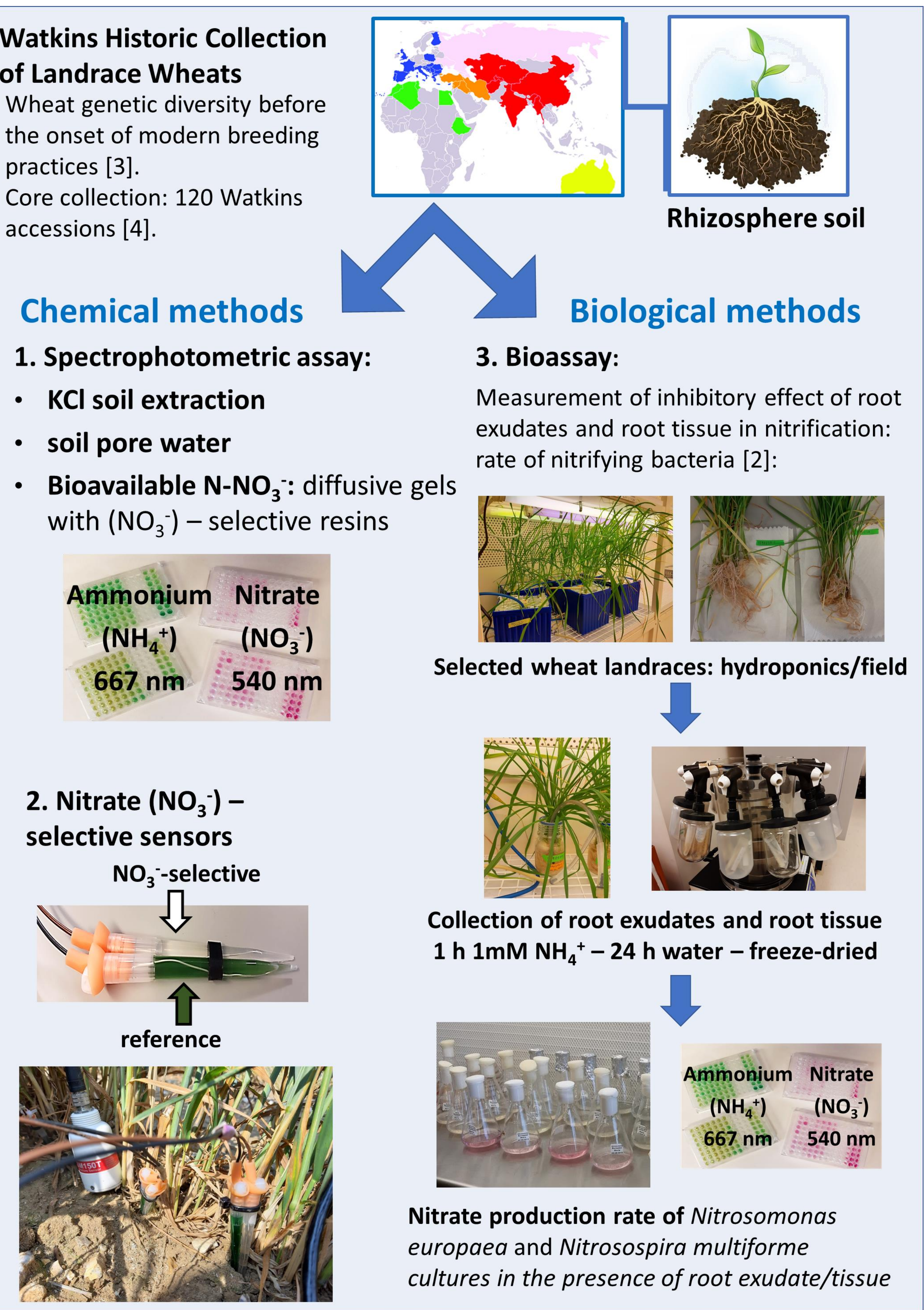


Figure 1. Potential nitrification rate (PNR) for rhizosphere soil collected from 7 historical wheat landraces [4] and a modern cultivar (Paragon) following addition of 100 mg N-NH₄ / kg soil.

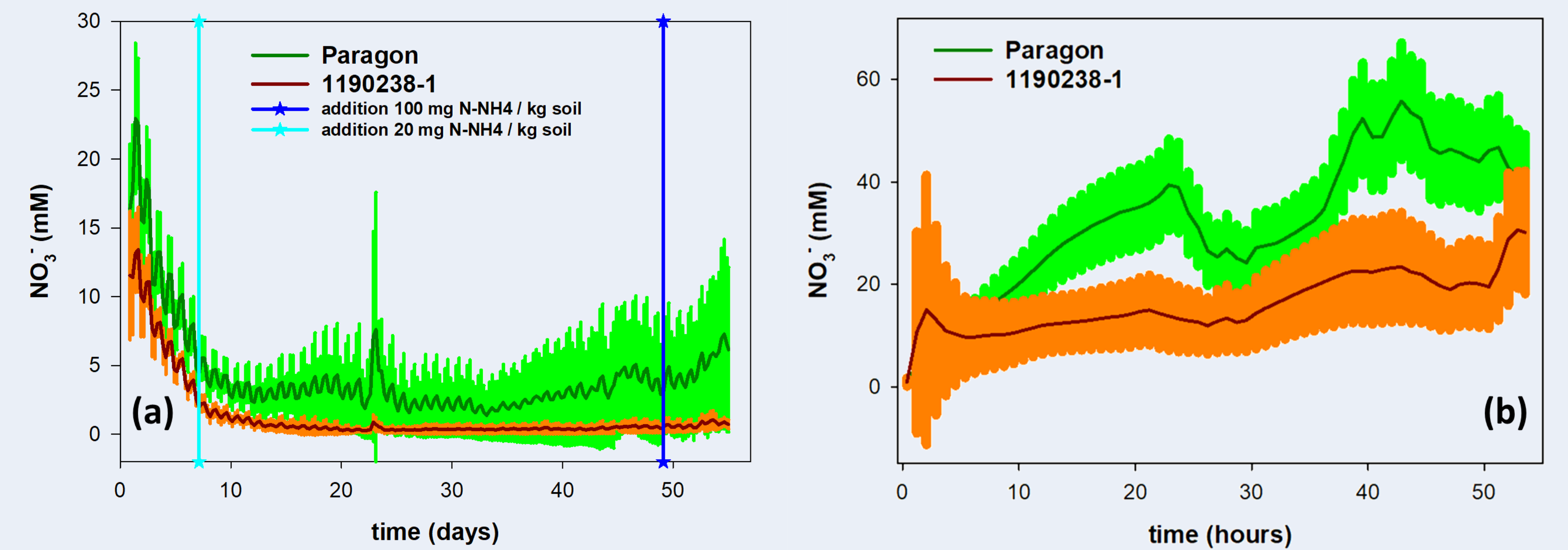


Figure 3. Data collected from nitrate (NO₃⁻) selective sensors for rhizospheres of a modern wheat cultivar (Paragon) and an historical landrace (1190238) grown in greenhouse (a) or in the field (b). Data variability is partly associated to the distance of the sensor to the rhizosphere area.

Preliminary results suggest that growing some of the historical wheat landraces can result in a decrease in nitrification rates in the soil rhizosphere. High variability is reported for geographic origin and replicates of modern cultivars.

Phase II

- Genome mining using the plantiSMASH platform [6] to identify genes clusters involved in BNIs biosynthesis pathways;
- Linkage analysis to identify trait loci (QTL) associated with BNI activity and ultimately identify potential breeding targets for BNI's [3]



Phenotyping population of 88 individuals of the wheat landrace candidate in a field experiment. The target trait, decrease of nitrification rate in rhizosphere soil, is being phenotyped for inhibition of nitrification by *N. europea* and pool of bioavailable NO₃ using selective resins.

