



Arthropod identification using DNA metabarcoding

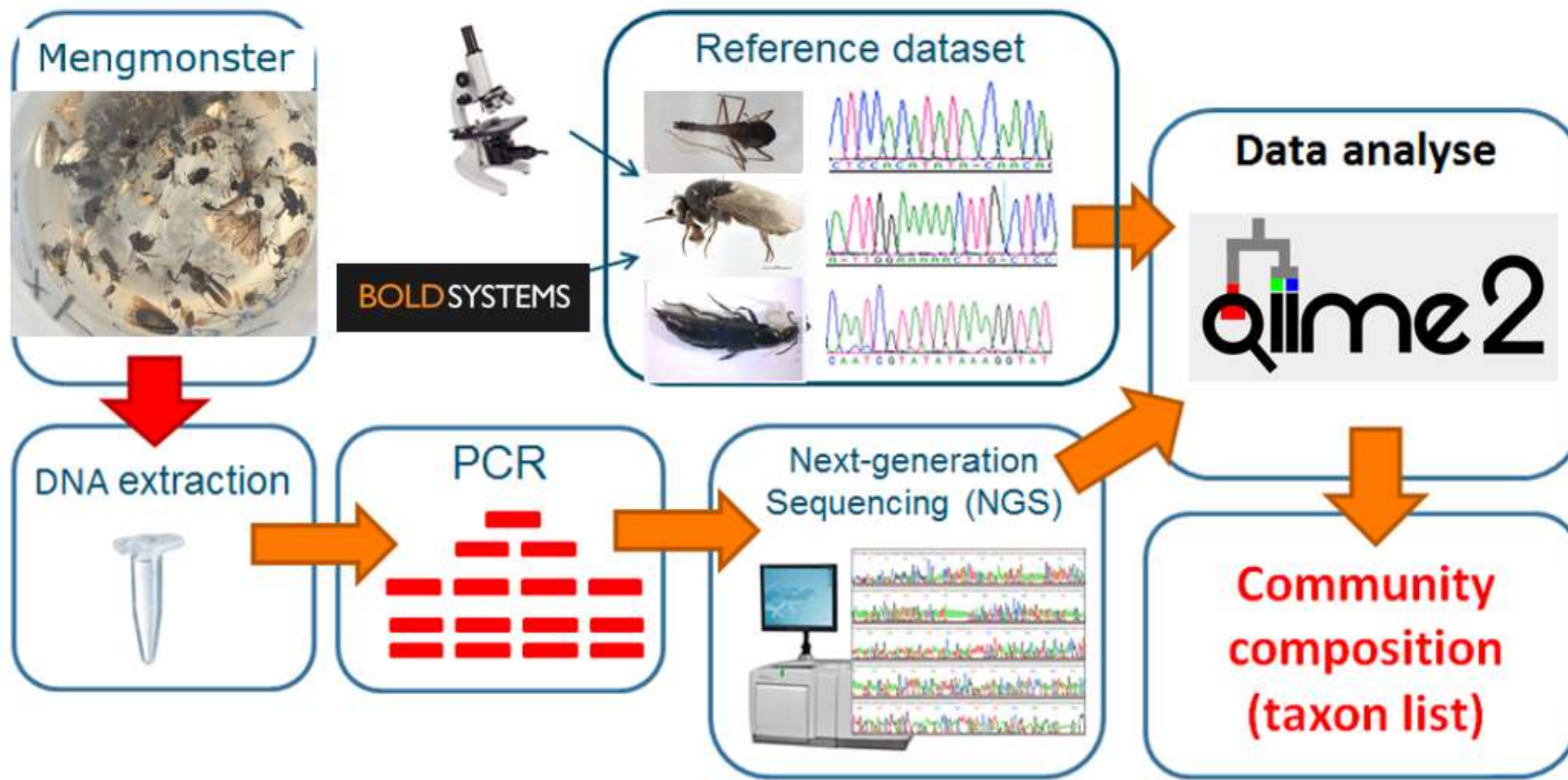
And combining results with AI and RS

Marcel Polling and Arjen de Groot

Team Animal Ecology, WENR

17/11/2022 RS&AI

DNA Metabarcoding



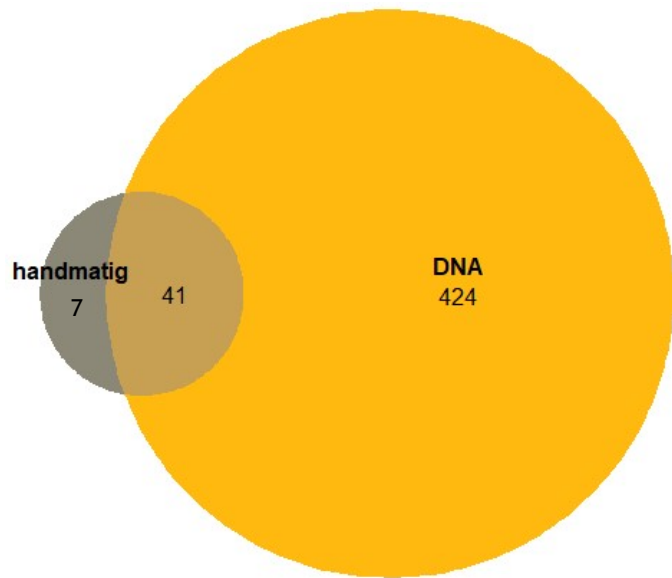
Insects collected using many different traps

- ▶ Projects on biodiversity, land use management, agriculture etc.



Projects: e.g. KB Efficiente monitoring for durable agriculture, OBN litter degradation micro-arthropods, PPS DZK Herb rich Grasslands, EcoCertified, SISEBIO (Univ. Barcelona)

DNA and morphology match very well,
but DNA finds many more species

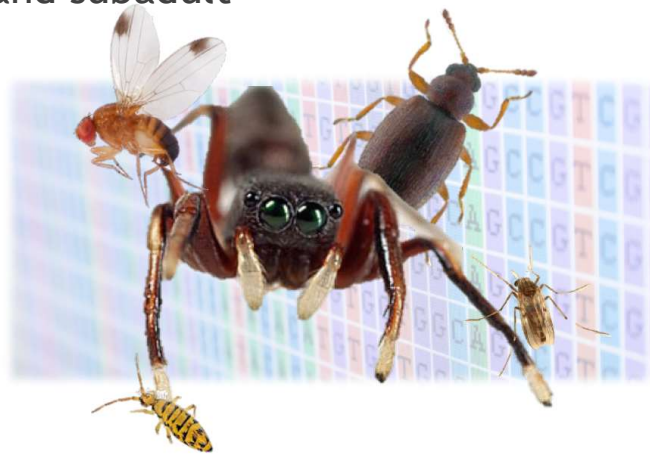


PPS Kruidenrijke graslanden

Staphorst Grassland – Dunnink 3 2009 Tube 40 (E201387)			Morphology		DNA	
Order	Family	Species	count	%	reads	%
Araneae	Lycosidae	<i>Trochosa ruricola</i>			12	0.2
Coleoptera	Staphylinidae		1	1.7	45	0.7
Coleoptera	Staphylinidae	<i>Philonthus cognatus</i>			135	2.2
Coleoptera	Staphylinidae	<i>Philonthus splendens</i>				
Diptera	Anthomyiidae		2	3.4		
Diptera	Anthomyiidae	<i>Delia florilega</i>			1757	28.8
Diptera	Anthomyiidae	<i>Delia platura</i>			155	2.5
Diptera	Cecidomyiidae		1	1.7		
Diptera	Cecidomyiidae	<i>Campylomyza flavipes</i>			16	0.3
Diptera	Ceratopogonida		1	1.7		
Diptera	Ceratopogonida	<i>Culicoides obsoletus</i>			14	0.2
Diptera	Ceratopogonida	<i>Culicoides punctatus</i>			273	4.5
Diptera	Chironomidae		5	8.6		
Diptera	Chironomidae	<i>Camptocladius stercorarius</i>			100	1.6
Diptera	Chironomidae	<i>Procladius culiciformis</i>			1219	20.0
Diptera	Chironomidae	<i>Pseudorthocladius</i>				
Diptera	Chironomidae	<i>Pseudorthocladius curtistylus</i>			9	0.1
Diptera	Chironomidae	<i>Pseudorthocladius filiformis</i>			24	0.4

Advantages:

- ▶ Faster and cheaper than morphological identification
- ▶ Objective: does not rely on taxonomic expertise
- ▶ Higher resolution (species level) allowing much more detailed ecological interpretations
- ▶ Semi-quantitative results possible
- ▶ Species ID for juvenile and subadult species

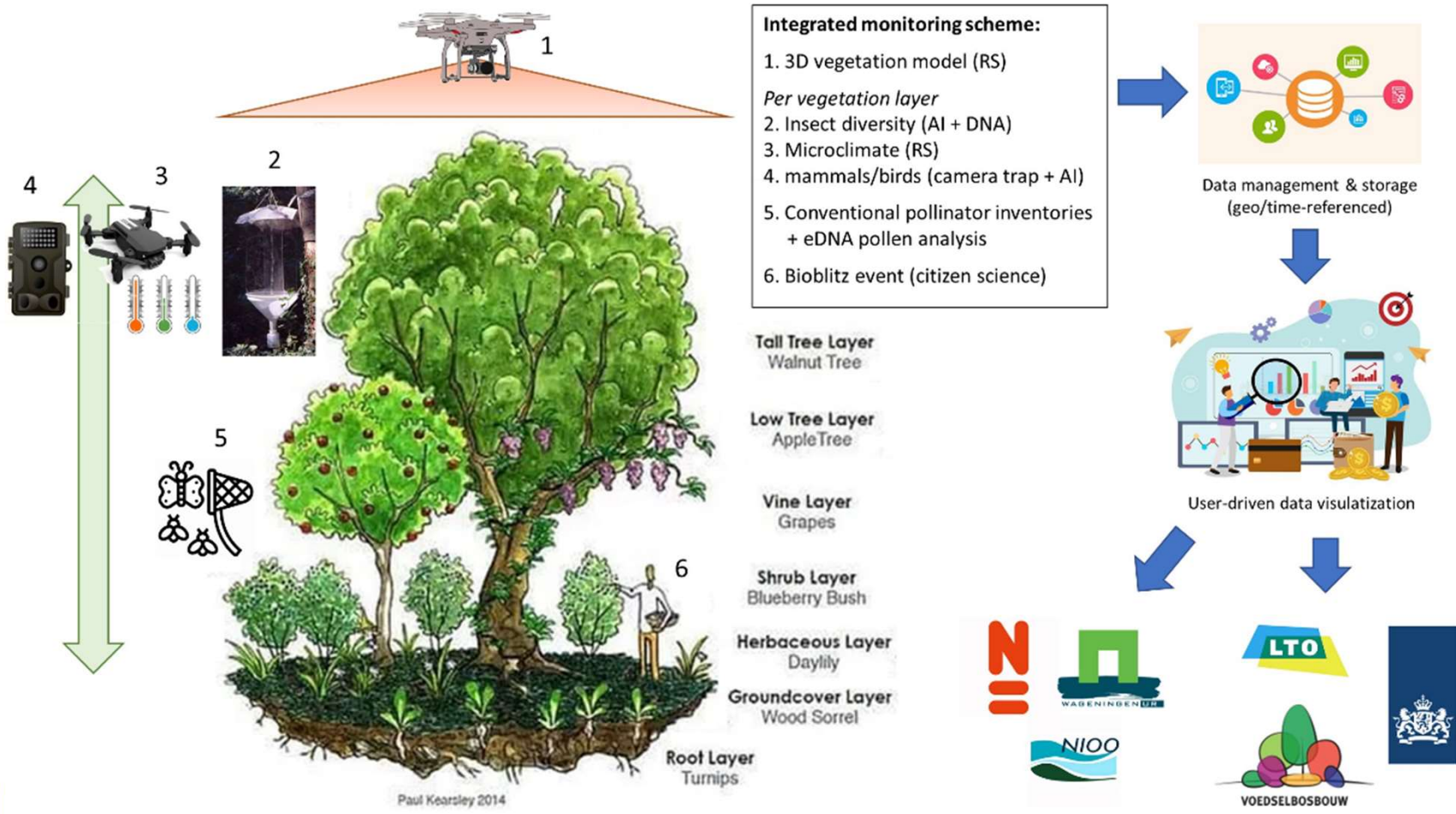


Disadvantages:

- ▶ No distinction between adults and juveniles possible
- ▶ Dependant on availability reference sequences
- ▶ No absolute quantities, abundance information lost

KB36 Cluster 5: Data Driven Biodiversity (food forests)

Collaboration with Earth Observation and Environmental Informatics group, Biodiversiteit en Beleid and FFC Post Harvest Technology



Integrating technologies showcase: AI and DNA for insect abundance estimations

