

***In Situ* CWR Management Planning and Implementation**

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Genetic reserve

- Area designated for the active management of the genetic diversity of one or more CWR populations in their natural habitat.
- Basic management unit for *in situ* CWR conservation

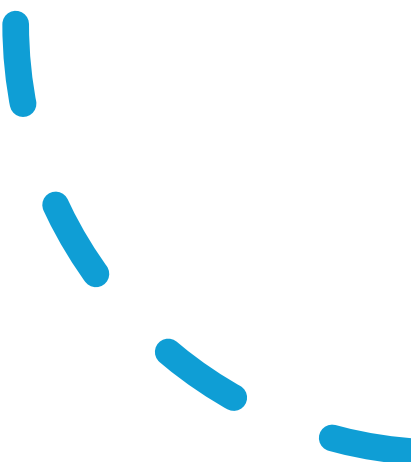


Genetic reserve management plan

- Management plan: planning tool that contains a set of prescriptions and actions to achieve the conservation goals

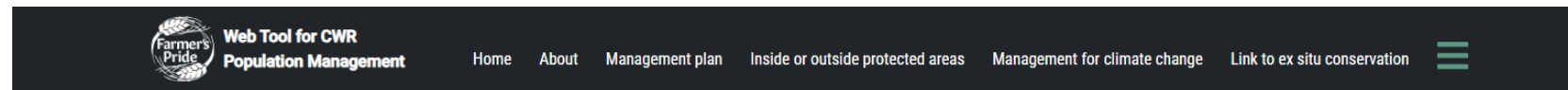


Elements of a genetic reserve management plan

- a. Site description
 - b. Target taxa population description
 - c. Habitat characterization
 - d. Objectives
 - e. Prescription
 - f. Monitoring and evaluation
 - g. Workplan and resources
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CWR population management guidelines

- Iriondo, J.M., Magos Brehm, J., Dulloo, M.E., Maxted, N. (eds.) (2021) Crop Wild Relative Population Management Guidelines. Technical Report. Farmer's Pride project.
- www.farmerspride.eu
- Web tool:
<https://cwrpopulation-toolkit.cropwildrelatives.org/>



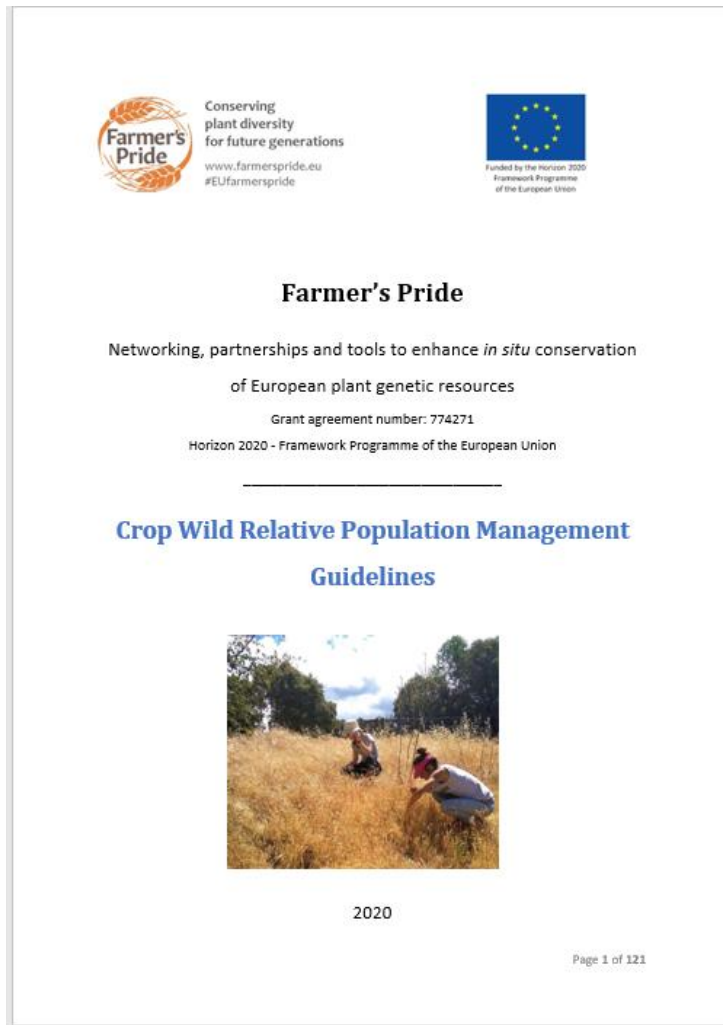
Web tool for CWR Population Management

Welcome!

This web tool for Crop Wild Relative Population Management provides **practical guidelines for the management of CWR populations *in situ*** and the sites in which they are being conserved.

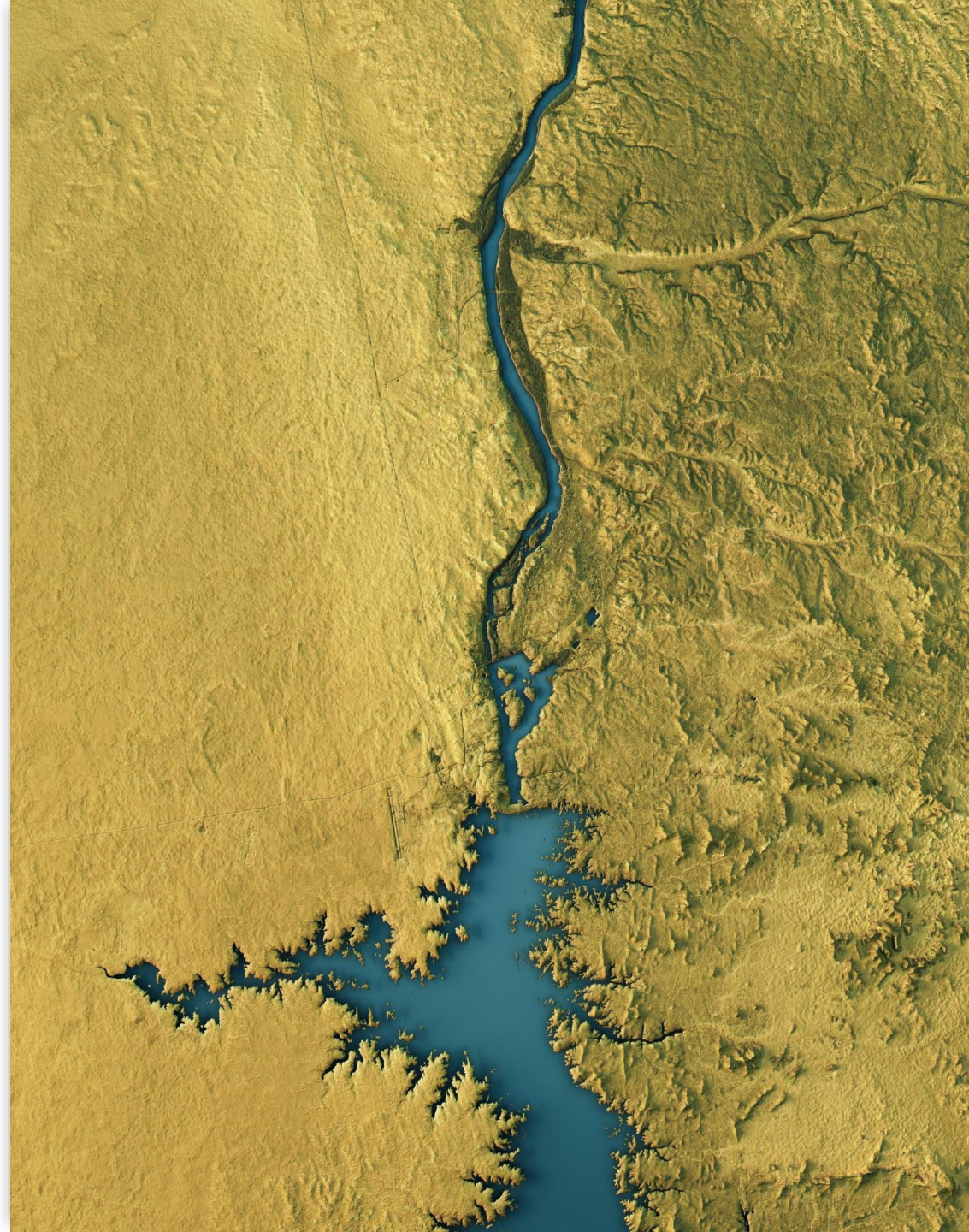
Bear in mind that there is **no single way to manage CWR populations** as it depends on the context in which the *in situ* conservation action has been established. However, this **user-friendly web tool** will guide you through the process of **developing and implementing an efficient CWR population management plan**, whether you are working in a protected area, on-farm, or other land use context. You will also learn how to **manage your CWR populations to address the impacts of climate change** and how **tomake these vital genetic resources available for use for the benefit of society** through complementary conservation in seed/genebanks.

The **community of people managing CWR populations across the globe is growing fast**. Do not hesitate, get started and be part of our network to conserve CWR in their natural habitats.



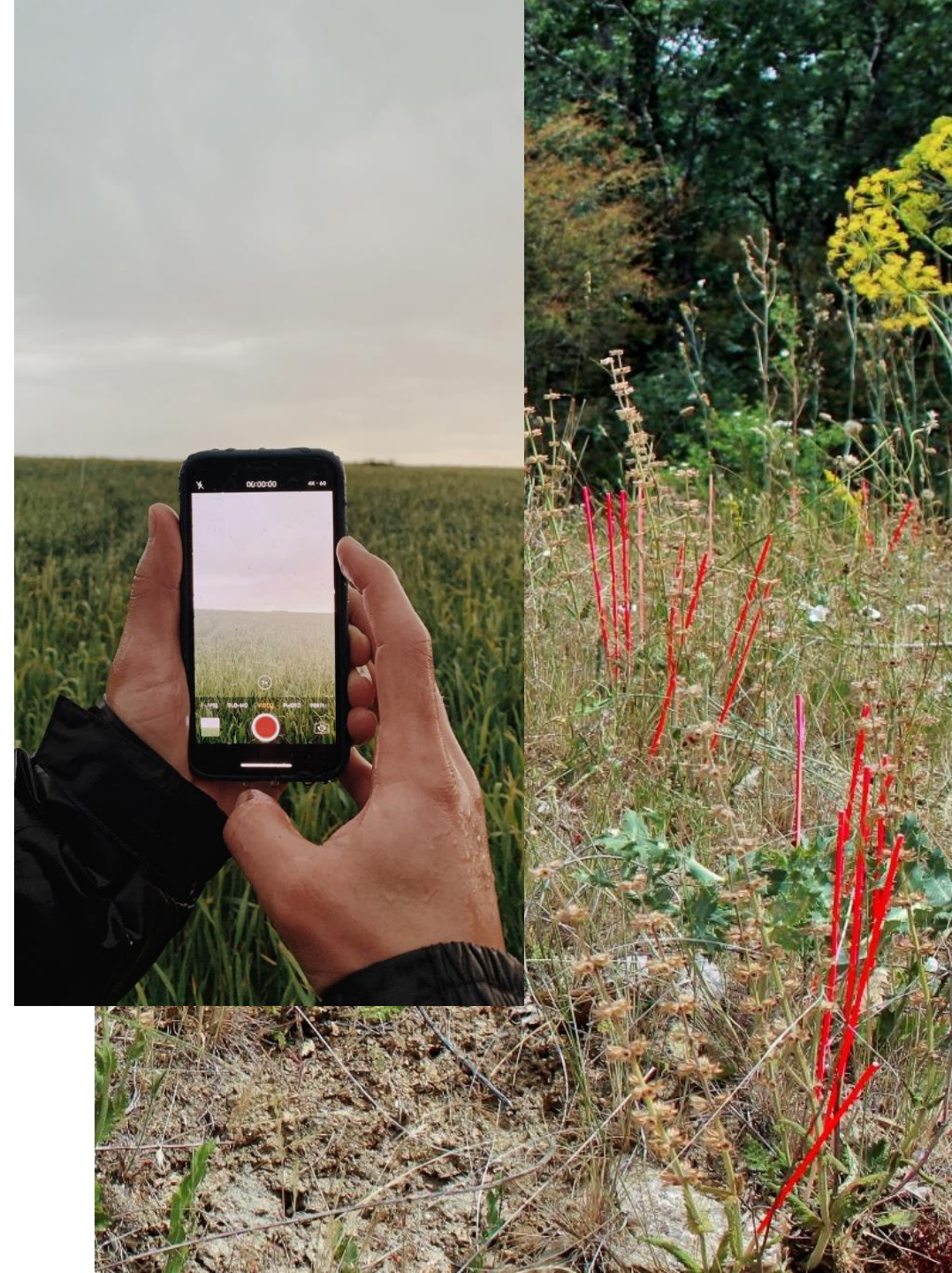
a. Site description

- geographic location, size
- legislation and policy measures
- land use and land tenure
 - public land: protected areas
 - private land: farms



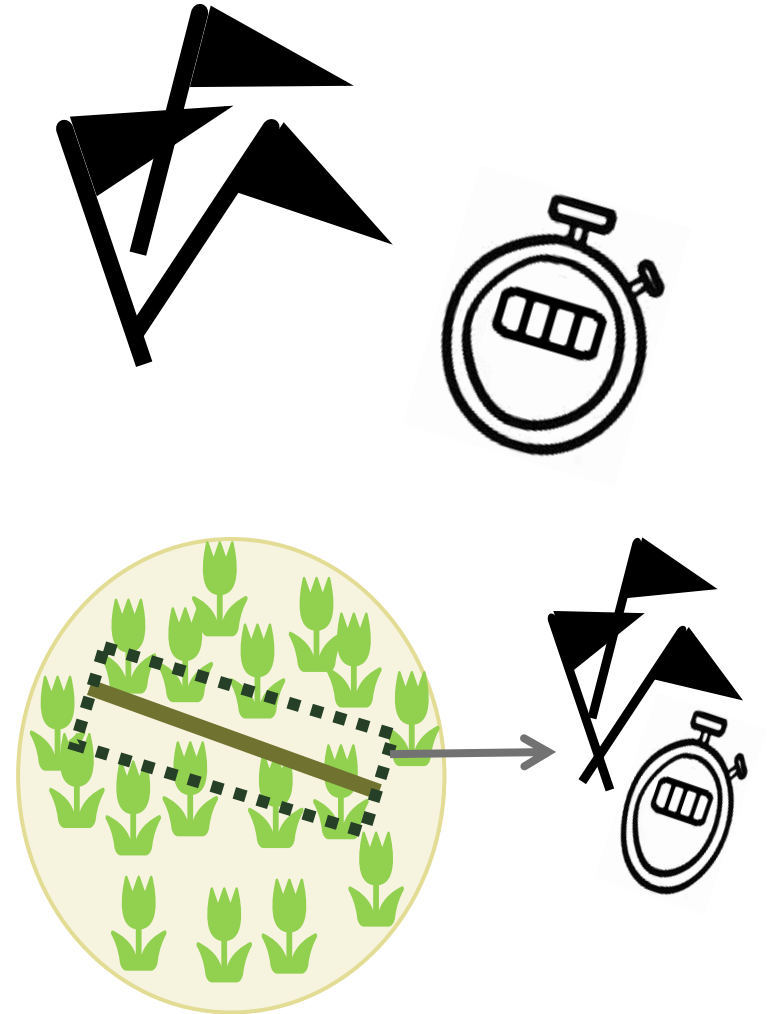
b. Target population description: geolocation and population size

- Georeferencing and delimitation of target CWR populations
- Population size: estimation of the number of reproductive individuals
 - Flowering time
 - Small populations: complete census
 - Large populations: estimation through sampling



b. Target population description: geolocation and population size

- Population trends:
 - Balance between resource investment and data quality.
 - Reasonable time investment per population: 2 hours
 - Simplified approach able to detect relevant demographic trends through time.



b. Target population description: **genomic characterization**

- **Goal: baseline characterization**

- Conservation
 - Genetic diversity and inbreeding
 - Population structure and gene flow
 - Signatures of selection
 - Introgression
- Use (Pre-breeding):
 - Identify genetically complementary populations for crossing
 - Genome-wide association studies (GWAS) for target traits
 - Genomic prediction
 - Trait introgression tracking
 - Marker-assisted selection

b. Target population description: genomic characterization

- Whole genome resequencing (WGS)
 - Assuming availability of reference genome of related crop species
 - 3-5 x coverage
 - Random sample of 30 individuals/population
 - 1500 € per population



b. Target population description: genomic characterization

- Goal: monitoring genetic diversity trends
 - Research on 'Genetic diversity' barcode
 - Genetic diversity estimators based on chloroplastic markers (e.g. 500 bp) optimized for each genus
 - 120 € per population (30 individual sample)



b. Target population description: phenotypic characterization

- Difficult to implement as a standardized practice in genetic reserves
- Only traits of interest by plant breeders
- Adapted standard descriptors of related crop
- Pilot studies to test usefulness and feasibility



b. Target taxa description:
threat assessment

THREAT ASSESSMENT			
Genetic reserve:			
Target CWR:			
Threat type	Timing	Scope	Severity
1 Residential and commercial development			
1.1 Housing and urban areas			
1.2 Commercial and industrial areas			
1.3 Tourism and recreation areas			
2 Agriculture and aquaculture			
2.1 Annual and perennial non-timber crops			
2.1.1 Shifting agriculture			
2.1.2 Small-holder farming			
2.1.3 Agro-industry farming			
2.1.4 Scale Unknown/Unrecorded			
2.2 Wood and pulp plantations			
2.2.1 Small-holder plantations			
2.2.2 Agro-industry plantations			

- Risks that could potentially threaten the genetic diversity and evolutionary potential of the population
- IUCN Threats Classification Scheme (version 3.2):
<https://www.iucnredlist.org/resources/threat-classification-scheme>

c. Habitat characterization

- **Physical characterization** (soil, geology, climate, hydrology)
- **Biotic characterization** (competitors, parasites, herbivores, pollinators, seed dispersers)





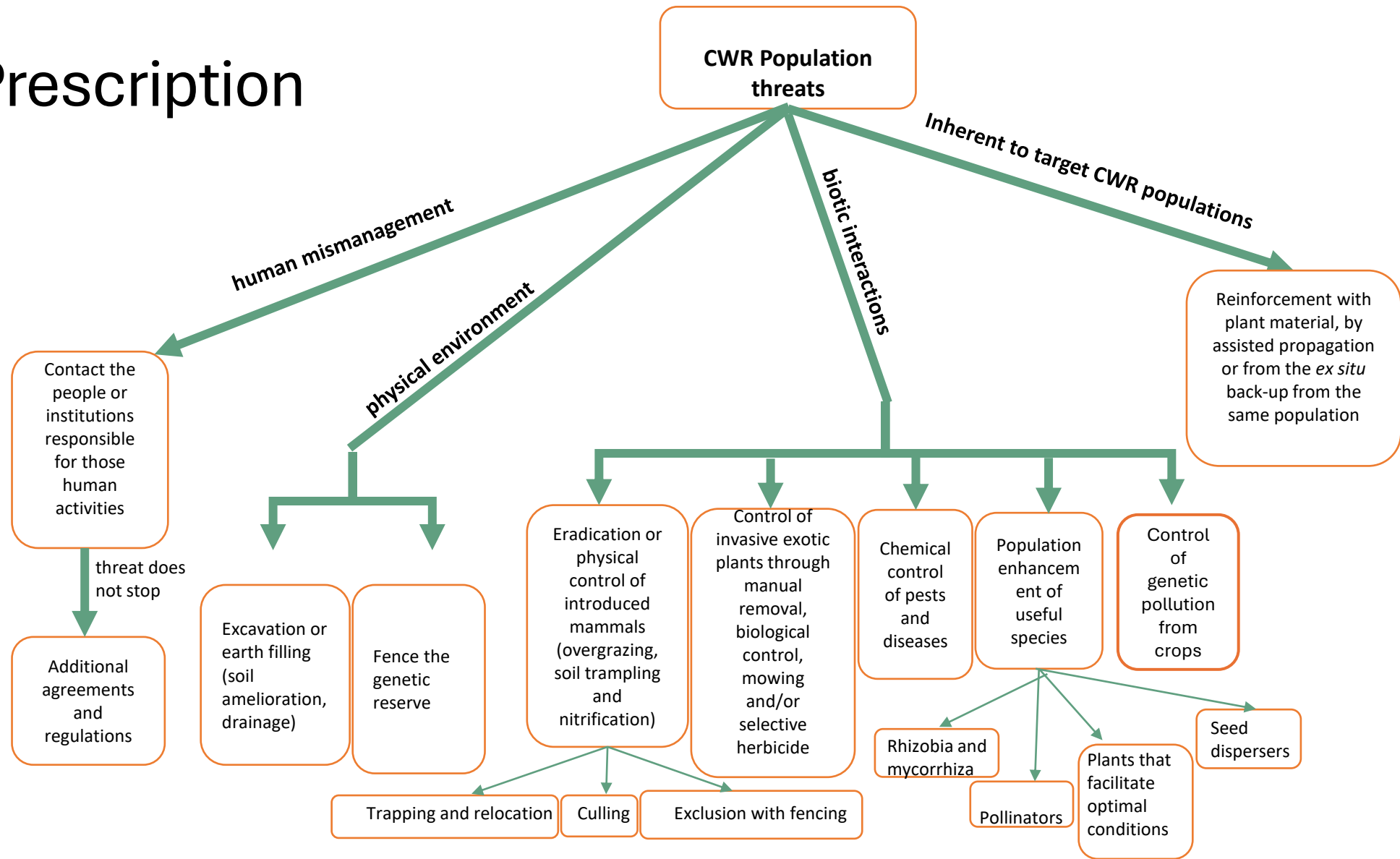
d. Management objectives

Overall objective: to maintain the viability and genetic diversity of the target CWR population(s)

Specific goals:

- Minimize risk of genetic pollution from nearby crops
- Support actions that promote genetic diversity
- Ensure access to populations for research, phenotyping, *ex-situ* backup for use in breeding, exploitation by local communities

e. Prescription



f. Monitoring and evaluation

- Aims:
 - to detect changes in population size
 - to assess trends in population genetic diversity
 - to detect changes in the physical and biotic components of the habitat
 - to determine the outcomes of management actions and to guide management decisions

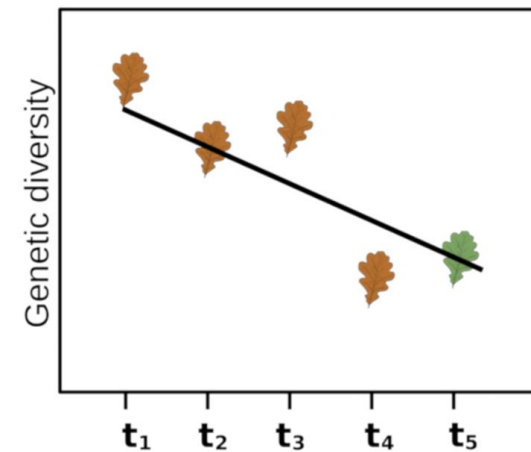
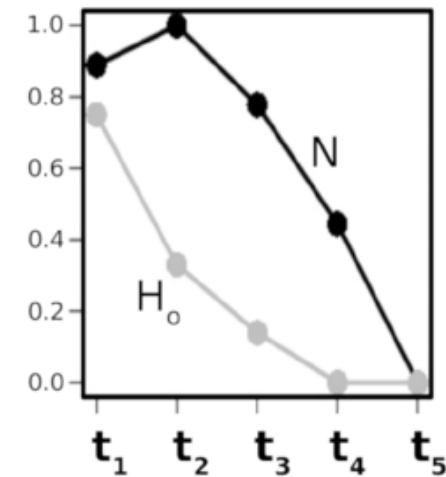
f. Monitoring and evaluation

- Qualitative check-ups throughout the year by protected area personel / farmers, managers
 - Alert about any significant perturbation
- Periodic monitoring and evaluation (e.g., every 5 years)
 - Demographic trends
 - Genetic diversity
 - Population threat assessment
 - Review of management plan



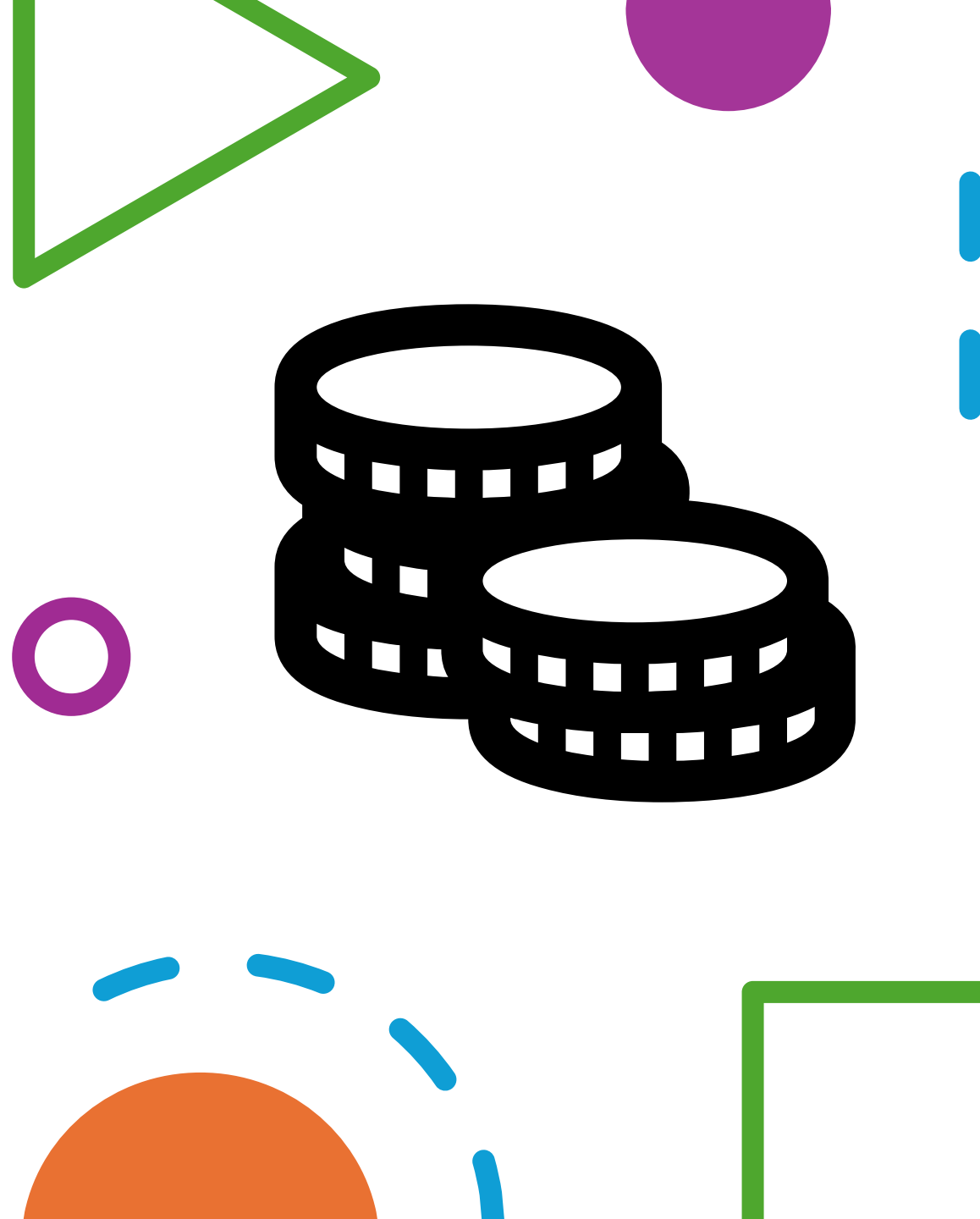
f. Monitoring and evaluation

- Identify demographic and genetic thresholds
- Life history traits (e.g. annuals vs. perennials)
- Trends through several time intervals



g. Workplan and resources

- Workplan by designated managers
- Management and monitoring resources:
 - Public sector:
 - **Agriculture** (CAP)
 - Environment (e.g., photovoltaic plants compensation measures)
 - Private sector:
 - Companies in the sector
 - Corporate responsibility



Integration of population information in the national *in situ* CWR database

- Simplify the work of genetic reserve managers
- Collect a variety of standardized data on:
 - Geolocation
 - Area of occupation
 - Demographic census
 - Phenotypic characterization
 - Population threat assessment
 - Habitat characterization
- Data gathering and delivery to cloud database







 KoboToolbox



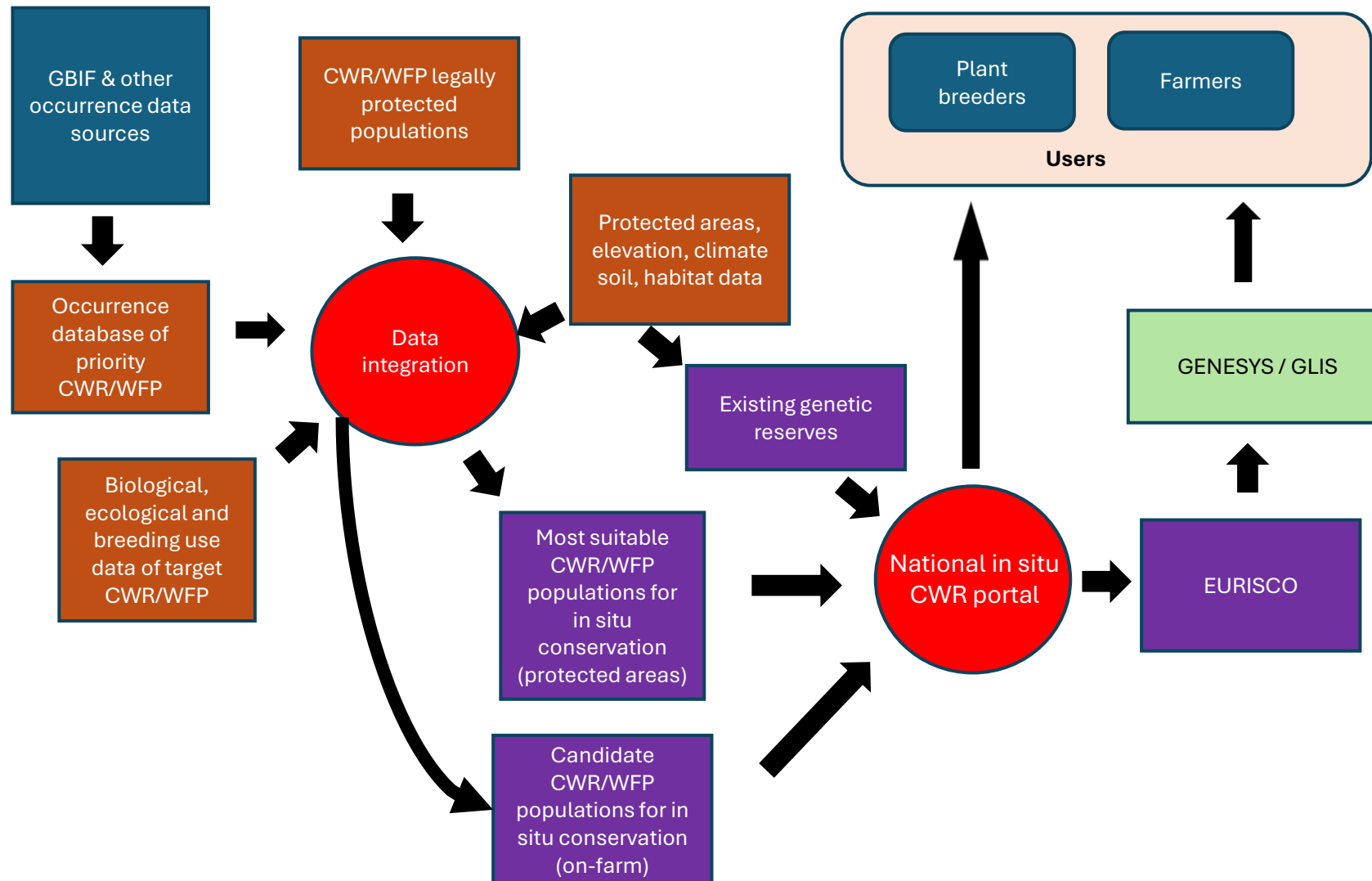
Ad-hoc app

Integration of population information in the national *in situ* CWR database

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RESUMEN FORMULARIO DATOS CONFIGURACIÓN											
ocultar campos											
1 - 3 3 resultados	Validación ▼	start ▼	end ▼	Genetic reserve polygon ▼	Name of the species ▼	Area of occupancy of the population ▼	Polygon of the core ▼	Population size ▼	Soil ▼	Habitat ▼	
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National Inventory of in situ conservation of CWR





Stakeholder
involvement
in CWR
conservation

