

# Rethinking Genebank Management – some critical thoughts and new approaches

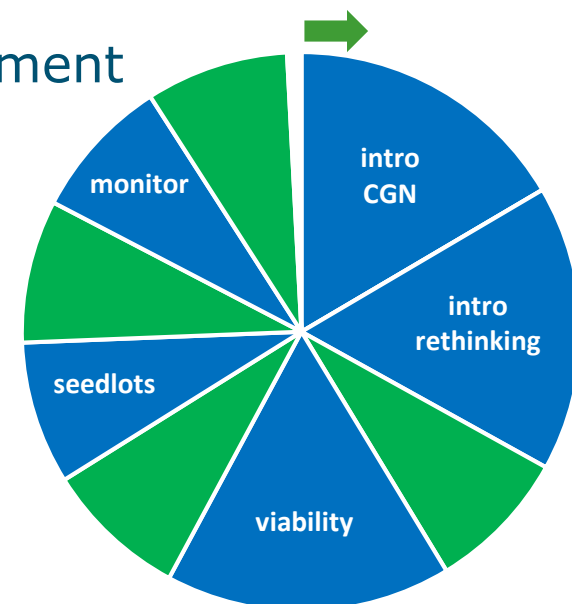
January 21<sup>st</sup>, 2021 – GROW webinar

*Theo van Hintum* (head CGN-Plant)  
Centre for Genetic Resources, The Netherlands (CGN)



# Rethinking Genebank Management

- overview webinar
  - short introduction CGN (10 min)
    - video
  - introduction rethinking genebank management (10 min)
  - viability monitoring (10 min)
  - principle seed lot management (5 min)
  - demo genebank monitor (5 min)
  - remaining 20 minutes used for discussion between the topics



# Short Introduction CGN

# Short Introduction CGN

- Centre for Genetic Resources, The Netherlands (CGN)
  - Dutch GRC for plants, animals and forest
    - this workshop is about **plant genetic resources (PGR)**
      - 2021 budget CGN-PGR is ± € 2 mln, 10.5 fte  
66% funded by governmental genebank program
  - CGN-PGR manages *ex situ* genebank
    - 23437 accessions
      - focus on **vegetables**
    - active collaboration with users
      - plant breeding industry
      - research community
    - ISO 9001 quality management system
      - since 2004



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# Short Introduction CGN

- Centre for Genetic Resources, The Netherlands (CGN)
  - CGN-PGR is involved in other PGR related activities
    - support NL **on-farm** actors
    - promotion **CWR** conservation to nature conservation
    - **policy** development
      - ITPGRFA
      - ABS & DSI debate
    - international **collaboration**
      - ECPGR
      - Crop Trust

# Short Introduction CGN

- Centre for Genetic Resources, The Netherlands (CGN)
  - as good as its **staff**



**Noor Bas**  
Curator /  
Communicatie



**Dione Bouchaut**  
Curator



**Martin Brink**  
Beleidsmedewerker



**Willem van  
Dooijeweert**  
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**Liesbeth de Groot**  
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Zaadbeheer



**Jarinka Heijink**  
Beleidsmedewerker  
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Afdelingshoofd /  
Onderzoeker



**Roel Hoekstra**  
Curator /  
Documentatie



**Chris Kik**  
Hoofdcurator /  
Verzamelaar



**Rik Lievers**  
Curator



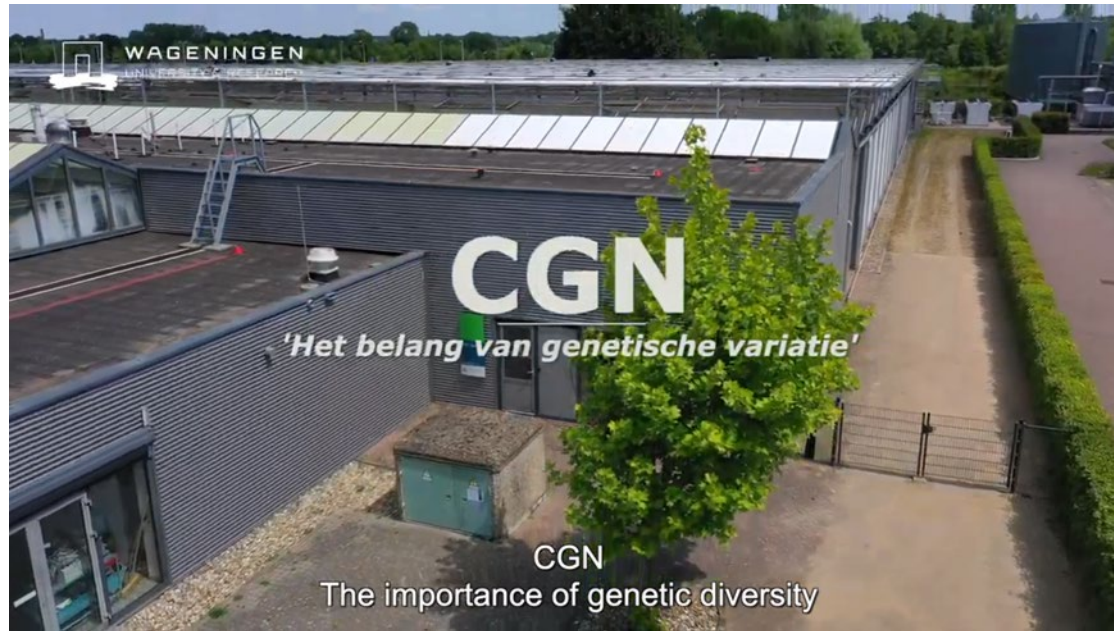
**Frank Menting**  
Documentatie /  
Communicatie



**Rob van Treuren**  
Curator /  
Onderzoeker



# Short Introduction CGN



<https://youtu.be/PGIpiMkhBGY>

# The Issues



# Rethinking Genebank Management

## ■ introduction

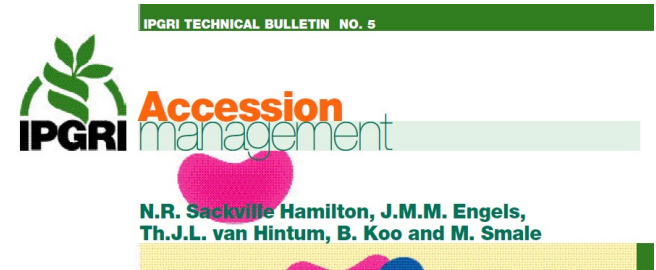
- genebanks arose **organically**
  - working collections for research and breeding grew
  - logistics became complicated due to size and use
  - legal and phytosanitary issues arose
- genebank management has become a **complicated** discipline

how would we set up a genebank from scratch?

- objective: **conserve PGR and make it available**
- funding is always the major limiting factor

# Rethinking Genebank Management

- genebank issues that could be re-thought
  - composition of the collection
    - what diversity to sample?
    - duplicate or complement other collections?
    - focus on cultivated or on CWR, and within these groups?
    - how much to invest to obtain material?
  - composition of the accessions
    - single lines or populations
      - splitting for self pollinators?
    - bulked accessions
      - bulking for cross pollinators?



*Genetic Resources and Crop Evolution* 43: 343–349, 1996.  
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## Reduction of duplication in a *Brassica oleracea* germplasm collection

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Received 7 March 1995; accepted in revised form 6 September 1995

**Key words:** duplication, bulking, *Brassica oleracea*, Brussels sprouts, white cabbage, genetic resources, isozymes

### Summary

To reduce the number of accessions in the *Brassica oleracea* collection of the Centre for Genetic Resources The Netherlands (CGN) groups of accessions were bulked. Accessions in a group were selections from the same landrace or old variety, and were chosen, with the help of crop experts, on the basis of their history and morphology.

# Rethinking Genebank Management

- genebank issues that could be re-thought
  - conservation method
    - clones or seeds or pollen or DNA
      - e.g. apple or potato
    - technical set-up
      - what temperature for seed storage?
      - small freezers or freezer rooms?
  - regeneration method
    - large numbers of accessions or high genetic integrity?
      - investment needed to maintain genetic integrity (during regeneration) can not be spend on increasing the collection size
  - viability monitoring
    - large numbers of accessions or high security?



# Rethinking Genebank Management

- genebank issues that could be re-thought
  - availability
    - how can (/should?) use be promoted?
    - what user groups should be targeted?
    - what conditions for access are applied?
    - what service level is targeted?
      - access to information / core selectors / bio-informatics interfaces
      - user consultancy
      - phytosanitary and import permits, non-GMO statements
  - quality management
    - invest in quality management?
    - what system and to what level of detail?

# Rethinking Genebank Management

- genebank issues that could be re-thought
  - automation
    - image recognition
    - material storage
  - monitoring regeneration
    - high-throughput techniques
  - integration bioinformatics
    - optimisation collection composition
    - selection material for use
  - integration *in situ* actors
    - become a true genetic resources centre

# Rethinking Genebank Management

## ■ messages

- genebank management decisions are **arbitrary**

- funding is limiting factor
- pro's and con's of decisions should be clear
  - sometimes more research is needed
- community consensus is difficult
  - optimal decision depends on purposes of genebank
  - genebank standards support collaboration



- whatever decisions taken – they need to be implemented correctly
  - quality management is the tool



# Viability Monitoring

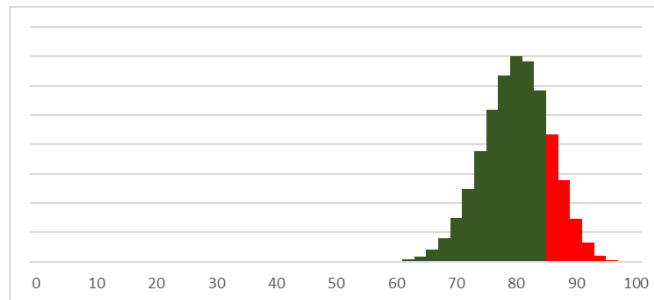
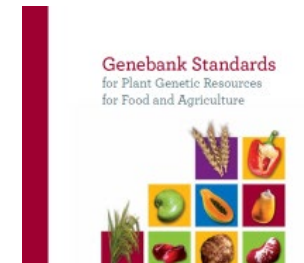


# Viability Monitoring

- to assure the **quality of the seeds** viability needs to be monitored

- **FAO (2014) standards** have been defined

- initial value > 85 %
- threshold for regeneration = 85%
  - or lower depending on the species or specific accessions of initial viability
- $N_{\text{seeds}} = ?$ 
  - “dependent upon the size of the accession but should be maximized to achieve statistical certainty; however, the sample size should be minimized to avoid wasting seed”



sampling effects:  
if true germination = 80%,  $N_{\text{seeds}} = 50$ ,  
then 19.0% of the tests will result in  
values over 85%

# Viability Monitoring

- **previous** CGN viability monitoring protocol
  - germination tested by ISTA certified labs
    - following ISTA protocols but 200 instead of 400 seed
  - reliability appeared **very low**
    - based on 5-10% blind doubles
    - too many wrong decisions were made
      - to regenerate – shortened generation span and unnecessary costs
      - not to regenerate – risk of losing the accession
- other genebanks appear to all do it in their own ways
  - based on a small inventory

van Dooijeweert, W., Menting, F. (2018) Procedures for Germination in Genetic Resources Conservation. CGN Internal Report. 33p. (available on request)

# Viability Monitoring

source: Hintum & Treuren (2012) PGRCU 10:134

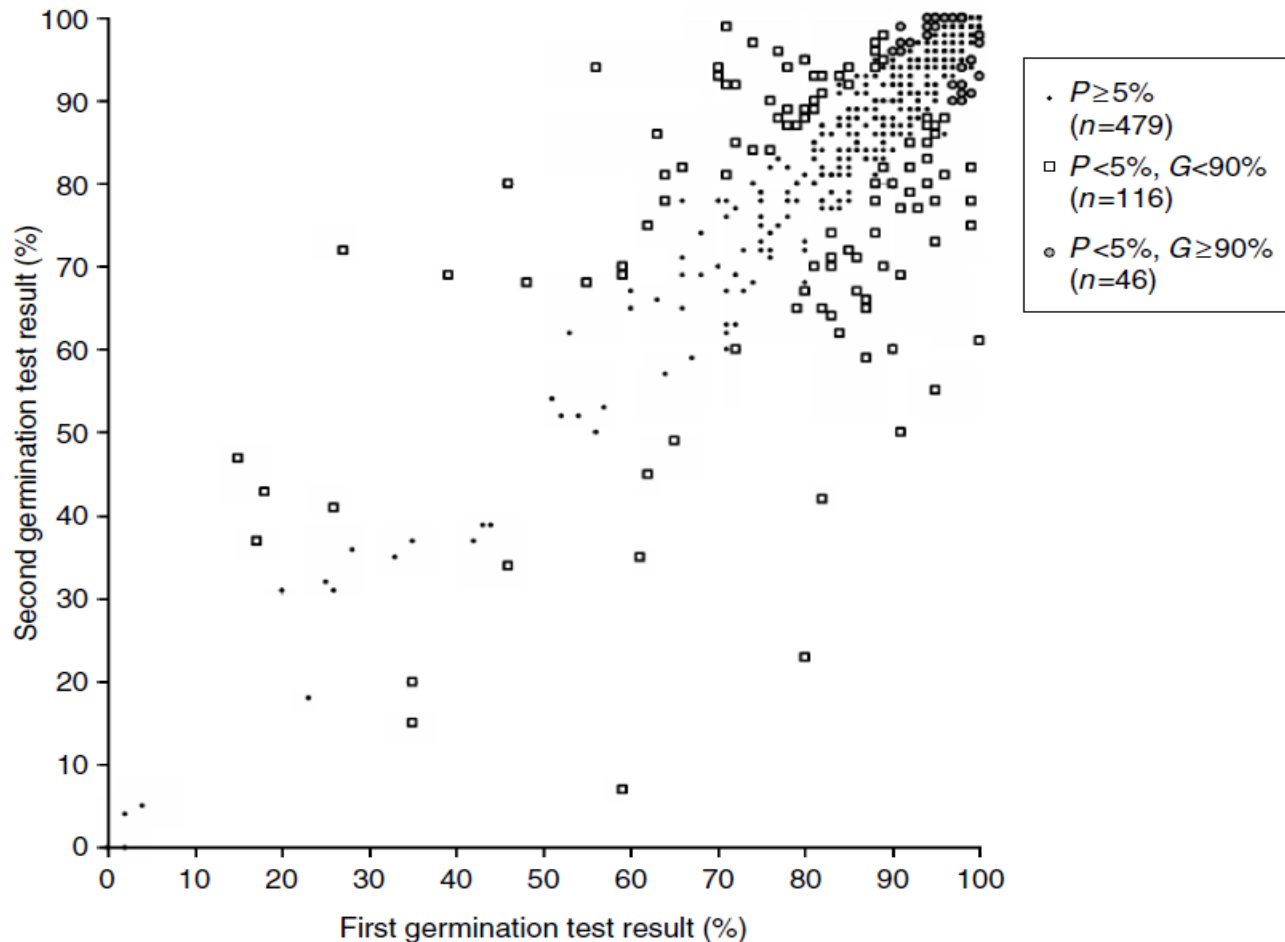


Fig. 1. Germination control tests of 641 seed lots.  $P$  indicates the probability of the results by chance alone, while  $n$  represents the number of included pairs. For data points with  $P < 5\%$ , a distinction is made for germination values ( $G$ ) lesser than 90% and values  $\geq 90\%$ .

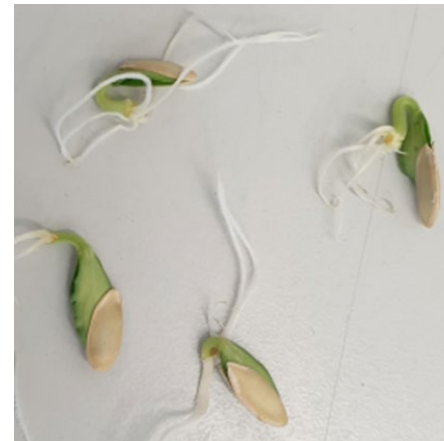
# Viability Monitoring

- grey zone between germinating and not germinating

- lettuce



- melon



# Viability Monitoring

- new viability monitoring protocol
  - still apply **same viability thresholds**
    - complying to FAO genebank standards
  - wait 25 years after **tested** successful regeneration
    - based on Treuren et al (2013)
  - determine quality of seeds the way you think best
    - use following scale
      1. **very good**, no worry, wait another 10 years (20 for small cereals)
      2. **good enough**, but pay closer attention, test within few years
      3. **dubious**, regenerate within few years
      4. **bad**, regenerate as soon as possible
  - further development of protocol is possible
    - blind doubles are being tested to determine repeatability
    - fate of 'bad seedlings' needs to be determined

# Seed Lot Management

# Seed Lot Management

- many genebanks maintain various seed-lots per accession
  - necessary complexity ?
- seed-lots serve two conflicting purposes
  - prolong generation span of material
    - every regeneration (rejuvenation) threatens genetic integrity
      - human error, genetic drift ( $N_e < \infty$ ), natural selection, etc.
  - always have seed available for distribution
    - seed stocks run out and need regeneration (multiplication)



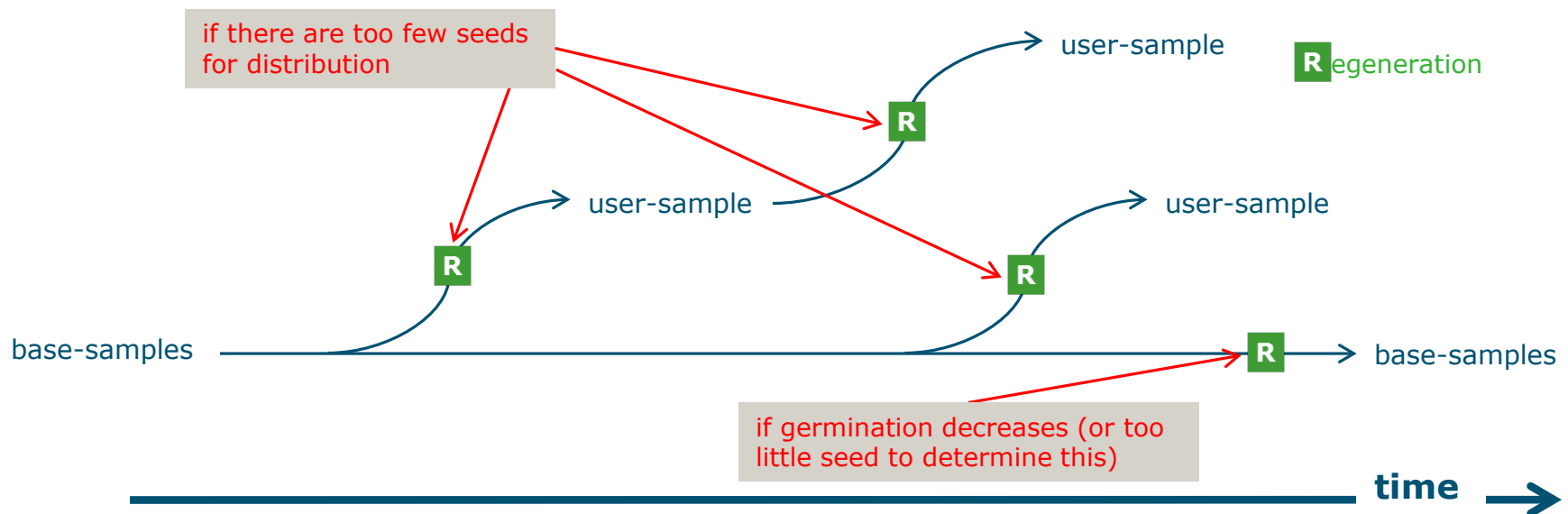


# Seed Lot Management

- CGN uses two types of seed-lots
  - base-seed-lots and user-seed-lots
  - every accession has a base-seed-lot
    - consisting of user-, regeneration-, germination- and rest-samples
      - viability is monitored
      - regenerated when viability drops or base-seed (for viability testing) runs out
  - some frequently used accessions have user-seed-lot
    - consisting of only user-samples
      - initial viability testing
      - no viability monitoring
      - not used for creating new base seed lots

# Seed Lot Management

- CGN uses two types of seed-lots
  - base-seed-lots and user-seed-lots



# Demo Genebank Monitor

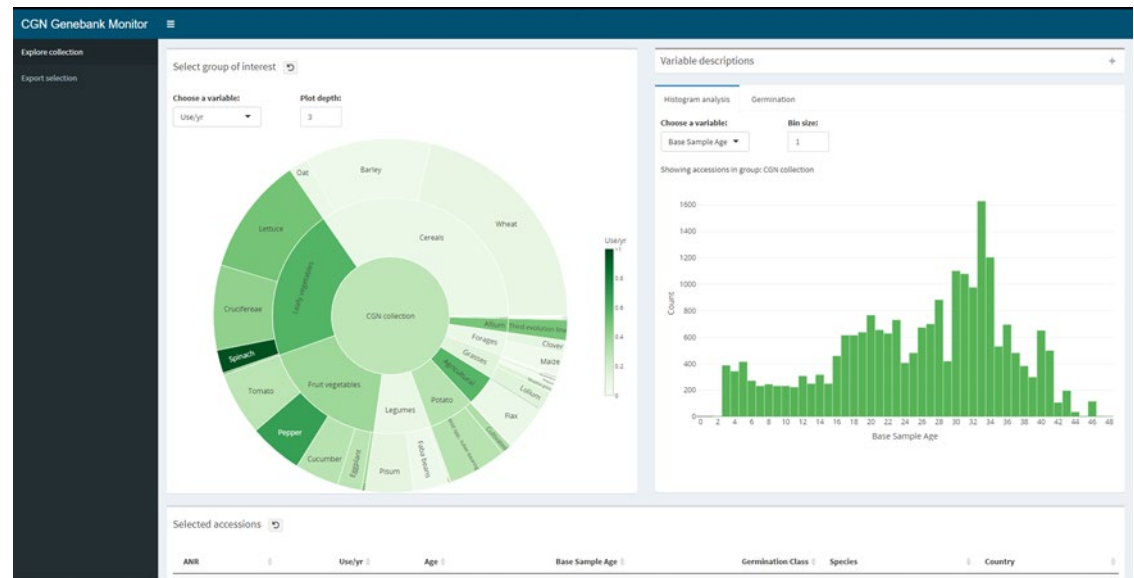
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# Demo Genebank Monitor

- genebank managers / curators need overview of their collection to make management decisions
  - composition
  - use
  - traits
  - seed age
  
- databases are often far from accessible
  - standard reports are not sufficient
  - more complex questions are difficult to answer

# Demo Genebank Monitor

- genebank monitor is based on
  - diversity tree of the collection
  - basic accession-based statistics
- prototype



# Demo Genebank Monitor

- **prototype** genebank monitor
  - available at <https://cgn-monitor.shinyapps.io/monitor/>
  - intended to convey idea of genebank management tool
    - software (R-package Shiny) possibly not the best
    - presented data very limited

we need concepts, knowledge, platforms and tools  
to professionalize genebank management

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# Thank you for your attention !

