



Strategic Research Center

GenSAP

Genomic Selection in Animals and Plants



GenSAP partners

- **University partners**
 - Aalborg University
 - Aarhus University
 - China Agricultural University
 - Copenhagen University
 - French National Institute for Agricultural Research
 - Melbourne University
 - North Carolina State University
 - Norwegian University of Life Science
 - University of Wisconsin-Madison
- **Industry partners**
 - DLF Trifolium
 - Kopenhagen Fur
 - LKF Vandel
 - NAV - Nordic cattle genetic evaluation
 - Nordic Seed
 - Sejet Plant breeding
 - VikingGenetics
 - VSP - pig research centre



Genomic selection

- Breeding for complex traits
 - Many genetic and non-genetic factors
 - Complex interactions among genes
 - Interactions between genes and environment
- Use genomewide markers in LD with causative genetic variants
 - Predict genetic merit
 - Breeding decisions and long term consequences of those
- Implemented in dairy cattle and pig breeding using 50K markers
 - Work very well for large homogeneous breeds
 - Works poorly across breeds and long genetic distances
- Large potential in other species
- New technologies available (sequence, OMICS)



Vision

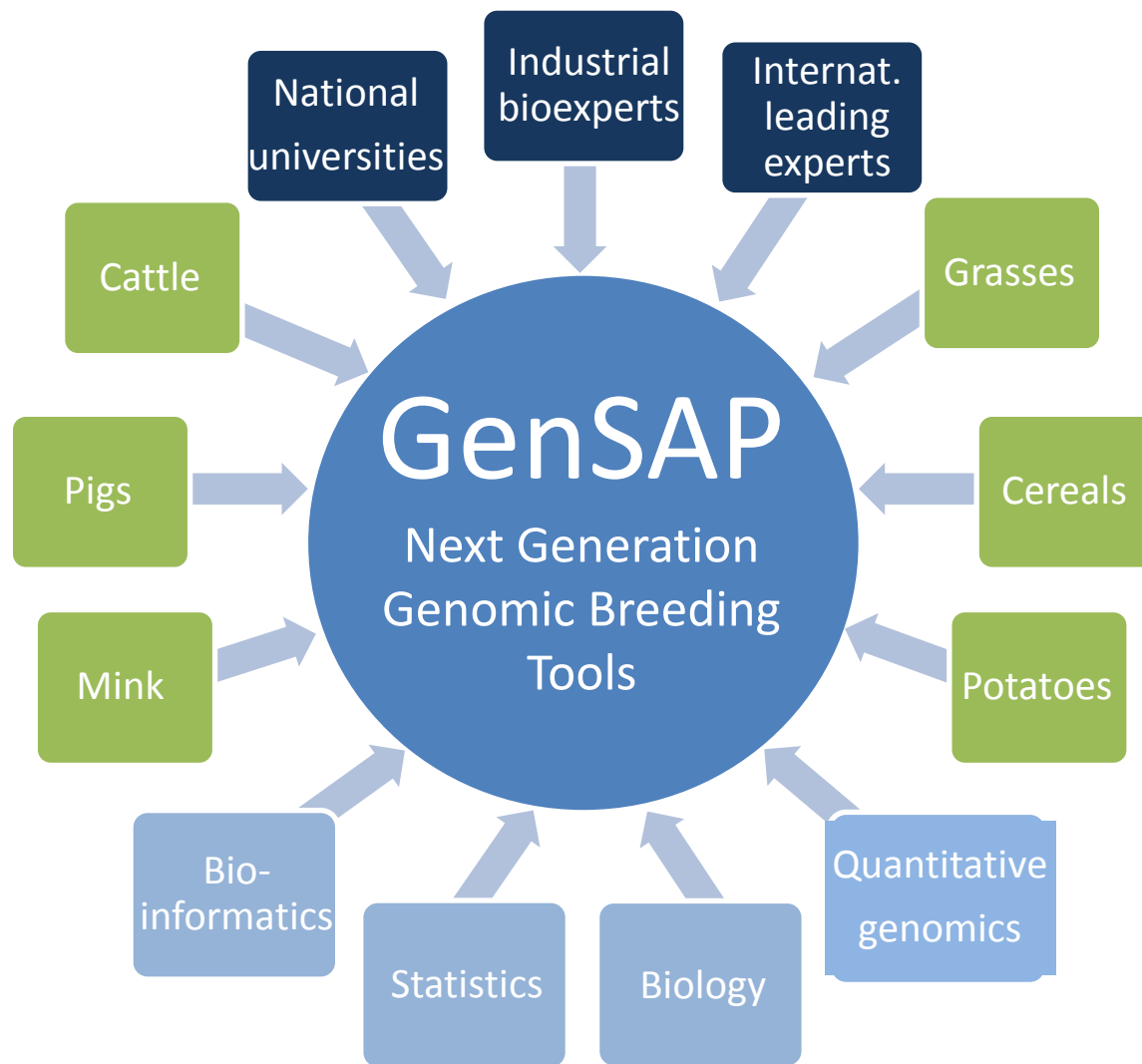
GenSAP

Next Generation
Genomic Breeding
Tools



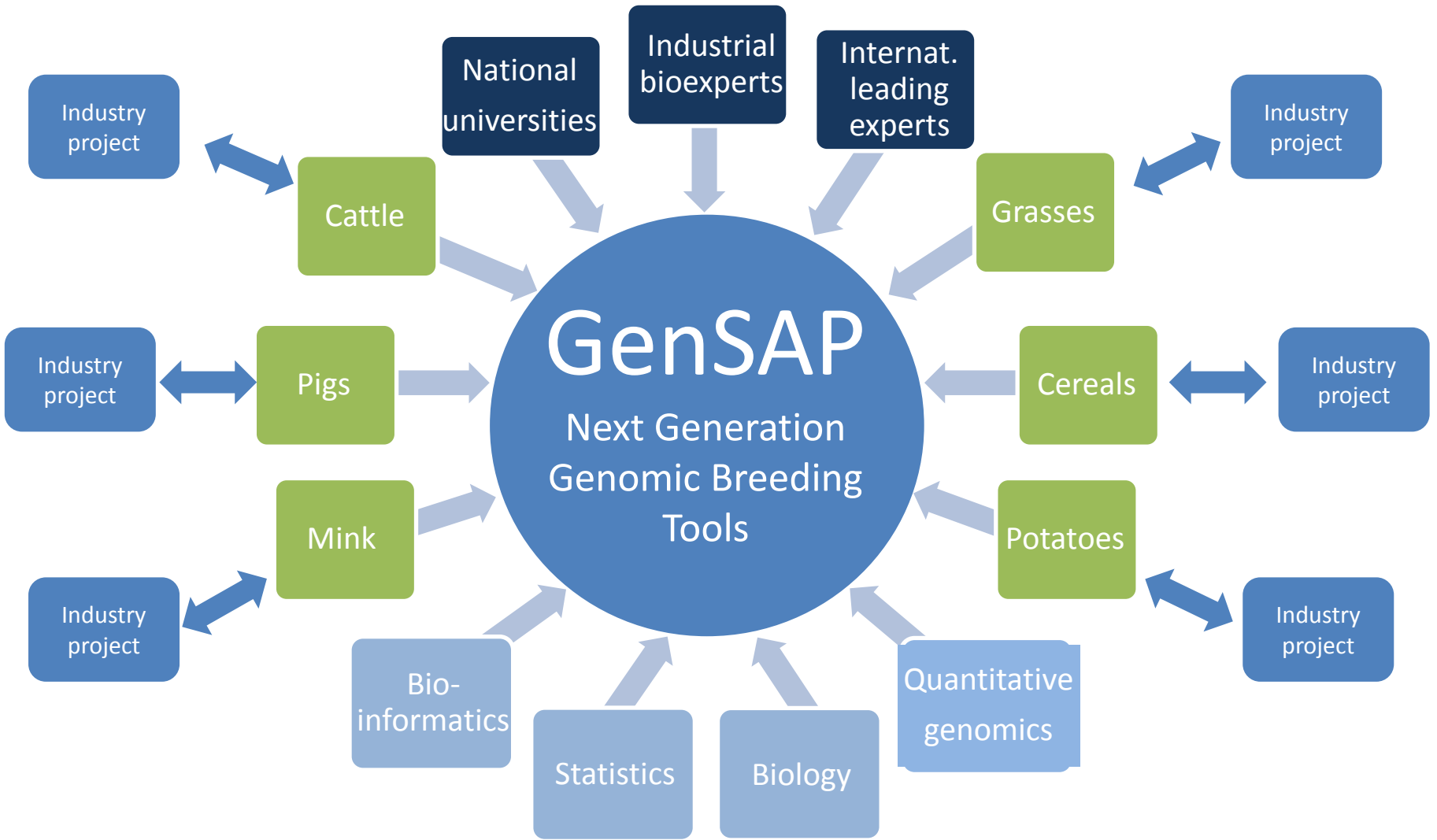
Animal and plant species







Synergy with other projects





Challenges

- Use full sequence information
- Genomic prediction in huge scale data
- Genomic prediction across breeds and populations
- Non-additive genetic effects
- Generalisation to plant species
- New genomic measure of inbreeding



GenSAP

SFA1

Genetic
architecture of
complex traits

SFA2

Genomic
prediction
models

SFA3

Assesment and
optimisation tools



GenSAP

Species

Universities

Industries

SFA1

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architecture of
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