



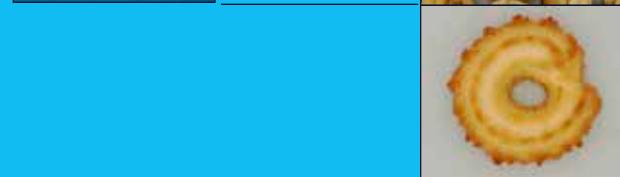
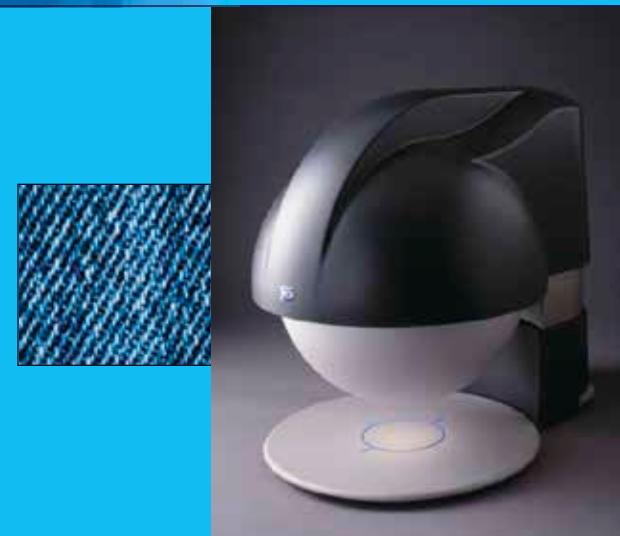
SpectraSeed: Seed phenotype database through spectral imaging

Jens Michael Carstensen

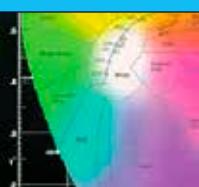
CTO, Videometer A/S

Assoc. Prof. DTU Compute

Affiliated professor of organism imaging,
University of Copenhagen



Coworker acknowledgement
Karsten Hartelius, Videometer A/S
Kåre Jensen, Videometer A/S



SpectraSeed



Funded by:



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UNIVERSITY



DLF TRIFOLIUM
SEEDS & SCIENCE



westrup



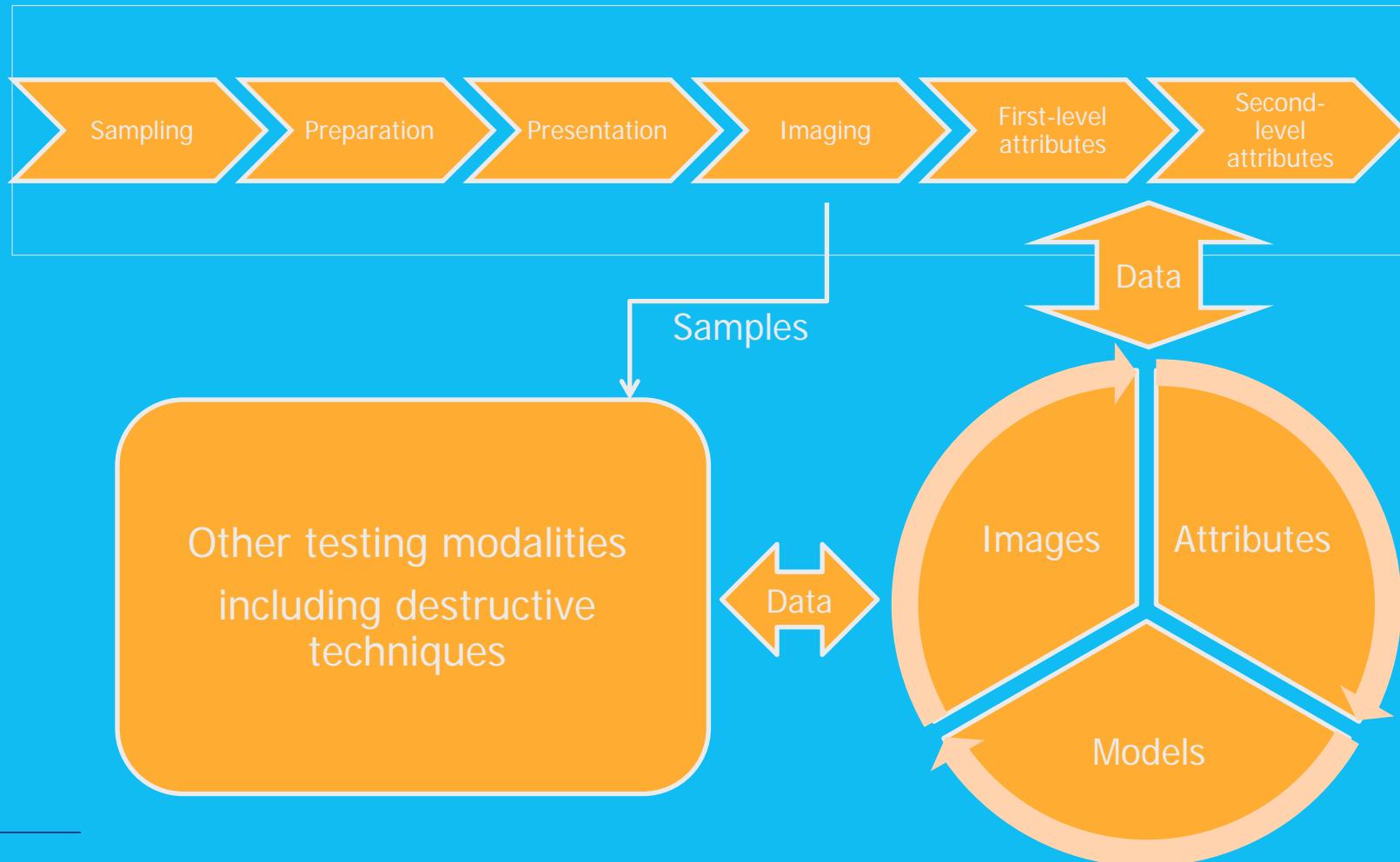
Jensen Seeds A/S





Concept

Spectral imaging seed testing protocol



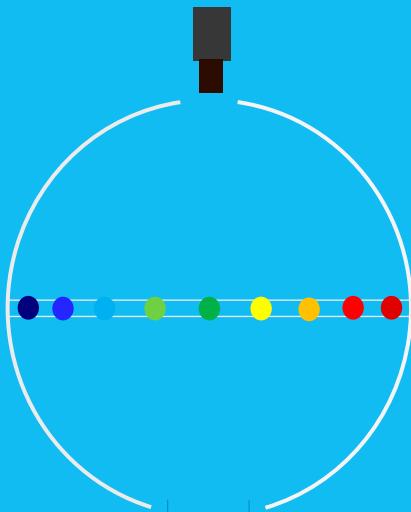


Use case in malting industry





Videometer BSQ Spectral Imaging



Camera
Lens
Integrating sphere
LEDs of multiple wavelengths
Sample is placed in target opening

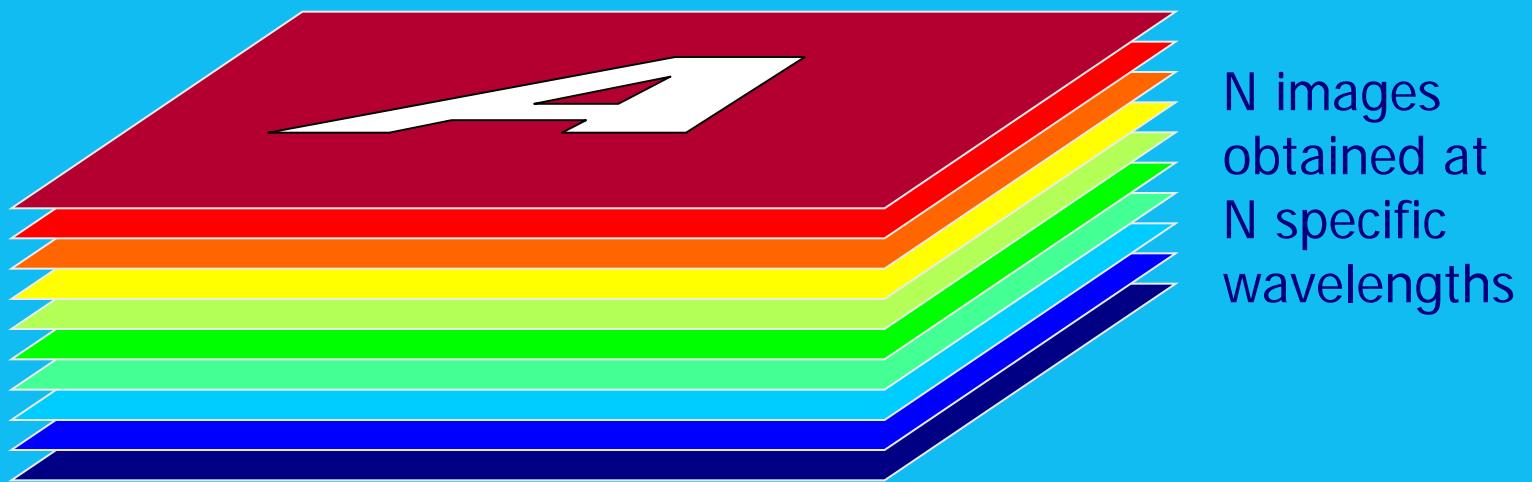
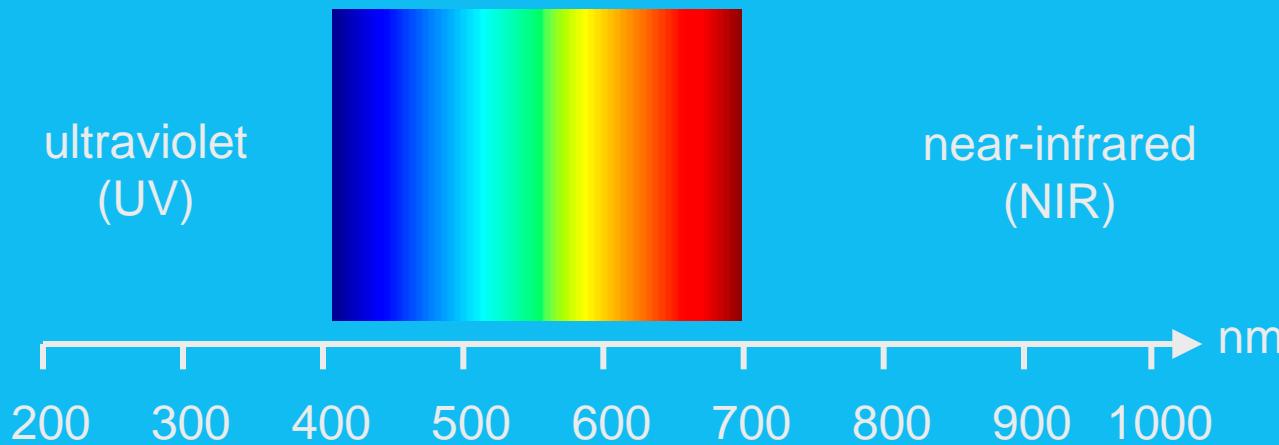


- LEDs: Stable, durable, large selection, rapidly developing technology
- Up to 20 different high-resolution bands acquired sequentially in 0.5-1.5 seconds depending on camera
- May be combined with emission filters, backlight, and darkfield illuminant





Band-sequential spectral imaging





Imaging

- Rapid, non-destructive measurement
- 10-20 seconds per sample including handling
- May easily be combined with other measurements – even destructive techniques
- High versatility measurement
- Focus on
 - Reproducibility
 - Traceability
 - Robustness
 - Transferability





First-level attributes

Spectral and morphological seed characteristics

- Size
- Shape
- Color
- Topographical texture
- Spectral texture
- Spectral components related to surface chemistry
- Count





Second-level attributes

- Seed purity
- Germination percentage
- Germination rate
- Seed vigour
- Seed health
- Seed maturity
- Seed longevity
- ...





On-line phenotype database

- Seed images
 - Tagged with origin data
- First-level attributes
 - Calculated from spectral images
- Second-level attributes
 - Measured by a number of different techniques including destructive techniques and subjective scoring
- Modelling algorithms to link first-level and second-level attributes
- Background services: sanity check, novelty detection etc.





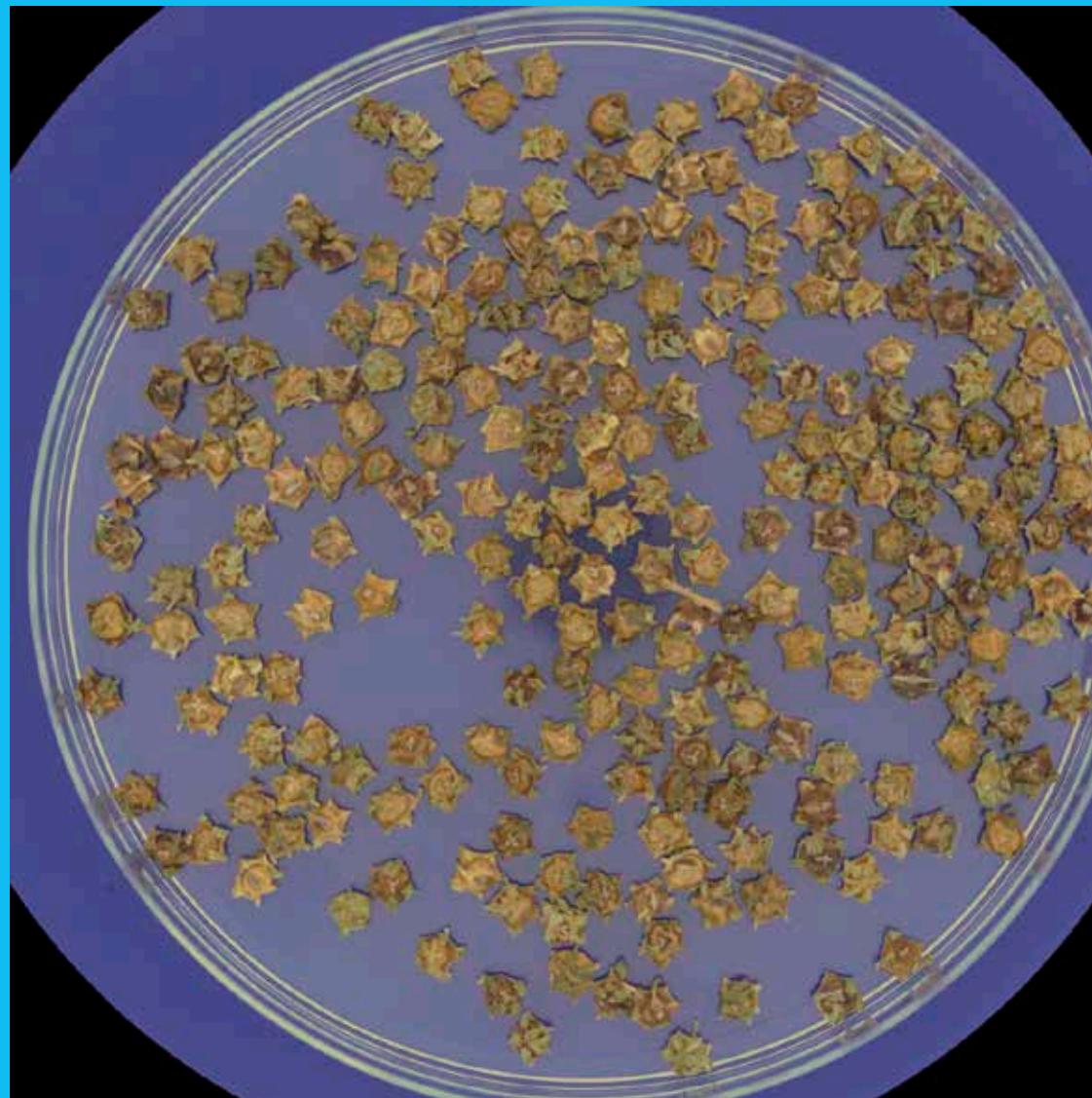
Database use cases

- Data support for parameterized generation of first-level attributes e.g.
 - Segmentation and identification of single seeds in a single layer presentation with touching seeds
 - Feature generation adapted to seed orientation like germ end orientation, dorsal/ventral/side presentation
- Dynamic generation and updating of prediction models for second-level attributes. These are based on a data support selected on e.g. crop, season, geographical region, instrument, data supplier and user.
- Data mining resource



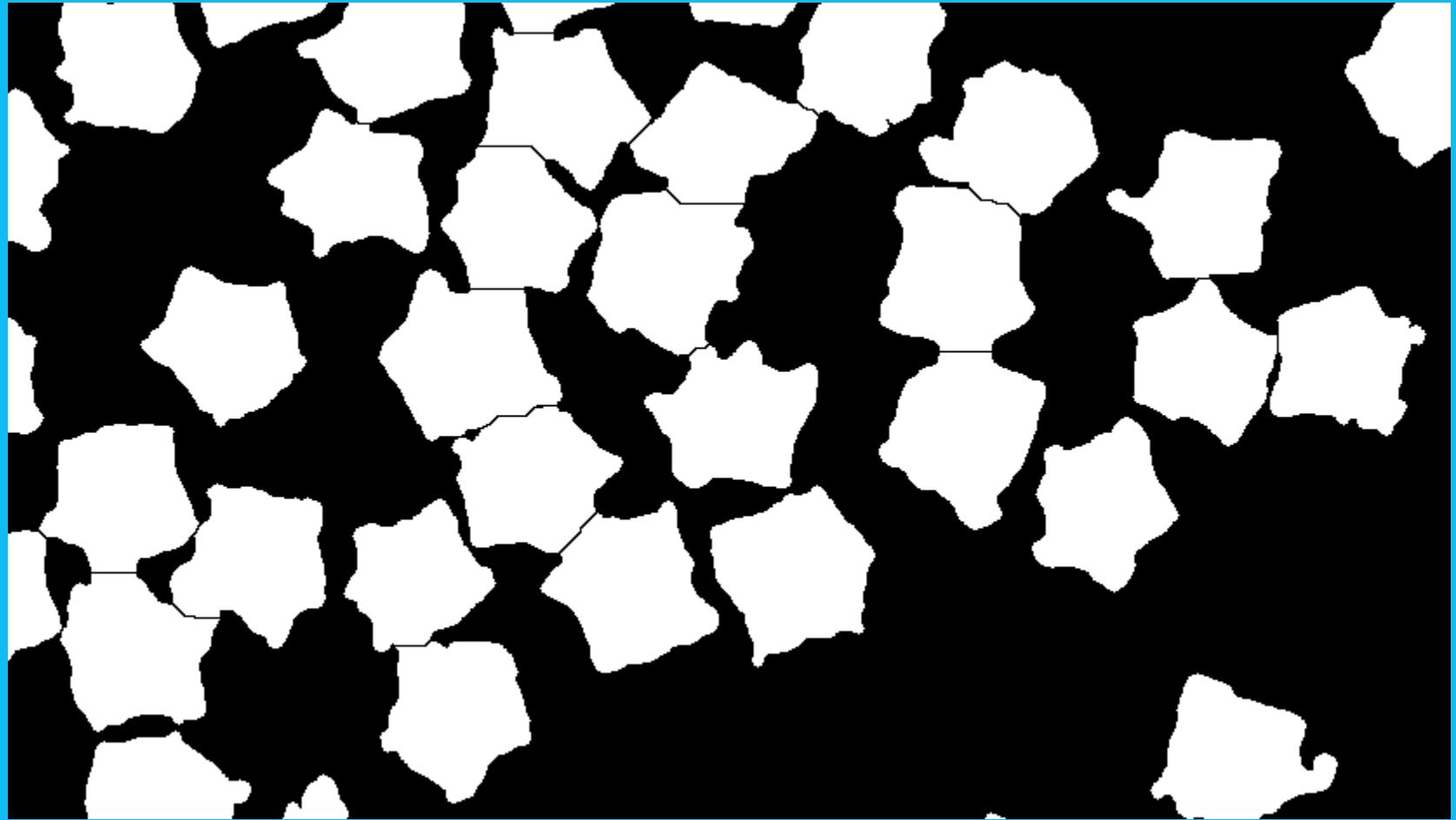


Sugar beet seed example





Sugar beet single seeds segmented

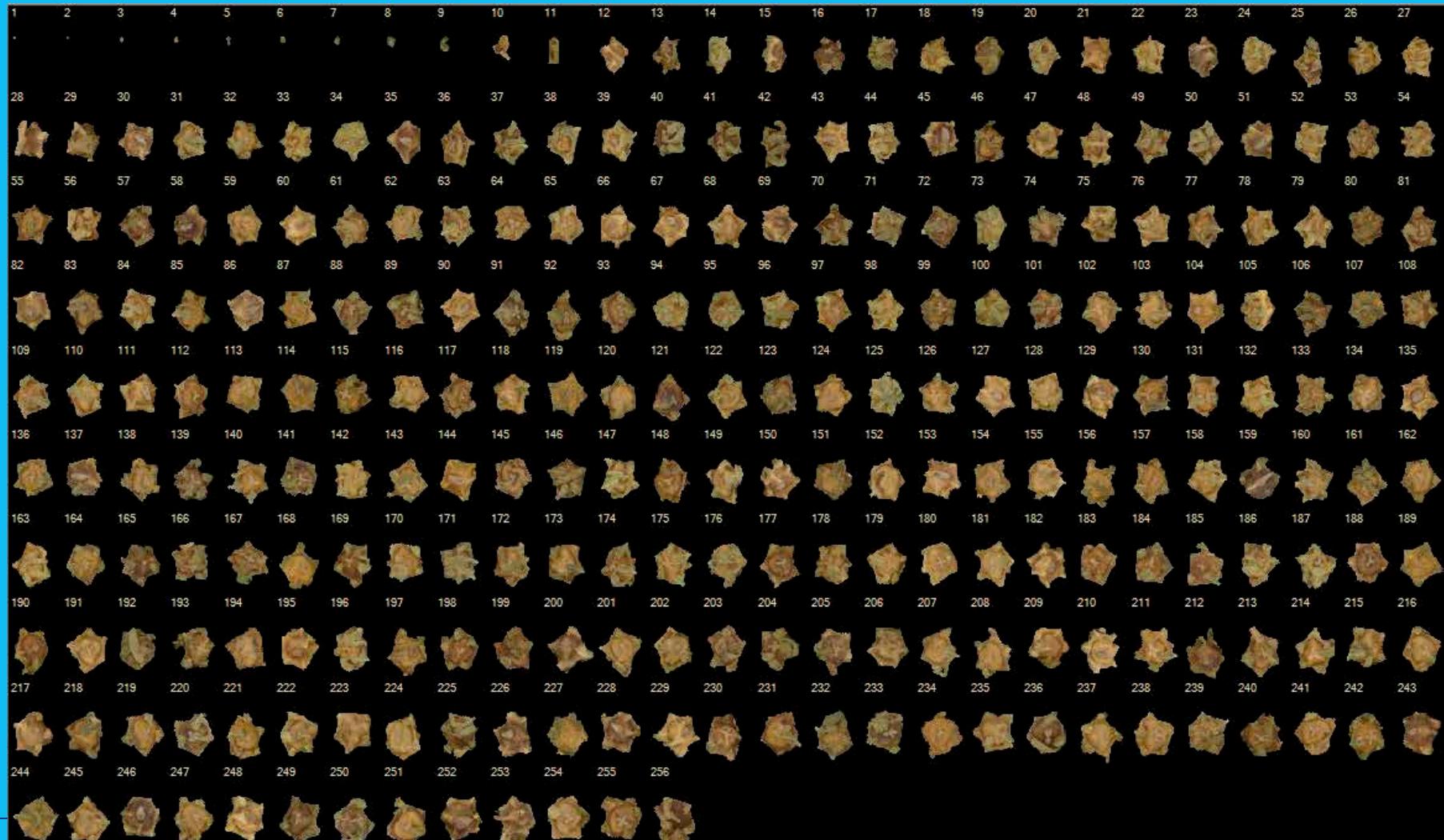


Segmentation is automated based on seed characteristics drawn from the database





Seeds sorted by area

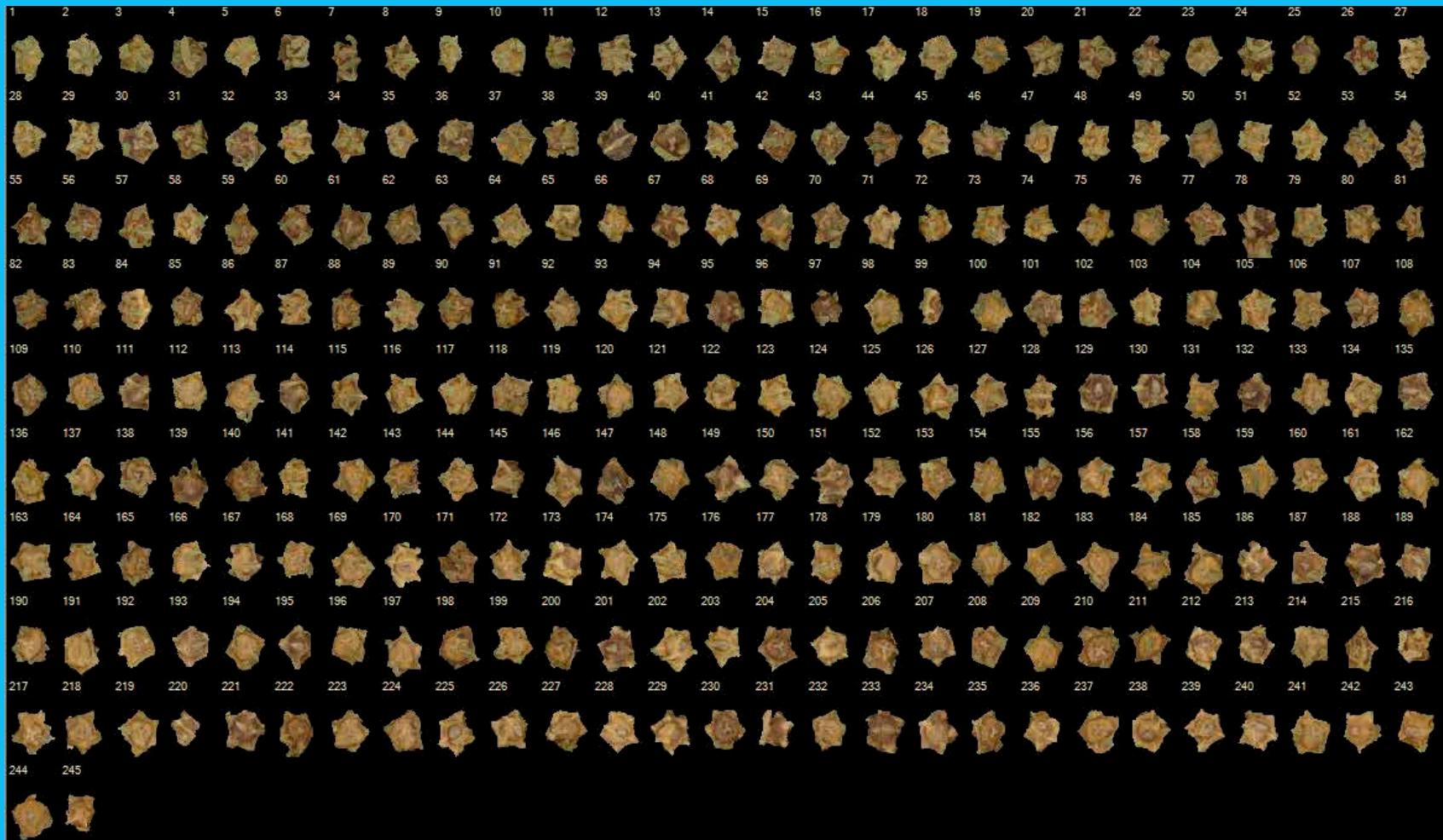


First-level attribute





Seeds sorted by chlorophyll/maturity

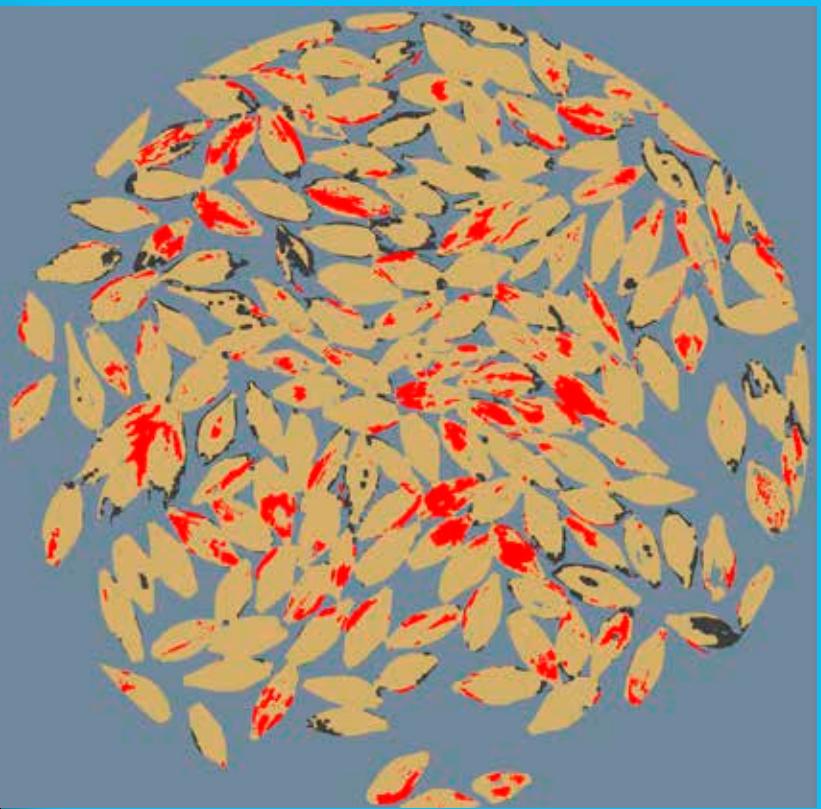


Second-level attribute





Seed health: Fusarium and gray molds detection



The red color = red, orange or purple area on kernels

The dark gray color = grey and black molds

Brownish = Barley without molds



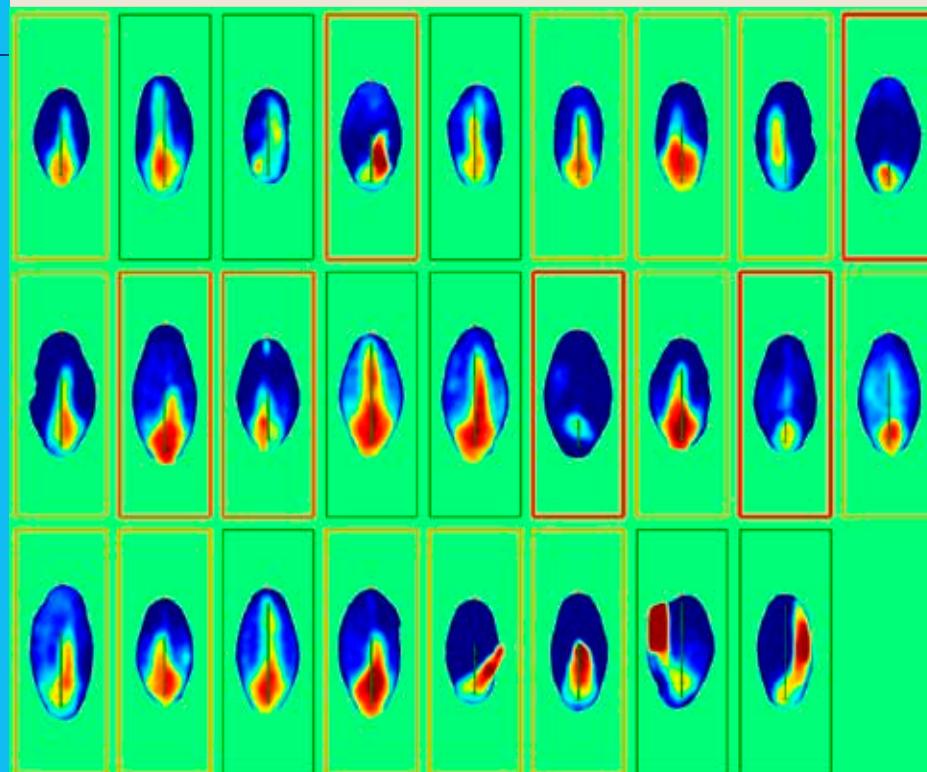


Seed germination: Acrospire length

Session: New Acrospire length bm 2003

Recipe: New Acrospire length bm 2

Plan: No Plan



| Sample ID | Note | #Total | Mean Relativ | #Group1 | #Group2 | #Group3 | #Group4 | #Gro |
|-----------|------|--------|--------------|---------|---------|---------|---------|------|
| Prøve0_1 | | 25 | 0,575 | 2 | 5 | 15 | 3 | 0 |
| Prøve0_2 | | 26 | 0,5576923 | 0 | 10 | 13 | 3 | 0 |
| Prøve0_3 | | 25 | 0,585 | 0 | 9 | 11 | 5 | 0 |
| Prøve0_4 | | 26 | 0,6298077 | 0 | 7 | 13 | 5 | 1 |
| Prøve1_1 | | 26 | 0,677884638 | 1 | 6 | 6 | 13 | 0 |
| Prøve1_2 | | 26 | 0,7692308 | 0 | 0 | 11 | 15 | 0 |
| Prøve1_3 | | 26 | 0,7355769 | 0 | 2 | 12 | 11 | 1 |
| Prøve2_0 | | 26 | 0,7019231 | 0 | 1 | 16 | 9 | 0 |
| Prøve2_1 | | 26 | 0,6923077 | 0 | 1 | 17 | 8 | 0 |
| Prøve2_2 | | 26 | 0,721153855 | 0 | 3 | 10 | 13 | 0 |
| Prøve2_3 | | 26 | 0,711538434 | 0 | 2 | 13 | 11 | 0 |
| Prøve2_4 | | 26 | 0,7355769 | 0 | 2 | 12 | 11 | 1 |



Resume

Finish Session





Conclusions and outlook

- Seed phenotyping is BIG DATA
- Systematic and automated generation and updating of models linking first-level and second-level attributes are necessary
- SpectraSeed plan:
 - 2013 Local databases running specific applications
 - 2014-2015 Parameterized analysis, automated modelling, more applications
- Launch of SpectraSeed on-line database

