

A plant breeding landscape

RESEARCH & DEVELOPMENT

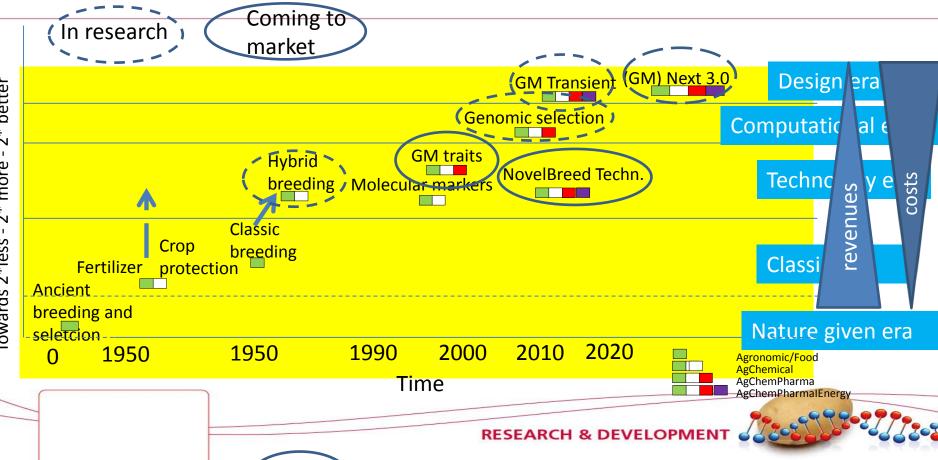
To come

- Crop improvement in time
- Breeder's limits and needs

RESEARCH & DEVELOPMENT

- Suggestions for science
- NBTs
- Durable resistance

Crop improvement in time



Introduction HZPC Group



- Since 1898; origin Netherlands
- > Number of employees: 262 (FTE)
- Sales of seed potatoes 652.000 ton
- > Export/license ca. 75 varieties to 82 countries
- Sales of table potatoes 180.000 ton
- ➤ Turnover €275 mio; gm <15%; profit ca €5 mio.</p>
- Shareholders: staff growers breeders

RESEARCH & DEVELOPME



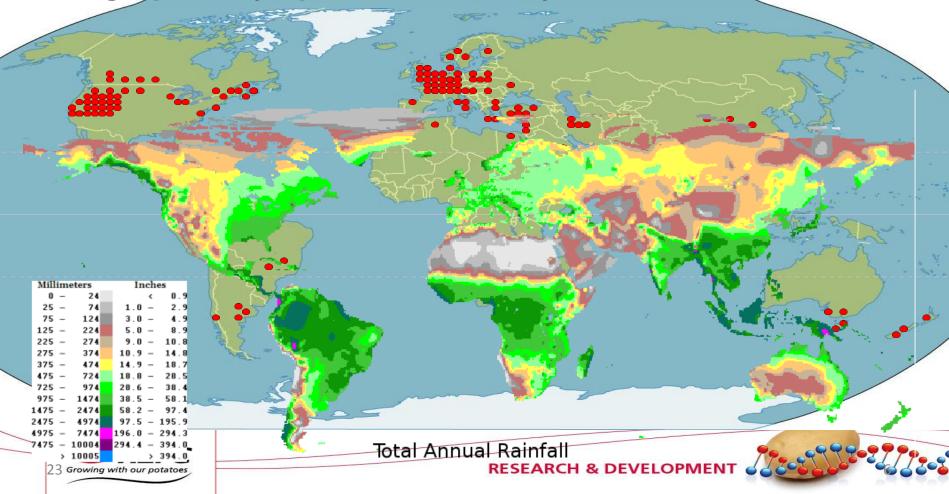
Market success is a 4-fold balance



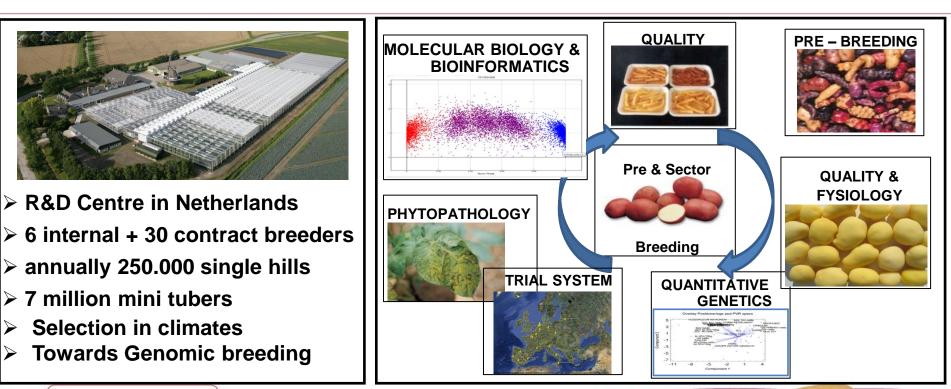
- Achieved by breeding and marketing (60/40; 40/60 ~ 50/50)
- Breeding: about control of heritable traits or performance and speed of recombination

RESEARCH & DEVELOPMEN

Geographically spread French Fry Factories



HZPC R&D





RESEARCH & DEVELOPMENT

Breeder's limitations

- 40.000 genes * 15 effective alleles is 600k of influences + epigenetical
- Challenge to make the best choice in 600k * 600k opportunities, fixed in series of quadruplex sets of performance.
- 100k potential crosses at present; <1000 are selected by phenotype, offspring, markers and statistical approaches.
- Potato breeding is still subjective at start; rely on yield measurement starts when <3% of initial genetic variation ('single hills') is left
- 8 yrs to identify the variety, 3 yrs to convince added value to customer and produce seed potato volume

RESEARCH & DEVELOPME

Breeder's needs

Become more objective with an effective start at crossing and single hill: Baseline is available

- Phase 1: improved by MAB for resistance Pi R genes, (cyst) nematodes, virus and more; supported by validation protocols for diseases, objective phenotyping, additive genes in a good germplasm background
 running and achievable, next,
- Phase 2: grow towards genomic breeding based on performance associated series of SNPs, control allelic contributions, supportive data systems.

RESEARCH & DEVELOPM

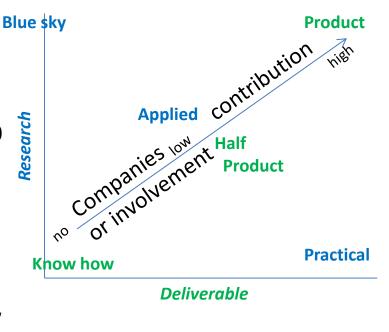
• Phase 3: ...

What's needed in/from collaboration with science:

Suggestions/challenges:

more

- Breeders that can handle the output
- Affordable FTO; more tools is less per unit
- Tuberosum reference genomeSSSSSS
- Broad Data handling systems
- Physiology know how (close the gap towards >100 structural tons/ha)
- Low-no mistake sequencers or an array for all important alleles, or both
- Non gm NBT's to solve/fixation simple traits
- Innovative approaches to unravel complex traits
- Potato companies understand your work better so Take big steps, Think big, Act collaborative, Focus, Achieve goals in time, Communicate active, Catch spinn-off



RESEARCH & DEVELOPMEN

Complexity to come to a logic predictable performance

60-70% GENOTYPE -	15-25% ENVIRONMENT	15-20% + MANAGEMEN ⁻	100% T => PHENOTYPE				
Controlled input	Variable component	modelling / DSS	Interactive result				
DNA sequence, genotyping	Climatic data Consumer	Physiology Control	Field performance				
by sequencing, genomic data	preferences	DSS	Plant architecture, growth rate, development, etc.				
Epigenome	Detailed metadata	QC's	Physiological status				
M. d. L. d			Proteins				
Methylation		Storage	Metabolites				
RNA: arrays \rightarrow RNA:	seq		Other compounds				
micro RNA's							

.....

RESEARCH & DEVELOPMENT

HZPC Challenges (at random)

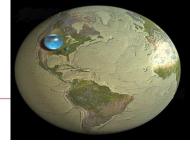
- Feed the world, locally (food security, carbon print)
- Availability sweet water (salinity)
- High production on poor soils (-2 input, +2 output, better food)
- Raw product storage at low risk (resistances, physiology)
- Low-no losses in the value chain (now 40% food waste)
- Climate extremes within growing season (physiolology)
- Secure an affordable legal ownership (build a world wide business)
- Fair profit for future developments (People Planet Profit, Corporate Governance)
- Consistent product on marketable yield & quality (measure, monitor)



Potato is a great carrier to serve all items!

Staple crop nr 4 now,

Potential to move up to 3! RESEARCH & DEVELOPMENT



Novel Breeding Technologies





Novel breeding technologies

- 1. Mutation 'like': SDN1-3, ODM, TALEN, CRISPr, RTDS; techniques can knock out 1 bp up to introduction of genes, protein based or mimic natural repair mechanisms
- 2. Virus induced gene silencing: (transient) gene silencing mostly used for scientific complementation
- 3. Cisgenesis: A. tum introduced, only native crossable genes, <20 bp new
- 4. Grafting: non gm grafted on gm

8.

- 5. Reverse genetics: redesign parental lines from a hybrid, contains a recombinant step to suppress the meiotic phase
- 6. RNA dependent DNA methylation; epigenetic change induced by dsRNA methylation

RESEARCH & DEVELOPME

 Agroinfiltration: local expression of foreign genes, parts of plant can be used for cuttings etc.

EU

- Important to know that EU2001/18/EC is process ÁND product based!!! Also the Cartagena protocol is Product and Process based
- Article 2(2) Definition GMO:
 - Part 1: GMO means an organism, with the exception of human beings, in which the genetic material has been altered in a way that does not occur naturally by mating and/or ...= PRODUCT based
 - Part 2: genetic modification occurs at least through the use of techniques listed in Annex 1A part 1 and the techniques listed in Annex 1A part 2 are not considered to result in genetic modification= PROCES based
- Article 3 is about the application of a gm technique to an organism but will not lead to a GM, listed in Article 1B.
- Legal landscape: scientific base is is <40%, EU lawyers must be comfortable when taken to court is >60%

RESEARCH & DEVELOPMEN

Durability of resistance

Definition: a resistance that holds over many years of agricultural use in various environments, challanged by pathogen recombination and selection in time.

CF9 (downey mildew tomato), Mlo (downey mildew wheat), H1 (Potato cyst nematode), Ve1 (Verticillium's tomato)

- Cf9: appears to be 2-3 genes, not one; mimic by Cisgenesis
- MIo: Susceptibility genes knocked out, hard to replace; reverse mimic by HIGS
- H1: highly effective, still to be cloned and learn why durable
- Ve1: apparently gene for gene response with Ave1 Achilles heel



Potato varieties with Pi R-genes

- Innovator, Santé, Escort, Toluca, Bionica, Carolus, Sarpo Mira, Biogold, Fresco, Connect, Raja, Athlete, Coquine, Voyager, Janine, etc..
- About 10 different R-genes from ca. 5 species are or have been present in listed potato varieties

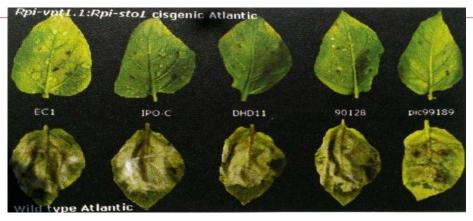
Aim: Durable Disease Resistance

"Resistance that remains effective over long periods of widespread agricultural use"



RESEARCH & DEVELOPMEN

Cisgenic ~ mimic durable Cf9 / wild species approach



2013, Kwang Ryong Jo, Unveiling and deploying durability of late blight resistance in potato, thesis.

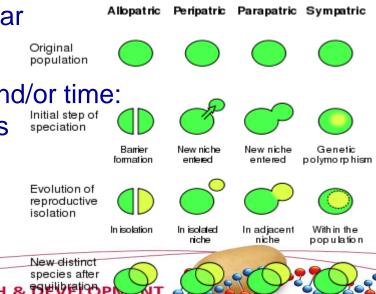
Year	Genotype	R-gene content	Delay in d	lavs	8	i r	ela	ati	VE	د
2011	Bintje	-	2	а	100		+	100	240	-
	3151-3	9b	9	6	ь	C	1	100	1	
	MaR2	2	12	-	b	C	d	25		
	3020-18	8	15			C	d	e		
	edn7727-104	edn2	19		10	-	d	e		100
	edn7727-148	10, edn2	22		46		d	e	+	14
	3025-53	3a, 3b, 4, 8, 9a, 9b?***	19		411		d	e		
	Toluca	blb2	18		+	+	d	e	+	
	3025-1	abpt, 3a, 3b, 4, 9a, 9b?***	23		2	-	d	e	-	-
	3025-43	3a, 3b, 8	19	¥65		- 30	d	e	10	
		3a, 3b, 4, 8, 9b?***	36			*	100	e	f	10
	Sarpo Mira	3a, 3b, 4, smira1, 8	44		-	-		e	F	
		3a, 3b, 4, 8	43		10		4	e	F	
			40			S.	1	e	f	-
			40		1		10	e	f.	-
	edn//2/-100	1, abpt, 3a, 3b, 4, 8, 9a, 9b	114	107			100		F	a
	MaR9 edn150-4	edn1, 4, edn2	No lesions**	*	4		1		-	g

Durable resistance: Nature will win, Varieties should hold long ..

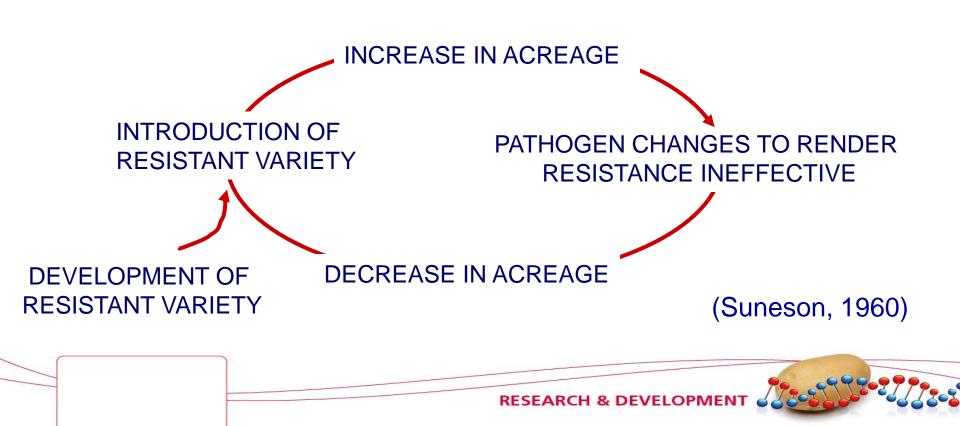
RESEAR

Strategies to enhance durability of resistance

- Pyramid/stack multiple genes increase evolutionary hurdle
- Utilize 'durable' or 'non-host' resistance
- Match life expectancy of resistance and cultivar
- Diversify selection pressure on pathogen
- heterogeneity of resistance genes in space and/or time:
- diversify resistance sources, host & non-hosts
- pipeline with different resistance genes,
- shaping syn- and allopatric evolution,
- multilines and cultivar mixtures.



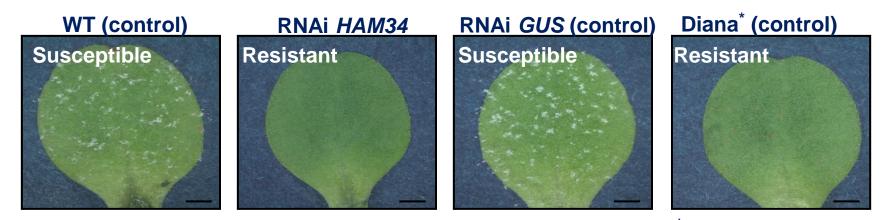
THE 'BOOM-AND-BUST' CYCLE OF DISEASE CONTROL The Agricultural Consequences of Pathogen Evolution



Host Induced Gene Silencing

Of six genes tested, two (HAM34 & Cellulose Synthase) resulted in inhibition of *B. lactucae*:

Transgenic T₃ lettuce seedlings expressing RNAi *HAM34* are resistant to *B. lactucae* (8 dpi)



Kindly provided by: Richard Michelmoore - Manjula Govindarajulu http://michelmorelab.ucdavis.edu/

