Rothamsted Research where knowledge grows

Nutrient use efficiency in wheat What are the challenges? Malcolm J. Hawkesford

Crop Science Workshop, May, 2014



- Yield and NUE
 - Link between yields and NUE
 - Genetic variation and scope for improvement
- Inverse relationship of grain protein and yield
- Quality: micronutrients
- Some opportunities
 - Exploiting more diverse germplasm diversity
 - Improved high throughput phenotyping





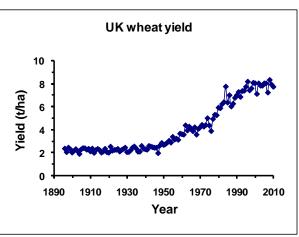


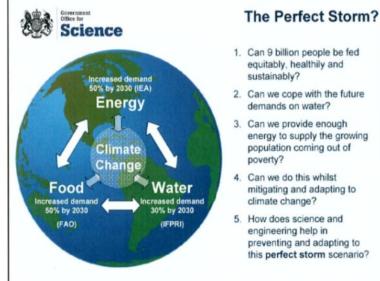


Goal: food security via sustainable increases in yields

- Double production by 2050
- Population increase
- Climate change
- Land pressure
- Water availability
- Cost/availability of fertiliser
- Plateauing yields



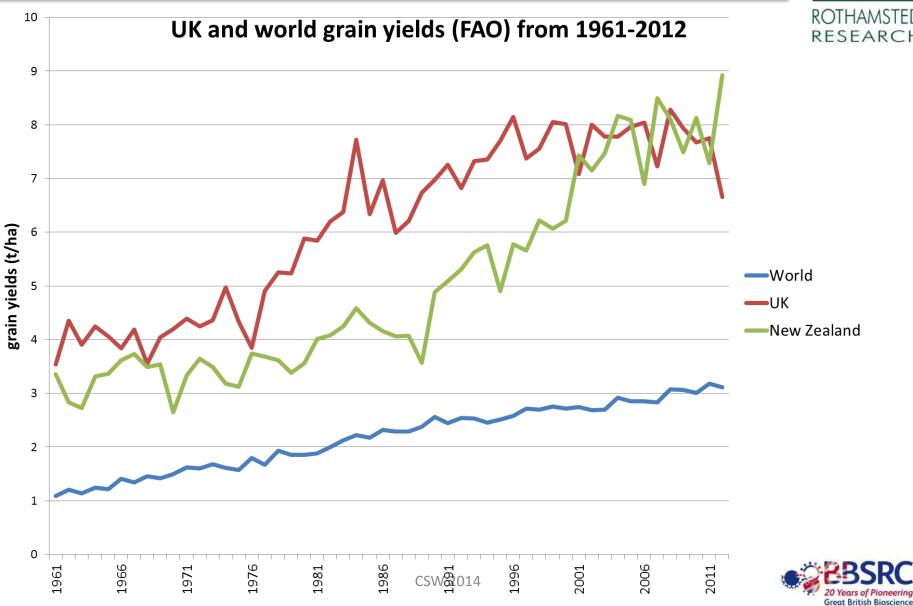






Content of Pioneering Great British Bioscience

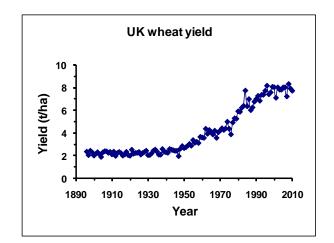
Trends in wheat yields (farm gate)





First green revolution

- Norman E. Borlaug
 - Dwarfing genes
 - More nitrogen
- Some issues
- Next green revolution?
 - ➢ Roots
 - Primary production



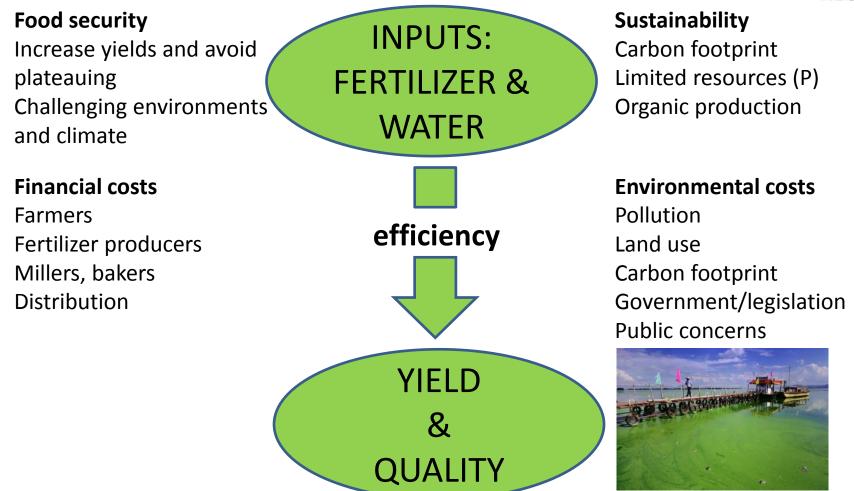






Why do we need NUE?







Target traits YIELD + ROTHAMSTED RESEARCH Nutrient use efficiency NUE= Carbohydrate & GPD NUpE x NUtE nutrient content) (yield/N_{av}) NH Senescence **BIOMASS** Nutrient-export Photosynthesis Canopy as a **NUtE** transient store = Yield/Nutrient taken up Root **Architecture** NUpE establishment Proliferation Activity = Nutrient taken up/Nutrient available

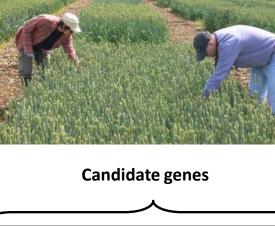
Great British Bioscience

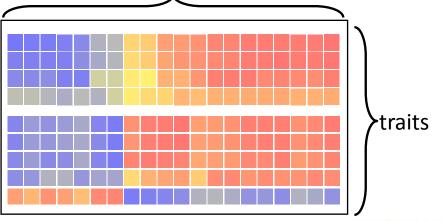
Key approach: genes correlating to NUE traits

- Trait de-convolution and prioritization
- Assessment of variation
 - Provision of data for breeding
 - Aid new gene discovery
- Identification of genes/markers
 - Transcriptomics
 - Correlation with traits
 - Mapping populations



• Breeding or biotechnology









NUE and diversity

Why are diversity studies needed?

- Assess current extent of variation (WGIN)
- Wider screening may be necessary if not enough variation in current germplasm
- Bottlenecks, recent and ancient
- Harness 'lost' genes and alleles, a particular problem for some sustainability traits
- Modern selection under high inputs

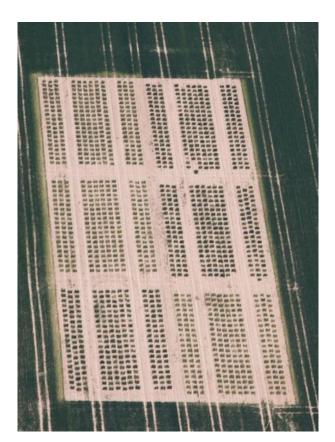
What are the problems?

- Finding the right germplasm
- Genetics and Phenotyping
- Metrics when comparing very different materials

WISP

- <u>http://www.wheatisp.org/</u>
- Watkins, Gediflux, SHW and modern cultivars
- Derived mapping populations









WGIN: Wheat Genetic Improvement Network







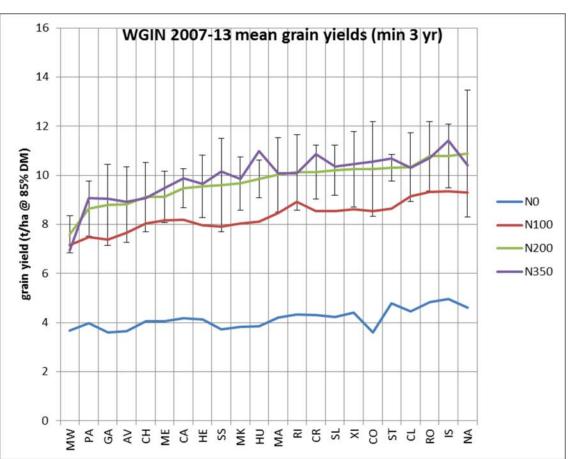
Wheat Genetic Improvement Network



WGIN: The Nitrogen-Diversity trial



- 2004-13
- 51 varieties
- 14 in at least 9 years
- All 4 groups
- 4 N levels in all except
 2 years
- Grain and straw, yield and %N
- Archived fresh grain
- Archived dry milled grain and straw
- Many spin-off projects



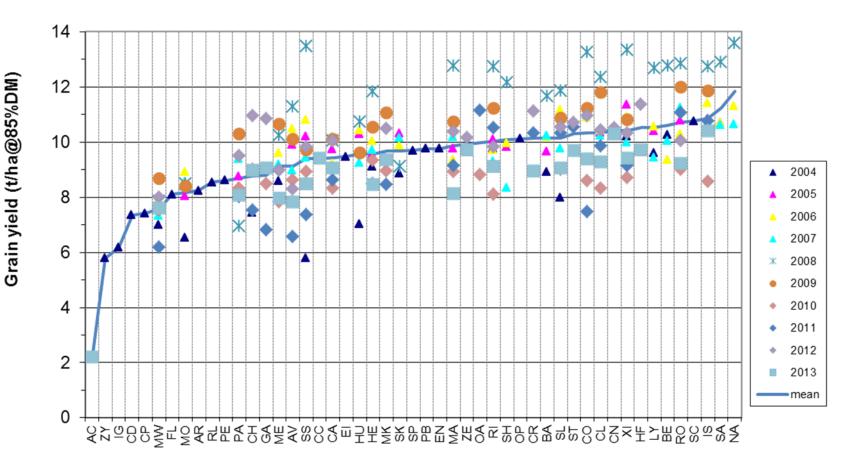


WGIN – yield stability

Rothamsted WGIN-N200

Combine Grain Yield (2004-13)

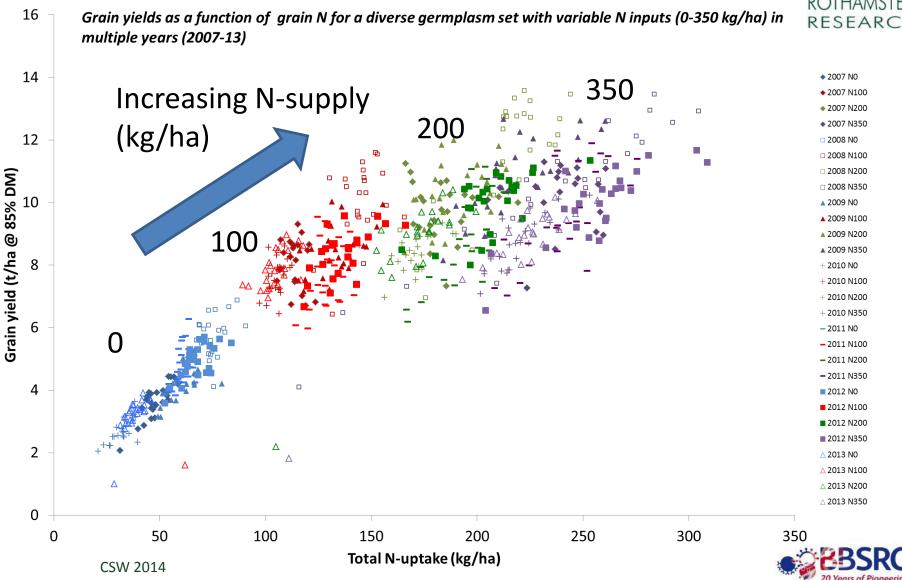






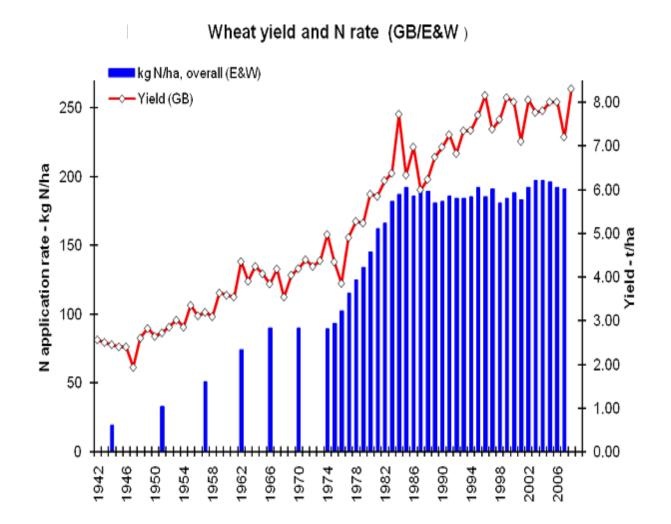
Variety

N-supply impacts on yields and quality





N fertiliser use trends in the UK

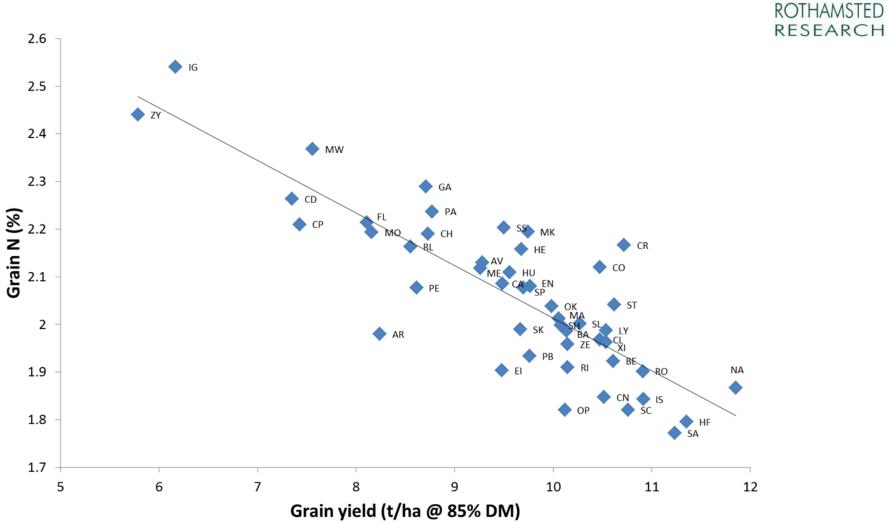


Hawkesford (2014) J Cereal Sci



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Grain protein deviation

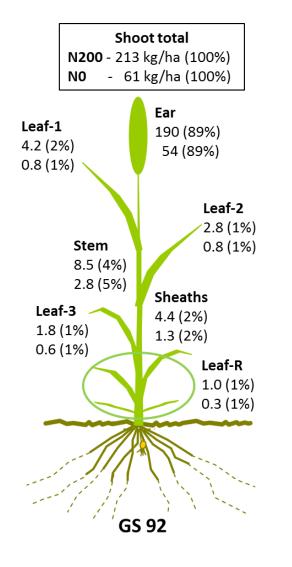


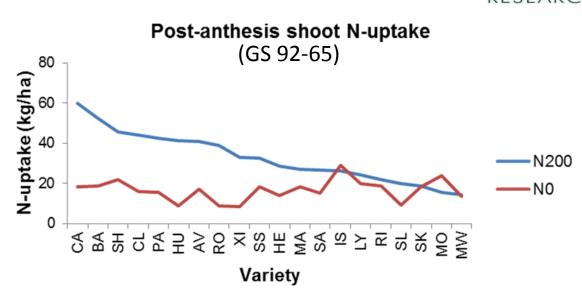


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Post anthesis N uptake







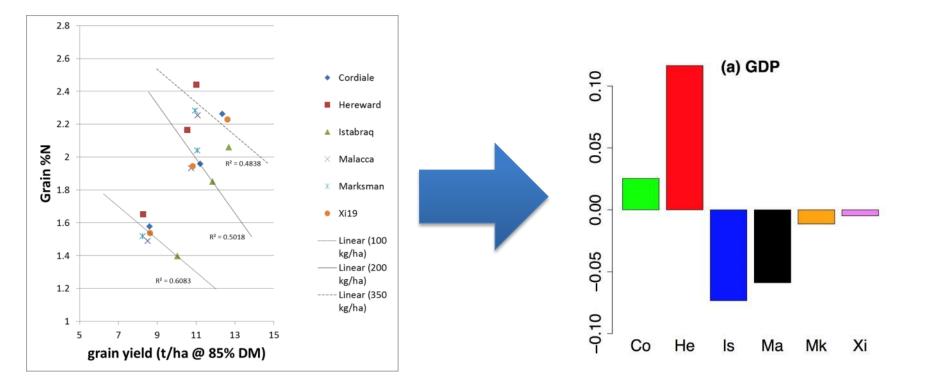


Genotypic variation in the uptake, partitioning and remobilisation of nitrogen during grain-filling in wheat $^{\rm th}$

Peter B. Barraclough*, Rafael Lopez-Bellido¹, Malcolm J. Hawkesford Plant Biology and Crop Science Department, Rothamsted Research, West Common, Harpenden, Hertfordshir e ALS 200, UK



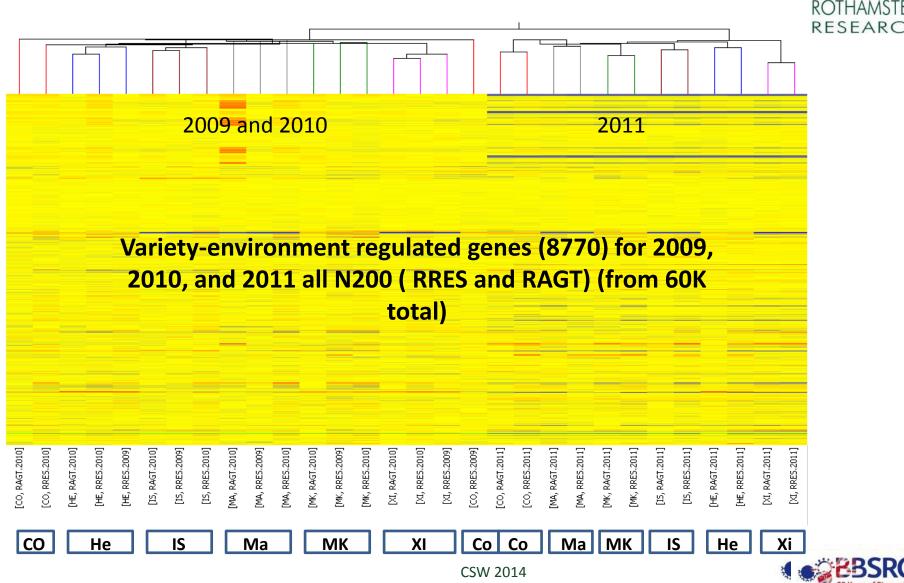
Normalising datasets







Transcriptomics

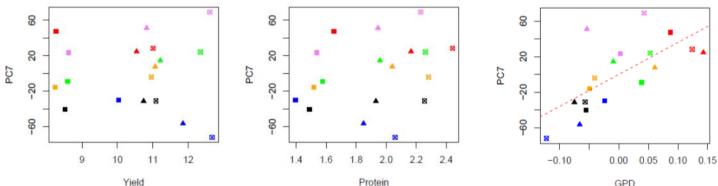




ANOVA – correlating the Affymetrix gene expression PC analysis to trait data for 2009



	Yield	Protein	Yield corr N	Protein corr N	Protein corr N & Y
	Tielu	Protein	Tielu con N	Protein com N	Protein con N & T
Mean PC1	0.078	0.031	0.740	0.519	0.581
Mean PC2	0.065	0.991	0.053	0.005	0.043
Mean PC3	0.152	0.460	0.008	0.006	0.129
Mean PC4	0.736	0.682	0.139	0.911	0.421
Mean PC5	0.362	0.037	0.000	0.015	0.643
Mean PC6	0.357	0.195	0.823	0.572	0.415
Mean PC7	0.853	0.063	0.199	0.000	0.000
Mean PC8	0.071	0.034	0.431	0.636	0.285
Mean PC9	0.314	0.446	0.080	0.067	0.297
Mean PC10	0.000	0.000	0.447	0.088	0.010

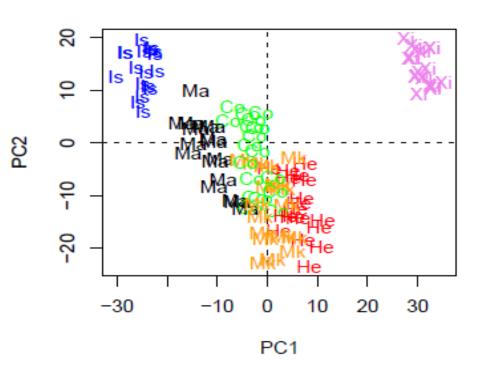


GPD



PCA of the selected GPD genes



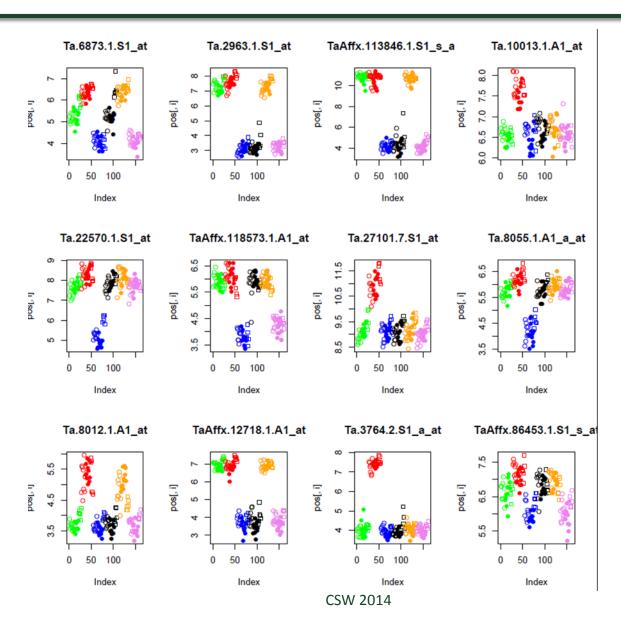


(a) Score plot

Partial Least Squares regression to further refine these list **resulting in 537 key genes defining differences in GPD between the varieties**



Validation of genes positively related to GPD







No varieties are perfect!

Variety Performance at 200 kg-N/ha (2004-08)

Variety	Code	Nabim	Years	Yield	%N	Uptake	Utilisation
Avalon	AV	1	5				
Flanders	FL	1	1				
Hereward	HE	1	5				
Hurley	HU	1	5				
Malacca	MA	1	5				
Mercia	ME	1	4				
Maris Widgeon	MW	1	5				
Shamrock	SH	1	4				
Solstice	SL	1	5				
Spark	SP	1	1				
Xi 19	XI	1	5				
Cadenza	CA	2	5				
Cordiale	со	2	3				
Einstein	EI	2	1				
Lynx	LY	2	5				
Rialto	RL	2	1				
Scorpion	SC	2	1				
Soissons	SS	2	5				
Beaver	BE	3	4				
Claire	CL	3	4				
Riband	RI	3	5				
Robigus	RO	3	4				
Istabraq	IS	4	4				
Napier	NA	4	3				
Savannah	SA	4	4				
Paragon (spring)	PA	1	5				
Chablis (spring)	СН	2	1				
Arche	AR	F	1				
Batis	BA	G	5				
Caphorn	СР	F	1				
Cappelle Desprez	CD	F	1				
Enorm	EN	G	1				
Isengrain	IG	F	1				
Monopol	MO	G	5				
Opus	OP	G	1				
PBis	PB	G	1				
Petrus	PE	G	1				
Sokrates	SK	G	5				
Zyta	ZY	Р	1				

Upper-Q Inter-Q Inter-Q Lower-Q

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Summary of variety performance (quartile rankings) based on 2004-07 WGIN datasets

EJA (2010) 33, 1-11

Europ. J. Agronomy 33 (2010) 1–11



Nitrogen efficiency of wheat: Genotypic and environmental variation and prospects for improvement

Peter B. Barraclough^{a,*}, Jonathan R. Howarth^a, Janina Jones^a, Rafael Lopez-Bellido^b, Saroj Parmar^a, Caroline E. Shepherd^a, Malcolm J. Hawkesford^a









Donor germplasm

Mapping populations

NILS

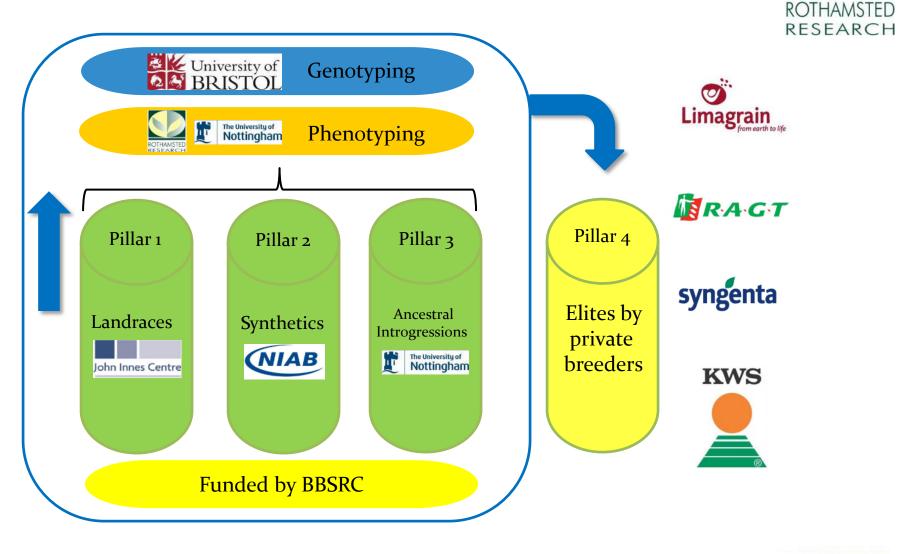
New synthetics

Introgressions





WISP (Wheat Improvement Strategic Program)

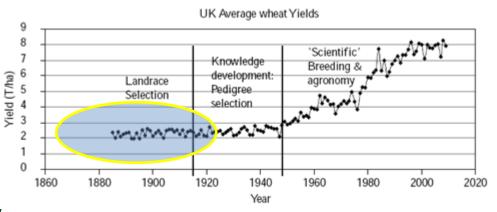




A E Watkins, University of Cambridge

- 1920's and 30's
- Board of Trade
- Farmers, markets and researchers
- Several thousand but now 136.
- 34 countries
- Held at JIC, duplicated in Australia
- Core genetic collection, c 120, plus other germplasm (Gediflux, synthetics) trialled at Rothamsted and Nottingham

Watkins diversity



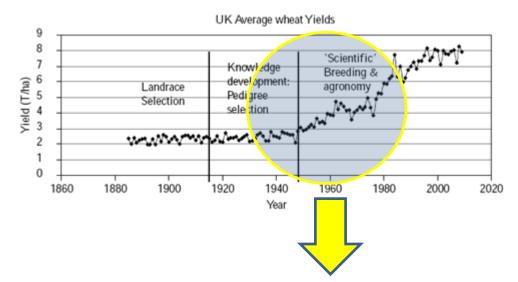


(graphics and map courtesy of S Griffiths & S. Orford, JIC)





Gediflux is designed to capture Western European winter wheat diversity since 1940



Country	Varieties	Years of release	examples
Austria	40	40-90	Tassilo (50s) Hubertos (90s)
Belgium	24	50-90	Norda (60s) Escorial (80s)
Germany	18	80-90	Calif (80s) Pegassos (90s)
E Germany	30	40-80	Mahndorf (50s) Kanzler (80s)
W Germany	19	50-90	Muck (50s) Borenos (90s)
Denmark	5	80-90	Anja (80s) Pepital (90s)
France	34	40-90	Vague d'epis (40s) Isengrain (90s)
UK	66	40-90	Holdfast (40s) Equinox (90s)
Netherlands	19	40-80	Lovink (40s) Nautica (80s)
Sweden	26	25-90	Jarl (20s) Meridien (90s)
UK NL	229		



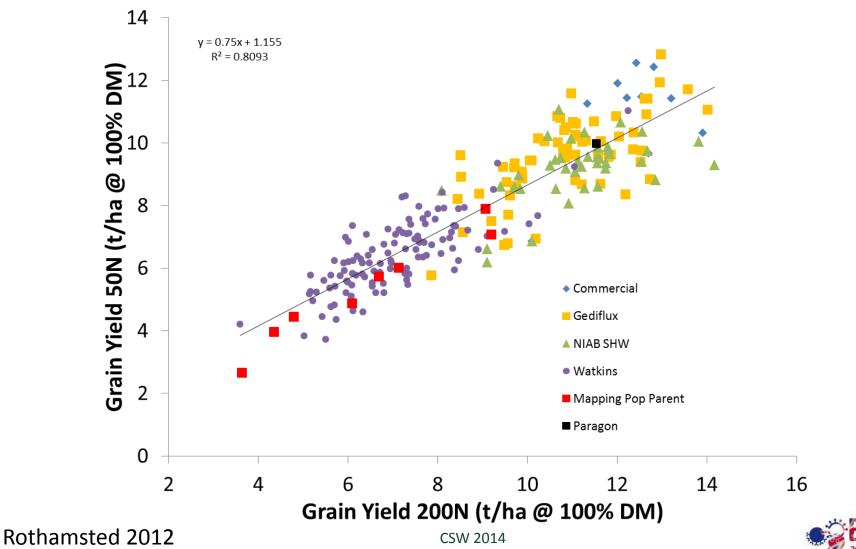
- Step changes
 - PBI
 - Rht1
 - 1BL.1RS
 - HMW glutenins
 - Claire, Robigus
- During each 'step change'
- What was being left behind?
- We can find out by looking into the 510 Gediflux varieties



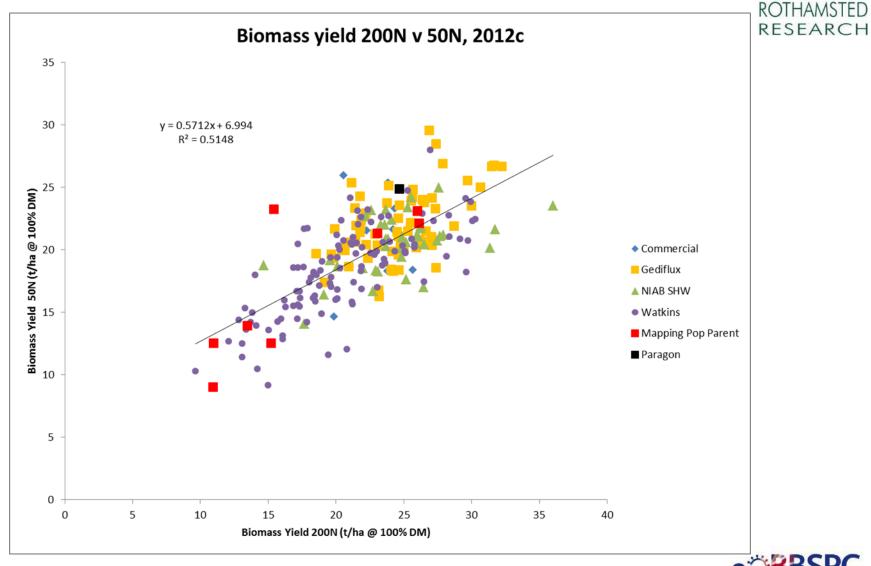
(slide from S. Griffiths, JIC)

Grain yields – response to N inputs



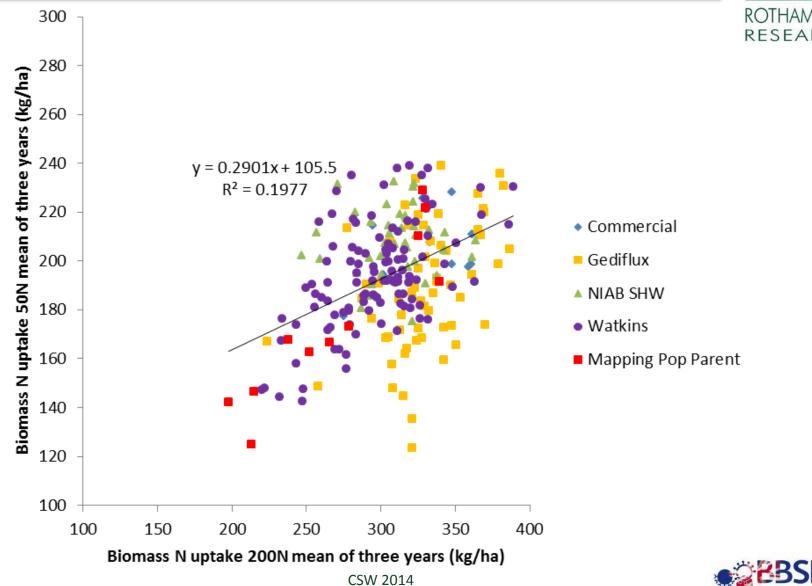


N-response: 2012 biomass



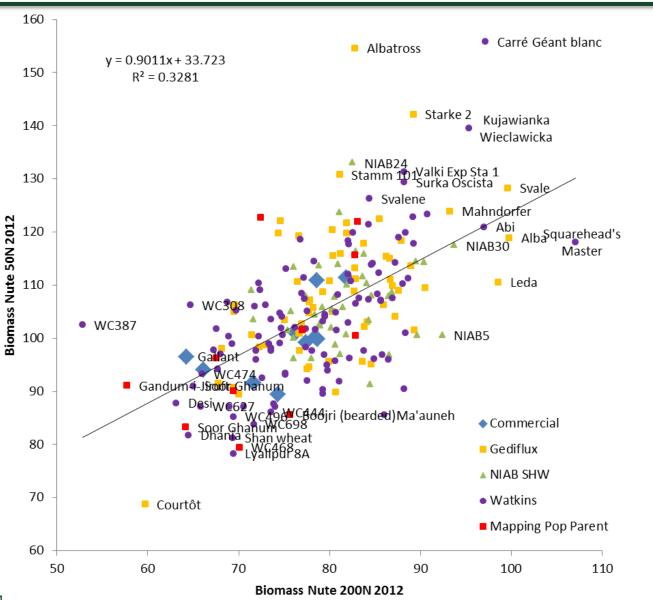
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N-uptakes, 3 year mean at RRes



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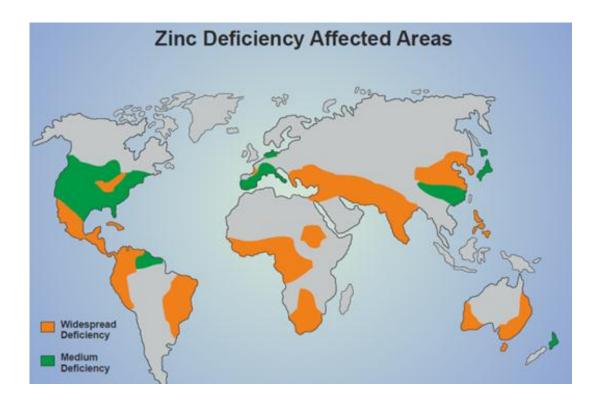
Biomass NUtE, 2012





Micronutrients: zinc

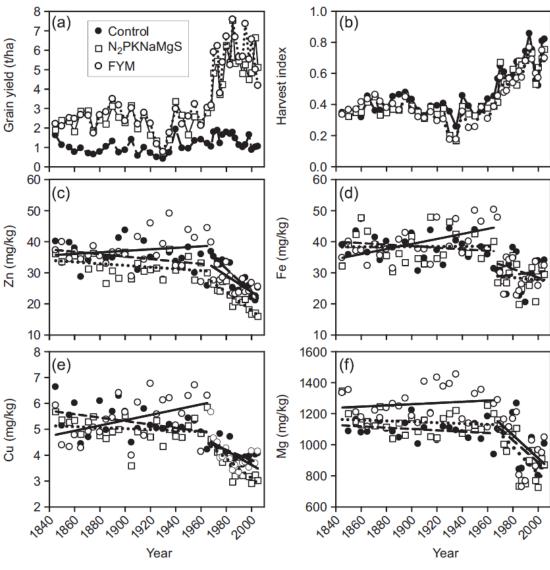
- Cereals are low in bioavailable essential micronutrients such as zinc
- Zinc deficiency affects 30% of the world's population
- Reduced zinc grain content also reduces yield
- What plant breeding strategies are there to overcome this?







Decreasing mineral content of wheat



ROTHAMSTED

Fan et al (2008) J trace Elements in Medicine &Biology; 22, 315-24.





Elevated CO₂ impacts on micronutrients

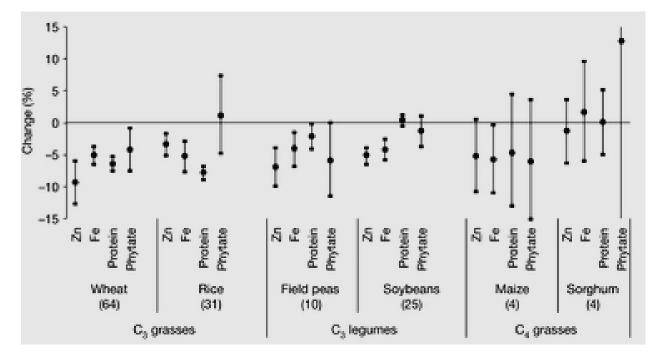
ROTHAMSTED

LETTER

doi:10.1038/nature13179

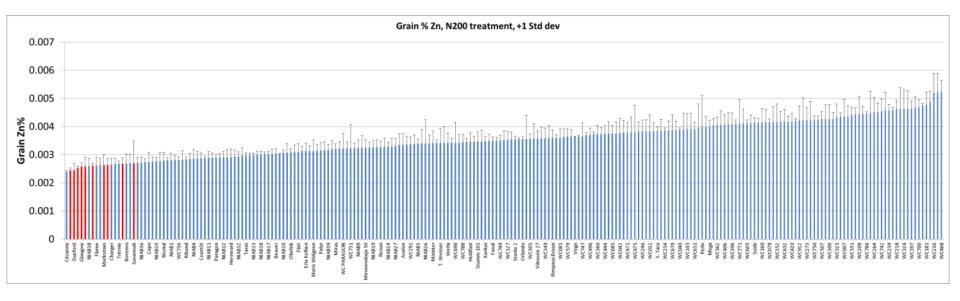
Increasing CO₂ threatens human nutrition

Samuel S. Myers^{1,2}, Antonella Zanobetti¹, Itai Kloog³, Peter Huybers⁴, Andrew D. B. Leakey⁵, Arnold J. Bloom⁶, Eli Carlisle⁶, Lee H. Dietterich⁷, Glenn Fitzgerald⁸, Toshihiro Hasegawa⁹, N. Michele Holbrook¹⁰, Randall L. Nelson¹¹, Michael J. Ottman¹², Victor Raboy¹³, Hidemitsu Sakai⁹, Karla A. Sartor¹⁴, Joel Schwartz¹, Saman Seneweera¹⁵, Michael Tausz¹⁶ & Yasuhiro Usui⁹





Grain Zn concentrations in WISP germplasm



24-52 mg/kg

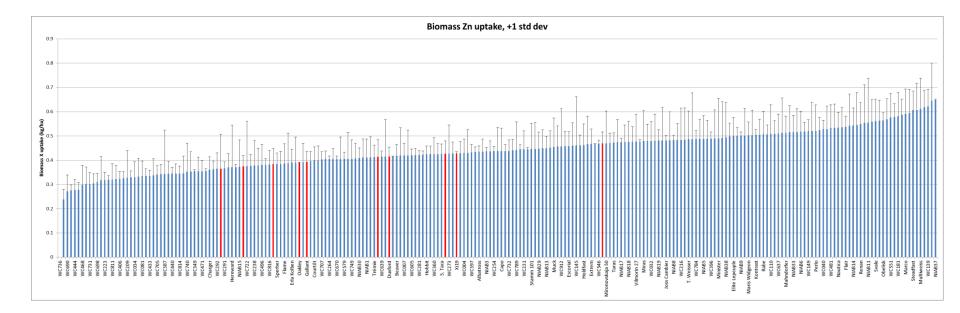


Red = modern cultivars 2012 Rothamsted High N



ROTHAMSTED

Total Zn uptake in biomass

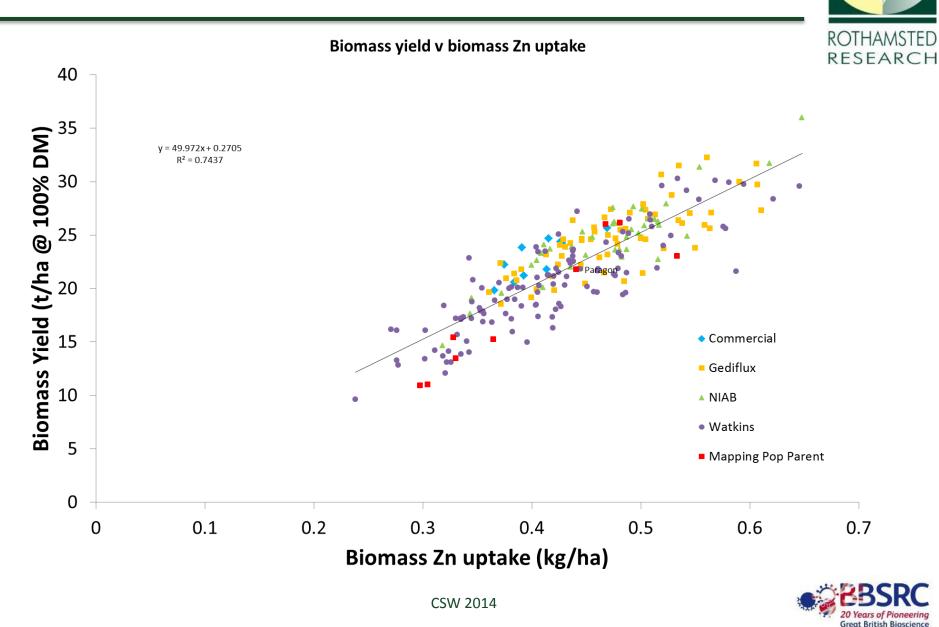


Range = 0.24-0.55 kg/ha Red = modern cultivars 2012 Rothamsted



ROTHAMSTED

Zn uptakes increases with biomass yield



What about roots?





Mapping populations

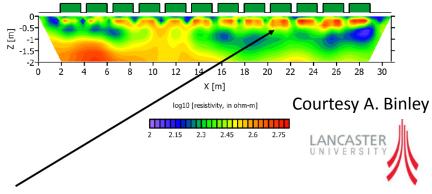


Penetrometer



Non-irrigated Absolute inversion 13-May-11

ERT Electrical resistance tomography



Appears to show significantly reduced water content down to 0.5m, especially in the plots towards the end of the line.



The University of Nottingham

CSW 2014

A QTL approach to dissect traits





RESEARCH PAPER

Identification of QTLs associated with seedling root traits and their correlation with plant height in wheat

Caihong Bai^{1,2}, Yinli Liang¹ and Malcolm J. Hawkesford^{2,*}

CSW 2014

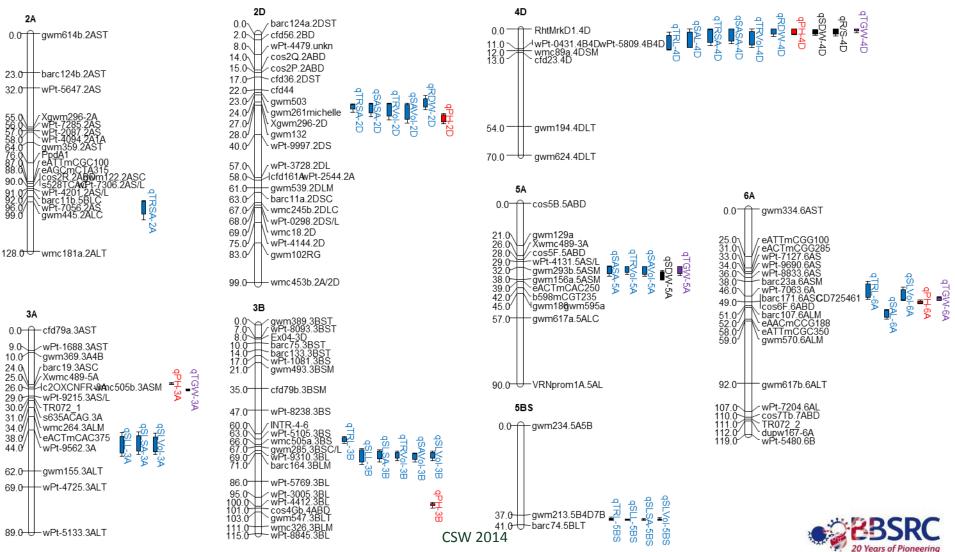


Botany www.jxb.oxfordjournals.org

Major root QTL and associated PH and TGW QTL



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The phenotyping challenge





Black Horse 2013

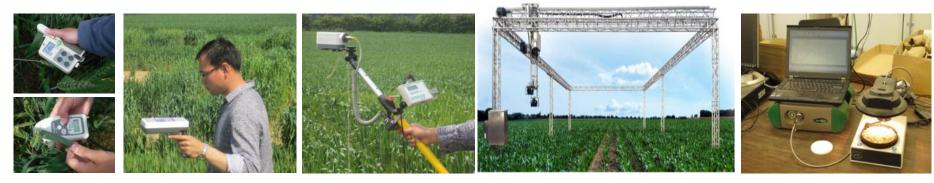


Scale

- Visual
- Hand held contact
- Hand held non contact
- UAV
- Phenomobiles
- Ground-based fixed site
- Post harvest high throughput analysis



8







Phenotyping from the air - UAV NDVI













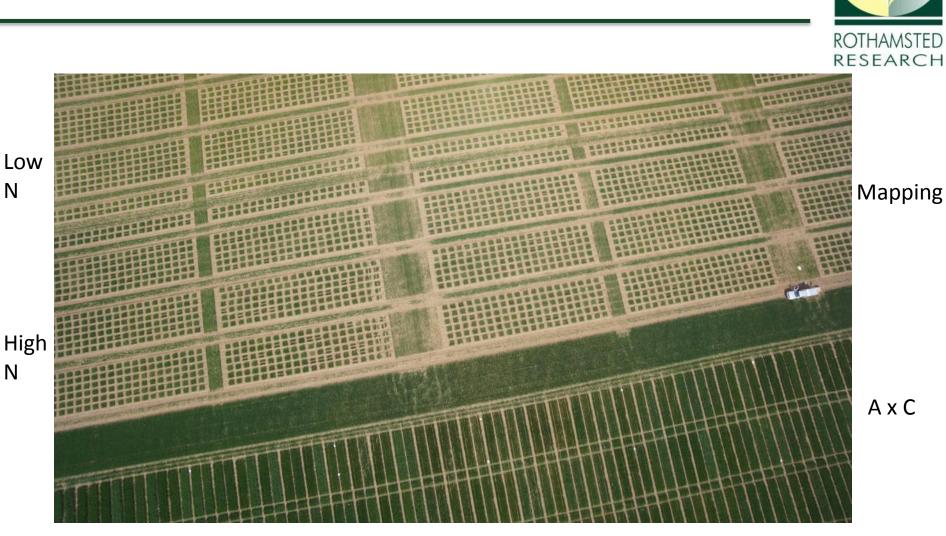
Mapping populations, Meadow 2014







Trials, Meadow 2014



24th April 2014



CSW 2014

Mapping population, Meadow 2014

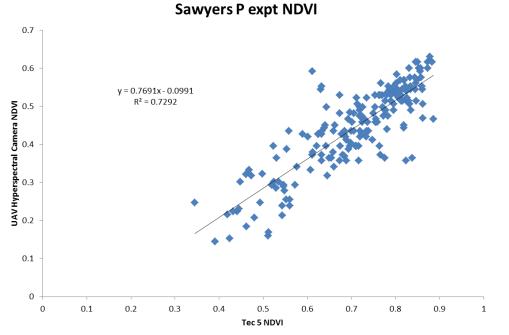




®

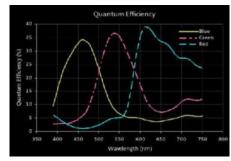
UAV capabilities

- Programed flight
- RGB
- Multi-spectral
- Thermal





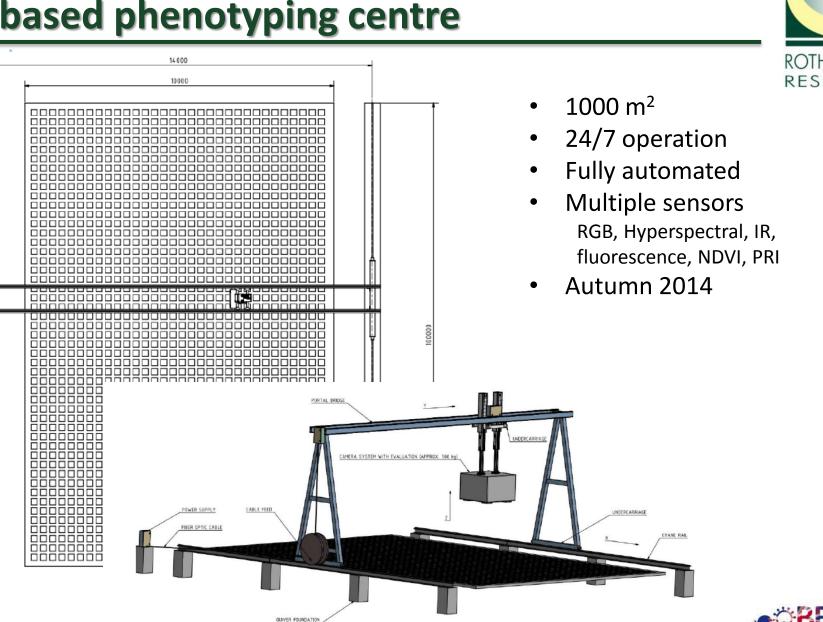








New Rothamsted automated groundbased phenotyping centre





20 Years of Pioneering Great British Bioscience

Summary

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- Substantial NUE diversity amongst modern germplasm
- BUT potential for much more by mining land races, diploid progenitors and wild relatives
- Essential to be precise about dissecting NUE traits



Rowland Biffen (1874–1949)



Thanks



- RRes Farm staff
- WISP, WGIN and 20:20 teams
- Peter Barraclough, Andrew Riche, Peter Buchner, Saroj Parmar, Yongfang Wan, Caihong Bai, Astrid Grün, Nick Evens
- Peter Shewry, Yongfang Wan, Ellen Mosletł (Nofirma), Gemma Chope (Campden BRI)

Wheat

Genetic Improvement

Network

- KWS, Limagrain, RAGT, Syngenta, bakers and millers
- Summer students and visitors













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