

DFW WP3 – TOOLKITS FOR BREEDING



The more elite varieties become the harder it gets to introduce diversity into them

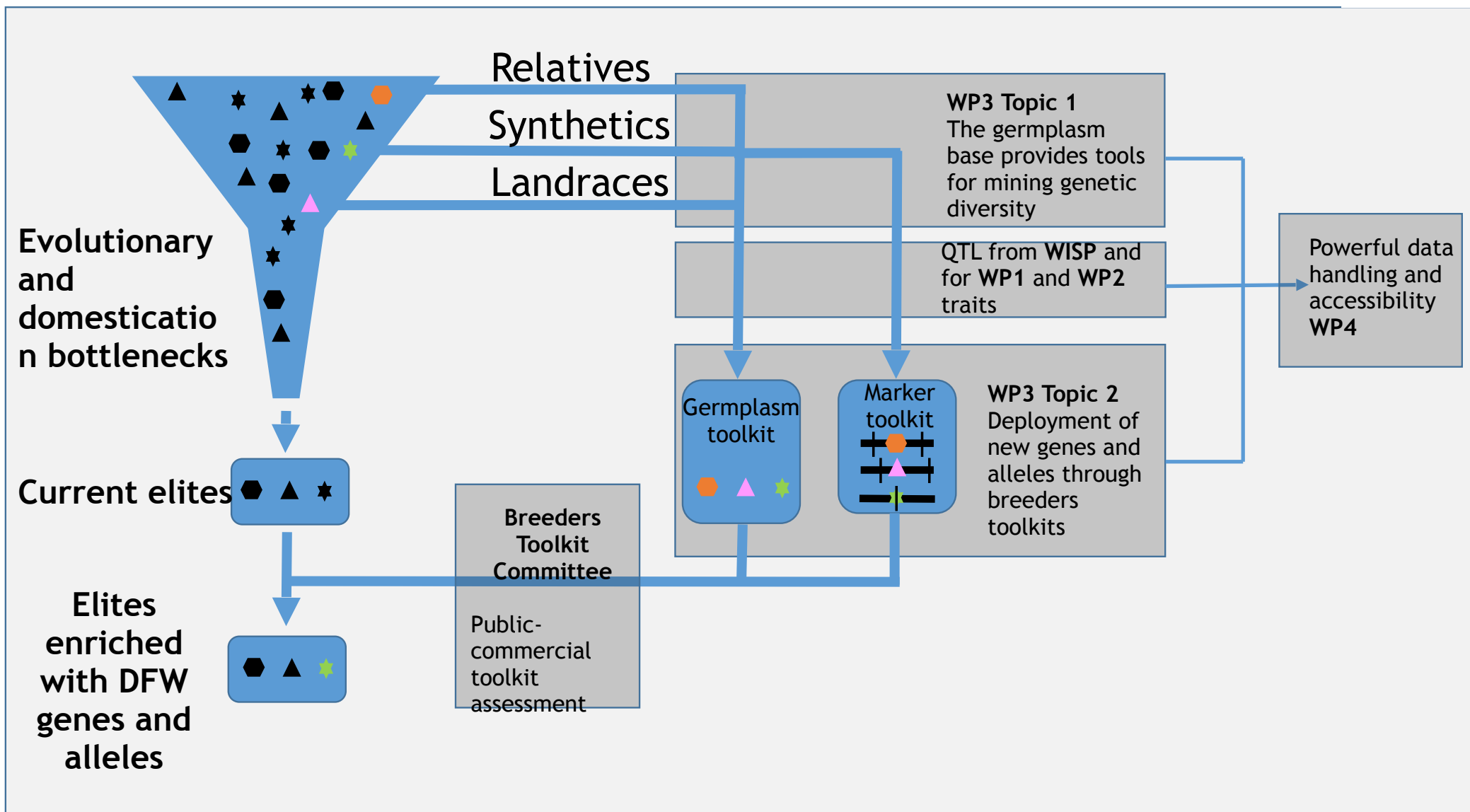


Sources of new and useful variation also carry a lot of undesirable variation

Eg 100%
lodging of the
AE Watkins
wheat
landrace
collection!



Strategy for allele mining and deployment



Breeders Toolkit Academic-Commercial Interface

Breeder Tool Kit Selection Committee



Chair
Simon Griffiths



Breeding Companies



KWS
Jacob Lage



RAGT
David Schafer



LSPB
Peter Jackson



Limagrain
Constance Lavergne



Syngenta
David Feuerhelm



DSV
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Work Package Leaders



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Rob
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Work Package 3 Institute Representatives



Bristol
Keith Edwards



NIAB
Keith Gardner



Nottingham
Julie King

GRU Representative
and BTK Coordinator
Simon Orford

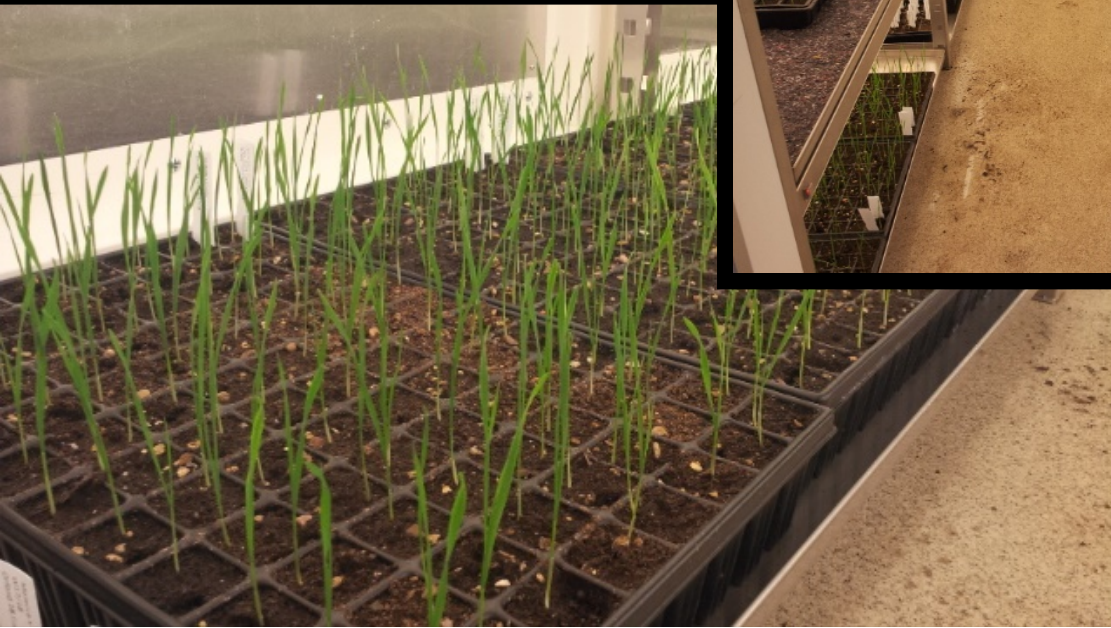


DFW Programme Manager
Julie Ellwood

<https://designingfuturewheat.org.uk/breeders-toolkit/>

BREEDERS TOOLKIT NOMINATION CRITERIA

- ▶ What is it? (e.g.QTL, alien introgression,mutation,)
- ▶ What trait does this relate to? (eg height, seed zinc content, lodging resistance)
- ▶ Work Package (please tick)
 - ▶ 1
 - ▶ 2.1
 - ▶ 2.2
 - ▶ 3.0
 - ▶ Material from outside DFW
- ▶ Describe the likely merit of the locus in breeding
- ▶ What evidence do you have to support this?
- ▶ What selection method can be used (marker, phenotypic)?
- ▶ How do you think this material should be assessed to validate its value in breeding? Quantity of seed available now
- ▶ Any other comments?





Michelle Leverington Waite



Field
multiplication
of Toolkit
germplasm



Academic Toolkit as of now

- 1742 lines ready
 - Deposited in GRU
 - Axiom 35 K genotyped
 - Grown at developer sites- mostly JIC, NIAB, Nottingham Uni
 - When sufficient seed is available they undergo central DFW testing
- 



Malcolm Hawkesfoed

Andrew Riche

Toolkit testing at Rothamsted

Harvest 2020 Breeders Toolkit

WBTK Code	TK Code	Genotype	BTK trial (Harvest year)	Description of Introgression	Recipient Cultivar	Comment
WBTK0020	WL0120	PW141-21-3-4-Q2A-GFP-W	BTK2020	Wat 1190141 QTL 2A-GFP	Paragon	JIC WP3
WBTK0021	WL0170	PW292-25-6-3-Q2A-MATU-W	BTK2020	Wat 1190292 QTL 2A-MAT	Paragon	JIC WP3
WBTK0022	WL0247	PW352-26-5-19-Q2A-DTMA-W	BTK2020	Wat 1190352 QTL 2A-MAT	Paragon	JIC WP3
WBTK0023	WL0296	PW468-84-4-2-Q5A-NDRE-W	BTK2020	Wat 1190468 QTL 5A-NDRE	Paragon	JIC WP3
WBTK0024	WS0095	SHW008 SYN Sel 51	BTK2020	NIAB SHW-008 BC1F6	Robigus	JIC WP3
WBTK0025	WS0105	SHW008 SYN Sel 61	BTK2020	NIAB SHW-008 BC1F6	Robigus	NIAB
WBTK0026	WS0117	SHW090 SYN Sel 70	BTK2020	NIAB SHW-090 BC1F6	Robigus	NIAB
WBTK0027	WR0032	D029 DHF1-29	BTK2020	Ae. mutica DH BC	Paragon	NIAB
WBTK0028	WR0154	D329 DHF1-329	BTK2020	Ae. mutica DH BC	Paragon	Nottingham
WBTK0029	WM0012	gw2-A / Par (+/-)	BTK2020	gw2-A	Paragon	Nottingham
WBTK0030	WM0013	Par / POL (-/+)	BTK2020	POL	Paragon	JIC WP1
WBTK0031	WM0014	gw2-A / POL (+/+)	BTK2020	gw2-A and POL	Paragon	JIC WP1

Multi site breeder trials coordinated by GRU



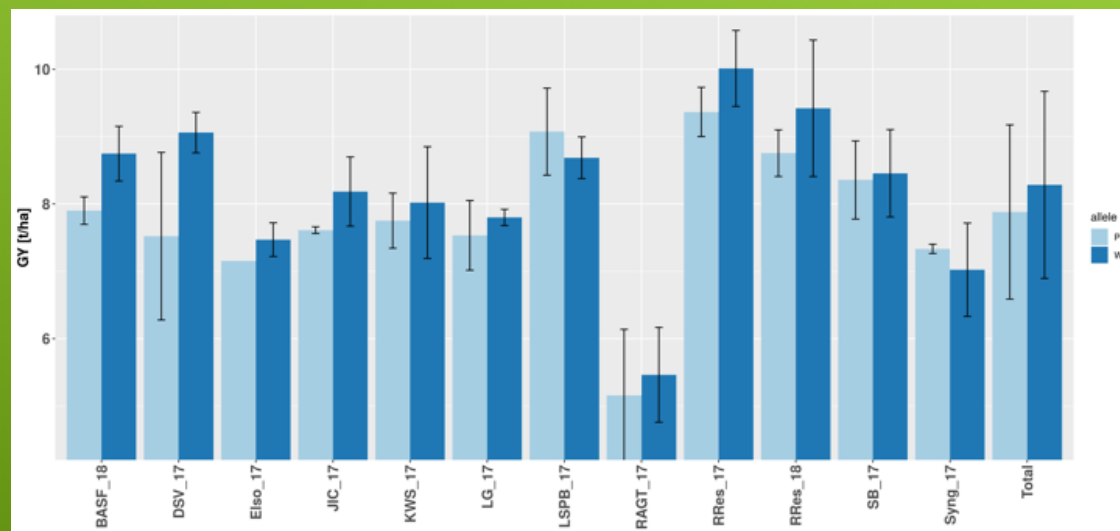
Simon Orford GRU

What does a successful BTK nomination look like?

Multi site trials



Consistent advantage conferred by new locus



Near isogenic lines carrying 7BL segment from Watkins landrace increase yield

SeedStor (Version 1.12) | SeedStor Home | Search Interface | Help Pages | GRU Information

SeedStor

<p>Reference panel of 376 European winter wheats used in association mapping research 2008-2009</p> <p>NIAB IAV IAV IAV IAV</p> <p>Triticaceae Genome Association panel</p>	<p>Wheat TILLING in silico wheat TILLING populations (Exome capture)</p>	<p>Vicia Faba</p>	<p>Wheat Pan Genome Pangenome Collection</p>	<p>Open Wild Wheat Consortium Aegilops tauschii Diversity Panel</p>
<p>DFW</p> <p>Designing Future Wheat Toolkit</p>	<p>BREEDERS TOOL KIT</p> <p>Designing Future Wheat Breeders Toolkit</p>	<p>GEDIFLUX Collection 1945-2000</p> <p>Gediflux Northern European Wheat Collection</p>	<p>Paragon X Chinese Spring SSD</p> <p>Paragon x Chinese Spring Mapping Population</p>	<p>Paragon γ Deletions</p> <p>Paragon Gamma Irradiated deletions</p>
<p>RenSeq lines</p> <p>Wheat Resistance gene enrichment (REN) sequencing collection</p>	<p>The Paragon Library</p> <p>Paragon Near Isogenic Line Library</p>	<p>DFW Nested Association Mapping Populations</p> <p>DFW Wheat Nested Association Mapping (DFW-NAM) panel</p>	<p>EMS Induced Mutations Paragon</p> <p>defra Wheat Genetic Improvement Network</p> <p>Paragon EMS Mutants</p>	

Toolkit
NILs

Breeders
Toolkit

RIL populations

RESOURCES IN GRU

