



Bringing a selection index into the IRRI programs

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Approach

1. Understand the targeted pipeline
 1. Breeding pipeline 1: market segments Hard White- Optimum Environment (HW-OE) and Hard White- Heat Tolerant Early Maturity (HW-HTEM).
 2. Breeding pipeline 2: market segments Hard White- Drought Tolerant Normal Maturity (HW-DTNM), Hard White- Drought Tolerant Early Maturity (HW-DTEM); Hard White- High Rainfall (HW-HR) & Hard Red
 3. Breeding Pipeline 3- Zn mainstreaming
2. Understand the targeted stage
 1. Stage 1
 2. Stage 2
 3. Stage n
3. Agree on the selection purpose
 1. Selection of parents
 2. Advancement of products
4. Understand the traits and selection procedure in an algorithm fashion.
5. Calculate retrospective weights
 1. $b = P-S$
6. Finetune weights.



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1. Understand the selection procedure in an algorithm fashion

- We identified that the process can be mapped back to a set of reduction and selection steps, each consisting in trait conditions (value and directionality):

As many traits as needed involved in each step

step	selectionUnit	stepType	trait	value	directionality	trait	value	directionality	valuesUsedAs	useTraitCovariance	TotalSelected
1	DESIGNATE	reduce	YLD_rel	0.4	>	ZNC_rel	0.3	>	culling	-	891
2	DESIGNATE	select	Xa21	R	=	YLD_BV	6.8	>	culling	-	20
3	DESIGNATE	select	Xa21	R	=	ZNC_BV	14	>	culling	-	20
4	DESIGNATE	select	Xa5	R	=	YLD_BV	5	top	culling	-	8
5	DESIGNATE	select	Xa5	R	=	ZNC_BV	5	top	culling	-	8
6	DESIGNATE	select	Pi54	R	=	YLD_BV	5	top	culling	-	8
7	DESIGNATE	select	Pi54	R	=	ZNC_BV	5	top	culling	-	8
8	DESIGNATE	select	Pita	R	=	YLD_BV	5	top	culling	-	6
9	DESIGNATE	select	Pita	R	=	ZNC_BV	5	top	culling	-	6
10	DESIGNATE	select	YLD_BV	10	top				culling	-	6
11	DESIGNATE	select	ZNC_BV	10	top				culling	-	6
12	DESIGNATE	select&reduce	PHT_BV	115	<	FLW_BV	90	<	culling	-	40
13	PARENTAGE	select	COUNT	3	random				culling	-	18

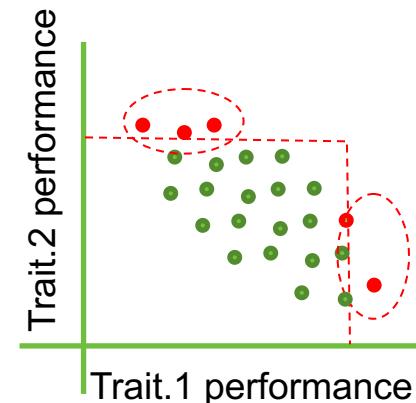
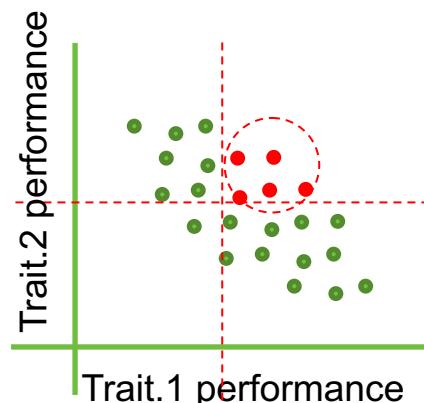
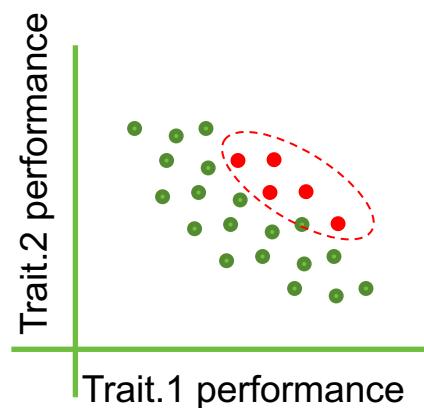
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Can we recreate or improve Josh's selections?

- We compared the selections made by Josh vs the algorithm and a selection index.
- Current method gives a strong weight to single-trait transgressive individuals.

Current method gives a strong weight to extreme value individuals NOT total merit

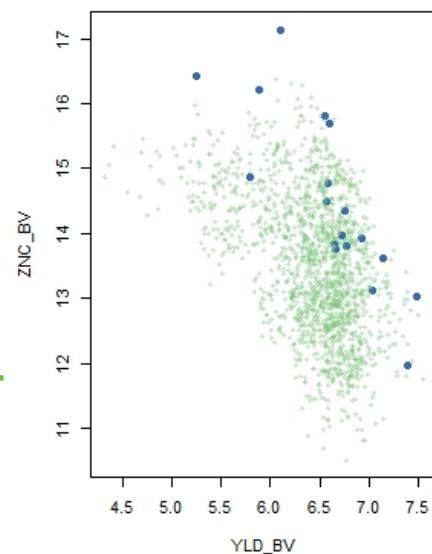
- Do simulations show that picking transgressive individuals is a good method to increase genetic gains?
 - T1: use an index to pick the best for total merit (10%)
 - T2: pick the best for yield and then best for zinc (31% > 31% = 10%)
 - T3: pick the best individuals for each trait (top 5% in each = 10%)



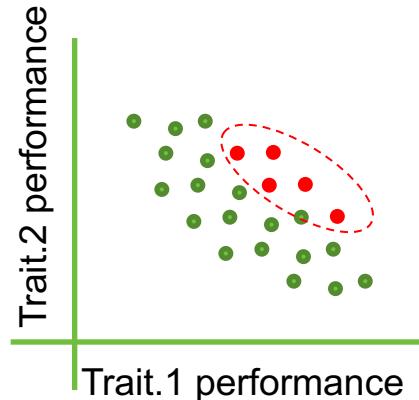
Selection index

Independent culling

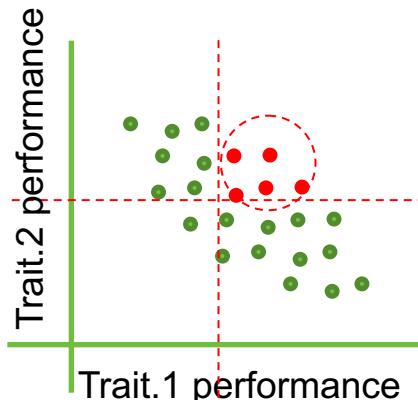
Tandem-type selection



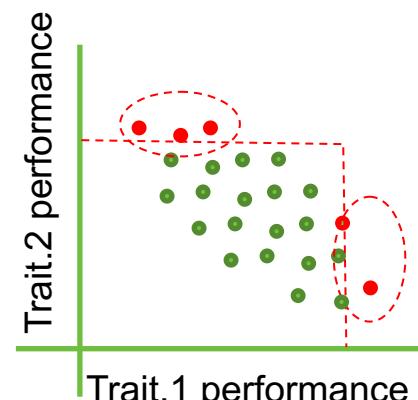
The best parents are not the extreme value individuals with lack of performance in other traits



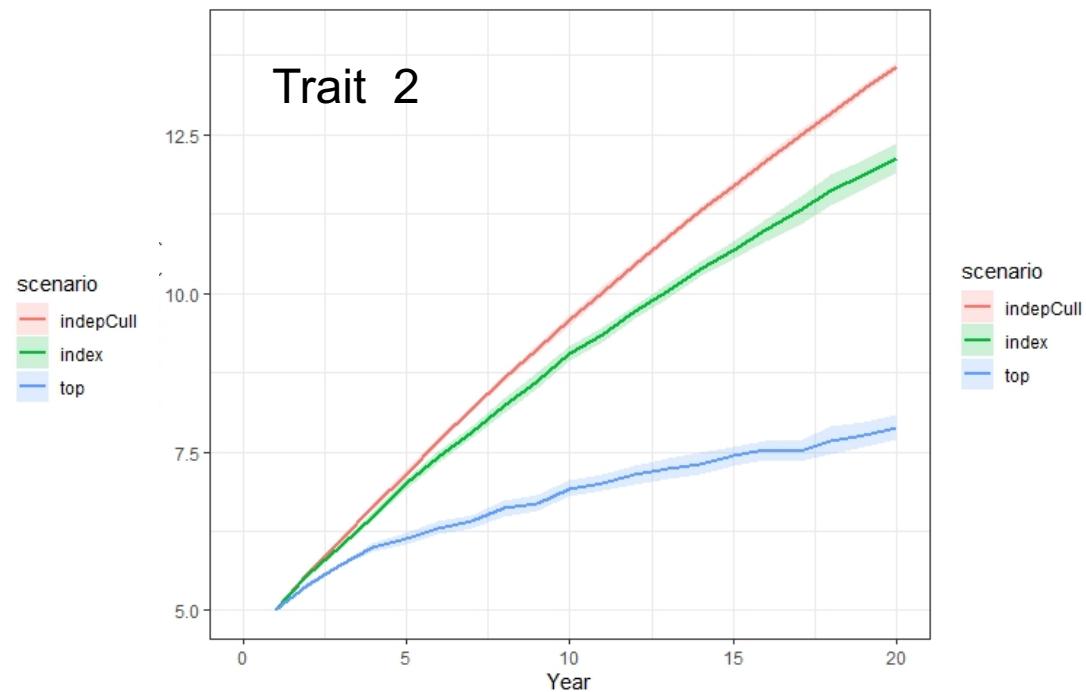
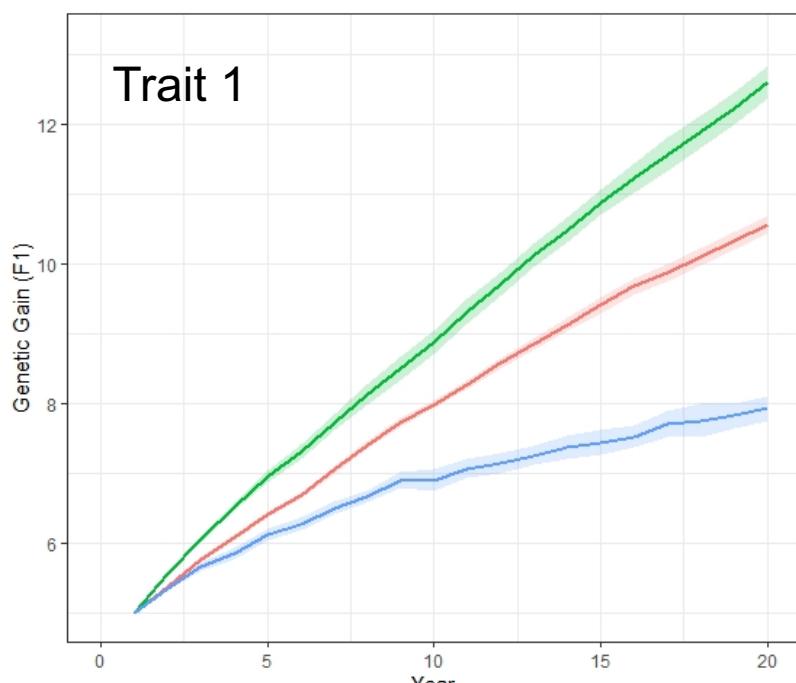
Selection index



Independent culling



Tandem-type selection



2. Identify which parts of the selection procedure can be replaced with an index

- We identified that the process can be mapped back to a set of reduction and selection steps, each consisting in trait conditions (value and directionality):

As many traits as needed involved in each step

step	selectionUnit	stepType	trait	value	directionality	trait	value	directionality	valuesUsedAs	useTraitCovariance	TotalSelected
1	DESIGNATE	reduce	YLD_rel	0.4	>	ZNC_rel	0.3	>	culling	-	891
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7	DESIGNATE	select	Pi54	R	=	ZNC_BV	5	top	culling	-	8
8	DESIGNATE	select	Pita	R	=	YLD_BV	5	top	culling	-	6
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10	DESIGNATE	select	YLD_BV	10	top				culling	-	6
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12	DESIGNATE	select&replace	PHT_BV	115	<	FLW_BV	90	<	culling	-	40
13	PARENTAGE	select	COUNT	3	random				culling	-	18

3. Build and refine an index: retrospective weights $b = P^{-1} s$

- If the selection differentials represent the breeder's goal, then the weights determines the merit of individuals selected.

Retrospective weights obtained using Josh's files

	Bangladesh	ESA	India	Philippines
YLD_BV	2.905	4.764	3.371	2.015
ZNC_BV	0.457	0.629	0.460	1.199
Xa21n	-3.148	1.003	-3.533	-0.844
Xa5n	0.751	-0.436	0.915	-0.492
Pitan	-0.496	-1.075	-1.153	-1.358
Pi54n	-1.051	-5.222	-2.889	-1.348

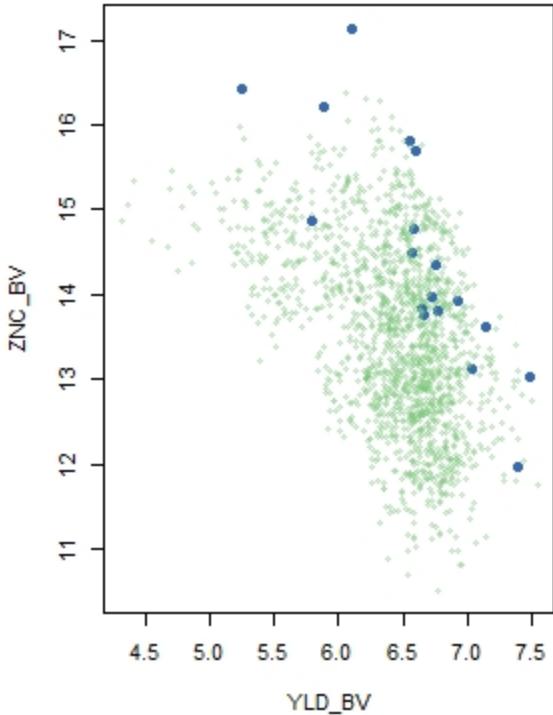


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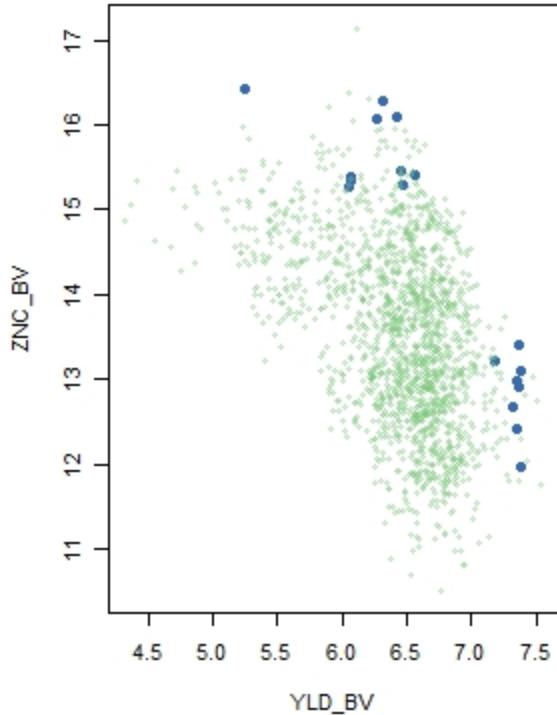
YLD_BV	2.015
ZNC_BV	1.199
Xa21n	-0.844
Xa5n	-0.492
Pitan	-1.358
Pi54n	-1.348

4. Compare to current approach using selection differentials

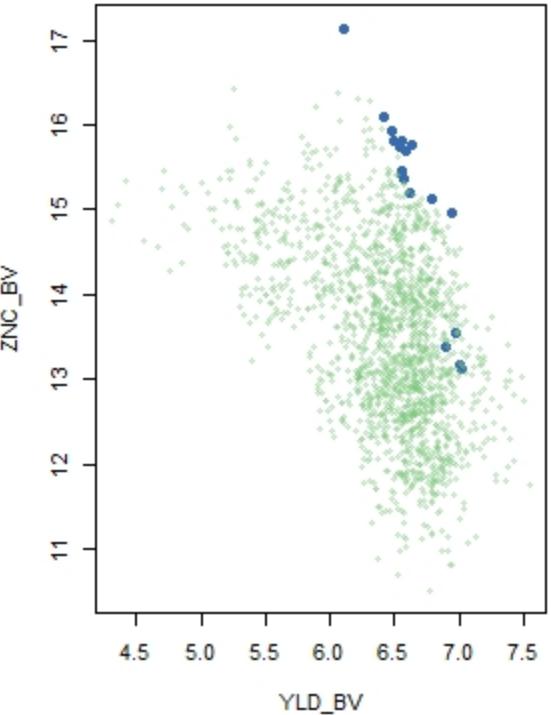
Josh selections



Algorithm selections



Index selections



	YLD_BV	ZNC_BV	Xa21n	Xa5n	Pitan	Pi54n	Final gain
Philippines.Josh	0.155	0.965	-0.078	-0.194	-0.174	-0.040	1.919
Philippines.Algorithm	0.255	0.906	-0.365	0.015	0.104	0.027	1.722
Philippines.Index	0.093	1.951	-0.564	-0.319	-0.174	0.027	3.361

Selection differential and total gain is higher for selection indices across all regions

	YLD_BV	ZNC_BV	Xa21n	Xa5n	Pitan	Pi54n	Final gain
Bangladesh.Josh	0.620	0.372	-0.188	-0.119	0.061	-0.032	2.477
Bangladesh.Index	0.671	1.447	-0.953	-0.208	-0.063	0.027	5.457
ESA.Josh	0.530	0.110	0.047	-0.165	-0.097	-0.127	3.481
ESA.Index	0.392	-0.484	0.047	-0.319	-0.174	-0.973	7.018
India.Josh	0.651	0.402	-0.188	-0.056	0.037	-0.078	3.178
India.Index	0.487	1.537	-0.953	-0.108	-0.069	0.027	5.617
Philippines.Josh	0.155	0.965	-0.078	-0.194	-0.174	-0.040	1.919
Philippines.Algorithm	0.255	0.906	-0.365	0.015	0.104	0.027	1.722
Philippines.Index	0.093	1.951	-0.564	-0.319	-0.174	0.027	3.361



You can consider that your weights were not the optimal and propose new ones

- You can finetune the weights to reflect better what you want.
- For example, we double the weight for yield:

	Bangladesh	ESA	India	Philippines	
YLD_BV	2.905	4.764	3.371	2.015	* 2 = <u>4.030</u>
ZNC_BV	0.457	0.629	0.460	1.199	
Xa21n	-3.148	1.003	-3.533	-0.844	
Xa5n	0.751	-0.436	0.915	-0.492	
Pitan	-0.496	-1.075	-1.153	-1.358	
Pi54n	-1.051	-5.222	-2.889	-1.348	



	YLD_BV	ZNC_BV	Xa21n	Xa5n	Pitan	Pi54n	Final gain
Philippines.Josh	0.155	0.965	-0.078	-0.194	-0.174	-0.040	2.230
Philippines.Index	0.425	1.078	-0.397	-0.319	-0.174	-0.029	3.772

Comparison of crosses selected by Josh vs the index

Cross	Freq.Josh	Freq.Index
BR 28/IR 50::C1	1	1
FEDEARROZ 50/IR 77298-14-1-2-10//IRRI 123/IR 45427-2B-2-2B-1-		
1///SANHUANGZHAN NO 2/...	1 NA	
IR 55182-3B-3-2-2-2/IR 10198-66-2	1 NA	
IR 98418-B-B-15/IR09A224	1	2
IR03A262/IR 50::C1	1	2
IR05N412/BRRI DHAN 55	1	1
IR09A116/IRRI 156	1	1
IR09N190/BRRI DHAN 55	1	1
IR09N190/IR09F436	1	1
IR09N190/IRRI 156	1	1
IR10N237/BRRI DHAN 55	1	1
IRRI 156/IR04A115	1 NA	
IRRI 156/IR11A293	1	1
IRRI 174/IR11A293	1 NA	
IRRI 174/IR12A330	1	1
MANAW THUKHA/IRRI 154	1	1
TAKANARI/IRRI 154	2 NA	
IR09A116/MANAW THUKHA	NA	1
IR10N225/BRRI DHAN 55	NA	1
IRRI 154/MOROBEREKAN	NA	1
IRRI 174/IR04A428	NA	1
PR 36921-B-6-1-3-4/IRRI 154//IR09A228	NA	1

Keeping selection of individuals per family balanced in the Philippines still favors the index

Selection differentials using index for across and within family selection

	YLD_BV	ZNC_BV	Xa21n	Xa5n	Pitan	Pi54n	FinalGain
Bangladesh.Josh	0.620	0.372	-0.188	-0.119	0.061	-0.032	2.477
Bangladesh.Index	0.649	0.478	-0.153	0.181	-0.074	-0.073	2.833
ESA.Josh	0.530	0.110	0.047	-0.165	-0.097	-0.127	3.481
ESA.Index	0.613	0.158	0.047	-0.052	-0.041	-0.173	4.038
India.Josh	0.651	0.402	-0.188	-0.056	0.037	-0.078	3.178
India.Index	0.706	0.211	-0.153	0.231	-0.024	-0.073	3.468
Philippines.Josh	0.155	0.965	-0.078	-0.194	-0.174	-0.040	1.997
Philippines.Index	0.252	0.989	-0.111	-0.056	-0.069	-0.026	2.069



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