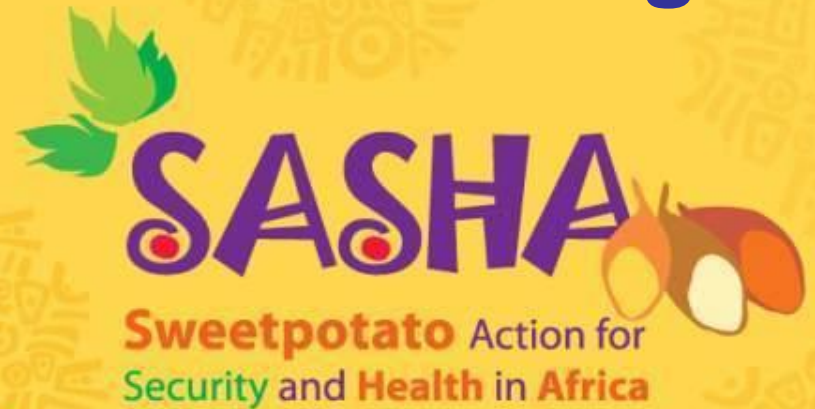


Improved Tools for Breeders: CloneSelector & AccuDataLog



Luka Wanjohi, Jacaranda Hotel, Nairobi
September 09, 2014

Introduction



- CloneSelector is a tool developed to help plant breeders carry out field trials, analyze the results and make selection decisions.
- Open source
- Based on MS excel & R statistical package

Features



- Design field trials – single and multiple location
- Generate a field book for each experiment
- Register metadata for each experiment – critical with Open Access
- Data collection in field and post-harvest
- Calculate derived variables (yield/ha etc)

Features



- Offer some options for statistical analysis
- CloneSelector now offers strong data management options for both breeding and non-breeding trials for clonally propagated crops

Highlights



- CloneSelector1.0 released in 2010
 - RCBD trial design and management
- CloneSelector2.0 released June 2011
 - MET analysis
- CloneSelector3.0 released August 2012
 - 2 more statistical designs to accommodate trials with large number of genotypes

Highlights

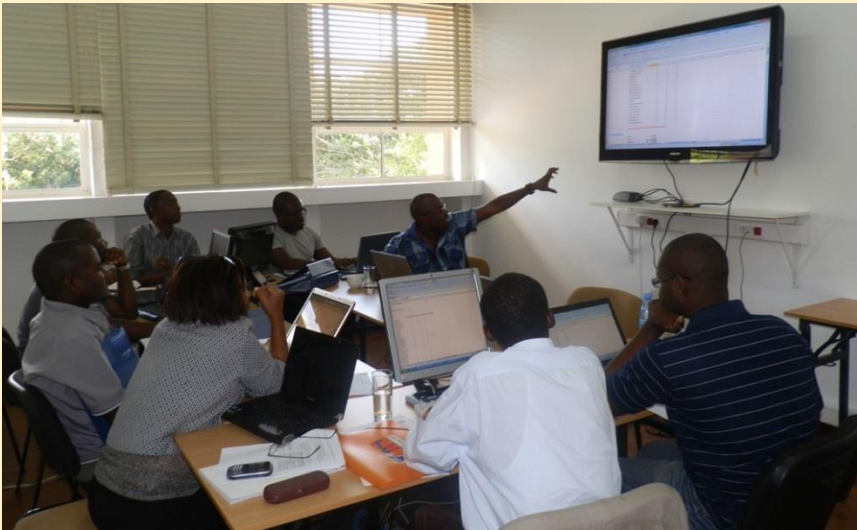


- CloneSelector3.0 released August 2012
 - Automatic import of NIR's data
- Training offered to SP breeders during annual meetings 2010-2012

Highlights



- Intensive in-country trainings in 2013
 - Over 10 countries



Highlights



- Work ongoing on integration of Elston and Pesek Baker Index into CloneSelector
 - Raul Eyzaguirre conducted a training on both indices in June during this year's breeders meeting
- AccuDataLog, a mobile Fieldbook app. with integrated Zebra Barcoding technology.

About AccuDataLog



- A mobile application for entering trial data into the CloneSelector Fieldbook while in the field
- Available on Windows and Android platforms

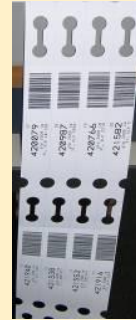
Main Features

- Automatic Import of CloneSelector Fieldbooks into mobile device
- Field based data entry



Main Features

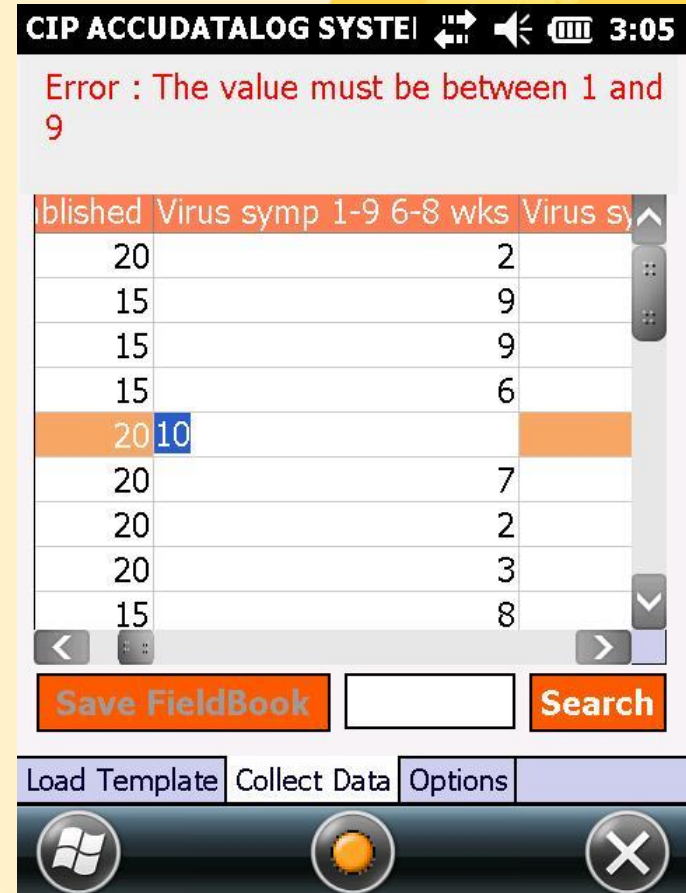
- Integrated barcode technology, 1D or 2D



Main Features



- Realtime data entry validation: numeric, date, string length, lower limits, upper limits, etc
 - User customizable



Main Features

- Print on demand (POD) of labels via mobile printing
 - Labeling of samples during harvest
 - Label regeneration



Main Features



- Easy transfer of data from PDA back to CloneSelector Fieldbook for analysis

SSP WA Case Study



- Trial: PT
- Genotypes: 31,
Reps:2,
Locations: 4
- Team consists
CIP and CSIR-
CRI colleagues



Planting

- Planting labels printed
 - white V-Max polyolefin 7.5 mil tag that provides tear strength and outdoor use up to 1-2 years. Offers good durability and chemical resistance



Data Collection



- Fieldbook uploaded on PDA's in readiness for field data collection
 - SSP-WA has 5 PDA's so far
 - Same Fieldbook uploaded on multiple PDA's as each technician will collect data of a unique trait
 - Data collected over breeding season and regularly backed up.
 - Previously data entered on printed copies of the Fieldbook

Harvesting

- Field data entry
 - Root count, foliage, etc
 - Complimentary paper data capture for backup



Harvesting

- NIR's samples labels printed in the field



NIR's Lab

- Fresh weight data entry into Fieldbook



Transfer back to CloneSelector



	B	C	D	E	G	H	AK	AL	AM	AN	AO	AP	AQ	AR	AS	AT	AU	AV	AW	AX	AY	AZ
1	Trial: PT3-2013 Site: Fumesua Country: Ghana Planting: 22/08/2013											fWeightRo	fVineWeig	fSkinColor	fFleshColo	fSizeRoots	fFormRoot	fRootDefe	fWeevilDa	fTrait4	fTrait5	fTrait6
10	PlotSize: 3 Harvest: 22/12/2013 Collaborator: Dr. Asafu-Agyei Institution:																					
11	CSIR-CRI/CIP																					
12																						
20	LowerLimit						0	0	0				1		1	1	1	1				
21	UpperLimit												9		9	9	9	9				
23	Rep	Bloc	Fla	Entry	Name	Pedgree	# plants Harvested	# plants w. Roots	# Roots Marketable	# Roots NonMarket	Weight Root Marketable	Weight Root NonMarket	Vine Weight kg	Skin Color 1-9	Flesh Color	Root Size 1-9	Root Form 1-9	Root Defects 1-9	Weevil Damage 1-9	Aldicides 1-9	Milipede 1-9	Trait6
25	1	1	1	1	UW11906-79		9	9	6	23	1.5	4.0	2.5			6	5	1	7	3.0	3.0	
26	1	1	2	2	Kemb10		10	8	4	12	2.0	2.0	3.1			5	5	1	7	6.0	3.0	
27	1	1	3	3	Ningshu1		10	5	0	9	0.0	0.5	3.3			3	3	1	1	2.0	1.0	
28	1	1	4	4	Mugande		10	10	6	30	2.0	2.5	3.0			3	3	1	2	2.0	1.0	
29	1	1	5	5	SPK004/616		10	8	4	10	2.0	1.5	2.0			2	2	1	2	1.0	1.0	
30	1	1	6	6	Apomuden		10	10	11	24	0.2	1.5	0.7			4	4	1	5	3.0	2.0	
31	1	1	7	7	199062		8	8	21	32	2.5	1.5	2.1			5	5	1	6	1.0	2.0	
32	1	1	8	8	Mohe		10	9	14	15	3.0	1.5	1.1			7	6	1	1	1.0	1.0	
33	1	1	9	9	Ejumula		10	10	11	20	2.5	1.0	2.1			5	4	1	5	2.0	3.0	
34	1	1	10	10	Uww11906-289		7	5	9	11	1.0	0.5	3.1			5	6	1	3	1.0	1.0	
35	1	1	11	11	Kemb37		10	10	11	24	2.0	1.0	1.7			5	5	1	4	2.0	2.0	
36	1	1	12	12	Cemsa78-326		10	8	8	4	2.5	0.1	1.0			5	4	2	4	3.0	2.0	
37	1	1	13	13	Maphutha-1		10	10	16	23	1.5	0.1	0.5			4	6	1	6	2.0	1.0	
38	1	1	14	14	Mugamba		10	10	30	15	4.5	0.5	1.1			6	5	2	4	2.0	2.0	
39	1	1	15	15	Zapallo		8	6	6	6	0.5	0.1	4.2			4	4	1	2	2.0	1.0	
40	1	1	16	16	Kamala Sundari		10	10	24	19	3.5	0.5	1.3			6	5	1	3	2.0	3.0	
41	1	1	17	17	UW119-15		10	10	20	14	5.5	0.5	2.3			6	5	1	3	3.0	2.0	
42	1	1	18	18	Ogyefo		10	10	11	30	2.5	1.5	3.0			5	5	1	3	3.0	1.0	
43	1	1	19	19	UW11906-175		10	9	11	21	2.5	1.0	1.3			5	5	1	5	2.0	1.0	
44	1	1	20	20	MUSG0616-18		10	9	11	26	1.5	1.0	1.1			4	4	1	7	1.0	2.0	
45	1	1	21	21	Jewll		10	10	16	18	3.0	0.5	1.1			5	4	1	8	3.0	3.0	
46	1	1	22	22	Ejumula25		10	10	13	24	3.5	1.5	1.6			7	5	1	6	3.0	3.0	
47	1	1	23	23	Tacna2		10	10	17	13	3.5	0.5	3.9			7	5	1	5	1.0	2.0	
48	1	1	24	24	Bauregard		10	9	22	18	4.0	1.0	11.7			5	4	2	5	2.0	1.0	
49	1	1	25	25	Jonathan		9	9	7	26	1.0	1.0	2.4			3	4	1	2	1.0	2.0	
50	1	1	26	26	Cemsa74-228		10	10	19	5	8.5	0.5	4.6			8	5	1	7	6.0	4.0	
51	1	1	27	27	Tamale Orange		10	10	11	8	2.5	0.5	0.6			5	5	1	6	3.0	2.0	
52	1	1	28	28	LO323-1		10	10	15	16	4.0	1.5	3.0			6	4	2	6	3.0	5.0	

- The rest of the NIR's data is imported back into CloneSelector via the CloneSelector routine for importing NIR's data
- On the first day of use data for 13 different harvest traits for a total of 62 genotypes in the PYT was entered into the Fieldbook in the field.

Challenges



- Limited informatics support in SSA
 - User support always key in success of any information systems

Getting Started



<http://sweetpotatoknowledge.org/germplasm/research-methods/cloneselector>

AccuDataLog:

<https://research.cip.cgiar.org/confluence/display/GIMS/CIP+AccuDataLog>

Files (5)

- About CloneSelector.pdf ▾
Mar 12, 2013 | Luka Wanjohi
- Advanced RExcel Setup Guide ▾
Jul 25, 2011 | Luka Wanjohi
- CloneSelector Users Guide - Multi Environment Trial Analysis ▾
Sep 19, 2013 | Luka Wanjohi
- CloneSelector Users Guide - Potato ▾
Sep 19, 2013 | Luka Wanjohi
- CloneSelector Users Guide - Sweetpotato ▾
Sep 19, 2013 | Luka Wanjohi

Subfolders under CloneSelector

Folders under this section are listed below. Click on the folder name to see

CloneSelector1-1

CloneSelector2-0

CloneSelector3-0

CloneSelector3-1

CloneSelector3-1

CloneSelector3-1 ▾
Sep 19, 2013 | File | Luka Wanjohi

See all...

There are no subfolders

Example of Query tool for CloneSelector

Acknowledgements



- Raul Eyzaguirre
- Edwin Rojas
- Carlos Velasquez
- Ted Carey
- Jan Low

