Tutorial of the Breeding Planner (BP) for Marker Assisted Backcrossing (MABC)

BP system consists of three tools relevant to molecular breeding.

- MARS: Marker Assisted Recurrent Selection
- MABC: Marker Assisted Backcrossing
- MAS: Marker Assisted Selection for pyramiding multiple genes
- This tutorial is designed for MABC

November 2013







Open the Breeder Planner (BP) software

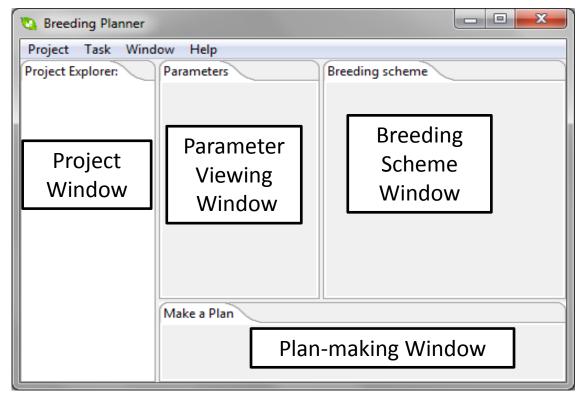
1. Open the software: The BP software can be opened by double click the software icon in your computer desktop

Breeding Planner

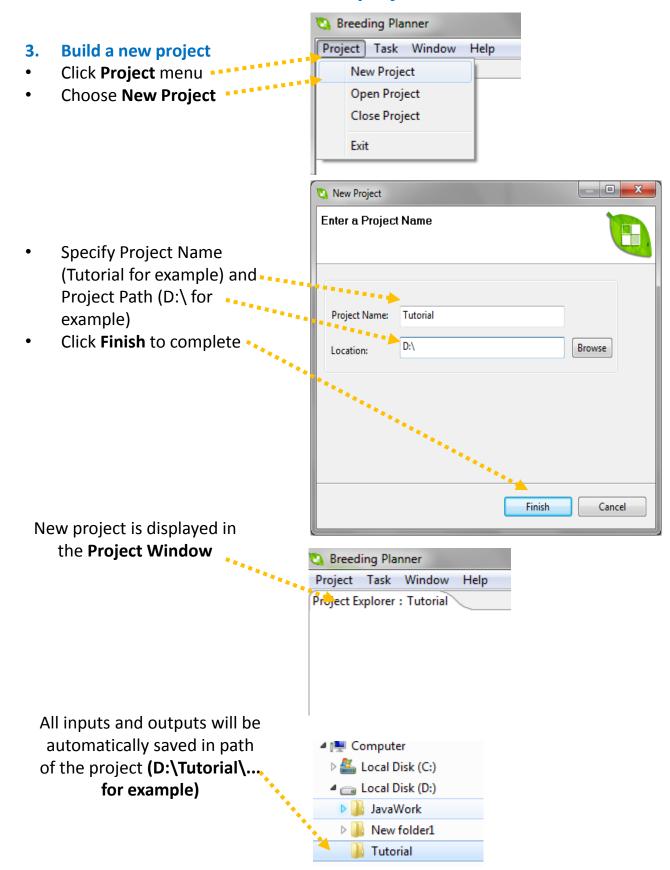
2. Overview of the software.

The screen is split into four windows.

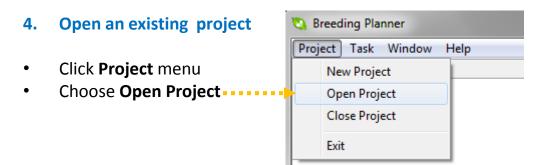
- Project Window: List all molecular breeding programs you have planned. Three distinct breeding programs can be considered: MARS, MABC and MAS.
- Parameter Viewing Window: You can view your breeding parameters in this window.
- Breeding Scheme Window: Once the breeding parameters are specified, a breeding flowchart will be demonstrated in this window.
- Plan-making Window: You can select the current stage/generation of your breeding programs in this window. A detailed plan for the near future will be made by the BP system.



Build a new project

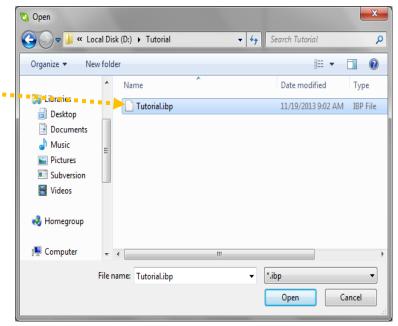


Open an existing project

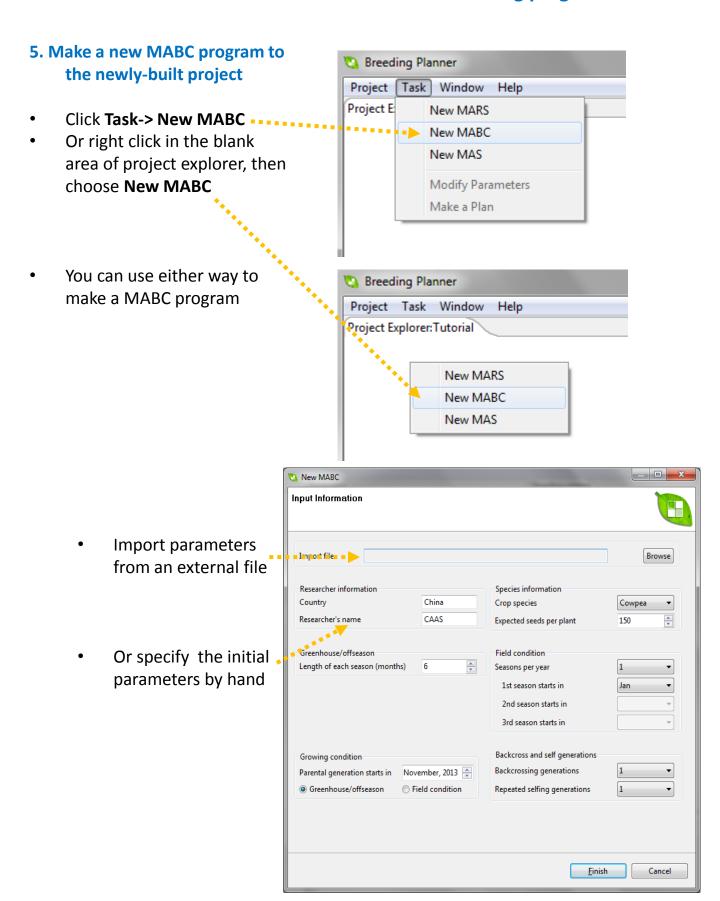


 Choose an existing project in your computer

• Please Note: The BP system is project-based. When you first use the system, you need to build a new project first. Then you can make various breeding programs. When you leave the system, the system automatically save the jobs you have done. The next time, you can start from a new project, or from an existing project.



PMABC Tutorial: Start a new MABC breeding program



PMABC Tutorial: Parameters required to define a MABC program

Users have to specify a set of parameters before a MABC scheme can given by the system. Below are more details on the required parameters.

1. Researcher information

Country;

Researcher's name;

2. Species information

Crop species: Select one crop species from Cowpea, Rice, Wheat, Maize, Groundnuts, and Cassava

Expected seeds per plant (or propagation rate) (more on the next page)

3. Greenhouse/offseason

Length of each season (months).

Note: We assume the crop can be grown across the whole year under the Greenhouse condition. That is, the next season can start in the same month when the previous season is harvested. So the planting time for each season is not needed.

4. Field condition

Seasons per year: the number of seasons per year, select from 1-3

1st season starts in: select a month

2nd season starts in: select a month, after the end of the 1st season 3rd season starts in: select a month, after the end of the 2nd season

Note: The crop cannot be grown across the whole year under the Field condition. So if multiple seasons are possible, the system asks for the planting time for each season.

5. Growing condition

Parental generation starts in: select the start time of the parental generation

Early generation growing condition: select one case "Greenhouse/offseason" or "Field condition"

6. Backcross and self generations

Backcrossing generations:1-3

Repeated selfing generations: 1-3

Minimum and maximum numbers of seeds per plant for each crop under optimum or normal conditions

Crop	Minimum number	Maximum number	Median (used as default in Breeding Planner)
Cowpea	10	300	50
Rice	50	300	200
Wheat	50	250	150
Maize	50	500	200
Groundnuts	30	200	80
Cassava	10	100	50

Notes:

- Expected seeds per plant in Breeding Planner will be used to calculate if there are enough seeds for phenotyping. If not, additional seed increase (by selfing) will be requested.
- The user input must fall into the min-max range for the selected crop! Otherwise, when the input number is smaller than the minimum number, the minimum number will be assumed. When the input number is greater than the maximum number, the maximum number will be assumed.
- The number of seeds required is calculated from settings for "Multi-locational phenotyping". Say, when genotyping is conducted in F2, multi-locational phenotyping is only possible when each F2 plant can give enough seeds. Otherwise, phenotyping will be delayed until the required seeds are produced.

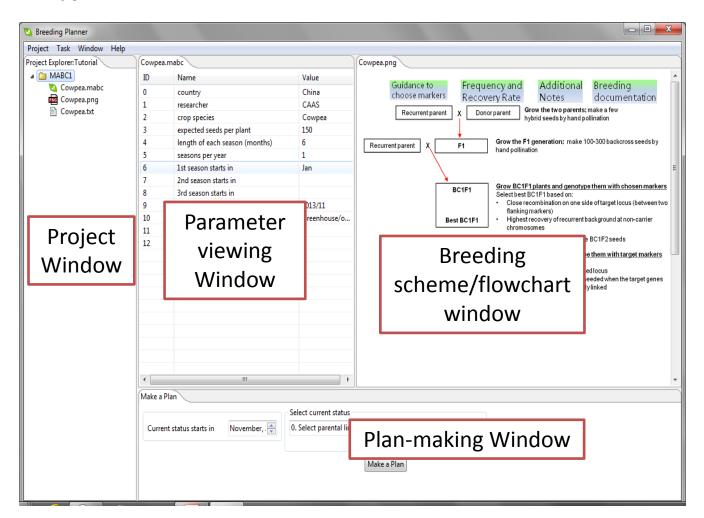
PMABC Tutorial: The interface

6. Overview of the MABC functionality:

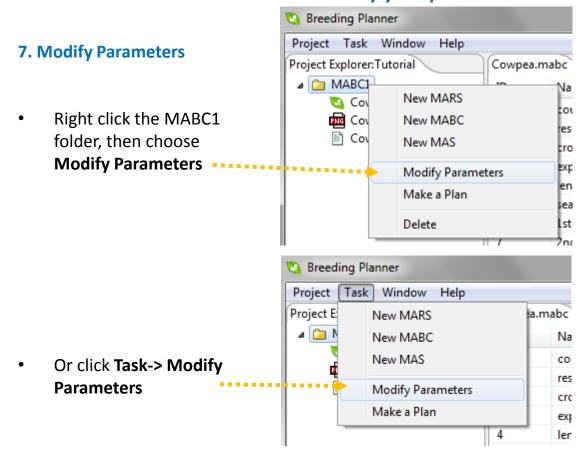
There are four windows in MABC functionality

- **Project Window:** List all molecular breeding programs you have planned. Three distinct MB programs can be considered: MARS, MABC and MAS.
- Parameter Viewing Window: You can edit/view your breeding parameters in this window.
- **Breeding Scheme Window:** Once the breeding parameters are specified, a breeding flowchart will be demonstrated in this window.
- **Plan-making Window:** You can select the current stage/generation of your breeding programs in this window. A detailed plan for the near future will be made by the BP system.

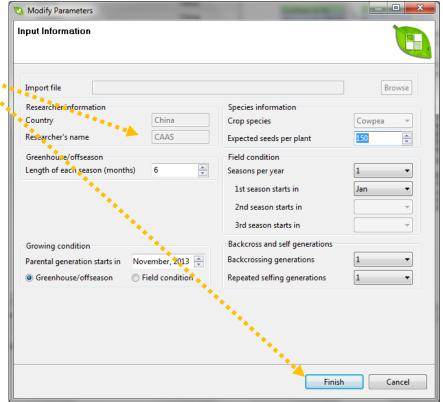
When the required parameters are set in the Parameter setting/viewing Window, the defined MABC breeding program is graphed in the Breeding Scheme/flowchart Window. A set of output files are listed in the Project Window.



PMABC Tutorial: Modify your parameters



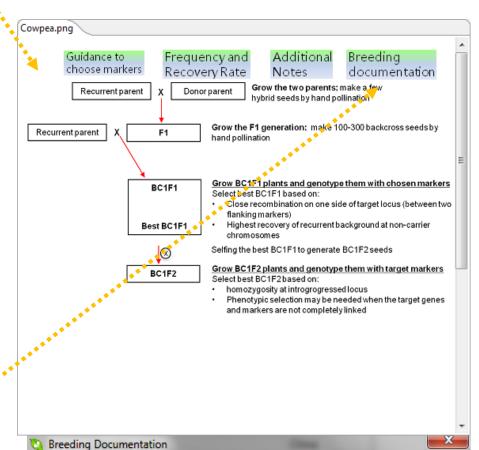
You may modify the parameters in the Dialog Window, and then click Finish button to save your modifications. The breeding scheme will be automatically modified.



PMABC Tutorial: The breeding scheme and documentation

8. View the flowchart

It is in the right window

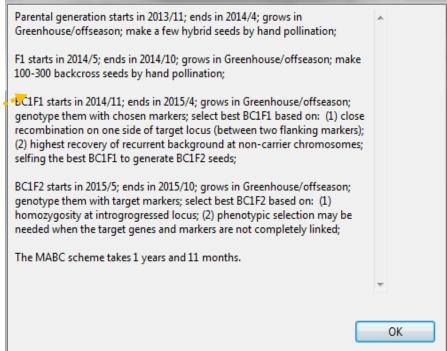


9. View breeding documentation

 Click the Breeding Documentation button

The Breeding

Documentation is in the pure text describing the whole flow of the MABC program.



PMABC Tutorial: Additional information

Guidance to choose markers

10. Additional information on the flowchart

 Click Guidance to choose markers Cowpea.png Frequency and Guidance to Additional Breeding choose markers Recovery Rate Notes documentation Grow the two parents; make a few Recurrent parent Donor parent hybrid seeds by hand pollination Grow the F1 generation; make 100-300 backcross seeds by Recurrent parent hand pollination

to choose markers will be popped up to help to choose markers to use in MABC process

 Click Frequency and Recovery Rate Choose markers to use in MABC process

2-4 well spread polymorphic markers per chromosome for background selection

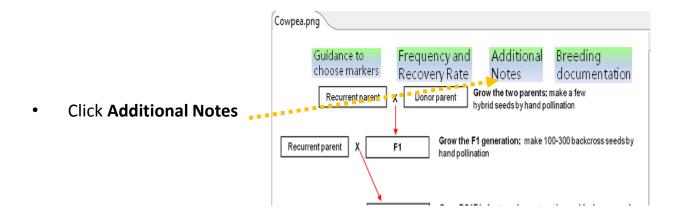
 2-3 flanking markers on each side of locus to introgress (2-5 cM apart if major gene, if QTL involved the whole confidence interval needs to be spanned (4-5 markers))

OK

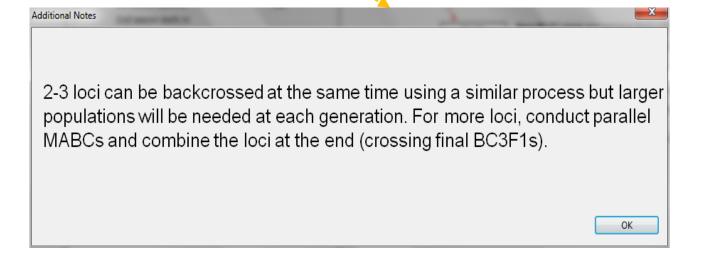
 The table for Frequency and Recovery Rate will be popped up to help to get the Rate (%) of homozygousity during repeated selfing

Rate (%) of h	nomozygousity d	uring repeated	d selfing					
Generation	Number of ind	ependent loci	(higher value	for linkage)				
	1	2	3	4	5	10	20	30
F2	50.00	25.00	12.50	6.25	3.13	0.10	0.00	0.00
F3	75.00	56.25	42.19	31.64	23.73	5.63	0.32	0.02
F4	87.50	76.56	66.99	58.62	51.29	26.31	6.92	1.82
F5	93.75	87.89	82.40	77.25	72.42	52.45	27.51	14.43
F6	96.88	93.85	90.91	88.07	85.32	72.80	52.99	38.58
F7	98.44	96.90	95.39	93.89	92.43	85.43	72.98	62.35
F8	99.22	98.44	97.67	96.91	96.15	92.46	85.48	79.03
F9	99.61	99.22	98.83	98.45	98.06	96.16	92.47	88.92
F10	99.80	99.61	99.42	99.22	99.03	98.06	96.17	94.30

PMABC Tutorial: Additional information



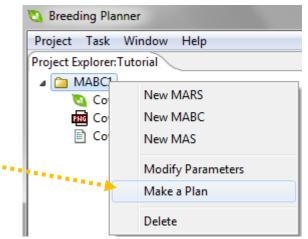
 The table for Additional Notes will be popped up to help to conduct the MABC process.

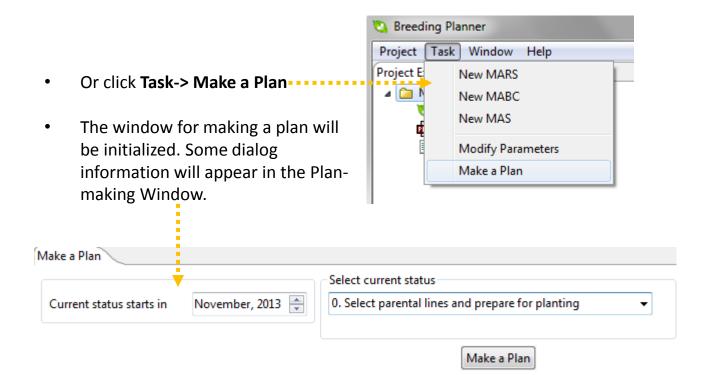


PMABC Tutorial: Make a plan from wherever you are

11. Make a plan

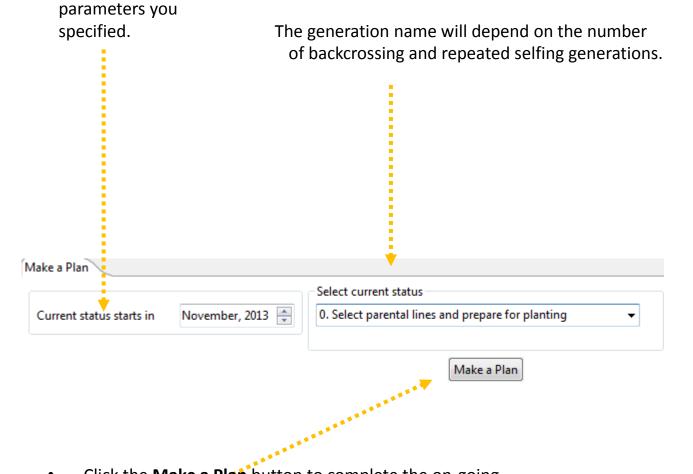
 Right click the MABC folder in the Project Window, then choose Make a Plan





PMABC Tutorial: Tell the system where you are

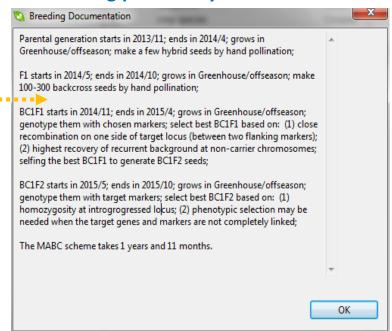
- Tell the system when the current season starts. The system will know the current season is grown in Greenhouse or in Field from the breeding
- Tell the system where you are by selecting:
 - 0. Select parental lines and prepare for planting
 - 1. Parental lines is growing
 - 2. F1 generation is growing
 - 3. BC1F1 generation is growing
 - 4. BC2F1 generation is growing
 - 5. BC3F1 generation is growing
 - 6. BC3F2 generation is growing
 - 7. BC3F3 generation is growing
 - 8. BC4F4 generation is growing



• Click the **Make a Plan** button to complete the on-going MABC breeding program.

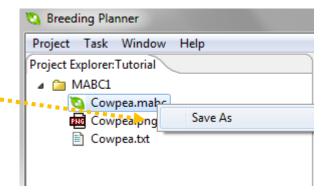
PMABC Tutorial: See the breeding plan the system makes

 New window will show the Breeding
 Documentation for the remaining status, and the time to complete the
 MABC breeding program

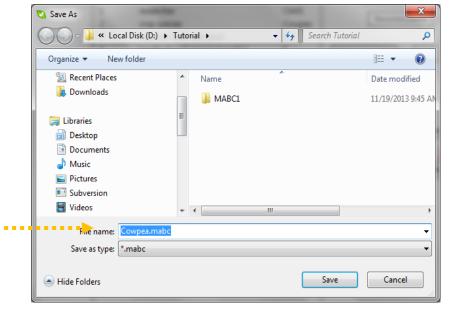


12. Save the result files

 Right click the file name, and then select Save As to save the breeding documentation on your computer



 Specify the path and rename of the output file



BP-MABC Tutorial: The contact information

- Any comments or suggestions? You may contact any one on the BP development team
 - Dr. Jiankang Wang, CIMMYT China and CAAS, <u>wangjiankang@caas.cn</u> or <u>jkwang@cgiar.org</u>
 - Dr. Luyan Zhang, CAAS, <u>zhangluyan@caas.cn</u>
 - Mr. Wencheng Wu, CAAS, <u>wuwencheng@caas.cn</u>
 - Dr. Xavier Delannay, GCP, <u>x.delannay@cgiar.org</u>
 - Dr. Hamer Paschal, GCP, <u>Hamerp99@gmail.com</u>





