Melon Trait and Germplasm Resources Survey 2011

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The Cucurbit Crop Germplasm Committee (CCGC), which operates under the auspices of the USDA-ARS National Plant Germplasm System (NPGS), is composed of ARS, university and industry scientists, and provides guidance to NPGS on matters relating to cucurbit crop and wild related species. The CCGC is responsible for all cultivated cucurbits (http://www.ars-grin.gov/ npgs/cgclist. html).

The CCGC Crop Report is available on-line and is periodically updated. The seed industry and public research communities have undergone major changes since the last update of the melon (*Cucumis melo* L.) section of the CCGC Crop Report. Moreover, changes in market demands and resource and regulatory constraints, further development of molecular technologies, and new production-limiting biotic and abiotic challenges warrant an update of the melon report.

Melon was introduced to North America after centuries of culture and selection throughout Europe, the Middle East, central, eastern and southern Asia, India, and Africa. Recent research supports an Asian origin (1-3) and reports the closest wild species, *C. picrocarpus*, to be in Australia (3).

A two-part survey instrument is, thus, available for input by any person in any country with knowledgeable interest in melon production, utilization, breeding, genetics, or botany (see following). Part 1 is concerned with fruit and plant traits of importance with emphasis on host plant resistance, fungal and viral pathogens in particular. Every region has specific, required combinations of fruit quality traits–they are not included. Part 2 is focused on Germplasm Resources, from "Germplasm," which consists of wild or feral accessions and land races, to varieties and cultivars including F_1 hybrids, and nine types of genetic stocks. Interest in markers and transgenics are also queried.

Download the survey from the website at the bottom of the survey and send via email, or photocopy and send via mail (address above), or fax to the number at the bottom of the survey. All responses will be useful for updating the melon section of the CCGC Crop Report. A summary of the survey will be prepared for inclusion in the next Cucurbit Genetics Cooperative Report.

Literature Cited

- Renner, S. S., Schaefer, H., and Kocyan, A. 2007. Phylogenetics of *Cucumis* (Cucurbitaceae): *C. sativus* (cucumber) belongs in an Asian/Australian clade far from *C. melo* (melon). BMC Evolut. Biol. 7:58-69.
- 2.Schaefer, H., Heibl, C., and Renner, S. S. 2009. Gourds afloat: a dated phylogeny reveals an Asian origin of the gourd family (Cucurbitaceae) and numerous oversea dispersal events. Proc. Royal Soc. Bot. 276:843-851.
- 3.Sebastian, P., Schaefer, H., Telford, I. R. H., and Renner, S. S. 2010. Cucumber (*Cucumis sativus*) and melon (*C. melo*) have numerous wild relatives in Asia and Australia, and the sister species of melon is from Australia. Proc. Natl. Acad. Sci. (U.S.A.) 107(32):14269–14273.

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Melon Survey 2011 Part 1: Fruit and Plant Traits of Interest/Needed

Fruit traits

- Increased shelf life
- Lightly processed characteristics
- ____ Increased yield
- ____ Nutritional value
- ____ New market types
- ____ Other: _____
- ____ Other: _____

Abiotic stress

- ____ Temperature
- ____ Salt excess
- ____ Mineral deficiency
- ____ Water
- ____ Other: _____
- ____ Other: _____

Disease: Fungal

- Powdery mildew
 - ____ Podosphaera xanthii
 - <u>Golovinomyces cichoracearum</u>
- ____ Downy mildew (*Pseudoperonospora cubensis*)
- ____ Anthracnose (*Colletotrichum lagenarium*
- ____ Fusarium wilt (*Fusarium oxysporum* f.sp. *melonis*)
- Verticillium wilt (Verticillium dahliae and V. albo-atrum)
 Phytopthora crown rot (Phytopthora capsici)
- Pythium (*Pythium* spp.) root and crown rot
- Alternaria (Alternaria cucumerina)
- Gummy stem blight (*Didymella bryoniae*)
- Vine decline (*Monosporascus cannonballus*)
- ____ Other: _____
- Other:

Disease: Bacterial

- ____ Fruit Blotch (Acidovorax avenae subsp. citrulli)
- ____ Bacterial wilt (Erwinia tracheiphila)
- ____ Angular leaf spot (Pseudomonas syringae pv. lachrymans) ____ Other: _____
- Other:

Disease: Viral

Aphid-transmitted

- ____ Cucumber mosaic virus (CMV)
- ____ Cucurbit aphid borne yellows virus (CaBYV)
- ____ Muskmelon yellow stunt virus (MYSV)
- ____ Papaya ringspot virus (PRSV) watermelon strain (= WMV 1)
- ____ Watermelon mosaic virus (WMV) (= WMV 2)
- ____ Watermelon mosaic Virus-Morocco
- ____ Zucchini yellow mosaic virus (ZYMV)

Disease: Viral (continued)

Whitefly-transmitted

- ____ Beet pseudo yellows virus (BPSYV)
- Cucurbit leaf crumple virus (CuLCrV)
- ____ Cucurbit yellow stunting disorder virus (CYSDV)
- ____ Lettuce infectious yellows virus (LIYV)
- ____ Melon chlorotic leaf curl virus (MCLCV)
- ____ Melon leaf curl virus (MLCV)
- ____ Squash leaf curl virus (SLCV)
- ____ Watermelon curly mottle virus (WCMoV)
- ____ Other _____
- ____ Other: _____

Soil-borne/ Seed-borne

- ____ Muskmelon necrotic spot virus (MNSV)
- ____ Squash mosaic virus (SqMV)
- ____ Other _____
- ____ Other: _____

Other

- ____ Cucumber green mottle mosaic (CGMMV)
- ____ Cucumber vein yellowing virus (CVYV)
- ____ Cucurbit latent virus (CLV)
- Curly top virus
- ____ Eggplant mottled dwarf rhabdovirus
- ____ Kyuri green mottle mosaic virus
- (KGMMV-YM)
- ____ Melon rugose mosaic virus (MRMV)
- ____ Melon vein-banding mosaic virus (MVbMV)
- ____ Melon yellow spot virus
- Melon yellowing-associated virus (MYaV) Melon yellows virus
- ____ Muskmelon yellow spot virus (MYSV)
- ____ Ourmia melon virus
- Tobacco ringspot virus (TrSV)
- Tomato leaf curl virus
- ____ Zucchini yellow fleck virus (ZYFV)
- Other ______ Other:

Insects and Nematodes

- _____ Sweetpotato whitefly, Bemisia tabaci Biotypes: A/B/Q
- ____ Greenhouse whitefly, Trialeuroides vaporariorum
- ____ Cucumber beetle, Acalymma trivittatum, Diabrotica undecimpunctata undecimpunctata, and D. balteata
- ____ Leafminer, Liriomyza sativae and Liriomyza trifolii
- ____ Green peach aphid, Myzus persiceae
- ____ Melon aphid, Aphis gossypii
- ____ Melon fly, *Myiopardalis pardalina*
- ____ Root knot nematode, *Meloidogyne* spp. ____ Other: _____
- ____ Other: ______ ____ Other ______
- Download from: http://www.ars-grin.gov/npgs/cgc_reports/melonsurvey

Return to: Jim McCreight, USDA, ARS, 1636 E. Alisal St., Salinas, CA 93905, U.S.A. Fax 01-831-755-2814; Email: jim.mccreight@ars.usda.gov

Melon Survey 2011 Part 2: Germplasm Resources

Germplasm

_ More germplasm is needed for various characters (as listed on the preceding page).

Germplasm exploration and exchange is:

____ high priority

____ moderate priority

____ low priority

If high or moderate priority, please identify potential countries and areas therein as well as names of in-country contacts who could collaborate in germplasm collection trips or /exchanges.

____ There are adequate germplasm resources available.

Comments on germplasm resources:

Genetic Stocks of interest Recombinant inbred lines (RILs) Near isogenic lines (NILs) Core Collections Test Arrays Mutant Stocks Pocket Collections Haploids Doubled Haploids Tetraploids	Transgenics high prio moderate low priori Transgenics high prio moderate low priori	Transgenics at present are high priority moderate priority high priority	
Molecular markers are			
high priority			
moderate priority			
low priority			
Additional comments			
Respondent information (optional, but h	nelpful for any needed clarification)		
Name:	Discipline area(s):		
	Breeding	Molecular	
Contact information:	Pathology	Physiology	

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____ Entomology

____ Other _____