A UNIQUE CYTOPLASMIC-NUCLEAR INTERACTION CAUSING SUNFLOWER PLANTS WITH REDUCED VIGOR AND THE GENETICS OF VIGOR RESTORATION

Zhao LIU¹, Wei GU², Chao-Chien JAN³

North Dakota State University
Heilongjiang Academy Of Agricultural Sciences, Harbin, Heilongjiang, China
Northern Crop Science Lab., USDA-ARS, Fargo, ND, USA

chaochien.jan@ars.usda.gov

ABSTRACT

Plants with pale yellow leaves and reduced vigor were observed in backcross progenies of inbred line HA 89 or HA 410 in the cytoplasms of 12 perennial *Helianthus* species, but not in the cytoplasms of annual *H. niveus*, *H. praecox*, *H. anomalus*, and *H. neglectus*. Segregation ratios of normal (N) to reduced-vigor (RV) plants in testcrosses and self-pollination of heterozygous normal plants, respectively, suggested that single dominant gene (V) controls vigor restoration. A high frequency of vigor restoration genes was found in 11 cultivated sunflower lines, with the exception of HA 89, HA 410, RHA 801, and Seneca. Testcross progenies of the half-diallel crossed F1s among HA 271, HA 234, VNIIMK, Armavir, Issanka, and HA 821 onto the RV cmsRIG1 were all normal, suggesting that all these lines possess the same V gene. Extensive use of H. tuberosus in early sunflower breeding programs might explain the presence of H. tuberosus V gene in many cultivated sunflowers, and the possible selective advantage of the V gene. A new V gene derived from H. giganteus was identified, which differed from the V gene commonly existing in cultivated lines. Other V genes derived from H. hirsutus and H. salicifolius will be compared among all the V genes. The V gene commonly existing in cultivated lines has been mapped to the linkage group 7 of the sunflower genome using SSR markers. The tightly linked markers will help select for normal vigor progenies when using perennial Helianthus cytoplasms in a breeding program

Key Words: sunflower, cytoplasmic-nuclear interaction, reduced vigor, wild perennial Helianthus