Inheritance of very high linoleic acid content and its relationship with nuclear male sterility in safflower

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Safflower (*Carthamus tinctorius* L.) possesses the highest amount of linoleic acid among the 10 major vegetable oil crops of the world. Very high linoleic acid content is controlled by recessive alleles at a single locus *Li*. However, deviated segregations from the expected monogenic inheritance have been observed in crosses involving nuclear male-sterile (NMS) lines. The present research was undertaken to study the inheritance of very high linoleic acid content in safflower and its relationship with nuclear male sterility. F₁, F₂, F₃, BC₁F₁ and BC₁F₂ seed generations were evaluated in a cross between CR-142 (a line with very high linoleic acid content, 88%) and CL1 (an NMS line with wild-type linoleic acid content, 74%). The genetics of linoleic acid content in male-sterile plants was determined by testcrossing with CR-142. The results confirmed monogenic inheritance. The analysis of the F₃ and BC₁F₂ to CL1 seed generations demonstrated a repulsion-phase linkage between *Li* and *Ms* loci, the latter conferring the NMS trait. The recombination rate between *Li* and *Ms* was estimated to be 0.09.