Molecular Phylogenetic Analyses of Geographical and Ecological Speciation of Anaphalis margaritacea

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Anaphalis margaritacea (Asteraceae) is a perennial species distributed widely from Asia to North America. In Japan, A. margaritacea consists of three varieties; var. margaritacea distributed mainly in the mountain areas in eastern Japan, var. angustifolia distributed in mountain areas in western Japan, var. yedoensis distributed in riverbanks covered with pebbles throughout Japan. These varieties are considered to have been derived via different speciation patterns such as geographical and ecological speciations. They seem to need taxonomic reexamination based on the molecular analysis because morphological intermediates between these varieties have been sometimes reported. In this study, we collected these three varieties from all over Japan and examined the genetic differentiation among the three varieties using the genome-wide SNPs by MIG-seq.

Analyses based on the SNPs showed that the populations of A. margaritacea var. angustifolia and var. yedoensis were monophyletic, respectively and these lineages were contained with A. margaritacea var. margaritacea, suggesting that they are of a single origin from A. margaritacea var. margaritacea. Population genetics analyses showed that the A. margaritacea complex comprised three clades as a whole: "A. margaritacea var. margaritacea in Hokkaido" clade, "A. margaritacea var. margaritacea in Honshu" clade, and "A. margaritacea var. yedoensis in Hokkaido" clade. Anaphalis margaritacea var. angustifolia was included in "var. margaritacea in Honshu" clade. On the other hand, it is revealed that A. margaritacea var. yedoensis in Honshu have a admixture of the two clades of A. margaritacea var. margaritacea in Honshu and A. margaritacea var. yedoensis in Hokkaido. Based on these results, we will discuss differentiation processes among the three varieties of A. margaritacea.