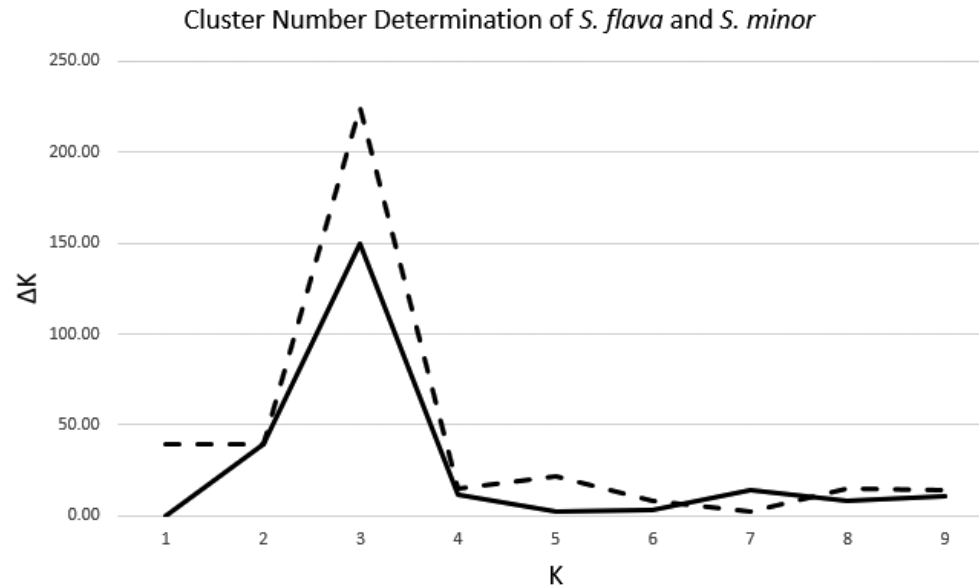


Population Genetic Structure and Natural Establishment of Hybrids Between *Sarracenia flava* and *Sarracenia minor* in Francis Marion National Forest

Jeremy D. Rentsch* and Ryan C. Holland

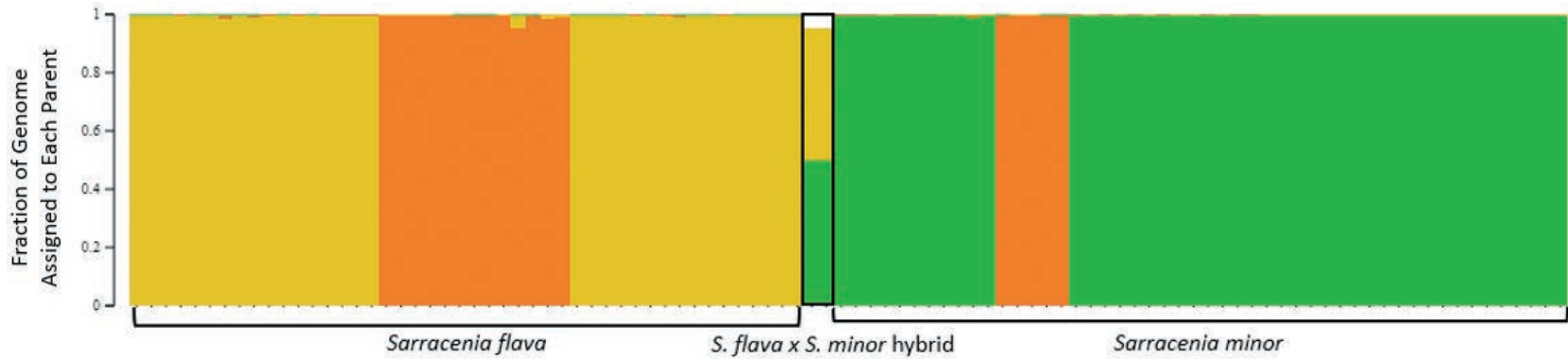
Francis Marion University, 4822 East Palmetto Street, Florence, SC 29505



Supplemental Figure 1. Estimation of cluster number for the 63 *Sarracenia flava* L. and 62 *Sarracenia minor* Walter individuals analyzed at eight nuclear microsatellite loci. Following the methods of Evanno et al. (2005), this ad hoc statistic uses ΔK to help identify the real number of clusters in a data set.

*email address: jrentsch@fmarion.edu

Received 7 April 2019; Accepted 31 January 2020



Supplemental Figure 2. Proportion of the putative F₁ hybrid's genome derived from each parental species; *Sarracenia flava* L. and *Sarracenia minor* Walter. Parental species are assigned a distinct cluster and the hybrid is analyzed as an individual with an unknown cluster or origin. Each allele in the hybrid is then assigned to a parental species of origin.