



The Evolution of Reproductive Systems and Sex-Determining Mechanisms Within *Rumex* (Polygonaceae) Inferred from Nuclear and Chloroplastial Sequence Data

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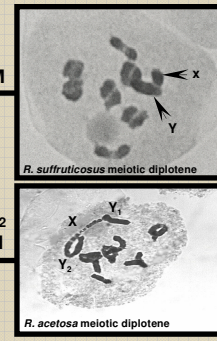
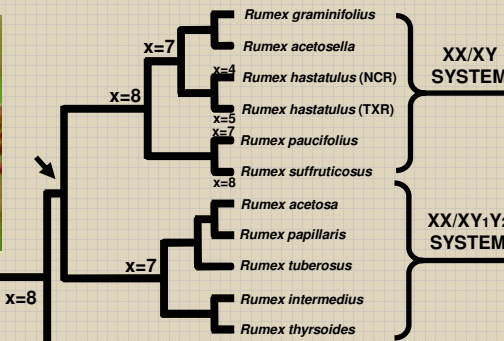
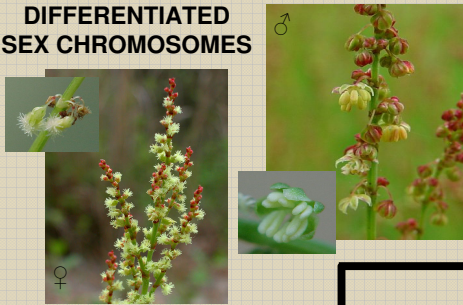
INTRODUCTION

The genus *Rumex* represents an exceptional case to test hypothesis regarding sex-chromosomes origin and evolution that still remain puzzling. In fact, *Rumex* constitutes a big group of species in which almost every mating system is present, comprising of hermaphrodite, polygamous, gynodioecious, monoecious and dioecious representatives. Here we present an evolutionary picture of the genus concerning basic chromosome number, sex-chromosomes and mating-systems evolution.

The genus *Rumex* is currently divided into four subgenera (Table 1): *Acetosella*, *Acetosa*, *Platypodium*, and *Rumex*. *Acetosella* contains two species, *R. acetosella* (which has several subspecies) and *R. graminifolius*. These species are dioecious and have a sex-determination mechanism based on the presence of an active Y and a simple chromosome system XXXY. Within the subgenus *Acetosa*, the section *Acetosa* is composed of *R. acetosa* and its relatives, which form an homogeneous group of species characterized by similar morphological and karyological characteristics, including a XXXY₂ sex-chromosome system plus a sex-determination mechanism based on the X/A balance. However, within the section *Americanae* of the subgenus *Acetosa*, there are two species: *R. paucifolius*, which has the XXXY system; and *R. hastatus*, which has two chromosomal races, one with the XXXY (called the "Texas race") and the other with the XXXY₂ (called the "North Carolina" race). Also, the second race has an X/A-based sex-determination mechanism, while the XXXY race has a Y-based one. Furthermore, the subgenus *Acetosa* contains four additional sections: *Scutarii*, *Vesicarii*, *Hastati*, and *Afroacetosa*. The first two are comprised of hermaphroditic and polygamous species. Strikingly, *Scutarii* has a dioecious species, *R. suffruticosus*, for which no chromosomal data were available until now. The sections *Hastati* and *Afroacetosa* are comprised of polygamous and gynodioecious species as well as a dioecious one, *R. sagittatus*, which lacks differentiated sex chromosomes. Meanwhile, the third subgenus, *Platypodium*, has one species (and several subspecies), *R. bucephalophorus*, which is hermaphroditic. Finally, the subgenus *Rumex* is composed of hermaphroditic species, although endemic Hawaiian species such as *R. giganteus* have evolved towards monoecy.

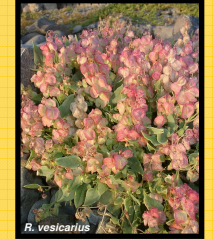
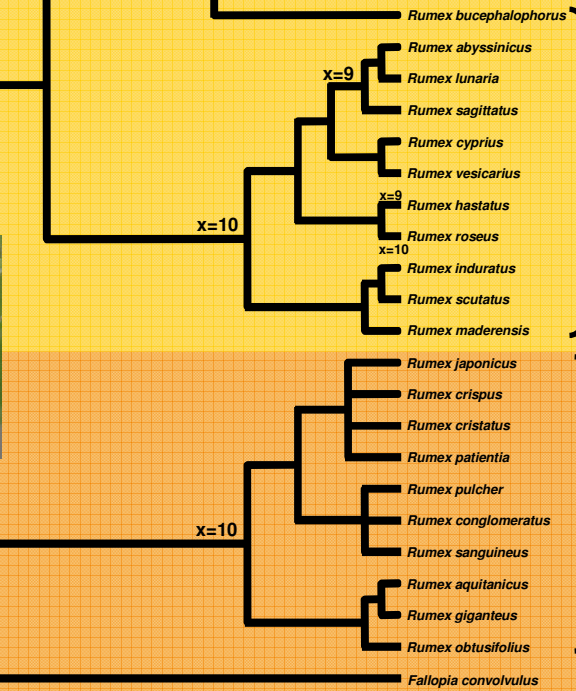
If this classification reflects the phylogeny of the genus, it implies that dioecy has appeared several times over the evolution of *Rumex* species directly from a hermaphroditic ancestor; also this classification involves that there has been no evolutionary constraint on the evolution of sexual systems and that forward and reverse evolution occur with equal probability. We test here whether these assumptions are correct or not by means of a phylogenetic analysis of the genus *Rumex* based on one nuclear and one chloroplastial marker (Table 1). Specifically, we want to address whether: i) dioecy has appeared once or several times in *Rumex*; ii) the Y-based sex-determination mechanism precedes to the X/A mechanism; iii) the multiple sex-chromosome system derived from an XXXY system; iv) a different infra-generic classification can be proposed for *Rumex* species consistent with their mating system and with their karyotype evolution.

DIFFERENTIATED SEX CHROMOSOMES



DIOECY

NON-DIFFERENTIATED SEX CHROMOSOMES



HERMAPHRODITISM
POLYGAMY
GYNODIOECY



HERMAPHRODITISM

RESULTS AND CONCLUSIONS

We found three phylogenetic clades within *Rumex*, implying a revision of the systematics for this genus. Thus, the subgenus *Rumex* appears coherent since all their species appear to form a well-supported (100% bootstrap) clade of closely related species. However, we find no evidence to maintain the subgenera *Platypodium* and *Acetosella*. Thus, *R. bucephalophorus*, as the only representative species of the subgenus *Platypodium*, is closely related to all the Eurasian and American dioecious species of subgenera *Acetosa* and *Acetosella*. On the contrary, our phylogeny finds support to include the species of these two subgenera within *Acetosa* and to separate this subgenus into two groups: one including *R. bucephalophorus* together with all the Eurasian and American dioecious species of the subgenera *Acetosa* and *Acetosella*; and a second group which would include the hermaphroditic/polygamous and the polygamous/gynodioecious species of the subgenus *Acetosa* together with the African dioecious species *R. sagittatus* (also, subgenus *Acetosa*). Within the main clade of dioecious species, there are two subclades. One of the subclades is composed by the XXXY species, which includes the Eurasian species *R. suffruticosus*, an endemic species of the Iberian Peninsula (for which we have described a XXXY system), *R. acetosella* and its relative *R. graminifolius* and the American dioecious species *R. paucifolius* and *R. hastatus*. The other subclade is composed of the species *R. acetosa* and its relatives, which form a homogeneous group of species characterized by similar morphological and karyological characteristics, including an XXXY₂ sex-chromosome system.

This new classification is consistent with the evolution of mating system and sex determination of *Rumex* species and of the karyotype. In contrast to the current view, this new phylogeny suggest a common origin for all Eurasian and American dioecious species of *Rumex* with gynodioecy as an intermediate state on the way to dioecy, since a second different lineage could be evolving from hermaphroditism towards dioecy via gynodioecy in Africa. Our results support the contention that sex determination based on the balance between the number of X chromosomes and the number of autosomes (X/A balance) has evolved secondarily from male-determining Y mechanisms and that multiple sex chromosome systems, XXXY₂, were derived twice from an XXXY system. Also, the resulting phylogeny is consistent with a classification of *Rumex* species according to their basic chromosome number, implying that the evolution of *Rumex* species might have followed a process of chromosomal reduction from x=10 (hermaphroditic species) toward x=7 (dioecious species) stages with x=9 and x=8 chromosomes (polygamous and gynodioecious species).

Table 1. Taxonomic and molecular data of Rumex species. The table lists species names, their subgenera and sections, chromosome numbers (x), and accession numbers for nuclear and chloroplast DNA sequences. It also includes a key for abbreviations and symbols used in the text.

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