

## **Data Management Plan**

This project will conform to NSF policy on the timely dissemination and sharing of research results. The major product of this proposal will be gene sequences and allele frequency data. All sequence data will be made publicly available at the time of publication through the NCBI database (GenBank), which is standard practice in the field of molecular ecology. Raw frequency data will be published as on-line supplemental files with the corresponding paper. Databases will include unique specimen identifiers, collection locality information, voucher ID # (if applicable), and the haplotypes or alleles associated with that individual. Vouchers of specimens from each location will be deposited in the malacology collection of the Los Angeles Natural History Museum (located ~5 miles from CSULA), where the PI has previously deposited type specimens.

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