

Rethinking phylogenetics using Caryophyllales (angiosperms), *matK* gene and *trnK*
intron as experimental platform

Sunny Sheliese Crawley

Dissertation submitted to the faculty of the Virginia Polytechnic Institute and State
University in partial fulfillment of the requirements for the degree of

Doctor of Philosophy
In
Biological Sciences

Khidir W. Hilu
Eric P. Beers
Carla V. Finkielstein
Jill C. Sible

December 2, 2011
Blacksburg, Virginia

Keywords: (phylogeny, missing data, caryophyllids, *trnK* intron, *matK*, RNA editing,
gnetophytes)

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