

GENOMIC RESOURCES NOTE

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GENOMIC RESOURCES DEVELOPMENT CONSORTIUM,¹ MATTHEW G. KING,² SÉBASTIEN RENAUT,³ LOREN H. RIESEBERG^{2,4} and HEATHER C. ROWE³¹Molecular Ecology Resources Editorial Office, 6270 University Blvd, Vancouver, BC V6T 1Z4, Canada, ²DuPont Pioneer, Box 1000, Johnston, IA 50131-0184, USA, ³Biodiversity Research Centre and Department of Botany, University of British Columbia, Vancouver, BC V6T 1Z4, Canada, ⁴Department of Biology, Center for Genomics and Bioinformatics, Indiana University, 1001 East Third Street, Bloomington, IN 47405, USA

Abstract

This article documents the public availability of three transcriptome sequences. These genomic resources were developed for the following species: *Helianthus anomalus*, *Helianthus deserticola* and *Helianthus paradoxus*.

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Table 1 contains information on the focal species, data type and resource developed, as well as access details for the data. The authors responsible for each genomic resource are listed in the final column. Full descriptions of how each resource was developed and tested are uploaded as Supplemental Material with the online

version of this manuscript; both Table 1 and the supplemental files are sorted by the generic name of the focal taxon.

Supporting Information

Additional Supporting Information may be found in the online version of this article:

Data S1 Renaut et al. GR Note.

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Species (no. of individuals)	Data type	Resources	Authors
<i>Helianthus anomalus</i> (3), <i>H. deserticola</i> (7), <i>H. paradoxus</i> (8)	Non-normalized cDNA	Sequence files: http://www.ncbi.nlm.nih.gov/sra/?term=PRJNA188794 Alignments: http://www.ncbi.nlm.nih.gov/sra/?term=PRJNA188794 SNPs: http://dx.doi.org/10.5061/dryad.fj594 Code: https://github.com/seb951/helianthus_hybrid_species_transcriptome	Renaut, Sébastien; King, Matthew G.; Rowe, Heather C.; Rieseberg, Loren H.