

GENOMIC RESOURCES NOTE

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Abstract

This article documents the public availability of (i) transcriptome sequence data and assembly for the rostrum dace (*Leuciscus burdigalensis*) naturally infected by a copepod ectoparasite (*Tracheliastes polycolpus*) and (ii) SNPs identified and validated from RAD sequencing for the Ugandan red colobus (*Procolobus rufomitratu tephrosceles*) using RAD sequencing.

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 Information with the online version of this manuscript.

Table 1 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

Species (no. of individuals)	Data type	Resources	Authors
<i>Leuciscus burdigalensis</i> (8)	Transcriptome sequencing and assembly	Transcriptome sequence data: NCBI BioProject PRJNA264971 Contig assembly: doi:10.5061/dryad.6365v Blast hits (with <i>Danio rerio</i> cDNA database): doi:10.5061/dryad.6365v	Olivier Rey, Géraldine Loot, Olivier Bouchez, Simon Blanchet
<i>Procolobus rufomitratu tephrosceles</i> (24)	SNP discovery and validation from RAD sequencing	RAD sequence data: NCBI Sequence Read Archive SRP050135 SNP information including validation results: http://dx.doi.org/10.5061/dryad.404fh	Maria Jose Ruiz-Lopez, Tony L. Goldberg, Colin A. Chapman, Patrick A. Omeja, James H. Jones, William M. Switzer, Paul D. Etter, Eric A. Johnson, Nelson Ting

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