

S4 Table. Amino acid sequence of riboflavin transporters used to construct the phylogenetic tree from Fig. 2B.

Classification	Species	Name	Accession number	Reference
Mammals	<i>Homo sapiens</i>	RFVT1	Q9NWF4.2	[1]
	<i>Homo sapiens</i>	RFVT3	Q9NQ40.4	[1]
	<i>Homo sapiens</i>	RFVT2	Q9HAB3.1	[1]
	<i>Mus musculus</i>	RFVT2	Q9D8F3.1	[2]
	<i>Pan paniscus</i>	member 1	XP_003810223.1	Predicted
	<i>Pongo abelii</i>	member 2	XP_002833179.2	Predicted
	<i>Macaca fascicularis</i>	member 2	XP_005564405.1	Predicted
	<i>Macaca nemestrina</i>	member 1	XP_011724203.1	Predicted
	<i>Equus caballus</i>	member 1	XP_014583611.1	Predicted
	<i>Tursiops truncatus</i>	member 2	XP_004319077.1	Predicted
	<i>Orcinus orca</i>	member 2	XP_012386875.1	Predicted
	<i>Bos mutus</i>	member 2	XP_014334325.1	Predicted
	<i>Ictidomys tridecemlineatus</i>	member 2	XP_005316095.1	Predicted
	<i>Cricetulus griseus</i>	member 2	XP_003507685.1	Predicted
	<i>Equus przewalskii</i>	member 2	XP_008508436.1	Predicted
	<i>Ceratotherium simum simum</i>	member	XP_004443067.1	Predicted
	<i>Ratus norvegicus</i>	member 3	NP_001032275.1	Predicted
	<i>Myotis brandtii</i>	member 3	XP_014402039.1	Predicted
	<i>Gorilla gorilla gorilla</i>	member 3	XP_004061693.1	Predicted
	<i>Pan troglodytes</i>	member 3	XP_009434927.1	Predicted
<i>Macaca mulatta</i>	member 2	NP_001181490.1	Predicted	
<i>Panthalops hodgsonii</i>	member 3	XP_005981822.1	Predicted	
Fungi	<i>Saccharomyces cerevisiae</i>		NP_014951.4	[3]
	<i>Arthroderma otae</i>		XP_002843476.1	Predicted
	<i>Colletotrichum fioriniae</i>		XP_007594293.1	Predicted
	<i>Colletotrichum gloeosporioides</i>		XP_007279280.1	Predicted
	<i>Eutypa lata</i>		XP_007792122.1	Predicted
	<i>Fibroporia radiculosa</i>		XP_012184704.1	Predicted
	<i>Gaeumannomyces graminis</i>		XP_009221625.1	Predicted
	<i>Magnaporthe oryzae</i>	MCH5	XP_003716530.1	Predicted
	<i>Marssonina brunnea</i>		XP_007296567.1	Predicted
	<i>Microsporium gypseum</i>		XP_003169176.1	Predicted
	<i>Ogataea parapolyomorpha</i>		XP_013933620.1	Predicted
	<i>Paracoccidioides lutzii</i>		XP_002797527.1	Predicted
	<i>Verticillium alfalfa</i>		XP_003000372.1	Predicted
	<i>Verticillium dahlia</i>		XP_009648470.1	Predicted
<i>Wickerhamomyces ciferrii</i>		XP_011271917.1	Predicted	
Nematodes	<i>Caenorhabditis elegans</i>		CCD69385.2	[4]
	<i>Brugia malayi</i>		XP_001896372.1	Predicted
	<i>Loa loa</i>	rft	XP_003145255.1	Predicted
	<i>Necator americanus</i>		XP_013300104.1	Predicted
	<i>Trichinella spiralis</i>		XP_003380128.1	Predicted

The sequences were obtained from the NCBI Protein database annotated as predicted riboflavin transporters. Function has not been established for the entries with the exception of transporters belonging to *H. sapiens*, *M. musculus*, *C. elegans* and *S. cerevisiae*.

Supplementary References

1. Yonezawa A, Inui KI. Novel riboflavin transporter family RFVT/SLC52: Identification, nomenclature, functional characterization and genetic diseases of RFVT/SLC52. *Mol Aspects Med.* 2013;34: 693–701. doi:10.1016/j.mam.2012.07.014
2. Yao Y, Yonezawa A, Yoshimatsu H, Omura T, Masuda S, Matsubara K. Involvement of riboflavin transporter RFVT2/Slc52a2 in hepatic homeostasis of riboflavin in mice. *Eur J Pharmacol.* 2013;714: 281–7. doi:10.1016/j.ejphar.2013.07.042
3. Reihl P, Stolz J. The monocarboxylate transporter homolog Mch5p catalyzes riboflavin (vitamin B2) uptake in *Saccharomyces cerevisiae*. *J Biol Chem.* 2005;280: 39809–39817. doi:10.1074/jbc.M505002200
4. Biswas A, Elmatari D, Rothman J, LaMunyon CW, Said HM. Identification and functional characterization of the *Caenorhabditis elegans* riboflavin transporters rft-1 and rft-2. *PLoS One.* 2013;8: e58190. doi:10.1371/journal.pone.0058190