

# **Sexual selection and the adaptive evolution of PKDREJ protein in primates and rodents**

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## **Supplementary data**

**Supplementary Figure S1** Amino acid alignment of PKDREJ in Muroidea.

**Supplementary Figure S2** Cladograms of rodents used for evolutionary analyses.

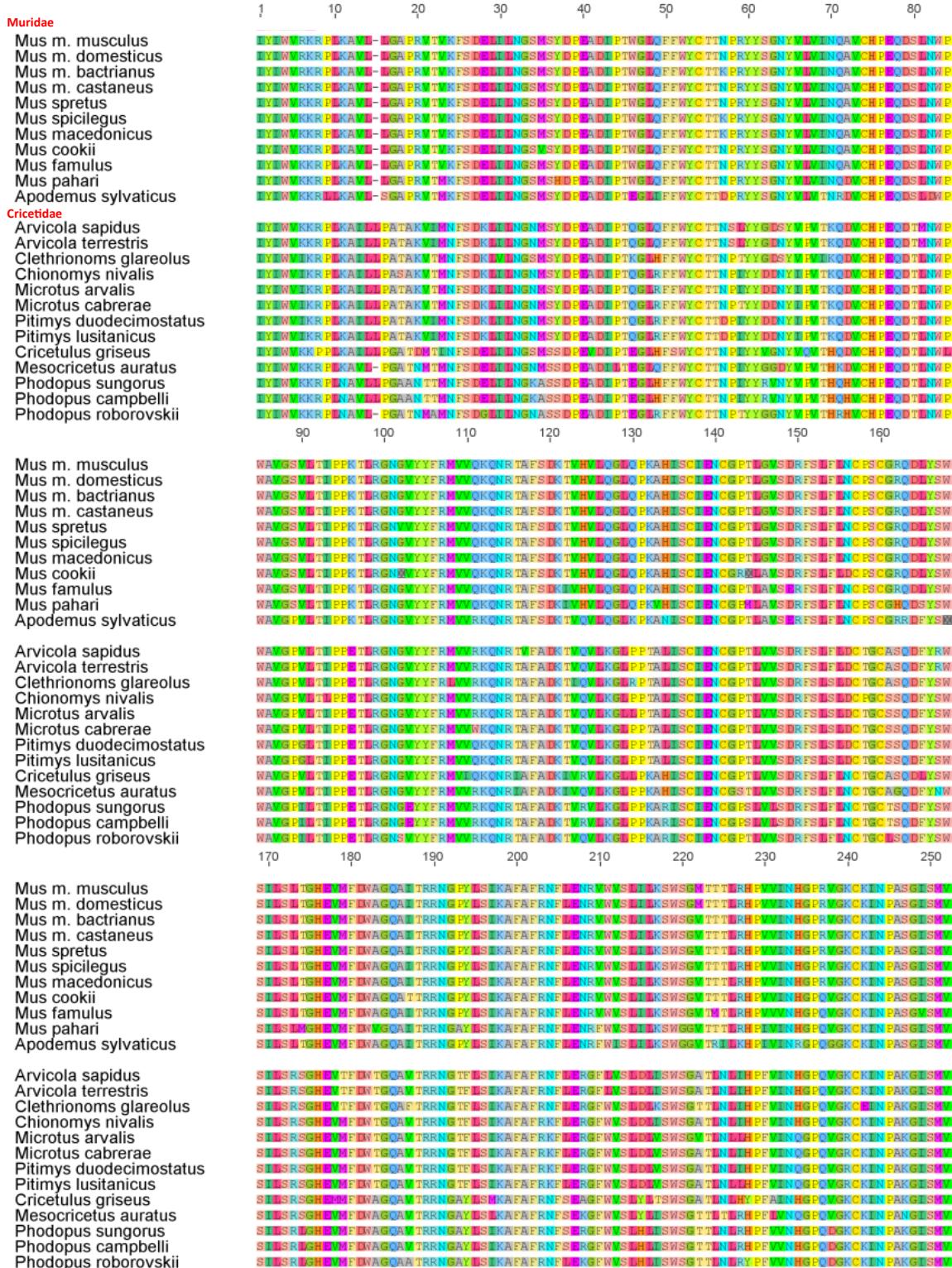
**Supplementary Table SI** Primers used for amplification of PKDREJ in rodent species.

**Supplementary Table SII** Body mass, testes mass and residual testes mass of primate species.

**Supplementary Table SIII** Body mass, testes mass and residual testes mass of rodent species.

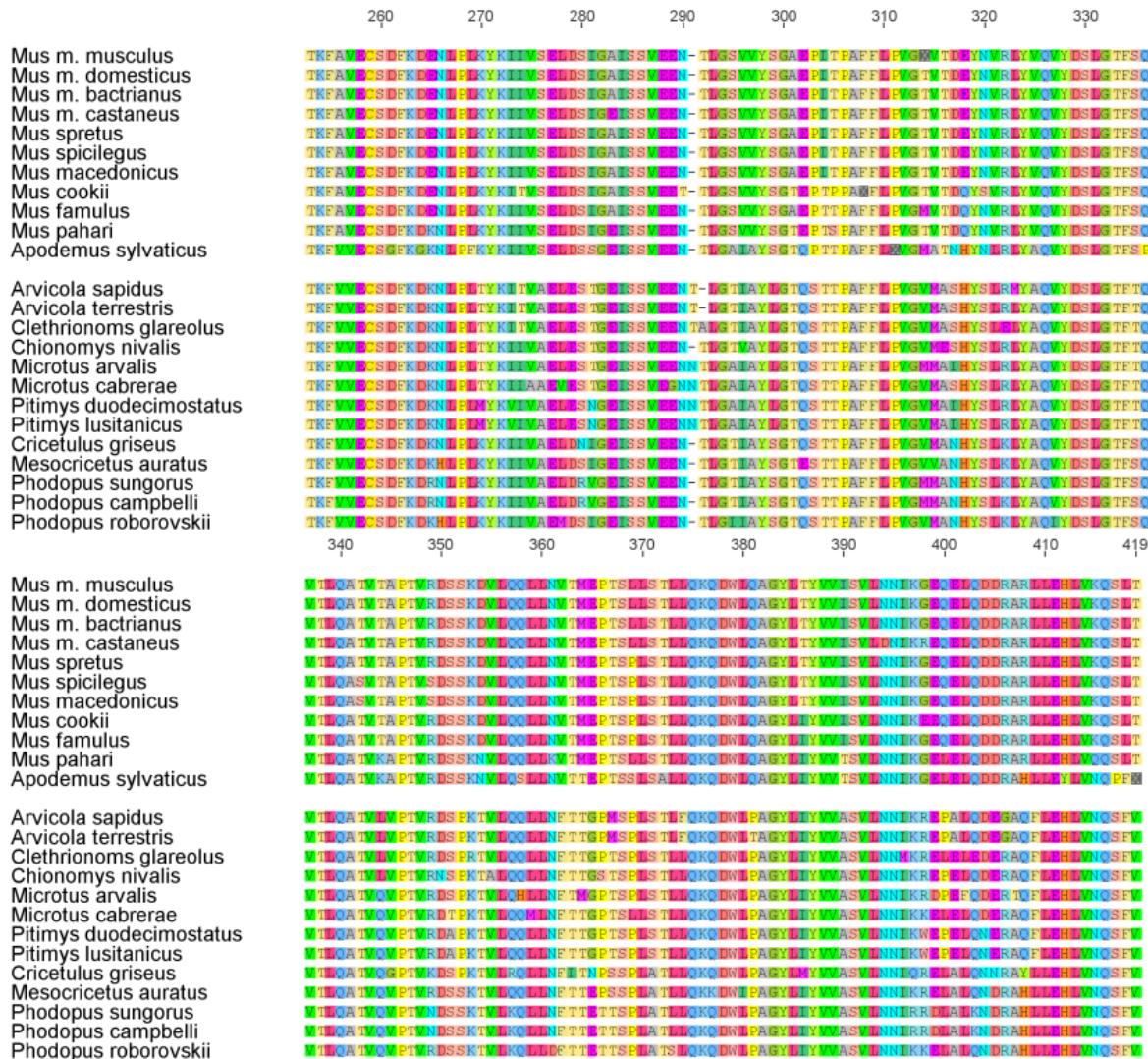
**Supplementary Table SIV** Lineage-specific evolutionary rates of PKDREJ in rodents.

**Supplementary Figure S1** Amino acid alignment of PKDREJ in Muroidea. Protein sequences of the REJ domain for the 24 muroid species used in the study are shown.

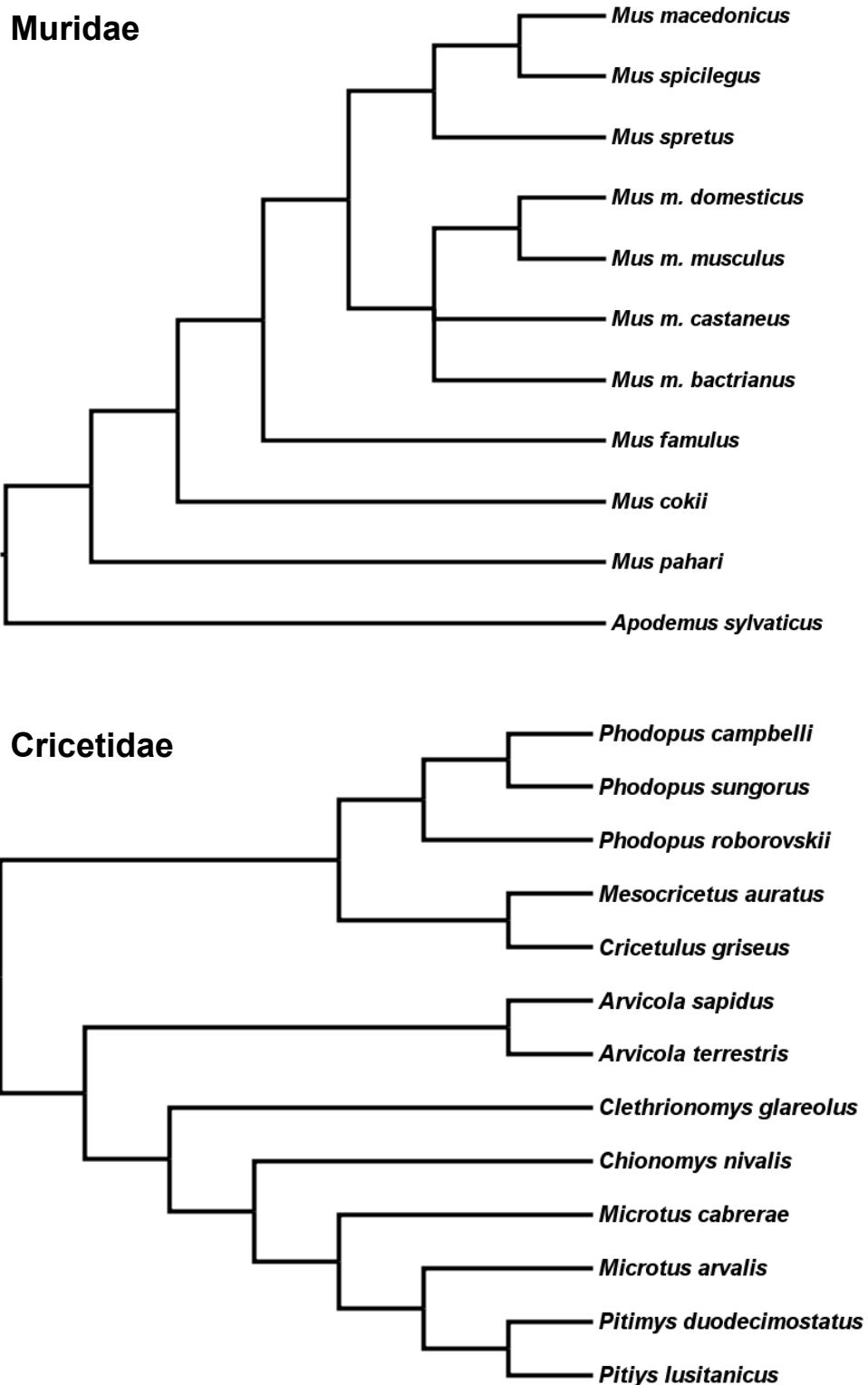


(cont.)

Supplementary Figure S1 (cont.)



**Supplementary Figure S2** Cladograms of rodents used for evolutionary analyses. The analyses were performed considering the two possible topologies derived from the politomy of *Mus musculus* subspecies.



**Supplementary Table SI** Primers used for amplification of PKDREJ in rodent species.

Primer	Sequence
PKDREJ 1-Fw	TTACGGGGTGTATGTGTTAAC
PKDREJ 1-Rv	CTGGGTTAATTGGTAATCAC
PKDREJ 3-Rv	TCGTCCCTGTGTTCATAGCAA
PKDREJ Vol 1-Fw	CTGTCCATCAACAGGTGGGA
PKDREJ Vol 1-Rv	AGTACCCCTGGACGAGATCTGC
PKDREJ 1 Ham-Fw	CCTGTAACATACAGTCCGTG
PKDREJ 1 Ham-Rv	ATCACACCCTGGCAGATCTG
Fw int PKDREJ	ATGTTGACTGGGCAGGGCAAG
Rv int PKDREJ	CCTGCCAGTCAAACATCACTTCA

Species	Primer pairs
<i>Mus m. musculus</i> , <i>Mus m. domesticus</i> , <i>Mus m. castaneus</i> , <i>Mus m. bactrianus</i> , <i>Mus spreitus</i> , <i>Mus spicilegus</i> , <i>Mus macedonicus</i> , <i>Mus famulus</i> , <i>Mus cokii</i> , <i>Apodemus sylvaticus</i> .	PKDREJ 1-Fw / PKDREJ 1-Rv
<i>Mus pahari</i> .	PKDREJ 1-Fw / PKDREJ 3-Rv
<i>Arvicola sapidus</i> , <i>Arvicola terrestris</i> , <i>Clethrionomys glareolus</i> , <i>Chinomys nivalis</i> , <i>Pitymys duodecimostatus</i> , <i>Pitymys lusitanicus</i> .	PKDREJ 1-Fw / PKDREJ Vol 1-Rv
<i>Microtus arvalis</i> , <i>Microtus cabrerae</i> .	PKDREJ Vol 1-Fw / PKDREJ Vol 1-Rv
<i>Cricetulus griseus</i> , <i>Mesocricetus auratus</i> , <i>Phodopus sungorus</i> , <i>Phodopus campbelli</i> , <i>Phodopus roborovskii</i> .	PKDREJ Ham 1-Fw / PKDREJ Ham 1-Rv

**Supplementary Table SII** Body mass, testes mass and log residual testes mass of primates.

Species	Body mass (g)	Testes mass (g)	log residual testes mass	Reference
<i>Macaca mulatta</i>	10430	76.0	0.569	Kenagy and Trombulak (1986)
<i>Macaca nemestrina</i>	9980	66.7	0.525	Kenagy and Trombulak (1986)
<i>Ateles geoffroyi</i>	9800	64.2	0.514	Anderson and Dixson (2004)
<i>Pan paniscus</i>	39100	135.2	0.456	Anderson and Dixson (2004)
<i>Pan troglodytes</i>	44340	118.8	0.366	Kenagy and Trombulak (1986)
<i>Homo sapiens</i>	63540	50.2	-0.107	Kenagy and Trombulak (1986)
<i>Callithrix jacchus</i>	320	1.3	-0.24	Kenagy and Trombulak (1986)
<i>Pongo pygmaeus</i>	74640	36.3	-0.292	Kenagy and Trombulak (1986)
<i>Saguinus labiatus</i>	501	1.5	-0.307	Kenagy and Trombulak (1986)
<i>Lemur catta</i>	27000	17.8	-0.322	Anderson and Dixson (2004)
<i>Erythrocebus patas</i>	13000	7.2	-0.515	Anderson and Dixson (2004)
<i>Gorilla gorilla</i>	134000	23.2	-0.648	Kenagy and Trombulak (1986)

**Supplementary Table SIII** Body mass, testes mass and log residual testes mass of rodents.

Family	Species	Body mass (g)	Testes mass (g)	log residual testes mass
Muridae	<i>Apodemus sylvaticus</i>	30.43	0.962	0.58
	<i>Mus spicilegus</i>	14.49	0.409	0.23
	<i>Mus spretus</i>	17.01	0.295	0.08
	<i>Mus macedonicus</i>	18.40	0.282	0.05
	<i>Mus cookii</i>	23.67	0.305	0.01
	<i>Mus m. bactrianus</i>	18.60	0.175	-0.06
	<i>Mus m. domesticus</i>	16.36	0.135	-0.07
	<i>Mus m. musculus</i>	21.13	0.134	-0.13
	<i>Mus m. castaneus</i>	19.50	0.1	-0.14
	<i>Mus pahari</i>	30.08	0.118	-0.26
	<i>Mus famulus</i>	27.40	0.053	-0.29
Cricetidae	<i>Mesocricetus auratus</i>	88.59	2.893	1.55
	<i>Cricetulus griseus</i>	36.37	2.27	1.41
	<i>Phodopus campbelli</i>	45.38	2.105	1.16
	<i>Phodopus sungorus</i>	39.42	1.211	0.32
	<i>Phodopus roborovskii</i>	21.07	0.706	-0.02
	<i>Chionomys nivalis</i>	43.64	0.851	-0.08
	<i>Arvicola sapidus</i>	209.75	2.144	-0.31
	<i>Clethrionomys glareolus</i>	25.65	0.401	-0.37
	<i>Microtus arvalis</i>	36.40	0.285	-0.58
	<i>Pitimys lusitanicus</i>	17.73	0.093	-0.6
	<i>Pitimys duodecimostatus</i>	29.76	0.076	-0.73
	<i>Microtus cabrerae</i>	44.27	0.142	-0.79
	<i>Arvicola terrestris</i>	91.56	0.411	-0.96

**Supplementary Table IV** Lineage-specific evolutionary rates of PKDREJ in rodents.  
 Parameters were calculated from the root of the phylogeny to the terminal branches.

Family	Species	dN	dS	$\omega$
Muridae	<i>Mus spicilegus</i>	0.0829	0.1654	0.5013
	<i>Mus spretus</i>	0.0794	0.1786	0.4449
	<i>Mus macedonicus</i>	0.0829	0.1654	0.5015
	<i>Mus cokii</i>	0.0844	0.1617	0.5217
	<i>Mus musculus bactrianus</i>	0.0818	0.1654	0.4944
	<i>Mus domesticus</i>	0.0806	0.1654	0.4873
	<i>Mus musculus musculus</i>	0.0806	0.1654	0.4873
	<i>Mus musculus castaneus</i>	0.0829	0.1680	0.4937
	<i>Mus pahari</i>	0.0775	0.1661	0.4663
	<i>Mus famulus</i>	0.0806	0.1641	0.4910
	<i>Apodemus sylvaticus</i>	0.0627	0.1478	0.4242
Cricetidae	<i>Arvicola sapidus</i>	0.1070	0.2850	0.3754
	<i>Arvicola terrestris</i>	0.1050	0.2930	0.3584
	<i>Clethrionomys glareolus</i>	0.1210	0.3010	0.4020
	<i>Chionomys nivalis</i>	0.1120	0.2810	0.3986
	<i>Microtus arvalis</i>	0.1190	0.2910	0.4089
	<i>Microtus cabrerae</i>	0.1210	0.2910	0.4158
	<i>Pitimys duodecimostatus</i>	0.1210	0.2910	0.4158
	<i>Pitimys lusitanicus</i>	0.1220	0.2940	0.4150
	<i>Mesocricetus auratus</i>	0.0990	0.2230	0.4439
	<i>Cricetulus griseus</i>	0.1210	0.2320	0.5216
	<i>Phodopus sungorus</i>	0.1080	0.2620	0.4122
	<i>Phodopus campbelli</i>	0.1080	0.2620	0.4122
	<i>Phodopus roborovskii</i>	0.1060	0.2880	0.3681