S4 Table. Likelihood ratio tests for positive selection using branch-site models

Gene	Models Compared	2AlnL	df	p value
MRPS5	Branch-site #1 (Mammals)	14.8240	1	p<0.001
	Branch-site #2 (Birds)	8.9825	1	p = 0.0027
	Branch-site #3 (Birds/Reptiles)	3.9416	1	p = 0.047
	Branch-site #4 (Amphibian)	12.7362	1	p<0.001
	Branch-site #5 (Fish)	10.0984	1	p = 0.0015
FOXO3a	Branch-site #1 (Mammals)	7.0982	1	p=0.008
	Branch-site #2 (Birds)	6.0542	1	p=0.014
	Branch-site #3 (Birds/Reptiles)	6.2140	1	p = 0.013
	Branch-site #4 (Amphibian)	21.9256	1	p<0.001
	Branch-site #5 (Fish)	15.2409	1	p<0.001
PARP1	Branch-site #1 (Mammals)	7.2486	1	p=0.007
	Branch-site #2 (Birds)	4.8149	1	p=0.03
	Branch-site #3 (Birds/Reptiles)	5.8271	1	p=0.016
	Branch-site #4 (Amphibian)	21.5609	1	p<0.001
	Branch-site #5 (Fish)	24.4764	1	p<0.001
PARP2	Branch-site #1 (Mammals)	11.3767	1	p<0.001
	Branch-site (Reptiles)	8.8035	1	p=0.003
	Branch-site #4 (Amphibian)	21.9734	1	p<0.001
	Branch-site #5 (Fish)	37.4631	1	p<0.001
PPARGCIA	Branch-site #1 (Mammals)	6.1631	1	p=0.013
	Branch-site #2 (Birds)	3.4102	1	p=0.06
	Branch-site #3 (Birds/Reptiles)	6.7810	1	p=0.009
	Branch-site #4 (Amphibian)	6.6429	1	p=0.01
	Branch-site #5 (Fish)	5.0601	1	p=0.024
SIRTI	Branch-site #1 (Mammals)	4.0928	1	p=0.04
	Branch-site #2 (Birds)	4.7832	1	p=0.03
	Branch-site #3 (Birds/Reptiles)	5.0012	1	p=0.025
	Branch-site #4 (Amphibian)	9.0935	1	p=0.003
	Branch-site #5 (Fish)	5.1349	1	p=0.023
SIRT2	Branch-site #1 (Mammals)	7.0923	1	p=0.008
	Branch-site #2 (Birds)	6.9963	1	p=0.008
	Branch-site #3 (Birds/Reptiles)	15.1935	1	p<0.001
	Branch-site #4 (Amphibian)	4.0173	1	p=0.045
	Branch-site #5 (Fish)	28.0972	1	p<0.001
SOD3	Branch-site #1 (Mammals)	6.0325	1	p=0.014
	Branch-site #2 (Birds)	6.4685	1	p=0.011
	Branch-site #3 (Birds/Reptiles)	8.0995	1	p=0.004
	Branch-site #4 (Amphibian)	9.1009	1	p=0.002
	Branch-site #5 (Fish)	12.1989	1	p<0.001
TP53	Branch-site #1 (Mammals)	8.9173	1	p=0.003
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	Branch-site #2 (Birds)	3.8893	1	p=0.048
	Branch-site #3 (Birds/Reptiles)	5.0916	1	p=0.024

p=0.001
p=0.001
p<0.001
p=0.002
p=0.003
p=0.008
p=0.003
p<0.001

Note: The p value adjusted by Bonferroni correction for multiple testing is 0.001.