

Table S3. Statistics and sample size from the test for allometry for each family, under the null hypothesis of isometry

	<i>N</i>	% predicted	<i>p-value</i>
<i>Tadpoles (shape ~ log(TL))</i>			
Alsodidae	30	21.19	<0.01
Batrachylidae	11	12.19	0.27
Cycloramphidae	26	12.32	0.04
Hylodidae	45	6.70	0.03
<i>Adults (shape ~ log(SVL))</i>			
Alsodidae	30	10.25	0.06
Batrachylidae	11	36.19	<0.01
Cycloramphidae	26	5.86	0.34
Hylodidae	45	12.10	0.01

¹ TL: total length; ² SVL: snout-vent length; *N*: number of individuals by family.