

Table A2 - Confusion matrix used for the discriminant analysis (misclassification error equals to 0.22) performed on the comparative sample pooled by genus (N=number of individuals per group).

	<i>Panthera</i> (N=9)	<i>Leptailurus</i> (N=1)	<i>Lynx</i> (N=3)	<i>Puma</i> (N=2)	<i>Acinonyx</i> (N=1)	<i>Felis</i> (N=3)
<i>Panthera</i>	8	0	0	1	1	0
<i>Leptailurus</i>	0	0	0	0	0	0
<i>Lynx</i>	0	0	3	0	0	0
<i>Puma</i>	1	0	0	1	0	0
<i>Acinonyx</i>	0	0	0	0	0	0