

Supplementary Table 2 Cross-species utility of 40 conserved microsatellite markers in three squirrel species.

Source species of primer set ^a	Squirrel species tested		
	<i>Hylopetes lepidus</i>	<i>Hylopetes phayrei</i>	<i>Menetes berdmorei</i>
<i>Hylopetes lepidus</i>	9/9	8/9	5/9
<i>Hylopetes phayrei</i>	8/10	10/10	7/10
<i>Menetes berdmorei</i>	0/8	1/8	8/8
<i>Glaucmys</i> / <i>Sciurus</i> species ^b	8/13	7/13	5/13

Fractions indicates the number of loci that amplified and were polymorphic in the test species relative to the number of loci that were designed from the stated source species

^a All primer set sequences match the source species stated and the thirteen-lined ground squirrel (*Spermophilus tridecemlineatus*)

^b *Glaucmys sabrinus*, *G. volans*, *Sciurus lis*, *S. niger* or *S. vulgaris* microsatellite sequences extracted from GenBank