

## Supplementary Materials

### Supplementary Text

#### Specimens Examined

N=469

Material studied: sn=skin, sl=skull, d=dental, p=penis

*Aeretes melanopterus*: AMNH45328 (sn, sl, d), 56933 (sn, sl, d). USNM219206 (sn, sl, d), 219205 (sn, sl, d). MNHNZM-MO-1874-551 (sl, d), ZM-MO-1867-554 (sn).

*Aeromys tephromelas*: FMNH90437 (sl, d). USNM291285 (sn, sl, d), 292649 (sn, sl, d), 292650 (sn, sl, d), 481192 (sn, sl, d). BMNH71.2736 (sn), 73.7.3.1 (sn, sl, d), 61.1166 (sl, d).

*Aeromys thomasi*: FMNH108895 (sl, d), 108896 (sl, d). USNM317237 (sn, sl, d).

*Belomys pearsonii*: AMNH87419 (sn, sl, d), 114889 (sn, sl, d), 167889 (sn, sl, d), 167890 (sl, d), 174853 (sl, d). FMNH35443 (sl, d), 76385 (sl, d). USNM257845 (sn, sl, d), 308160 (sn, sl, d), 358354 (sn, sl, d), 358355 (sn, sl, d), 358356 (sn, sl, d), 258345 (p). BMNH15.5.5.43 (sn, sl, d), 79.11.21.381 (sl, d), 85.8.1.136 (sl, d). KIZ034940 (sn, sl, d), 0305143 (sn, sl, d).

*Biswamoyopterus biswasi*: ZSI20705 (sn, sl, d).

*Biswamoyopterus gaoligongensis*: KIZ035622 (sn, sl, d), 034924 (sn, sl, d, p).

*Biswamoyopterus laoensis*: NUoLFES.MM.12.163 (sn, sl, d).

*Eoglaucomys fimbriatus*: USNM173361 (sn, sl, d), 173367 (sn, sl, d), 353239 (sn, sl, d), 353241 (sn, sl, d), 353243 (sn, sl, d), 410953 (sn, sl, d), 429353 (p). BMNH7.8.1.2 (sn, sl, d).

*Eupetaurus cinereus*: BMNH88.9.28.1 (sn), 17.4.8.1 (sn), 14.2.10.1 (sn), 20.1.17.6 (sn). ZSI9492 (sn, sl, d).

*Eupetaurus tibetensis*: ZSI19103 (sn). BMNH88.9.29.1 (sn), 23.11.10.2 (sn). NML19524 (sn, sl, d).

*Eupetaurus nivamons*: KIZ034192 (sn, sl, d), 034189 (sn, sl, d), 034190 (sn, sl, d), 035087 (sn, sl, d, p), 035088 (sn, sl, d).

*Glaucomys sabrinus*: AMNH95193 (p), USNM129708 (sn, sl, d), 140192 (sn, sl, d), 235940 (sn, sl, d), 19909 (sn, sl, d), 24271 (sn, sl, d), 68753 (sn, sl, d), 83152 (sn, sl, d), 87310 (sn, sl, d), 94310 (sn, sl, d), 136137 (sn, sl, d), 256993 (sn, sl, d), 260420 (sn, sl, d), 292278 (sn, sl, d).

BMNH7.7.7.3994 (sn, sl, d).

*Glaucomys volans*: AMNH188251 (p). USNM64682 (sl, d), 252203 (sn, sl, d), 364560 (sn, sl, d), 132833 (sn, sl, d), 136400 (sn, sl, d), 261694 (sn, sl, d), 261695 (sn, sl, d), 329704 (sn, sl, d).

*Glaucomys oregonensis*: USNM141952 (sn, sl, d), 264921 (sn, sl, d), 561687 (sn, sl, d).

*Hylopetes alboniger*: AMNH114884 (sl, d), 114885 (sl, d), 114886 (sl, d). FMNH76379 (sn, sl, d), 76381 (sn, sl, d), 76382 (sn, sl, d), 40996 (sn, sl, d), 37889 (sn, sl, d), 76374 (sn, sl, d), 76377 (sn, sl, d), 76378 (sn, sl, d), 76384 (sn, sl, d), 105547 (sn, sl, d), 105548 (sn, sl, d), 76383 (sn, sl, d), 82841 (sn, sl, d), 32313 (sn, sl, d), 82842 (sn, sl, d). USNM37875 (sl, d), 20887 (sn, sl, d), 253608 (sn, sl, d), 253609 (sn, sl, d). KIZ034950 (sn, sl, d), 034970 (sn, sl, d), 031849 (sn, sl, d), 033840 (sn, sl, d), 034945 (sn, sl, d), 019675 (sn, sl, d), 034946 (sn, sl, d), 0305016 (sn, sl, d), 034949 (sn, sl, d),

034942 (sn, sl, d), 032543 (p).

*Hylopetes bartelsi*: USNM501690 (sl, d).

*Hylopetes nigripes*: AMNH203309 (sl, d). USNM477994 (sn, sl, d), 477995 (sn, sl, d), 477998 (sn, sl, d), 478009 (sn, sl, d). FMNH63031 (sn).

*Hylopetes phayrei*: AMNH163556 (sl, d), 163558 (sl, d), 163561 (sl, d), 163562 (sl, d), 167891 (sl, d), 167892 (sl, d), 167893 (sl, d), 58161 (sl, d), 58171 (sl, d), 58179 (sl, d). FMNH25588 (sn, sl, d), 25589 (sn, sl, d), 82836 (sn, sl, d), 82837 (sn, sl, d), 1074 (sn). USNM235580 (sn, sl, d), 259761 (sn, sl, d), 260623 (sn, sl, d), 260624 (sn, sl, d), 294889 (sn, sl, d), 294894 (sn, sl, d), 297086 (sl, d), 297089 (sn, sl, d), 355126 (sn, sl, d), 584420 (sn, sl, d).

*Hylopetes platyurus*: AMNH101834 (sl, d), 101837 (sl, d), 101839 (sl, d), 101431 (sl, d), 101711 (sl, d), 106641 (sl, d), 106737 (sl, d), 101440 (sl, d), 101442 (sl, d), USNM488618 (sn, sl, d), 488619 (sn, sl, d), 488626 (sn, sl, d), 488627 (sn, sl, d).

*Hylopetes sagitta*: FMNH82834 (sn, sl, d), 88340 (sn, sl, d). USNM301023 (sn, sl, d).

*Hylopetes sipora*: BMNH47.1475 (sn).

*Hylopetes spadiceus*: AMNH54822 (sn, sl, d), FMNH37886 (sn, sl, d), 37888 (sn, sl, d), 46646 (sn, sl, d), 46644 (sn). USNM104627 (sn, sl, d), 123931 (sn, sl, d), 123933 (sn, sl, d), 292651 (sn, sl, d), 292652 (sn, sl, d), 308155 (sn, sl, d), 357017 (sn, sl, d), 481109 (sn, sl, d), 481115 (sn, sl, d), 535203 (sn, sl, d), 122883 (sn, sl, d). BMNH94.9.28.42 (sn, sl, d).

*Iomys horsfieldii*: AMNH185169 (sl, d, p), 185170 (sl, d). FMNH10889 (sn, sl, d), 108900 (sn, sl, d). USNM151792 (sn, sl, d), 292653 (sn, sl, d), 292654 (sl, d), 317240 (sn, sl, d), 153684 (sn, sl, d), 301024 (sn). BMNH55.12.24.102 (sn, sl, d), 47.1474 (sn, sl, d).

*Iomys sipora*: AMNH103151 (sn, sl, d), 103313 (sn, sl, d). USNM252321 (sn, sl, d). BMNH47.1473 (sn).

*Petaurillus hosei*: BMNH99.12.9.85 (sn, sl, d).

*Petaurillus kinlochii*: USNM488708 (sn, sl, d), 488709 (sn, sl, d), 488710 (sn, sl, d), 488711 (sn, sl, d).

*Petaurista albiventer*: AMNH83441 (sn, sl, d), 163570 (sn, sl, d), 163573 (sn, sl, d), 32643 (sn, sl, d), 55837 (sn, sl, d). USNM257713 (sn, sl, d), 297078 (sn, sl, d), 297080 (sn, sl, d), 297082 (sn, sl, d), 174079 (sn, sl, d), 353202 (sn, sl, d), 410952 (sn, sl, d), 35495 (sn, sl, d), 37877 (sl, d). BMNH114c (sn, sl, d), 5.11.19.3 (sn, sl, d), 85.8.1.121 (sn, sl, d). MNHNZM-MO-2000-608 (sn, sl, d).

*Petaurista alborufus*: MNHNZN-MO-1870-42 (sn, sl, d). BMNH10.10.19.4 (sn, sl, d), 14.12.1.5 (sn), 95.7.4.1 (sn, sl, d), 23.4.1.93 (sn).

*Petaurista caniceps*: AMNH163577 (sl, d). BMNH22.9.1.44 (sn, sl, d), 1937.6.14.1 (sn, sl, d), 558a (sn, sl, d), 79.11.21.531 (sn, sl, d), 47.1472 (sn, sl, d). KIZ034932 (sn, sl, d), 034962 (sn, sl, d), 034931 (sn, sl, d), 034933 (sn, sl, d), 034963 (sn, sl, d).

*Petaurista elegans*: AMNH101780 (sl, d), 101781 (sn, sl, d), 102468 (sl, d), 163578 (sl, d), 57372 (sl, d). USNM84422 (sn, sl, d), 292647 (sn, sl, d), 292648 (sn, sl, d), 300107 (sn, sl, d), 307574 (sn, sl, d), 240857 (sn). BMNH12.7.25.33 (sn, sl, d), 46.3.4.8 (sn, sl, d), 16.3.26.14 (sn, sl, d), 47.1471 (sn). KIZ034937 (sn, sl, d), 034938 (sn, sl, d), 034972 (sn, sl, d), 034973 (sn, sl, d), 034939 (sn, sl, d).

*Petaurista hainana*: AMNH58201 (sl, d), 58209 (sl, d), 58210 (sl, d), 58211 (sl, d), 58212 (sn, sl, d),

58213 (sn, sl, d), 58214 (sl, d), 58200 (sl, d).

*Petaurista lena*: AMNH183145 (sl, d), 184534 (sl, d), 184916 (sl, d), 184918 (sl, d). BMNH8.4.1.39 (sn, sl, d).

*Petaurista leucogenys*: USNM1470879 (sl, d), 149880 (sl, d), 140881 (sl, d). BMNH5.3.3.17 (sn, sl, d), 5.1.4.50 (sn, sl, d), 3.5.18.1 (sn, sl, d), 80.3.30.3 (sl, d), 23.3.8.2 (sn).

*Petaurista magnificus*: USNM290079 (sl, d), BMNH43.1.12.47 (sn, sl, d) .

*Petaurista nobilis*: FMNH105543 (sn, sl, d), 105544 (sn, sl, d), 105545 (sn, sl, d), 105546 (sn, sl, d), 114365 (sn, sl, d), 114643 (sn, sl, d), 112560 (sn, sl, d), 114364 (sn, sl, d), 114366 (sn, sl, d), 114367 (sn, sl, d). BMNH79.11.21.529 (sn, sl, d).

*Petaurista petaurista petaurista*: USNM121499 (sn, sl, d), 121500 (sn, sl, d), 156386 (sn, sl, d).

*Petaurista petaurista batuana*: USNM121742 (sn, sl, d).

*Petaurista petaurista cicur*: BMNH49.431 (sn, sl, d).

*Petaurista petaurista grandis*: AMNH184922 (sl, d), 184924 (sl, d), 184927 (sl, d), 184929 (sl, d), BMNH62.12.24.10 (sl, d), 62.12.24.9 (sn, sl, d).

*Petaurista petaurista marchio*: USNM143341 (sn, sl, d). BMNH0.8.2.24 (sn, sl, d).

*Petaurista petaurista melanotus*: BMNH116a (sn, sl, d).

*Petaurista petaurista nigrescens*: USNM19177 (sn, sl, d).

*Petaurista petaurista nigricaudatus*: BMNH49.432 (sn, sl, d).

*Petaurista petaurista nitidula*: BMNH94.9.28.10 (sn, sl, d).

*Petaurista petaurista penangensis*: BMNH49.433 (sn, sl, d).

*Petaurista petaurista rajah*: BMNH99.12.9.31 (sn, sl, d).

*Petaurista petaurista stellaris*: BMNH47.1470 (sn, sl, d).

*Petaurista petaurista terutaus*: USNM123934 (sn, sl, d).

*Petaurista philippensis annamensis*: BMNH79b (sn).

*Petaurista philippensis cineraceus*: MNHN1960-3676 (sn), 1962-2174 (sn).

*Petaurista philippensis lylei*: MNHNZM-MO-1948-373 (sn). KIZ019674 (sn, sl, d).

*Petaurista philippensis miloni*: MNHN1982-843 (sn).

*Petaurista philippensis*: AMNH87417 (sl, d), 87418 (sl, d), 55836 (sl, d). FMNH82830 (sl, d), 82831 (sl, d), 96323 (sl, d), 98266 (sl, d), 99406 (sl, d), 99407 (sl, d), 82828 (sl, d), 82829 (sl, d). BMNH6.11.6.14 (sn, sl, d), 25.1.1.42 (sn, sl, d), 98.10.5.41 (sn, sl, d), 14.4.3.1 (sn, sl, d), 22.8.21.1 (sn, sl, d), 23.1.6.53 (sn, sl, d), 23.1.6.50 (sn, sl, d), 115b (sn, sl, d), 115d (sn, sl, d), 96.11.7.5 (sn, sl, d), 198a (sn, sl, d), 96.3.27.1 (sn, sl, d).

*Petaurista xanthotis*: AMNH85063 (sl, d). FMNH34208 (sl, d). USNM144021 (sn, sl, d), 255136 (sn, sl, d). BMNH23.4.1.29 (sl, d), 27.6.24.1 (sn), 23.4.1.27 (sn).

*Petaurista yunanensis*: MNHN1896-2069 (sn). KIZ034928 (sn, sl, d), 034930 (sn, sl, d), 034927 (sn, sl, d), 034966 (sn, sl, d), 034964 (sn, sl, d), 034967 (sn, sl, d), 034926 (sn, sl, d), 034925 (sn, sl, d), 034923 (sn, sl, d), 034929 (sn, sl, d), 031250 (p).

*Petinomys crinitus*: FMNH87439 (sn, sl, d), 92787 (sn, sl, d). USNM239217 (sn, sl, d).

*Petinomys fuscocapillus*: AMNH150063 (sn, sl, d), 240844 (sn, sl, d), 240845 (sl, d). BMNH52.5.9.19 (sn).

*Petinomys genibarbis*: USNM3930 (sn, sl, d), 488671 (sn, sl, d), 488672 (sn, sl, d). BMNH60.5.4.83 (sn), 99.12.9.35 (sn).

*Petinomys hageni*: FMNH47126 (sn, sl, d). USNM143344 (sn, sl, d), 143345 (sn, sl, d).

*Petinomys lugens*: AMNH103149 (sl, d), 103318 (sn, sl, d). USNM252319 (sn, sl, d), 252320 (sn, sl, d), 121531 (sn, sl, d), BMNH95.1.9.7 (sn, sl, d).

*Petinomys mindanensis*: AMNH207540 (sl, d), 207541 (sn, sl, d). USNM254651 (sn, sl, d), 267975 (sn, sl, d).

*Petinomys setosus*: AMNH113031 (sl, d). FMNH89461 (sn, sl, d). USNM301022 (sn, sl, d), 481132 (sn, sl, d), 481140 (sn, sl, d), 481141 (sn, sl, d), 535204 (sn, sl, d).

*Petinomys vordermanni*: USNM124986 (sn, sl, d), 481153 (sn, sl, d), 481168 (sn, sl, d), 481173 (sn, sl, d). BMNH14.12.8.243 (sn).

*Priapomys leonardi*: KIZ034951 (sn, sl, d), 034952 (sn, sl, d), 034953 (sn, sl, d), 034954 (sn, sl, d), 034971 (sn, sl, d), 0410099 (sn, sl, d), 035090 (p). BMNH20.8.8.2 (sn, sl, d). DUE110001 (sn), E110002 (sn), E110003 (sn), E110005 (sn), E110006 (sn), E110007 (sn).

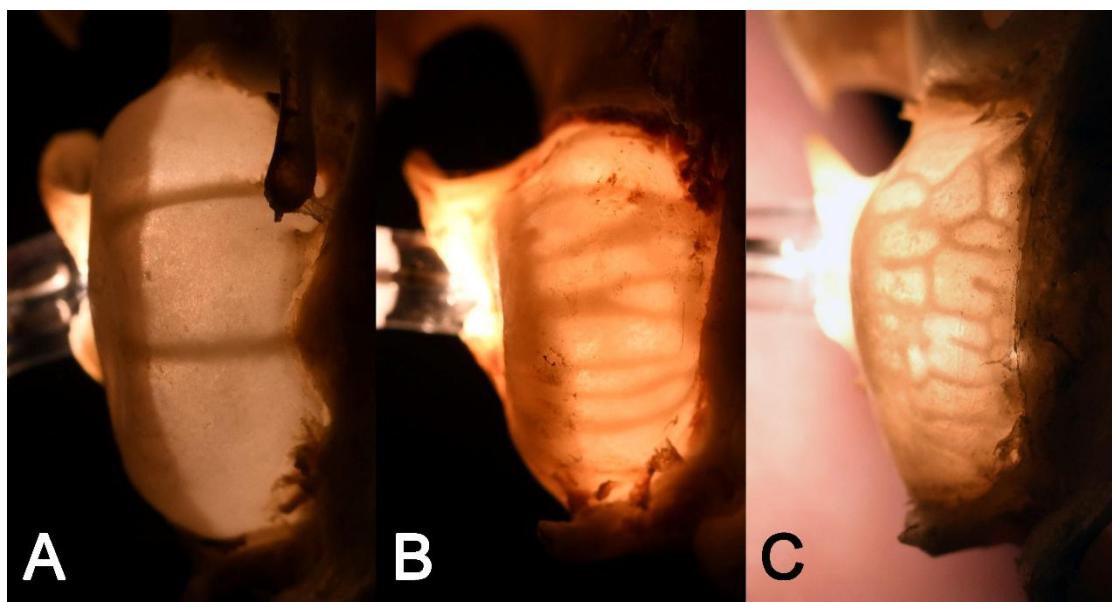
*Pteromys momonga*: USNM38075 (sn, sl, d). BMNH6.1.4.122 (sn), 6.1.4.125 (sl, d).

*Pteromys volans*: AMNH19526 (sl, d), 19534 (sl, d), 85466 (sl, d), 85487 (sl, d), 178821 (sl, d). FMNH92938 (sn, sl, d). USNM172624 (sn, sl, d), 172625 (sn, sl, d), 172626 (sn, sl, d), 270545 (sn, sl, d), 237587 (sl, d), 237588 (sl, d), 254934 (sn, sl, d), 200613 (sn, sl, d). BMNH1937.6.12.1 (sn), 20.7.4.21 (sl, d). KIZ022998 (p).

*Pteromyscus pulverulentus*: AMNH217576 (p). USNM481184 (sn, sl, d), 481186 (sn, sl, d), 489512 (sn, sl, d), 489514 (sn, sl, d). BMNH28.7.14.2 (sl, d), 91.8.28.10 (sn, sl, d), 71.1534 (sn, sl, d), 73.7.3.2 (sn).

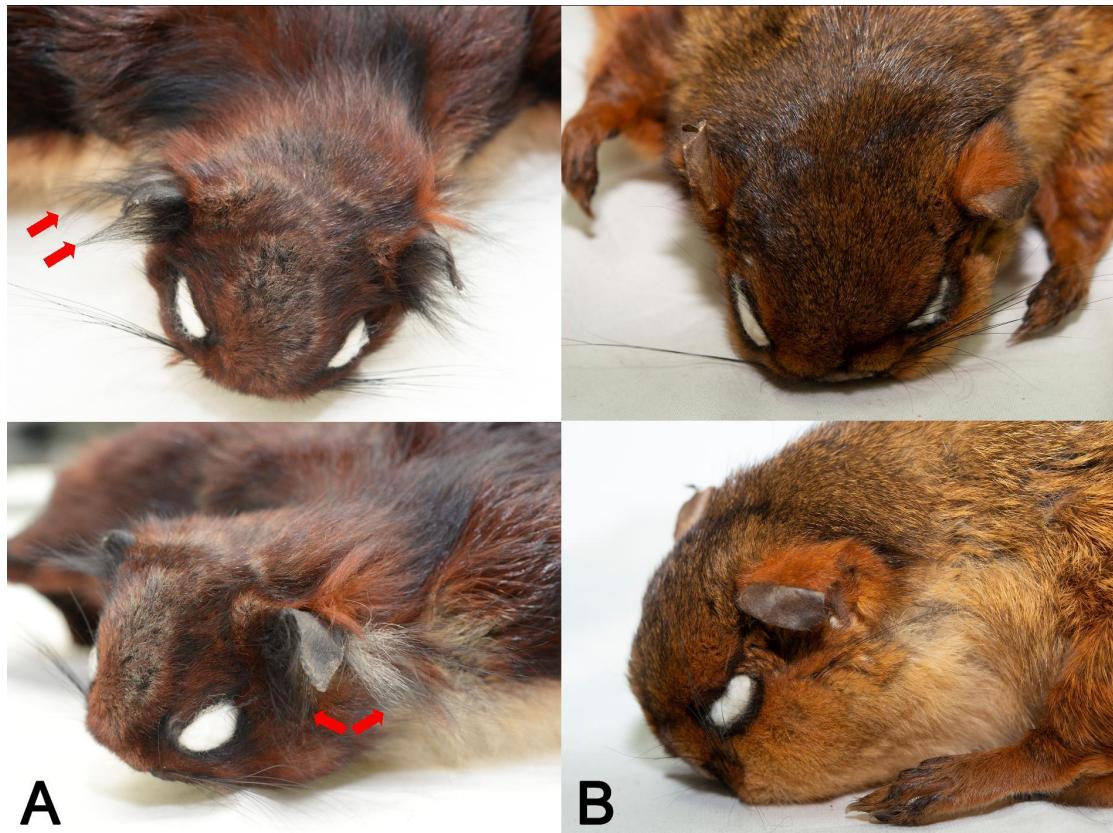
*Trogopterus xanthipes*: FMNH39834 (sn, sl, d), 39835 (sn, sl, d). USNM241271 (sn, sl, d), 254807 (sn, sl, d), 258520 (sn, sl, d), 258980 (sn, sl, d), 268872 (sn, sl, d). BMNH48.309 (sn, sl, d), 23.4.1.32 (sl, d), 14.6.24.1 (sl, d), 9.7.21.4 (sl, d), 95.7.5.1 (sl, d). MNHNZM-MO-1867-555 (sn, sl, d), ZM-MO-1867-556 (sn, sl, d). KIZ034969 (sn, sl, d), 034956 (sn, sl, d), 034957 (sn, sl, d, p), 034958 (sn, sl, d), 034959 (sn, sl, d), 022612 (sn, sl, d), 034968 (sn, sl, d), 028831 (sn, sl, d), 034960 (sn, sl, d).

**Supplementary Figure S1**



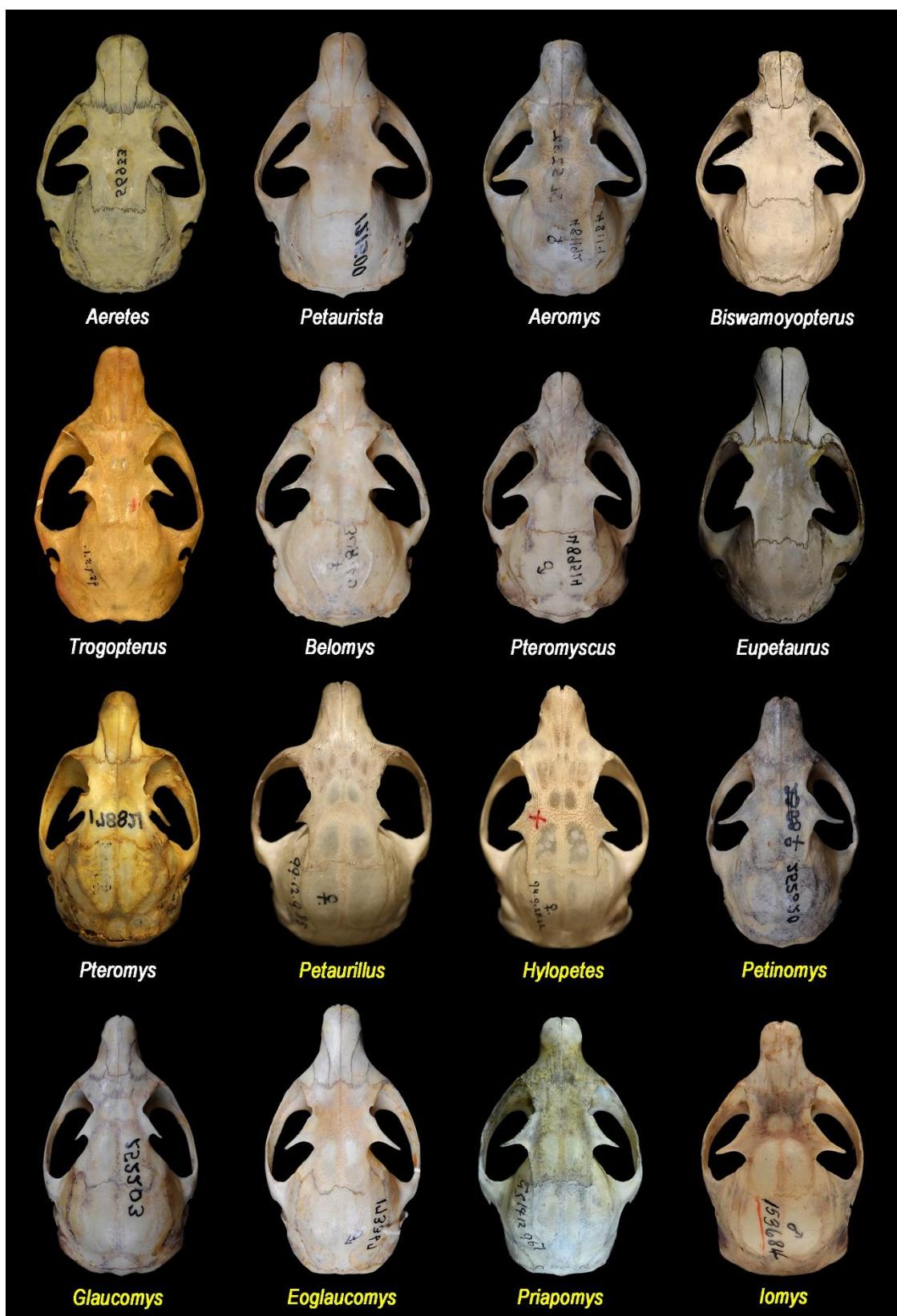
Bulla septae of A) Common form,  $\text{septae} < 4$  (*Aeretes*, *Aeromys*, *Eoglaucomys*, *Eupetaurus*, *Glaucomys*, *Hylopetes*, *Iomys*, *Petaurillus*, *Petaurista*, *Pteromys*, *Pteromys (Hylopetes) leonardi* (=*Priapomys* gen. nov.)); B) honeycomb form,  $\text{septae} > 4$ , not crossing each other (*Belomys*, *Biswamoyopterus*, *Trogopterus*, *Pteromyscus*, *Petynomys setosus*); and C) cobweb form,  $\text{septae} > 4$ , crossing each other (*Petynomys* excluding *Petynomys setosus*).

**Supplementary Figure S2**



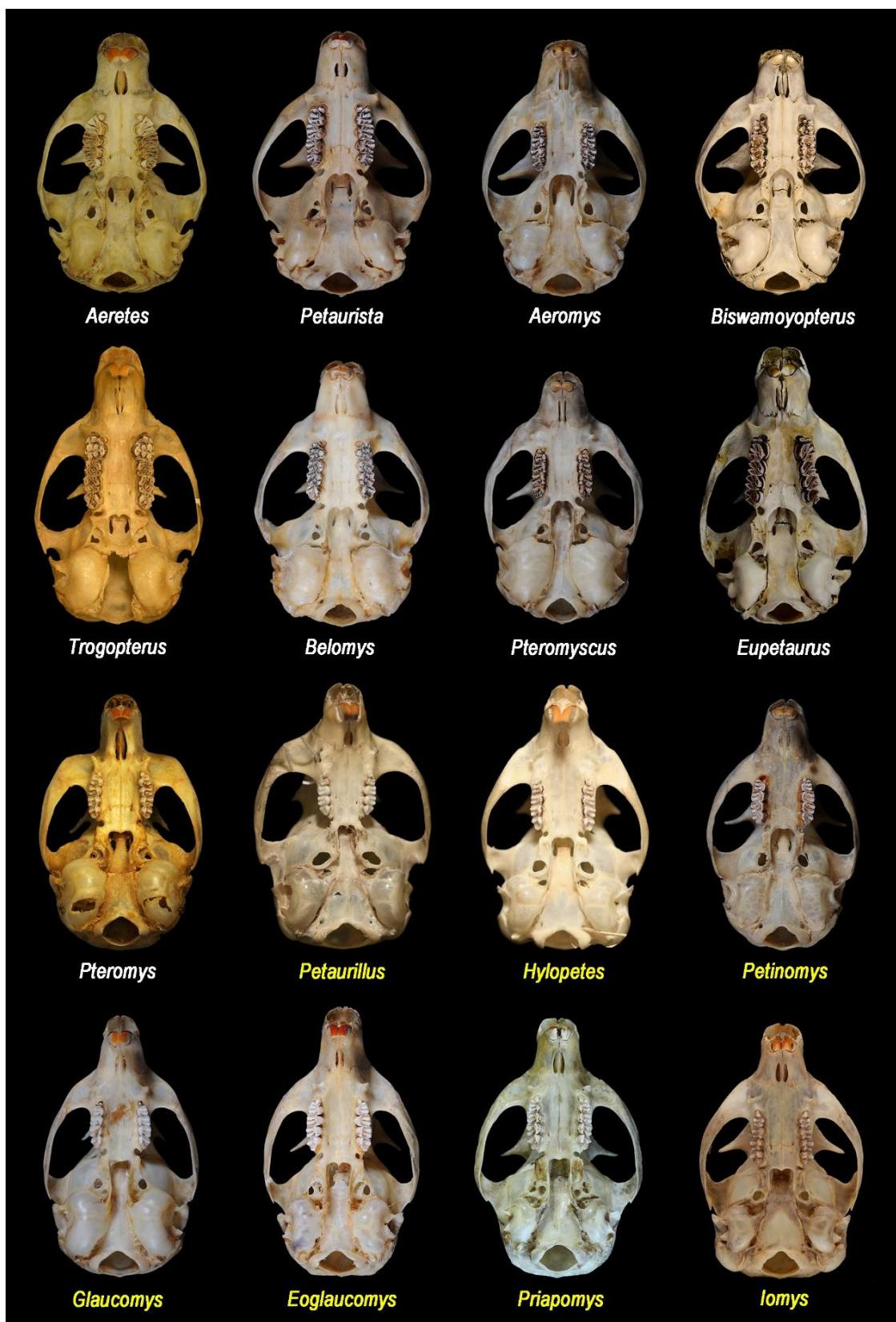
Ear tufts, A. present (red arrow), B. absent.

Supplementary Figure S3A



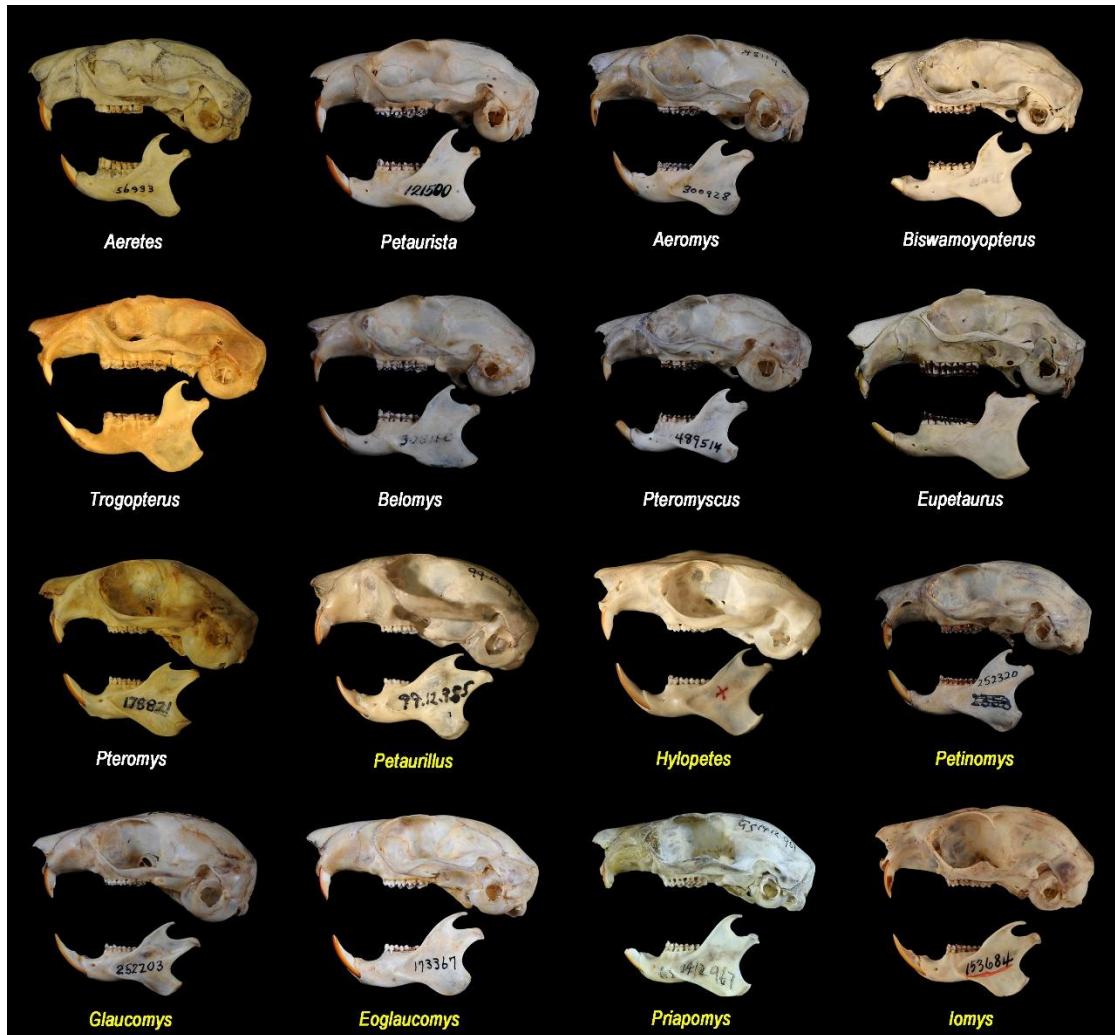
Dorsal view of skulls of sixteen extant flying squirrel genera, scaled to similar length. Refer to the legend of Supplementary Figure S3C for the detailed information of the specimens.

Supplementary Figure S3B



Bottom view of skulls of sixteen extant flying squirrel genera, scaled to similar length. Refer to the legend of Supplementary Figure S3C for the detailed information of the specimens.

Supplementary Figure S3C

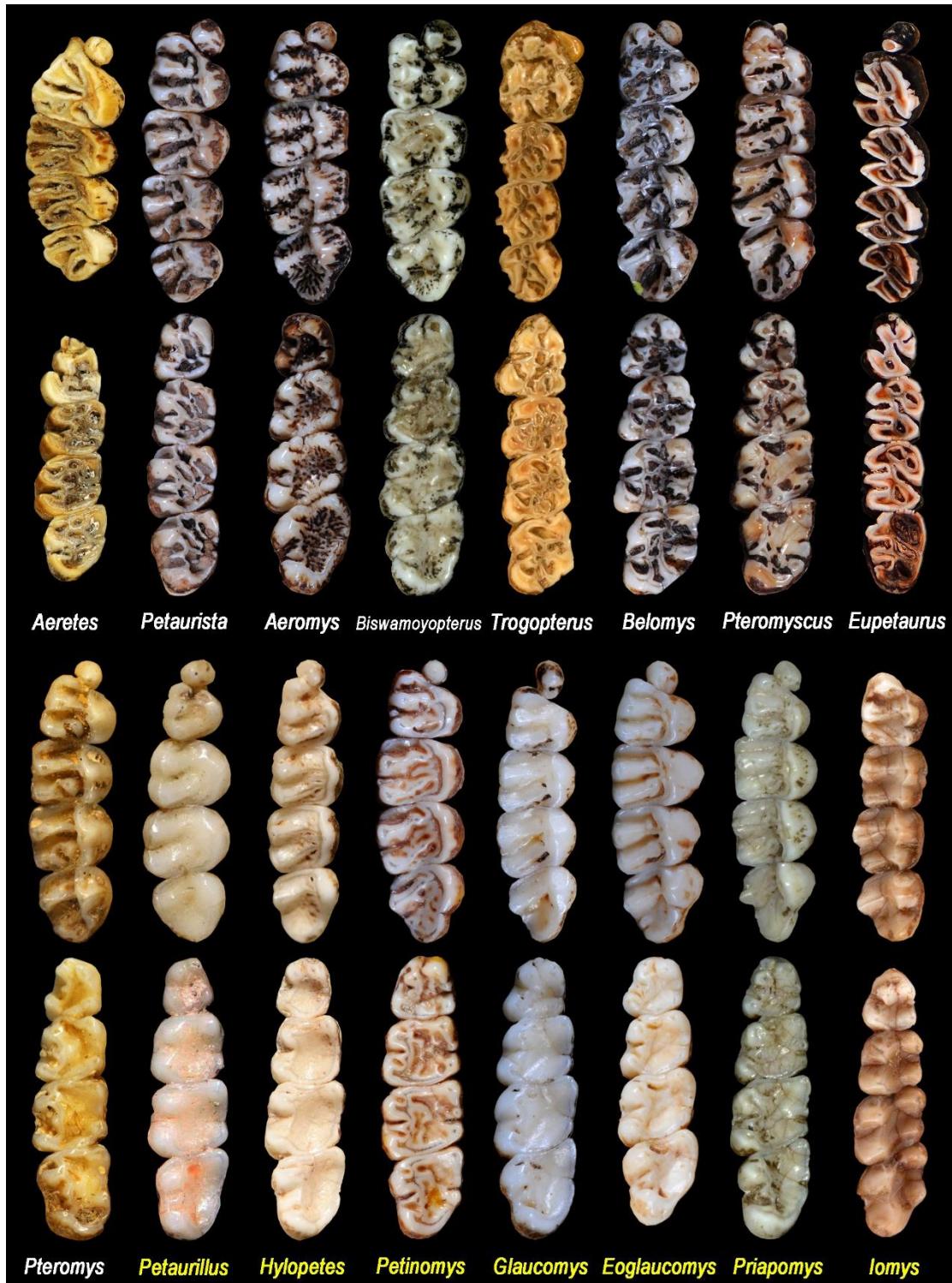


Lateral view of skulls and mandibles of sixteen extant flying squirrel genera, scaled to similar length.

**Detailed information of the specimens in Supplementary Figure S3ABCD:**

*Aeretes*: *Aeretes melanopterus* (AMNH56933), *Petaurista*: *Petaurista petaurista* (USNM121500), *Aeromys*: *Aeromys tephromelas* (USNM481192), *Biswamoyopterus*: *Biswamoyopterus gaoligongensis* (KIZ034924), *Tropopterus*: *Tropopterus xanthipes* (BMNH95.7.5.1), *Belomys*: *Belomys pearsonii* (USNM308160), *Pteromyscus*: *Pteromyscus pulverulentus* (USNM489514), *Eupetaurus*: *Eupetaurus nivamons* (KIZ034190), *Pteromys*: *Pteromys volans* (AMNH178821), *Petaurillus*: *Petaurillus hosei* (BMNH99.12.9.85), *Hylopetes*: *Hylopetes spadiceus* (BMNH94.9.28.42), *Petinomys*: *Petinomys lugens* (USNM252320), *Glaucomys*: *Glaucomys volans* (USNM252203), *Eoglaucous*: *Eoglaucous fimbriatus* (USNM173367), *Priapomys*: *Pteromys (Hylopetes) leonardi* (=*Priapomys leonardi*) (KIZ034951), and *Iomys*: *Iomys horsfieldii* (USNM153684). White letters indicate members of Pteromyina, yellow indicate Glaucomyina.

Supplementary Figure S3D



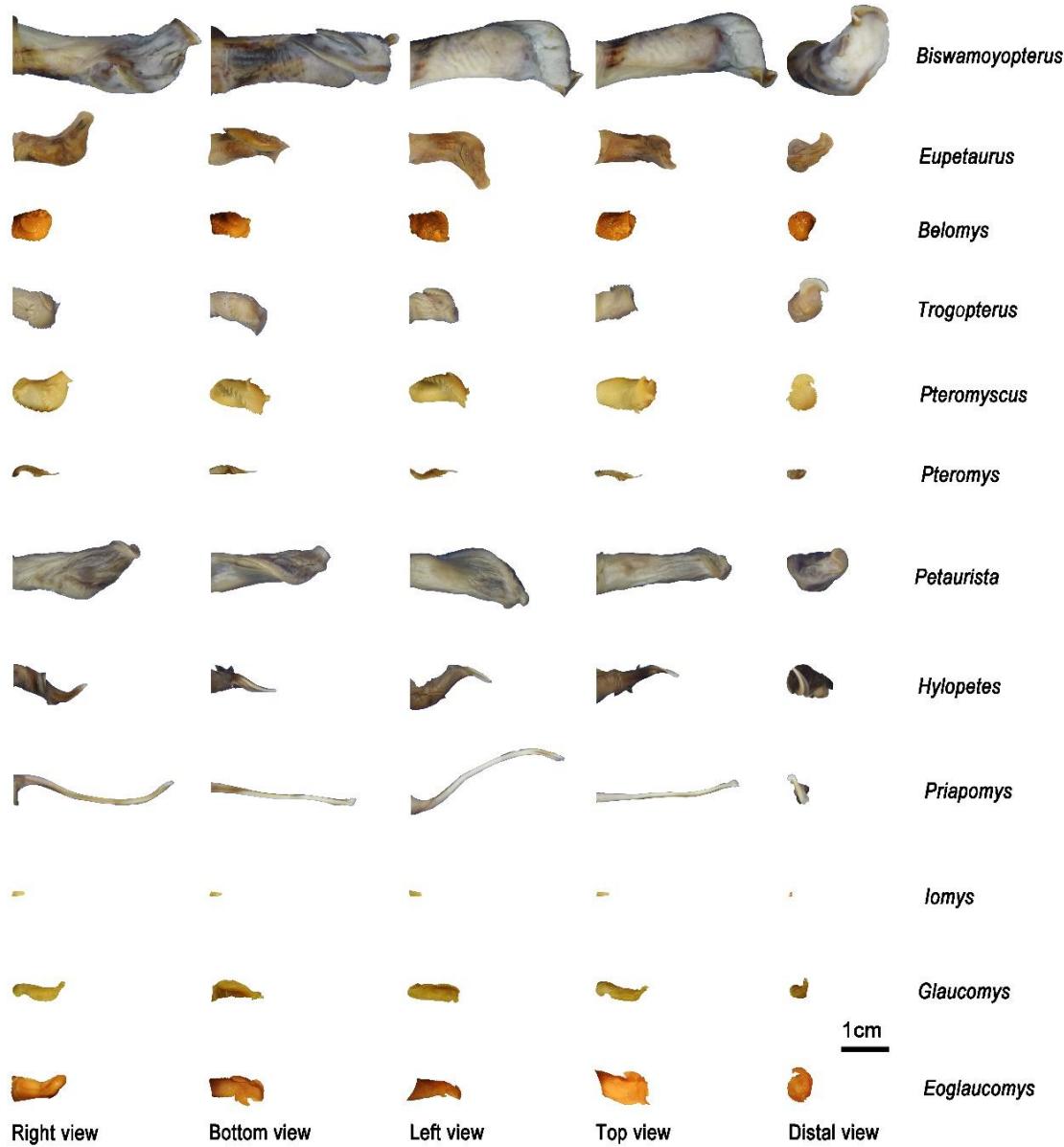
Upper cheek teeth (1<sup>st</sup> & 3<sup>rd</sup> row) and lower cheek teeth (2<sup>nd</sup> & 4<sup>th</sup> row) of sixteen extant flying squirrel genera, scaled to similar length. Refer to the legend of Supplementary Figure S3C for the detailed information of the specimens. For all teeth, the left side is buccal side.

**Supplementary Figure S4**



Buccal view of the upper cheekteeth of *Pteromys (Hylopetes) leonardi* (=*Priapomys leonardi*) (left, three cusps, BMNH20.8.8.2), *Hylopetes alboniger* (middle, four cusps, FMNH76377), and *Petinomys lugens* (right, four cusps, USNM252319).

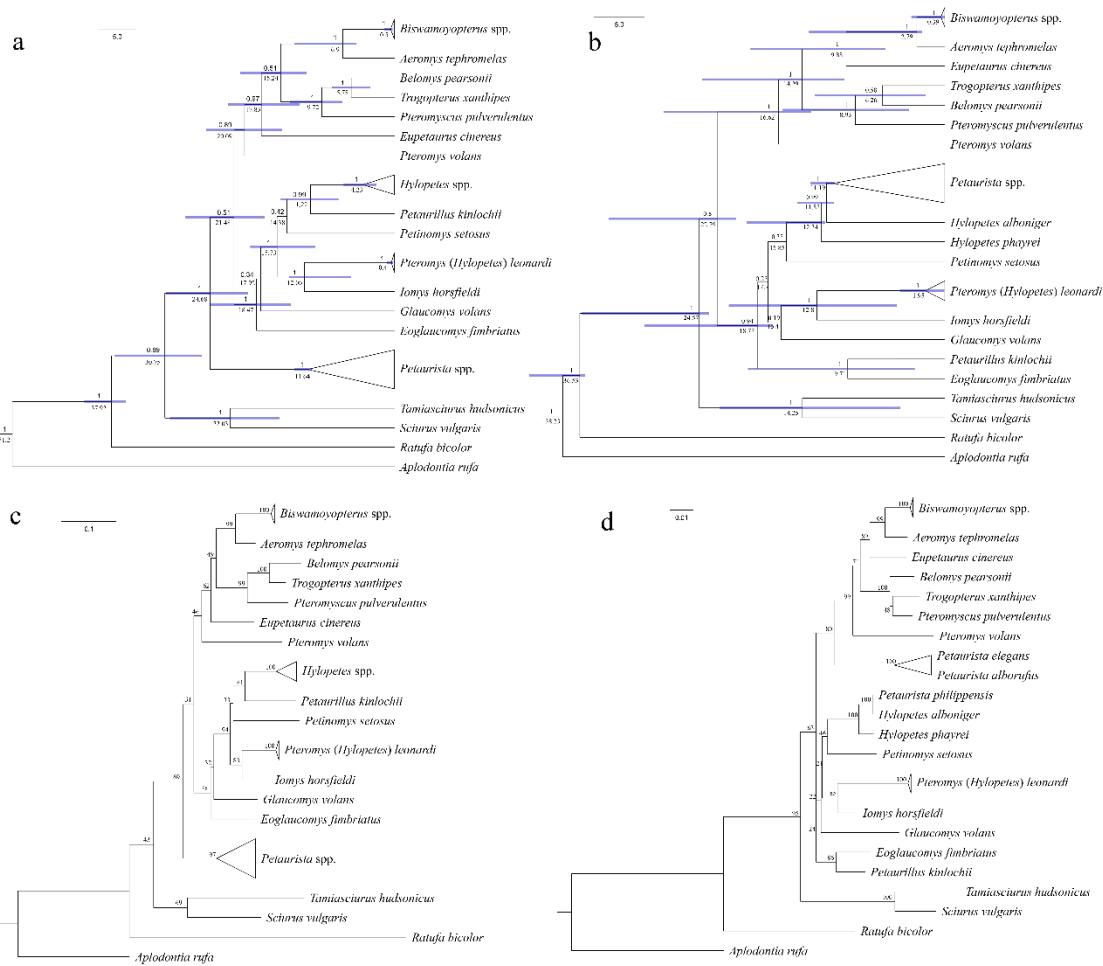
Supplementary Figure S5



Glans penis of some extant flying squirrel genera:

*Biswamoyopterus*: *Biswamoyopterus gaoligongensis* (KIZ034924, Body mass=1 370 g),  
*Eupetaurus*: *Eupetaurus nivamons* (KIZ035087, Body mass=1 555 g), *Belomys*: *Belomys pearsonii* (USNM258345, Body mass=156 g\*), *Trogopterus*: *Trogopterus xanthipes* (KIZ034957, Body mass=600 g), *Pteromyscus*: *Pteromyscus pulverulentus* (AMNH217576, Body mass=122g), *Pteromys*: *Pteromys volans* (KIZ022998, Body mass=138 g\*), *Petaurista*: *Petaurista yunanensis* (KIZ031250, Body mass=2 120 g), *Hylopetes*: *Hylopetes alboniger* (KIZ032543, Body mass=251 g), *Priapomys*: *Pteromys (Hylopetes) leonardi* (=*Priapomys leonardi*) (KIZ035090, Body mass=209 g), *Iomys*: *Iomys horsfieldii* (broken, AMNH185169, Body mass=165 g\*), *Glaucomys*: *Glaucomys sabrinus* (AMNH95193, Body mass=142 g\*), and *Eoglaucous*: *Eoglaucous fimbriatus* (USNM429353, Body mass=734 g\*). The asterisk indicates that the body mass data comes from the summary of Thorington *et al.* (2012) rather than from the specimen itself. Scale bar shown.

**Supplementary Figure S6**



Flying squirrel phylogeny (includes all genera except *Aeretes*) and node dating estimates utilizing different datasets: a. (12S+16S) Bayesian tree, b. IRBP Bayesian tree, c. (12S+16S) Maximum likelihood tree, d. IRBP Maximum likelihood tree.

**Supplementary Table S1**

Terms of dental morphology used in this paper with notes on the equivalents by different authors

position	Terms used in this paper		Equivalents
	Terms	Abbrev	
Upper cheekteeth	paracone	pa	
	metacone	me	
	protocone	pr	
	hypocone	hy	
	parastyle	past	
	mesostyle	msst	
	paraconule	pacu	protoconule (Qiu <i>et al</i> 2019, Tong 2007)
	metaconule	mecu	
	anteroloph	anl	anterior cingulum (James, 1963)
	paraloph	pal	protoloph (Qiu <i>et al</i> 2019, McKenna 1962), anterior transverse ridge (Allen, 1940) , postparaconule crista + preprotocrista (Zhou <i>et al</i> 1975)
	metaloph	mel	posterior transverse ridge (Allen, 1940), premetaconule crista + postprotocrista (Zhou <i>et al</i> 1975)
	posteroloph	pol	posterior cingulum (James, 1963)
	endoloph	enl	
	anterolophule	anlu	
	mesolophule	mslu	protolophule (Qiu <i>et al</i> 2019)
	preparaconule crista	prepalc	
	postmetaconule crista	pomelc	
	anterior valley	av	anterofossette (Tong 2007), paraflexus+ anterofossette (McKenna 1962), paraflexus (Kawamura 1988)
	central valley	cv	medifossette (Tong, 2007), posterobuccal flexus (McKenna 1962, Kawamura 1988), trigon basin (Zhou <i>et al</i> 1975)
	posterior valley	pv	posterofossette (Tong, 2007, McKenna 1962, Kawamura 1988)
	anterior flexus	af	
	central flexus	cf	
	posterior flexus	pf	posterolingual (diagonal) flexus (Tong, 2007, McKenna 1962, Kawamura 1988)
Lower cheekteeth	protoconid	prd	
	hypoconid	hyd	
	metaconid	med	parametaconid (James, 1963)

	entoconid	end	
	mesoconid	msd	
	paraconid	pad	anterobuccal cingulum (Qiu <i>et al</i> 2019), anteroconulid (Tong, 2007)
	ectostyloid	ecstd	
	metastyloid	mestd	mesostyloid (Kawamura 1988)
	hypoconulid	hycud	metaconulid (Tong, 2007)
	anterolophid	anld	protolophid (James, 1963)
	protolophid	prld	metalophulid (Qiu <i>et al</i> 2019)
	entolophid	enld	
	ectolophid	ecld	
	posterolophid	pold	
	anterobuccal sinusid	asd	
	hypoflexid	hyfd	buccal valley (Qiu <i>et al</i> 2019)
	anterior fossettid	afd	trigonid basin (Qiu <i>et al</i> 2019) trigonid or anterior valley (James, 1963)
	central flexid	cfid	talond basin (Qiu <i>et al</i> 2019) “lingual-flexid+medifossettid” (Tong, 2007), lingual flexid+metafossettid (Kawamura 1988), lingual diagonal flexid (McKenna 1962)
	posterior fossettid	pfd	talond basin (Qiu <i>et al</i> 2019) talond basin remnant (McKenna 1962, Kawamura 1988)

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**Supplementary Table S2**

GenBank accession numbers for 12S ribosomal RNA (12S), 16S ribosomal RNA (16S), and interphotoreceptor retinoid-binding protein (IRBP) sequences utilized in this study.

Species	12S rRNA gene	16S rRNA gene	IRBP
<i>Aeromys tephromelas</i>	AY227536.1	AY227482.1	AY227594.1
<i>Belomys pearsonii</i>	AY227537.1	AY227483.1	AY227595.1
<i>Biswamoyopterus biswasi</i>	MK105526.1 MK105527.1	MK105519.1 MK105520.1	MK105534.1 MK105535.1
<i>Eoglaucomys fimbriatus</i>	AY227562.1	AY227485.1	AY227597.1
<i>Eupetaurus cinereus</i>	AY227538.1	AY227484.1	AY227596.1
<i>Glaucomys volans</i>	AY227559.1	AY227486.1	AY227598.1
<i>Hylopetes phayrei</i>	AY227539.1	AY227487.1	AY227599.1
<i>Hylopetes alboniger</i>	MW929356	MW929360	MW929363
<i>Iomys horsfieldii</i>	AY227540.1	AY227488.1	AY227600.1
<i>Petaurillus kinlochii</i>	AY227542.1	AY227490.1	AY227602.1
<i>Petaurista albiorufus lena</i>	AY227541.1	AY227489.1	AY227601.1
<i>Petaurista elegans</i>	MW929357	MK105524.1	MK105539.1
<i>Petaurista philippensis</i>	MK105528.1	MK105521.1	MK105536.1
<i>Petinomys setosus</i>	AY227544.1	AY227492.1	AY227604.1
<i>Pteromys volans</i>	AY227545.1	AY227493.1	AY227605.1
<i>Pteromyscus pulverulentus</i>	AY227543.1	AY227491.1	AY227603.1
<i>Trogopterus xanthipes</i>	AY227546.1	AY227494.1	AY227606.1
<i>Priapomys leonardi</i>	MW929354 MW929355	MW929358 MW929359	MW929361 MW929362
<i>Aplodontia rufa</i>	AY227508.1	AY227451.1	AY227564.1
<i>Ratufa bicolor</i>	AY227548.1	AY227496.1	AY227608.1
<i>Tamiasciurus hudsonicus</i>	AY227555.1	AY227504.1	AY227622.1
<i>Sciurus vulgaris</i>	AY227553.1	AY227501.1	AY227620.1

**Supplementary Table S3**

Primers and PCR conditions for amplification and sequencing used in the genetic analyses.

Gene	Primer Sequence		Annealing temp	Remarks
	Forward	Reverse		
12S rRNA	rRNA-L613-hk1:  GGCGGGCGAGCAAAGCACTGAAAATG	rRNA-H1478-hk1:  TGATTGGTGGAGGGTGACGAGCGGTGTGT	56	(He et al. 2010)
16S rRNA	16Sar-L: CGCCTGTTATCAAAAAACAT	16Sbr-H: CCGGTCTGAACTCAGATCACGT	52	(He et al. 2010)
IRBP	IRBPA:  ATGGCCAAGGTCCCTTGGATAACTACTGCTT	IRBPB:  CGCAGGTCCATGATGAGGTGCTCCGTGTCCTG	61	(Jansa et al. 2009)

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**Supplementary Table S4**

Character loading and percentage of variance explained on the components of the principal component analysis, morphological measurements from *Pteromys (Hylopetes) leonardi* (=*Priapomys leonardi*) and related species ("similar sized + sympatric/parapatric" or "genetic similarity").

Variables	PC 1	PC 2	PC 3	PC 4	PC 5	PC 6	PC 7	PC 8	PC 9	PC 10	PC 11	PC 12	PC 13	PC 14	PC 15	PC 16	PC 17	PC 18	PC 19	PC 20
GLS	0.229	-0.011	0.014	-0.004	-0.032	-0.195	0.038	0.053	-0.001	-0.139	-0.084	0.072	0.026	0.082	0.030	0.012	-0.048	0.052	0.855	0.359
CBL	0.250	-0.001	0.019	0.038	-0.012	-0.157	0.034	0.050	-0.171	-0.160	-0.005	0.049	0.029	0.062	0.103	0.340	-0.012	-0.138	0.195	-0.814
LN	0.275	-0.128	-0.053	-0.044	-0.232	-0.105	0.688	0.217	0.447	-0.040	-0.107	0.221	0.008	-0.074	-0.101	-0.118	0.003	-0.013	-0.161	-0.028
ABN	0.234	0.148	0.069	-0.162	-0.108	0.630	0.005	0.310	0.035	-0.004	-0.249	-0.414	0.106	0.305	-0.041	0.047	-0.187	0.091	0.032	-0.028
PBN	0.149	0.073	-0.701	-0.177	0.614	-0.053	0.134	0.076	-0.061	-0.049	0.072	-0.137	0.075	-0.017	-0.021	-0.072	-0.062	0.023	-0.024	0.006
IOB	0.225	0.019	-0.071	-0.149	0.033	0.446	-0.199	-0.391	0.345	-0.443	0.222	0.287	-0.187	-0.101	-0.047	0.119	0.085	-0.114	0.019	0.024
ZB	0.264	0.021	0.075	-0.006	-0.048	0.057	0.059	-0.219	-0.195	0.048	-0.014	0.001	-0.060	-0.145	0.533	-0.485	-0.442	-0.293	-0.043	-0.009
BOC	0.242	0.063	-0.015	-0.097	-0.047	-0.275	-0.128	-0.421	0.057	-0.060	-0.384	-0.175	-0.068	0.411	-0.150	-0.364	0.346	0.068	-0.099	-0.099
BPP	0.227	0.049	-0.069	0.020	-0.128	-0.007	0.102	-0.459	-0.047	0.452	0.243	0.085	0.363	0.204	-0.289	0.268	-0.309	-0.008	-0.025	0.077
MB	0.178	-0.015	0.041	0.248	0.009	0.199	0.220	-0.225	0.009	0.157	0.170	-0.351	0.186	-0.360	0.270	-0.005	0.492	0.305	0.123	-0.048
LIF	0.215	-0.688	0.382	-0.144	0.476	0.033	-0.119	0.073	0.100	0.186	0.004	0.055	0.082	0.102	0.020	-0.002	0.029	-0.008	-0.035	0.006
PL	0.263	-0.060	-0.046	0.090	-0.089	-0.230	0.027	0.041	-0.050	-0.132	0.148	-0.165	-0.340	0.300	0.407	0.451	-0.015	0.141	-0.303	0.313
PPL	0.233	0.093	0.180	-0.005	0.111	0.092	0.171	-0.011	-0.607	-0.262	-0.309	0.270	0.183	-0.192	-0.175	0.139	0.158	-0.022	-0.215	0.238
LAB	0.011	0.483	0.254	0.404	0.486	0.066	0.120	-0.030	0.248	0.238	-0.203	0.200	-0.222	0.129	0.069	0.092	-0.022	-0.050	0.030	-0.010
BIF	0.220	0.125	0.258	0.215	0.075	-0.062	0.058	0.189	-0.164	-0.190	0.604	-0.163	-0.172	0.112	-0.401	-0.339	-0.028	0.020	-0.021	-0.031
LP4M3	0.231	-0.023	-0.146	0.411	-0.080	-0.081	-0.311	0.210	0.191	-0.017	-0.059	-0.154	0.316	-0.081	-0.044	0.045	0.163	-0.607	-0.093	0.154
IBPP4	0.203	0.358	0.296	-0.483	0.065	-0.339	-0.178	0.046	0.223	0.049	-0.001	-0.245	0.047	-0.444	-0.060	0.148	-0.106	0.031	-0.079	0.034
IBPM3	0.158	0.251	-0.054	-0.331	-0.110	0.066	-0.140	0.300	-0.099	0.351	0.258	0.414	0.064	0.207	0.242	-0.128	0.421	-0.056	0.001	0.008
lp4m3	0.225	0.019	-0.102	0.297	-0.049	-0.067	-0.389	0.151	0.119	-0.101	-0.076	0.293	0.254	-0.086	0.060	-0.151	-0.237	0.613	-0.129	-0.084
HM	0.286	-0.151	-0.222	0.082	-0.142	0.096	-0.165	0.067	-0.152	0.413	-0.175	0.014	-0.606	-0.315	-0.283	0.035	0.000	0.022	0.066	-0.035
Eigenvalue	0.028	0.006	0.003	0.002	0.001	0.001	0.001	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
% variance	65.504	13.251	6.311	5.265	3.431	1.359	1.192	0.836	0.757	0.479	0.418	0.358	0.272	0.238	0.118	0.092	0.069	0.025	0.019	0.006

**Supplementary Table S5**

Measurements recorded from *P. (H.) leonardi* (=*Priapomys leonardi*) and related species ("similar sized + sympatric/parapatric" or "genetic similarity"). Abbrev:

B=broken, N=not measured, LP4M3=length from the fourth upper premolar to the third molar, lp4m3=length from the fourth lower premolar to the third molar.

ID	species	GLS	CBL	LN	ABN	PBN	IOB	ZB	BOC	BPP	MB	LIF	PL	PPL	LAB	BIF	LP3M3	LP4M3	IBPP4	IBPM3	BP4	BM1	BM3	lp4m3	HM
FM 108900	<i>lomys horsfieldii</i>	45.23	40.91	14.32	6.67	3.87	8.56	26.82	9.73	15.35	19.25	2.93	24.17	13.51	6.77	10.83	N	8.63	5.36	5.97	N	N	N	8.54	17.28
FM 108898	<i>lomys horsfieldii</i>	44.83	41.17	14.29	7.06	4.16	9.62	26.73	9.64	15.19	19.86	3.08	23.65	14.17	7.04	10.89	N	8.69	5.43	5.94	N	N	N	8.66	17.14
FM 108901	<i>lomys horsfieldii</i>	44.03	40.99	13.63	7.12	4.12	9.29	27.12	9.74	14.87	19.71	2.99	24.00	13.92	6.55	10.29	N	8.93	5.04	5.80	N	N	N	8.68	17.61
FM 108899	<i>lomys horsfieldii</i>	46.33	42.21	15.32	7.03	4.03	9.63	28.68	10.07	15.80	19.85	3.08	24.26	14.67	6.80	10.96	N	8.61	5.64	6.54	N	N	N	8.80	17.73
FM 108897	<i>lomys horsfieldii</i>	45.69	41.87	15.06	8.13	4.20	10.46	28.67	9.98	15.84	20.65	3.34	23.17	15.25	7.13	B	N	9.11	5.65	6.28	N	N	N	9.23	18.77
AMNH 185170	<i>lomys horsfieldii</i>	44.55	41.53	13.84	6.98	3.76	9.03	28.14	9.70	16.80	21.40	2.90	23.97	14.55	7.79	10.60	N	9.23	5.07	5.69	N	N	N	9.02	18.16
AMNH 185169	<i>lomys horsfieldii</i>	42.91	39.46	12.77	6.90	4.28	9.21	26.99	9.37	14.95	20.72	3.11	22.80	13.32	7.33	10.42	N	8.81	5.45	5.85	N	N	N	8.70	16.84
AMNH 103313	<i>lomys sipora</i>	45.36	41.79	13.92	7.10	4.51	9.49	26.66	9.36	14.98	20.22	3.14	24.62	13.59	7.40	11.10	N	10.38	4.97	5.73	N	N	N	10.03	19.71
AMNH 103151	<i>lomys sipora</i>	B	B	13.48	6.32	4.53	9.06	26.64	B	B	B	3.17	24.50	B	7.78	11.08	N	10.28	4.78	5.84	N	N	N	10.12	19.35
AMNH 163556	<i>Hylopetes phayrei</i>	37.68	34.93	9.57	5.74	4.26	8.21	23.16	8.75	13.20	17.25	2.02	19.12	13.09	8.76	9.74	7.85	7.46	5.49	5.71	N	N	N	7.66	14.06
AMNH 163555	<i>Hylopetes phayrei</i>	39.50	35.61	11.01	6.75	3.71	8.11	24.20	8.67	13.88	17.80	2.12	19.73	13.32	8.67	10.17	7.92	7.63	5.53	6.18	N	N	N	7.95	14.10
AMNH 163557	<i>Hylopetes phayrei</i>	40.75	37.54	12.01	6.67	3.89	7.97	24.72	8.55	13.80	18.00	2.36	20.91	13.70	8.93	10.66	8.15	7.98	5.87	6.46	N	N	N	8.18	14.60
AMNH 163554	<i>Hylopetes phayrei</i>	39.76	35.95	11.38	6.66	4.56	8.62	24.06	8.59	13.97	18.34	2.24	20.06	12.45	8.98	10.35	8.29	7.89	5.77	6.02	N	N	N	7.92	14.13
AMNH 163553	<i>Hylopetes phayrei</i>	39.36	36.37	11.02	6.97	3.52	8.20	24.11	8.44	13.50	17.55	2.19	20.19	13.26	8.56	10.03	8.24	7.63	5.25	6.22	N	N	N	7.97	14.49
AMNH 167892	<i>Hylopetes phayrei</i>	41.31	37.85	12.62	6.33	3.93	8.38	24.93	8.81	14.14	18.46	2.49	20.77	B	8.65	9.84	8.80	8.42	5.59	5.86	N	N	N	8.57	15.15
AMNH 167891	<i>Hylopetes phayrei</i>	40.96	37.39	12.54	6.57	4.06	9.25	25.24	9.29	14.48	18.75	3.00	21.08	13.64	8.89	10.31	7.81	7.43	5.71	5.73	N	N	N	7.62	14.98
AMNH 176893	<i>Hylopetes phayrei</i>	41.40	38.16	12.11	7.25	4.41	9.75	25.51	9.18	14.29	18.82	2.64	21.37	13.87	8.82	10.73	8.64	8.04	5.98	6.35	N	N	N	8.32	15.19
FM 82836	<i>Hylopetes phayrei</i>	38.56	33.74	11.80	6.69	3.38	8.04	23.06	8.67	13.24	18.12	2.13	18.67	12.06	8.80	9.73	8.37	7.97	5.53	5.71	N	N	N	7.87	13.85
FM 82837	<i>Hylopetes phayrei</i>	40.29	36.51	11.84	6.30	3.35	8.07	23.39	8.78	12.34	17.83	2.69	20.84	12.87	8.58	10.02	8.31	8.02	5.37	5.33	N	N	N	8.23	13.91
FM 105548	<i>Hylopetes albioniger</i>	49.01	45.57	14.94	8.37	4.76	9.39	29.28	10.76	15.74	20.48	2.82	25.55	16.68	8.72	12.33	9.71	9.21	7.16	7.13	N	N	N	9.20	19.10
FM 105547	<i>Hylopetes albioniger</i>	46.77	43.64	13.63	8.67	3.49	10.40	30.17	10.53	16.67	20.83	2.51	24.65	15.43	8.60	12.20	10.13	9.51	7.37	7.61	N	N	N	9.53	18.77
AMNH 114886	<i>Hylopetes albioniger</i>	52.62	49.05	16.06	8.71	5.38	10.64	32.92	11.70	17.98	22.50	3.36	27.83	17.32	9.16	12.98	11.18	10.08	7.29	7.30	N	N	N	10.09	19.83
AMNH 114885	<i>Hylopetes albioniger</i>	51.46	47.91	16.05	8.23	4.74	10.68	33.28	11.30	17.19	23.02	2.84	26.93	17.01	9.11	12.93	10.72	9.96	7.03	6.96	N	N	N	10.09	19.57
AMNH 114884	<i>Hylopetes albioniger</i>	51.09	48.32	16.09	8.97	4.42	11.09	32.60	11.54	18.40	22.32	2.53	27.21	16.79	8.78	13.60	10.85	10.43	6.77	7.26	N	N	N	10.46	19.82
KIZ 034949	<i>Hylopetes albioniger</i>	50.53	46.74	15.43	7.65	3.90	9.79	31.14	11.35	17.08	21.25	2.54	27.20	16.00	8.66	12.97	N	10.11	6.78	6.48	N	N	N	10.23	19.08
KIZ 019675	<i>Hylopetes albioniger</i>	50.30	47.20	15.11	8.23	4.34	10.56	31.19	10.77	17.16	21.51	2.48	26.18	17.27	9.29	13.08	N	9.99	7.31	7.09	N	N	N	10.28	17.57
KIZ 034946	<i>Hylopetes albioniger</i>	50.10	46.75	13.46	8.26	4.51	10.68	32.11	11.55	16.75	21.85	3.14	25.93	17.35	8.64	12.30	N	10.48	6.25	6.49	N	N	N	10.67	19.51
KIZ 034948	<i>Hylopetes albioniger</i>	49.16	45.05	15.01	7.44	4.33	10.24	30.66	11.14	17.33	21.91	2.80	25.33	16.12	9.77	12.23	N	9.89	6.11	6.53	N	N	N	10.12	19.04
KIZ 034942	<i>Hylopetes albioniger</i>	48.82	45.06	13.69	7.94	4.18	10.56	30.13	11.23	16.81	21.09	3.15	26.44	14.97	9.35	12.80	N	10.24	6.55	6.61	N	N	N	10.34	18.49
KIZ 034952	<i>Priapomys leonardi</i>	46.14	42.31	13.82	7.24	3.54	9.15	29.05	10.53	15.61	20.22	3.40	24.25	14.74	7.65	11.43	9.18	8.59	6.71	6.04	2.16	2.51	2.54	8.55	17.34
KIZ 034951	<i>Priapomys leonardi</i>	46.68	43.11	14.01	7.64	4.31	9.33	29.65	10.09	15.12	19.33	3.56	23.91	15.83	7.95	11.44	9.46	8.80	6.08	6.30	2.19	2.61	2.46	8.96	18.14
KIZ 034954	<i>Priapomys leonardi</i>	44.98	40.89	13.20	6.92	3.54	9.39	28.09	9.95	14.93	18.96	3.16	23.35	14.18	7.68	11.46	9.07	8.61	6.39	6.33	2.13	2.53	2.40	8.91	16.32
KIZ 410099	<i>Priapomys leonardi</i>	45.99	42.24	13.75	7.78	3.73	10.24	29.87	9.78	15.58	20.21	3.60	23.57	15.62	7.46	11.74	8.92	8.23	6.46	6.51	N	N	N	8.93	18.43
BMNH 20.8.8.2	<i>Priapomys leonardi</i>	B	B	12.87	6.81	3.74	8.93	26.59	B	B	18.78	3.50	22.52	B	B	11.35	8.80	8.54	6.66	6.33	N	N	N	8.82	15.73
KIZ 034940	<i>Belomys pearsonii</i>	44.55	41.60	13.11	7.12	3.09	8.90	27.68	8.77	13.98	21.67	3.49	23.01	15.84	9.66	13.25	N	8.93	5.44	5.33	N	N	N	8.71	15.44
KIZ 034965	<i>Belomys pearsonii</i>	42.16	40.05	12.42	6.84	3.53	8.15	27.27	9.13	15.40	22.00	2.76	22.57	14.85	9.94	12.60	N	9.44	5.18	5.60	N	N	N	9.24	16.22

### Supplementary Table S6

Morphological comparison: when the phenotype of the examined specimens of the taxa are consistent, no additional modifiers are added; when there are a few exceptional individuals, "mostly" is added before the phenotype of most individuals; if the two phenotypes appear almost in a 1:1 ratio, recorded as "A or B".

Abbrev: present (P), absent (A), developed (D, very clear), reduced (R, obsolete, but recognizable); Anterior flexus (af), Central flexus (cf), Posterior flexus (pf).

Genera	Skin	Skull		Glans	
	Ear tufts	Auditory bulla septae	Coronoid process	Glans length /HB	Pattern
<i>Aeretes</i>	A	<4	D	unknown	unknown
<i>Aeromys</i>	A	<4	D	unknown	unknown
<i>Belomys</i>	P	>4, honeycomb	D	5%	short and stout, two crests on the right side with thorns on the edge
<i>Biswamoyopterus</i>	P	>4, honeycomb	D	4%	long and stout, two smooth crests on the right side
<i>Eoglaucomys</i>	A	<4	D	4%	short, two smooth crests on the right side, two prominent spiniform processes on the left side
<i>Eupetaurus</i>	A	<4	R	4%	short, three crests on the right side with thorns on the edge
<i>Glaucomys</i>	A	<4	D	7-16%	<i>G. sabrinus</i> : long, three crests on the right side with thorns on the edge <i>G. volans</i> : very long (baculum is only half the length of the glans) and slim, straight, two crests on the right side with thorns on the edge
<i>Hylopetes</i>	A	<4	D	7%	long, two smooth crests on the right side, a prominent spiniform process on the left side
<i>Iomys</i>	A	<4	D	unknown	The only sample available is a broken base, very slim.
<i>Petaurillus</i>	A	<4	D	unknown	unknown
<i>Petaurista</i>	A	<4	D	2-6%	short and stout, one smooth crest on the right side
<i>Petinomys</i>	P	>4, Cobweb*	D	unknown	unknown, baculum can refer to Pocock 1923

<i>Priapomys</i>	A	<4	D	17%	very long (baculum as long as the glans) and slim, "S" shaped, two faint smooth crests on the right side
<i>Pteromys</i>	A	<4	R	6%	long, two crests on the right side with thorns on the edge, a prominent spiniform process on the left side
<i>Pteromyscus</i>	A	ca. 5, honeycomb	D	5%	resembles <i>Belomys</i> , short and stout, two crests on the right side with thorns on the edge
<i>Trogopterus</i>	P	>4, honeycomb	D	3%	resembles <i>Belomys</i> , short and stout, two crests on the right side with thorns on the edge

Genera	Upper cheekteeth								Lower cheekteeth	
	P <sup>3</sup>	mesostyle	mesolophule	anterolophule	Paraconule	Metaconule	Metaloph on M <sup>3</sup>	Lingual flexus	Hypoconulid	protolophid & entolophid
<i>Aeretes</i>	P	mostly A	A	A	D	D	D	D pf, R af & cf	mostly A	D
<i>Aeromys</i>	P	P, isolated	A	A	R	R	A	R af & cf	A	A
<i>Belomys</i>	P	P, connect to paracone	A	A	D	D	R	R af & cf	D	D
<i>Biswamoyopterus</i>	P	P, connect to paracone	R	A	A	R	A	R af & cf	mostly A	A
<i>Eoglaucomys</i>	P	mostly A	A	mostly A	R	R	R	R af	A	mostly A
<i>Eupetaurus</i>	P	A	A	A	very D	A	D	D cf	A	D
<i>Glaucomys</i>	P	mostly A	mostly R	A	A	A	A	R af	A	A
<i>Hylopetes</i>	P	mostly P,	mostly P,	mostly R	mostly A	mostly A	A	R af	A	A

		connect to mesolophule	connect to mesostyle							
<i>Iomys</i>	A	mostly A	A	A	A	A	A	R af & cf	A	D
<i>Petaurillus</i>	mostly P	A	A	A	A	A	mostly A	R af	A	A
<i>Petaurista</i>	P	mostly isolated or connect to paracone	mostly A	A	D	D	D	D pf, R af & cf	A or R	D
<i>Petinomys</i>	P	mostly P, connect to mesolophule	mostly P, connect to mesostyle	R or A	A	mostly R	A	A	A	A or R
<i>Priapomys</i>	P	mostly A	R	R or A	A	R	A	A	A	A
<i>Pteromys</i>	P	mostly A	A	mostly R	A	D	R	R af & cf	A	R
<i>Pteromyscus</i>	mostly P	P, connect to paracone	A	A	R	D	R	R af & cf	D	R
<i>Trogopterus</i>	mostly P	P, connect to paracone	A	A	D	D	D	R af & cf	D	D

\*exception: *Petinomys setosus* is honeycomb.

Pocock, R. I. 1923. The Classification of the Sciuridae. *Proceedings of the Zoological Society of London*, 1923: pp. 244, fig. 28.

**Supplementary Table S7A**

Average genetic distances (%) for 12S ribosomal RNA sequences between the groups of studied flying squirrel species

	<i>Aeromys</i>	<i>Belomys</i>	<i>Biswamoyopterus</i>	<i>Eoglaucomys</i>	<i>Eupetaurus</i>	<i>Glaucomys</i>	<i>Hylopetes</i>	<i>Iomys</i>	<i>Petaurillus</i>	<i>Petaurista</i>	<i>Petinomys</i>	<i>Pteromys</i>	<i>Pteromyscus</i>	<i>Trogopterus</i>
<i>Belomys</i>	0.094													
<i>Biswamoyopterus</i>	0.073	0.119												
<i>Eoglaucomys</i>	0.088	0.109	0.089											
<i>Eupetaurus</i>	0.075	0.104	0.108	0.096										
<i>Glaucomys</i>	0.087	0.123	0.09	0.082	0.091									
<i>Hylopetes</i>	0.12	0.152	0.115	0.123	0.126	0.103								
<i>Iomys</i>	0.099	0.115	0.117	0.088	0.1	0.091	0.107							
<i>Petaurillus</i>	0.125	0.138	0.124	0.116	0.118	0.116	0.1	0.102						
<i>Petaurista</i>	0.109	0.139	0.114	0.106	0.106	0.099	0.117	0.120.117						
<i>Petinomys</i>	0.127	0.15	0.113	0.111	0.134	0.105	0.1	0.094	0.124	0.122				
<i>Pteromys</i>	0.102	0.117	0.116	0.099	0.102	0.108	0.136	0.109	0.144	0.131	0.13			
<i>Pteromyscus</i>	0.089	0.09	0.103	0.105	0.087	0.112	0.148	0.118	0.134	0.131	0.134	0.112		
<i>Trogopterus</i>	0.09	0.054	0.093	0.098	0.102	0.112	0.134	0.113	0.134	0.128	0.127	0.103	0.074	
<i>Priapomys</i>	0.124	0.142	0.107	0.089	0.116	0.095	0.104	0.069	0.112	0.126	0.097	0.124	0.124	0.122

**Supplementary Table S7B**

Average genetic distances (%) for 16S ribosomal RNA sequences between the groups of studied flying squirrel species

	<i>Aeromys</i>	<i>Belomys</i>	<i>Biswamoyopterus</i>	<i>Eoglaucomys</i>	<i>Eupetaurus</i>	<i>Glaucomys</i>	<i>Hylopetes</i>	<i>Iomys</i>	<i>Petaurillus</i>	<i>Petaurista</i>	<i>Petinomys</i>	<i>Pteromys</i>	<i>Pteromyscus</i>	<i>Trogopterus</i>
<i>Belomys</i>	0.12													
<i>Biswamoyopterus</i>	0.052	0.093												
<i>Eoglaucomys</i>	0.117	0.121	0.081											
<i>Eupetaurus</i>	0.085	0.111	0.077	0.11										
<i>Glaucomys</i>	0.099	0.121	0.098	0.068	0.114									
<i>Hylopetes</i>	0.109	0.13	0.094	0.091	0.121	0.079								
<i>Iomys</i>	0.094	0.106	0.086	0.085	0.107	0.078	0.072							
<i>Petaurillus</i>	0.104	0.109	0.093	0.075	0.112	0.066	0.068	0.059						
<i>Petaurista</i>	0.123	0.136	0.122	0.127	0.126	0.124	0.104	0.107	0.102					
<i>Petinomys</i>	0.109	0.127	0.11	0.078	0.132	0.099	0.079	0.097	0.08	0.116				
<i>Pteromys</i>	0.109	0.109	0.098	0.096	0.096	0.106	0.115	0.094	0.073	0.118	0.101			
<i>Pteromyscus</i>	0.109	0.085	0.11	0.131	0.114	0.121	0.147	0.121	0.108	0.133	0.146	0.121		
<i>Trogopterus</i>	0.114	0.055	0.086	0.108	0.101	0.109	0.11	0.104	0.094	0.123	0.119	0.096	0.073	
<i>Priapomys</i>	0.113	0.12	0.092	0.093	0.109	0.084	0.065	0.056	0.06	0.113	0.098	0.105	0.133	0.102

