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A new acaremyid rodent (Caviomorpha, Octodontoidea) from Scarritt Pocket, Deseadan (late Oligocene) of Patagonia (Argentina)

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CHARACTER LIST S1. Morphological character list based on the dataset of Vucetich & Kramarz (2003). The following multistate characters are treated as ordered: 1, 6, 14 and 15.

- (1) Degree of hypsodonty: brachydont (0); slightly hypsodont (1); mesodont (2); protohypodont (3); euhypsodont (4).
- (2) Cusps differentiable from respective crests: yes (0); no (1).
- (3) Figure eight shaped lower cheek teeth: absent (0); present (1).
- (4) Crests obliquity in lower teeth: no (0); yes (1).
- (5) Deciduous premolar: normal replacement (0); retained (4).
- (6) Hypoflexus in the upper premolar: absent or incipient (0); poorly developed (1); well developed (2).
- (7) Number of crest of upper premolar: trilophodont (0); tetralophodont (1).
- (8) Quadrangular protocone area in M1-M2: no (0); yes (1).
- (9) Flexid on the anterior wall of the p4: present (0); absent (1).
- (10) Development of metalophulid II in the p4: short (0); long (1).
- (11) Lingual end of metalophulid II in p4: free (0); join to the metaconid (1); join to the metalophulid I.
- (12) Ectolophid length in p4: long (0); short (1).
- (13) Hypolophid in p4 in intermediate ontogenetic stage: absent or incipient (0); well developed (1).

(14) Development of the metalophulid II in m1: reaching the lingual wall (0); not reaching the lingual wall (1); absent (2).

(15) Development of the metalophulid II in m2: reaching the lingual wall (0); no reaching the lingual wall (1); absent (0).

(16) Length of the posterolophid in m1-m2: short (0); long (1).

(17) Orientation of nMpi: oblique AD-PV (0); horizontal (1); oblique AV-PD (2).

(18) Anterodorsal limit of the mandibular masseteric fossa: absent (0); present (1).

(19) Mental foramen: present (0); absent (1).

(20) Figure eight shaped upper cheek teeth: absent (0); present (1).

(21) AP-L of anterior lobe in m2 with respect to AP-L of posterior lobe: subequal (0); about 75% (1); less than 50% (2).

(22) Origin of metalophulid II in m1: from protoconid (0); begin the protoconid (1).

(23) Origin of metalophulid II in m2: from protoconid (0); begin the protoconid (1).

(24) Discontinuity between the masseteric crest and the nMpi: absent (0); present (1).

MATRIX S1. Character-taxon matrix used for phylogenetic analysis. Characters between brackets represent polymorphic or uncertain scorings.

Taxon	10	20	24
<i>Phiomys andrewsi</i>	0000000010	2001010000	0110
<i>Deseadomys arambourgi</i>	1101000100	00011[01]0000	1110
<i>Platypittamys brachyodon</i>	0000000100	0000[01]00000	1110
<i>Migraveramus beatus</i>	11000??11	0110[01]1000?	0000
<i>Galileomys antelucanus</i>	101001[01]001	1001100101	1110
<i>Acaremys murinus</i>	101001[01]000	0000110101	1110
<i>Sciamys principalis</i>	2110021001	1000011101	1110
<i>Protacaremys prior</i>	10001??1??	???0011000	0110
<i>Chasichimys bonaerense</i>	31111??0??	???221101?	2??1
<i>Chasicomys octodontiforme</i>	31101??0??	???221?011	2??1
<i>Octomys mimax</i>	41111??0??	???2212011	0??1
<i>Massoiamys obliquus</i>	21111??0??	???2210010	2??0
<i>Eumysops laeviplicatus</i>	21001??0??	???2210010	2??0
<i>Stichomys regularis</i>	21011??1??	???[12][12]10000	20?0
<i>Changquin woodi</i>	2110???????	???00101??	0110
<i>Draconomys verai</i>	01000??111	0111110100	0000
<i>Sallamys pascuali</i>	100000[01]100	0010010000	1000
<i>Prospaniomys priscus</i>	11011??0??	???1100000	0000
<i>Spaniomys riparius</i>	21011??0??	???0001000	0000
<i>Willidewu steparius</i>	1000000010	0111111000	2000