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A comparative analysis of the molars of *Mus booduga*, *Mus dunnii* and fossil *Mus* of the Indian subcontinent: Phylogenetic and palaeobiogeographic implications

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Upper molars of *Mus booduga* collected from Varanasi and *Mus dunnii* (chromosome type-I) show slightly derived features compared to those of *Mus booduga* collected from Mysore and *Mus dunnii* (chromosome type-II). *Mus auctor*, *Mus elegans*, *Mus flynnii*, *Mus sp. indet.*, *Mus jacobsi*, *Mus sp.*, *Mus booduga* and *Mus dunnii* are closely related to each other in having moderately elongated and rather strongly distorted M¹s and moderately reduced M³s (except for *Mus elegans*, *Mus sp.* and *Mus jacobsi*). Representation of *Mus*, both in terms of taxonomic diversity and considerable numbers, in the Plio-Pleistocene deposits of the Indian subcontinent is indicative of an early diversification event in the history of *Mus*.

In all murids, M¹ (first upper molar) is relatively longer than M² (second upper molar), which in turn is longer than M³ (third upper molar). According to Misonne¹ lengthening of M¹ in a forward direction accompanied by reduction of M³, brings the whole series forward and occupies the place left by P⁴ (fourth upper premolar). The length of M¹ and M³ is given in relation to M², which has a standard relative length of 100% (ref. 1). Generally ancient murids have broad molars, thus Misonne¹ suggested that broad molars correspond to the more generalized types. Jacobs² proposed that in primitive murids, labial cingulum on lower molars are poorly developed and anterostyle on M¹ relative to lingual anterocone, but not on the posterior extremity.

During the preliminary studies on Late Pliocene *Mus*, it was observed that it resembles *Mus booduga*, the Indian Pigmy Field mice³, most closely. In the light of evolutionary trends of murids suggested by Misonne¹ and generalized dental characters of murids proposed by Jacobs² an attempt has been made here to study *Mus booduga* (extant taxon, collected from Varanasi and Mysore), *Mus dunnii* (extant taxon, chromosome types I, II and III), *Mus auctor* and *Mus sp.* reported from Late Miocene and Early Pleistocene Siwalik deposits, respectively², *Mus flynnii* and *Mus sp. indet.* recovered from Late Pliocene (around 2.5 m.y.) Siwalik sediments

of India⁴, *Mus jacobsi* discovered from Late Pliocene (around 2.4 m.y.) deposits of Karewas (Kashmir, India)⁵ and *Mus elegans* reported from Pliocene deposits of Paul-I-Charkhi, Afghanistan⁶.

Eleven skulls (with 132 teeth) each of adult *Mus booduga* and *Mus dunni* housed at Cytogenetics Laboratory, Centre of Advanced Study in Zoology, Banaras Hindu University, were examined. Forty isolated teeth of fossil *Mus* housed at Vertebrate Palaeontology Laboratory, Centre of Advanced Study in Geology, Panjab University have been used here for comparison.

Measurements were made using a microscope fixed with a reticule. Dental terminology of murid molars proposed by Jacobs² is followed here.

Matthey and Petter⁷ were first to distinguish between *Mus booduga* and *Mus dunni* on the basis of divergent karyotypes but they also observed a slight difference in shape of molar and colour of underparts. The colour of underparts however is at times misleading. The pygmy field mice *Mus booduga* and *Mus dunni*, which are endemic to the Indian subcontinent, are morphologically very similar and they inhabit the same ecological fields.

The diploid number of chromosomes is 40 in them but while the karyotype of *Mus booduga* with all acrocentric chromosomes is identical to *Mus musculus*, that of *Mus dunni* is distinct due to presence of composite submetacentric X and large acrocentric Y chromosomes. The works carried out in the Cytogenetics Laboratory of BHU over the last several years have revealed that *Mus dunni* is in active phase of speciation. In sharp contrast to the extreme conservatism observed in the karyotype of *Mus booduga* throughout India, *Mus dunni* populations of different places have divergent karyotypes. Three chromosome types (I, II and III), which are apparently parapatric in distribution, have been found so far, and on conducting mate-preference behaviour in them have shown some degree of ethological isolation⁸. Recently, on hybridization between chromosome types I and III, it has been observed that they are also to some extent reproductively isolated since varying conditions of hybrid sterility of males and inviability of hybrid females have been observed⁹. The stable karyotypic difference among the three chromosome types of *Mus dunni* has been achieved by establishing homozygosity for heterochromatin variation at the centromeric regions and as prominent short arms of the autosome pairs 1, 3 and 6. The chromosome types are incipient biological species.

Marshall¹⁰ examined specimens of *Mus booduga* collected from Uttar Pradesh, Madhya Pradesh, Karnataka and Tamil Nadu states, and of *Mus dunni* collected from Haryana, Uttar Pradesh, Madhya Pradesh, Maharashtra and Tamil Nadu states of India. According to him, *Mus booduga* has a long and slender

first upper molar with a long anterior cusp (lingual anterocone) surmounted by an accessory cusp (prestyle) and an inconspicuous antero-external cusp (labial anterocone). During the present investigation, the above-mentioned features were clearly observed on *Mus booduga* collected from Varanasi but specimens of *Mus booduga* collected from Mysore have relatively shorter lingual anterocones on M¹ and lacked a prestyle (Figure 1 a, b).

Incisive foramina of *Mus booduga* collected from Varanasi penetrate deeper (reaching between anterostyle and enterostyle of the M¹) compared to those of *Mus booduga* collected from Mysore and *Mus dunni* (chromosome types I, II and III), where it reaches the level of anterostyle of M¹ (Figure 1 f-j). Upper incisors of *Mus booduga* collected from Mysore are more curved compared to those of *Mus booduga* (Varanasi) and *Mus dunni* (types I, II and III) (Figure 1 k-o).

Mus is characterized by having a reduced M₃ and M³, anterostyle on M¹ anteroposteriorly compressed and



Figure 1. Occlusal views of upper and lower molars of a, *Mus booduga*, (Varanasi specimen no. Mbv-3); b, *Mus booduga* (Mysore sp. no. M3); c, *Mus dunni* (type I, sp. no. Mdv-4), d, *Mus dunni* (type II, sp. no. M6); e, *Mus dunni* (type III, sp. no. Mdm-1); f-j and k-o, skulls showing position of incisive foramina and curvature of upper incisors (sp. nos. Nbv 3, M3, Mdv-4, M6 and Mdm-1 respectively). Bar = 1 mm.

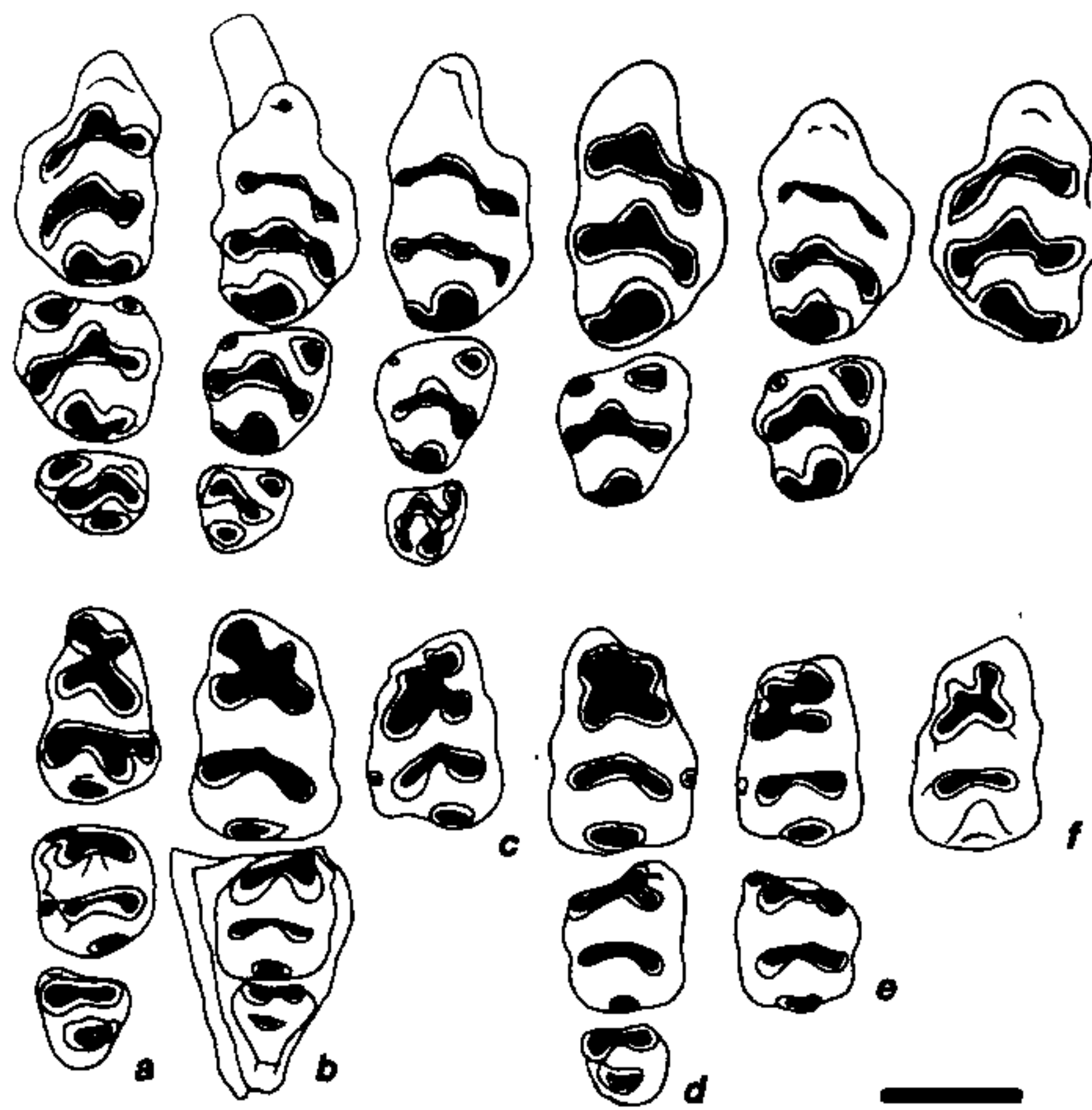


Figure 2. Occlusal views of upper and lower molars of a, *Mus auctor* (sp. nos. M¹-DP210, M²-DP325, M³-DP 326, M₁ DP294, M₂-DP 335, M₃-DP 337) redrawn from Jacobs²; b, *Mus flynni* (all VPL/RP numbers. M¹SM-2, M²-Sm, -32, M³-Sm-45, M₁SM-10, M₂&M₃-SM-38), redrawn from Patnaik⁴; c, *Mus sp. indet.* (all VPL-RP nos. M¹-SM-5, M²-SM-33, M³-SM-44, M₁-M-6-A), redrawn from Patnaik⁴; d, *Mus jacobsi* (M¹-KHM/11, M²-KHM/104, M₁-KHM/37, M₂-KHM/85, M₃-KHM/120, redrawn from Kotlia⁵; e, *Mus elegans* (M¹-AFG 932, M²-AFG-936, M₁, AFG-415, M₂-AFG-421), redrawn from Sen⁶; f, *Mus sp.* (M¹-DP-210, M₁-DP-202), redrawn from Jacobs². Bar = 1 mm.

posterior relative to the lingual anterocone and posterior cingulum reduced or absent. Further, M₁ has an 'X' pattern formed by the joining of labial and lingual anteroconids, protoconid and metaconid. *Mus auctor* is considered to be the most primitive form of *Mus* recovered so far. It differs from *Mus flynni*, *Mus sp. indet.*, *Mus jacobsi*, *Mus elegans*, *Mus sp.*, *Mus booduga* and *Mus dunni* in having a reduced M¹ with distinct precingulum, M² with a conspicuous labial anterocone and a reduced posterior cingulum and M³ with a labial anterocone (Figure 2a). Possible phylogenetic relationships between *Mus flynni* and *Mus auctor* and *Mus jacobsi* and *Mus auctor* have been suggested^{4, 5}.

Molars of *Mus dunni* (type-II), *Mus booduga* (Mysore) and *Mus flynni* resemble each other in having moderately elongated lingual anterocones and M₁s lacking labial cingulum (Figure 1b, d and 2b). On the other hand, molars of *Mus dunni* (type I and III), *Mus booduga* (Varanasi), *Mus sp. indet.* and *Mus jacobsi* are quite similar to each other in having considerably elongated lingual anterocones, prestyles (variably present) M₁s with distinct labial cingulum and medial anteroconids (Figure 1a, c, e, 2c and d).

Misonne¹ suggested that in advanced murids, M¹ is extremely elongated relative to M² and M³ and some

Table 1. Measurements (in mm) of upper molars of various *Mus* species examined. Mean lengths of *Mus auctor*, *Mus flynni*, *Mus elegans*, *Mus jacobsi* and *Mus sp. indet.* taken from Jacobs², Patnaik⁴, Sen⁶, Kotlia⁵, and Patnaik⁴, respectively. Length of M² has been taken here as a standard.

Species	Element	Mean length	Percentage
<i>Mus auctor</i>	M ¹	1.699	162
	M ²	1.047	100
	M ³	0.63	60.17
<i>Mus flynni</i>	M ¹	1.83	179
	M ²	1.02	100
	M ³	0.66	64
<i>Mus elegans</i>	M ¹	1.876	179
	M ²	1.044	100
<i>Mus jacobsi</i>	M ¹	2.149	200.27
	M ²	1.073	100
<i>Mus sp. indet.</i>	M ¹	2.13	206
	M ²	1.03	100
	M ³	0.56	54.36
<i>Mus booduga</i> (Mysore)	M ¹	1.75	190.21
	M ²	0.92	100
	M ³	0.52	56.52
<i>Mus booduga</i> (Varanasi)	M ¹	1.93	198.96
	M ²	0.97	100
	M ³	0.58	59.79
<i>Mus dunni</i> (type I)	M ¹	1.84	197.84
	M ²	0.93	100
	M ³	0.56	60.21
<i>Mus dunni</i> (type-II)	M ¹	1.78	189.36
	M ²	0.94	100
	M ³	0.58	61.70
<i>Mus dunni</i> (type-III)	M ¹	1.87	190.80
	M ²	0.98	100
	M ³	0.55	56.12

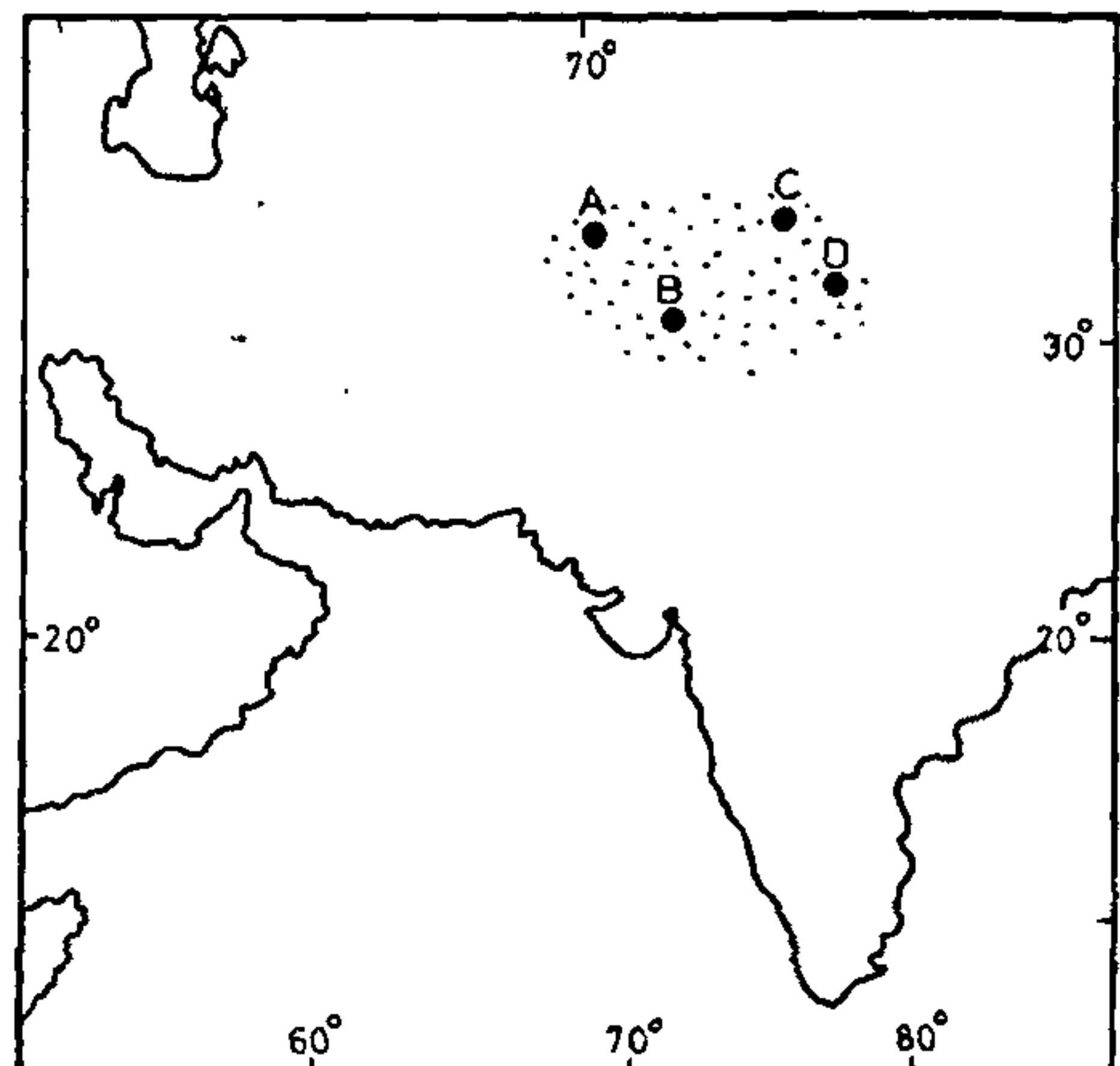


Figure 3. Plio-Pleistocene biogeographic province based on the fossil *Mus* occurrences in the Indian subcontinent. A, Paul-I-Charkhi, Afghanistan; B, Pabbi Hills, Pakistan; C, Kashmir Valley, India; D, Saketi Village (Himachal Pradesh, India)

forms of *Mus* even have M¹ with length over 240% of M² and M³ around 40%; this gives a total of

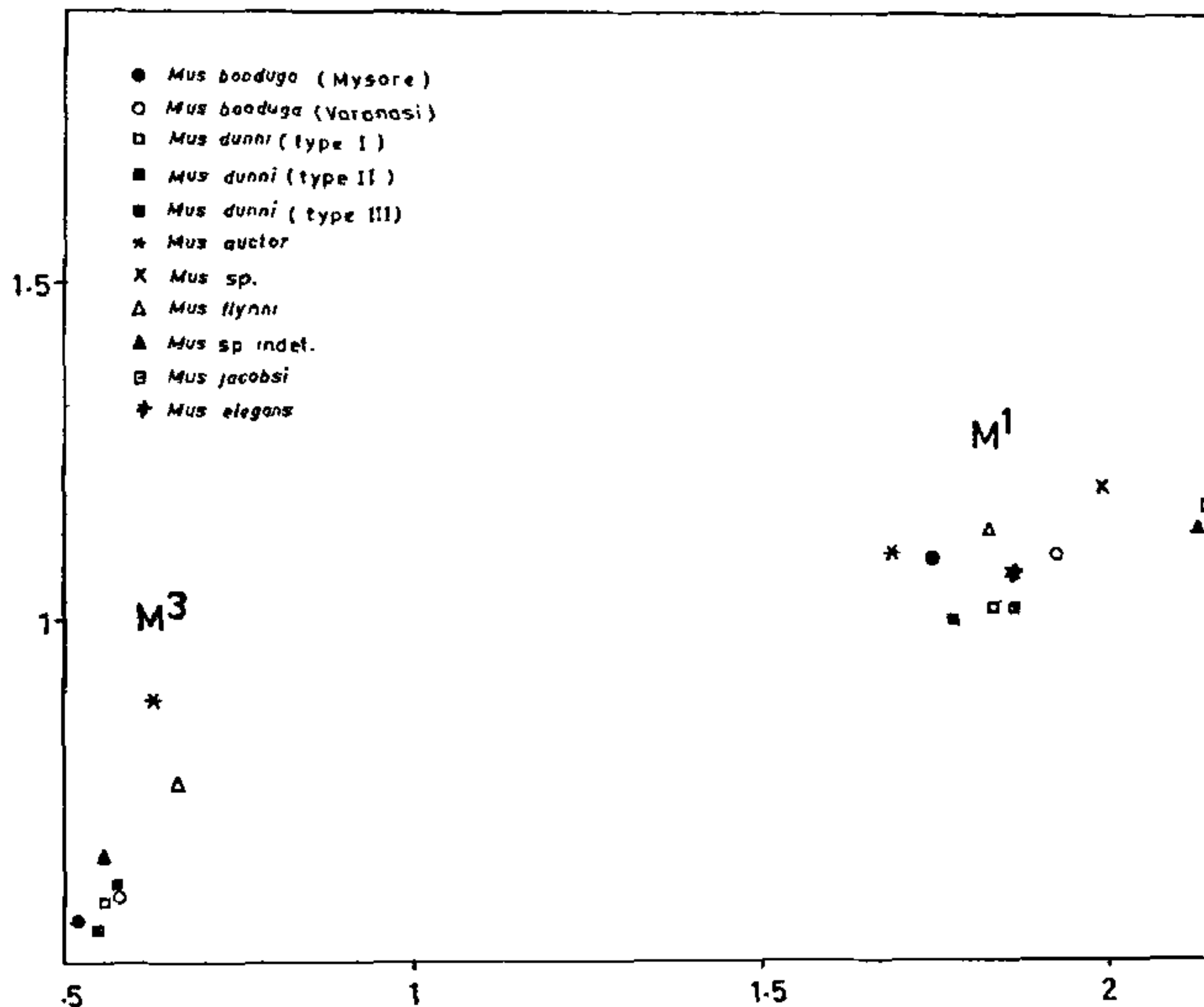


Figure 4. Scatter diagram of length and width of M^1 and M^3 's of various *Mus* species examined. Here plots are made by taking mean values of length and width of *Mus auctor* and *Mus sp.* (from Jacobs²) *Mus flynni* and *Mus sp. indet.* (from Patnaik⁴) *Mus jacobsi* (from Kotlia⁵) and *Mus elegans* (from Sen⁶).

240 + 100 + 40 = 380% which implies that in spite of having three teeth, such specimens would have a surface equal to that of four teeth of dental series of primitive rodents. Such calculations were carried out on fossil and extant taxon compared herein. The results show that *Mus auctor* should be most generalized of all in this character followed by *Mus flynni* and *Mus elegans* (Table 1). *Mus sp. indet.* and *Mus jacobsi* are considered here to be relatively derived in this character. Among the extant forms, *Mus booduga* (Mysore) and *Mus dunni* (type II) are generalized in this character compared to *Mus booduga* (Varanasi) and *Mus dunni* (type-I), whereas *Mus dunni* (type-III) falls somewhere in between these two types.

Three specimens of *Mus sp.* have been reported from the Early Pleistocene Siwalik deposits, near Pabbi Hills, Pakistan². Forty-three specimens of *Mus jacobsi* come from Late Pliocene deposits of Karewa, Kashmir⁵. Twenty-nine specimens of *Mus flynni* and 11 specimens of *Mus sp. indet.* have been collected from Late Pliocene, Siwalik sediments, near Saketi Village, Himachal Pradesh, India⁴. Such a representation of *Mus* species may suggest an early diversification in the history of *Mus*. On the basis of occurrences of *Mus* in Pliocene and Pleistocene deposits, a palaeobiogeographic province has been constructed (Figure 3).

Apart from the Indian subcontinent, Plio-Pleistocene *Mus* is known from Hadar Formation, Ethiopia (represented by four molars)¹¹ and Lake Turkana, Kenya (represented by six molars)¹². As this collection from Africa lacks M^1 's, it is at the moment difficult to compare it with that of the Indian subcontinent from the phylogenetic point of view.

In Figure 4, values (length and width) of M^1 's and M^3 's of all the *Mus* specimens fall in one domain. This observation encourages us to propose that on the whole *Mus booduga*, *Mus dunni*, *Mus flynni*, *Mus auctor*, *Mus elegans*, *Mus sp. indet.*, *Mus jacobsi* and *Mus sp.* are very close to each other and further recovery of fossil material, particularly from Pleistocene deposits may provide a better picture on the evolutionary lineages of *Mus*.

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Prostatic inhibin has a predominantly anti-parallel β -sheet structure

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Prostatic inhibin (94 amino acid residues, $M_r = 10,540$) is a protein isolated from the human and animal prostate glands. Three-dimensional structure of this cysteine-rich (10/94) protein has been studied by NMR spectroscopy. Preliminary investigations provide valuable information on the secondary structure of this protein. It is found to acquire a predominantly anti-parallel β -sheet structure and possibly the molecule is locked into several such sheets through disulphide linkages.

PROSTATIC inhibin is a protein with 94 amino acids and molecular weight of 10 kDa. It has been isolated from the human and animal prostate glands¹. More than a decade of research has established a wide range of its biological activities, ranging from preventing pregnancy to curing prostate cancer. Inhibin prevents pregnancy by modulating the level of circulating follicle-stimulating hormone (FSH) in mammals¹. It suppresses prolactin, a hormone that promotes lactation². Therefore, neutralizing inhibin through active immunization has been found to increase milk production. Although the primary structure of this molecule³ has been determined, no information is available so far about the three-dimensional structure. Our preliminary NMR investigations throw light on the three-dimensional structure of prostatic inhibin – an information of great value, in view of the useful biological activities of this molecule.

About 20 mg of HPLC pure protein⁴ was dissolved in 0.5 ml of an appropriate solvent (approximately 3 mM) and buffered with 100 mM acetate buffer. The pH was

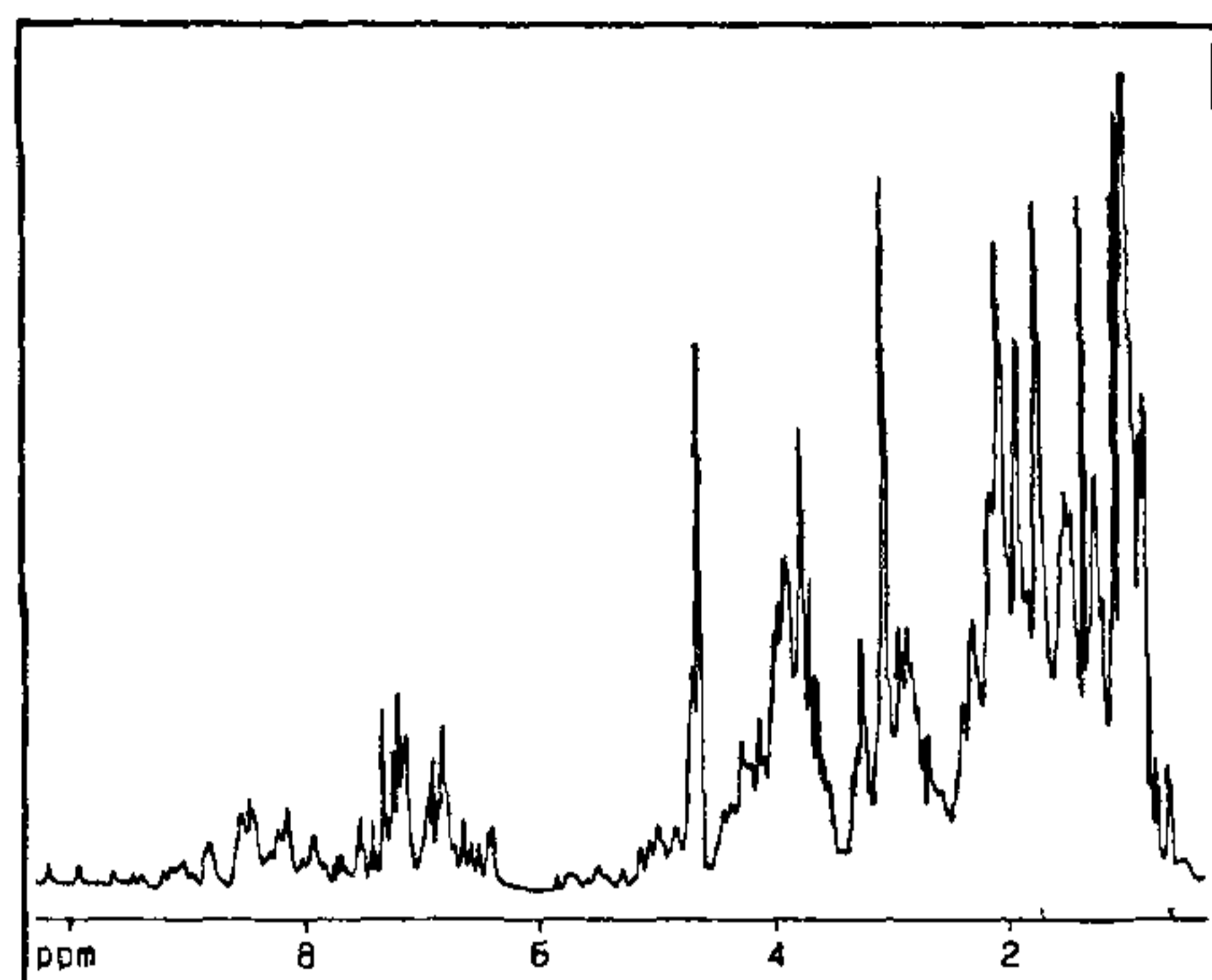


Figure 1. 500 MHz ^1H NMR spectrum of prostatic inhibin in a mixed solvent of 90% H_2O + 10% $^2\text{H}_2\text{O}$ at 310 K and pH 4.2 (acetate buffer).

adjusted to 4.2. NMR measurements were carried out in 99.9% $^2\text{H}_2\text{O}$ and in a mixed solvent consisting of 90% H_2O and 10% $^2\text{H}_2\text{O}$. Temperature was optimized for the best possible resolution and all spectra were recorded at 310 K. NMR experiments were carried out on a Bruker AMX 500 spectrometer with a ^1H frequency of 500 MHz and involve (i) two-dimensional (2D) two-quantum-filtered correlation spectroscopy (2QF COSY)⁵, (ii) 2D clean total correlation spectroscopy (clean TOCSY)⁶ with a mixing time of 100 ms, and (iii) 2D nuclear Overhauser enhancement spectroscopy (NOESY)⁷.

Prostatic inhibin has about 600 observable protons. The 500 MHz NMR spectrum shows reasonably well resolved features. Figure 1 shows the 1D ^1H spectrum of the protein in a mixed solvent of 90% H_2O and 10% $^2\text{H}_2\text{O}$. Figure 2A and 2B show 2QF COSY and NOESY spectra, respectively, in 99.9% $^2\text{H}_2\text{O}$. A detailed analysis of these and other spectra has enabled us to identify several spin systems. These include eight threonines, five valines, two glycines, one leucine, one isoleucine and twenty-one AMX spin systems (belonging to Cys, Ser, Asp, Asn, Tyr, Trp, His and Phe). The subspectral features are well dispersed. For example, the $\text{C}_\beta\text{H}-\text{C}_\gamma\text{H}_3$ correlations for all the eight threonine residues present in the prostatic inhibin are shown in Figure 3. Thus, at this stage, we have been able to identify almost half of the spin systems. Even in the absence of sequential resonance assignments, the NMR data provide valuable information on the secondary structure of this protein.

In the first instance, the 1D spectrum (Figure 1) indicates that the protein has a well-defined and ordered structure. There are several downfield-shifted C_αH protons as well as several upfield-shifted methyl resonances. The downfield shift of the C_αH protons is a