Localization of monoamine oxidase A and B genes on the mouse X chromosome

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SOURCE/DESCRIPTION: A 700bp Sacl fragment of the 2.1 kb human monoamine oxidase A(MAOA) cDNA, cloned in the EcoRl site of pUCl9, and a 2.5 kb cDNA of the human monoamine oxidase B gene(MAOB) in the EcoRl site of pSP6T719 (1).

<u>POLYMORPHISM</u>: The MAOA probe recognises a 14 kb S allele and two M alleles at 6.6 and 4.4 kb in Taq digests. The MAOB cDNA recognises two S alleles at 6.6 and 4.8 kb and two M alleles at 9.0 and 5.1 kb in Taq digests (Fig.1).

FREQUENCY: The M alleles are detected in the inbred <u>Mus musculus</u> (C57BL/10) strain, and the S alleles in the <u>Mus spretus</u> mouse.

NOT POLYMORPHIC FOR: Pstl, Mspl.

<u>CHROMOSOMAL LOCALISATION:</u> The mouse genes homologous for MAOA and MAOB have been mapped to the <u>Cybb</u> - <u>Timp</u> interval of the proximal mouse X chromosome by genetic breakpoint analysis. This predicts a human localisation of Xp21.1p11.21 for both genes, and is in agreement with published human mapping data (2) showing conservation of gene order.

<u>MENDELIAN INHERITANCE:</u> Segregation of the restriction fragments was followed in 60 recombinant backcross progeny, resulting from an interspecific backcross between <u>Mus musculus x Mus spretus</u> mice. 6 animals in this pedigree recombine in the <u>Cybb</u> - <u>Timp</u> interval and both <u>Maoa</u> and <u>Maob</u> cosegregate with <u>Timp</u> in 5 of 6 recombinant animals. This suggests <u>Maoa</u> and <u>Maob</u> lie close together on the mouse X-chromosome and centromeric to <u>Timp</u>. PROBE AVAILABILITY: Contact Dr J.C. Shih.

<u>OTHER COMMENTS:</u> We propose the gene symbols <u>Maoa</u> and <u>Maob</u> to identify the mouse equivalents of MAOA and MAOB.

REFERENCES: 1. Bach et al., Proc.Natl.Acad.Sci. USA 85, 4934-4938 (1988). 2. Lan et al., Genomics 4, 552-559 (1989).

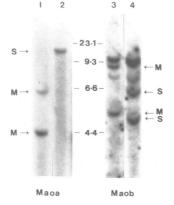


Fig. 1. Autoradiogram of a southern blot of Taq 1 digested murine DNA probed with a 700bp Maoa cDNA probe (lanes 1 and 2), and a 2.5 kb Maob cDNA probe (lanes 3 and 4). Lanes 1 and 3 show the musculus alleles (M) and lanes 2 and 4 the spretus alleles (S). Common bands are seen in lanes 3 and 4 at 9.5 kb and 7.5 kb.