

## The gene coding for variant hepatic nuclear factor 1 (*Tcf-2*), maps between the *Edp-1* and *Erba* genes on mouse Chromosome 11

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Variant Hepatic Nuclear Factor 1 (vHNF1), also called TCF2, HNF1 $\beta$ , or LFB3, is a nuclear protein that binds to proximal promoter sequences of  $\alpha$  and  $\beta$  fibrinogen and albumin. The protein was first identified in dedifferentiated rat hepatoma cell lines (Baumhueter et al. 1988; Cereghini et al. 1988). The mRNA is found in significant levels in adult rat and mouse kidney and in low levels in liver, intestine, and lung (Rey-Campos et al. 1991; De Simone et al. 1991; Mendel et al. 1991). The cDNA corresponding to the rat mRNA has been isolated and sequenced from the dedifferentiated hepatoma H5 cell line (Rey-Campos et al. 1991). The human and mouse cDNAs have also been isolated and share approximately 93% sequence homology with the rat cDNA (Bach et al. 1991; De Simone et al. 1991; Mendel et al. 1991). The gene codes for a homeoprotein homologous to Hepatic Nuclear Factor 1 (HNF1) or LFB1 in regions important for DNA binding and dimerization while the sequence is highly divergent in the transactivation domain. Both HNF1 and vHNF1 act as transcriptional activators of the albumin promoter in transient co-transfection experiments, and these proteins form heterodimers (Rey-Campos et al. 1991). Though HNF1 and vHNF1 are two closely related homeoproteins, they are localized on different chromosomes. HNF1 has been previously mapped to mouse Chr 12 and human Chr 5 (Bach et al. 1990).

Transcription factor 2 (*Tcf-2*) has been assigned to mouse Chr 11 and human Chr 17 by analysis of somatic cell hybrids (Milatovich et al. 1991). In situ hybridization has localized the gene to human Chr 17q11.2-q21.1 and mouse Chr 11B4-D (Bach et al. 1991). We report chromosomal location of mouse *Tcf-2* relative to other previously mapped Chr 11 loci using an intersubspecific mouse backcross. This (DF/B-*df/df*  $\times$  CASA/Rk) $F_1$   $\times$  DF/B-*df/df* backcross has been typed for 12 loci on Chr 11 (Buckwalter et al.

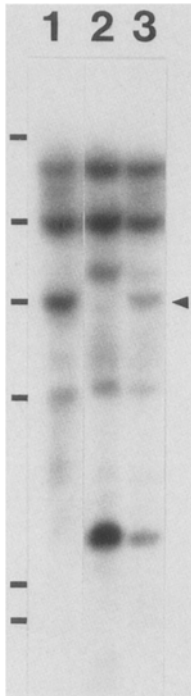
1991). Southern blotting and probe labeling was done by standard methods.

An informative *CASA/Rk BamHI* polymorphism was detected with the *Tcf-2* probe (Fig. 1). A DF/B-*df/df* specific fragment is located at 7.5 kb and a *CASA/Rk* specific fragment at 6.8 kb. The results of typing 86 animals for endothelial cell-derived protein (*Edp-1*; Wolf et al. 1992; Buckwalter et al. 1991), *Tcf-2*, and Avian erythroblastosis virus oncogene A (*Erba*) (Lazar et al. 1988; Buckwalter et al. 1991) are given in Table 1. Five recombinants were found between *Edp-1* and *Tcf-2*, placing *Tcf-2*  $5.8 \pm 2.5$  cM (centimorgans  $\pm$  standard deviation) distal to *Edp-1* on Chr 11. Fourteen recombinants were found between *Tcf-2* and *Erba*, placing *Tcf-2*  $16.3 \pm 4.0$  cM proximal to *Erba* on this chromosome. This mapping generates the following gene order: *Edp-1*-5.8 cM-*Tcf-2*-16.3 cM-*Erba*.

This localization corrects the preliminary mapping of *Tcf-2*, also known as *DII Pas3*, in an interspecific backcross with *Mus spretus* performed in the laboratory of J.-L. Guénet (Buchberg et al. 1991). Further analysis of those data places *Tcf-2* between *Csfgm* and *Gfap*, confirming our mapping of *Tcf-2*. In 22 progeny, there was one recombinant between *Csfgm* and *Tcf-2*, and there were four between *Tcf-2* and *Gfap*.

Homeobox-2, (*Hox-2*) a cluster of homeotic genes on Chr 11, has been mapped  $0.7 \pm 0.7$  cM proximal to *Erba* (Buchberg et al. 1989). The 95% confidence limits for the *Erba-Hox-2* and *Erba-Tcf-2* distances are not overlapping. Thus, *Tcf-2* is localized outside this cluster of other homeotic genes. This was further confirmed by detection of four recombinants out of 22 progeny typed for both *Tcf-2* and *Hox-2* in the interspecific backcross with *M. spretus* mentioned above.

This multipoint molecular genetic localization of *Tcf-2* is consistent with the in situ hybridization of *Tcf-2* to mouse 11B4-D. Our localization of *Tcf-2* corresponds to 11B4 (Buchberg et al. 1991). Our mapping is also consistent with the in situ localization of the human gene to 17q11.2-21.1 because the genes that we



**Fig. 1.** Southern blot identification of a unique *M. castaneus* (*CASA/Rk*) RFLP in *Tcf-2*. An arrowhead signifies the 6.8-kb fragment present in genomic DNA from *CASA/Rk* (lane 1) and from an animal heterozygous for *DF/B-df/df* and *CASA/Rk* (lane 3), but absent in an animal homozygous for *DF/B-df/df* at this locus (lane 2). Digests were performed with the restriction endonuclease *Bam*HI. Molecular size standard is  $\lambda$  DNA-*Hind*III fragments.

mapped proximal (*Edp-1*) and distal (*Erba*) to *Tcf-2* also map to human Chr 17. *Edp-1* maps to human 17q22-23 (Wolf et al. 1992), and *Erba* maps to human 17q11.2-21 (Buchberg et al. 1991). The localization of *Tcf-2* near *Edp-1* is consistent with the synteny con-

**Table 1.** Segregation of three loci on Chr 11 among 86 backcross offspring from the cross (*DF/B-df/df*  $\times$  *CASA/Rk*)F<sub>1</sub>  $\times$  *DF/B-df/df*. The symbols D and C represent *DF/B-df/df*-derived and *CASA/Rk*-derived alleles, respectively.

	Loci inherited from F <sub>1</sub> parent			Number of progeny
	<i>Edp-1</i>	<i>Tcf-2</i>	<i>Erba</i>	
Nonrecombinants	D	D	D	34
	C	C	C	33
Single recombinants	D	C	C	4
	C	D	D	1
	D	D	C	5
	C	C	D	9

servation and lack of linkage conservation observed between mouse Chr 11 and human Chr 17 (Buchberg et al. 1989).

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