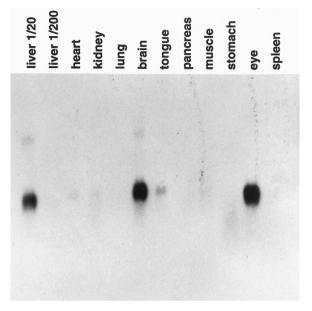
Erratum





Northern blot analysis of the TTR mRNA from tissues of human wild-type transthyretin transgenic mice in which 10 μ g of mRNA from the noted tissues of the mice were treated with glyoxal, separated on a 1% agarose gel, transferred onto nitrocellulose paper and probed with ³²P-cDNA encoding human TTR. This figure replaces the previous Figure 2 that represents a similar analysis in mice transgenic for a single copy of human mutant TTR Leu55Pro and was included in the paper in error. The present figure shows the greater density of the mRNA bands, as expected in animals integrating the greater number of intact gene copies, found in the same tissues as in the single copy transgenics, reflecting the fact that both constructs were under the same tissue-specific transcriptional control.