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Response: In the note included in reference 11 of our report (1) we did not indicate that clone 4D-12 described by Witney and Furano (2) cannot be regarded as a member of the "ID" repeat family in the rat genome. In fact we stated that Furano regards 4D-12 as a typical representative of the ID repeat family. We added the comment on 3' end divergency only because in comparison to clone p2A120 and the common or consensus 82-nucleotide sequence arrived at by Sutcliffe *et al.* (3) the sequence of clone 4D-

12 is different at the 3' end. Within the 3' region of the ID clone p2A120 and the 82-nucleotide common sequence the sequence CCCAGCTCCGAAAAA occurs (3, 4). In clone 4D-12, beginning after alignment of the common triplet CCC, the sequence ...CAGCATCACAGG is found (2), which is divergent from the clone p2A120 that we used in our measurements. Perhaps we were overly cautious in raising this minor point for readers to consider. Of more importance and as we mentioned twice in our report is the observation by Witney and Furano that the 4D-12 cloned repeat hybridizes to nuclear RNA from several sources including liver and kidney (2), just as we found using the ID repeat in clone p2A120. Hence the determinations made by Witney and Furano and those contained in our report both

point to rather ubiquitous transcription of the ID repeat in various rodent cells and organs.

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