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## References

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Tables should be numbered and carry headings enabling them to be understood without reference to the text. Each Table should be typed on a separate sheet. Symbols for footnotes should be in the order : †, ‡, §, ¶, ††, etc. The use of \*, \*\*, etc, should be limited to indicating levels of significance.

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Printed originals of figures and photographs should be provided as best possible quality. Figures such as graphs must be supplied in an editable file format, such as Excel. The use of bar graphs and histograms should be restricted, as the information can often be better presented in a table. In the presentation of results, experimental points should be indicated by symbols, used in order: ○, ●, △, ▲, □, ■, ×, +. Scale marks should be on the inside of the axes. Each Figure should be provided with a legend such that with the Figure it is comprehensible without reference to the text. Figure legends should be typed on a separate sheet or sheets, beginning Fig. 1.

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Individual results should not normally be given. The methods of statistical analysis should be clearly described; a suitable reference is adequate. Authors should make it clear whether they are quoting (e.g.) SD or SE. Any statement that two groups of values are different should be supported by the level of significance involved, as a single or range of *P* value: (*P* = 0.008) or (*P* < 0.01). Differences should not be claimed or implied if *P* > 0.05.

## Gene Sequences

Original DNA sequences reported in JDR must also be submitted to GenBank. Instructions can be found at <http://www.ncbi.nlm.nih.gov/Genbank/index.html>

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