



Modeling of B-Box domain. Substitutions at P369S, and R408Q highlighted, neither are predicted to have a significant effect on protein conformation. See p 1386.

Editor

Tore K Kvien

Associate Editors

Francis Berenbaum
Hans Bijlsma
Dimitrios Boumpas
Gerd Burmester
Mary Crow
Iain McInnes

Editorial office

Annals of the Rheumatic Diseases
BMJ Publishing Group Ltd
BMA House
Tavistock Square
London WC1H 9JR, UK
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F: +44 (0)20 7383 6668
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