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Resolution improvement by modulation of analog-to-digital converters

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A program for template matching of protein sequences
ANTHEPROT: protein sequence analysis
A simple method for predicting the secondary structure of globular proteins: implications and accuracy
Discriminant analysis of promoter regions in E.coli sequences
A microcomputer program to evaluate the saturation of complex solutions with respect to biominerals
POEMS: program for outliers elimination in multidimensional space
An algorithm for comparing multiple RNA secondary structures
The use of various properties of amino acids in color and monochrome dot-matrix analyses for protein homologies

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