



# Results

**MOLECULAR MODELING & COMPUTATIONAL CHEMISTRY**

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Editorial and News

I am the new Editor for MMCC Results, the Molecular Modeling and Computational Chemistry News Letter. I took this project responsibility from Prof. David Busath starting from this year. I have worked two years as an Assistant Editor of MMCC Results under the previous Editor, Prof. David Busath. We have covered more than 25 journals every month and selected the more interesting articles for each issue of this newsletter.

I sincerely hope that all the scientists from academic and industrial background will give their full support to this project.

R.Nageswar, Editor  
RR Labs Inc.,  
8013 Los Sabalos Street  
San Diego, CA 92128



## MMCC Results

R.Nageswar, Editor  
8013 Los Sabalos Street  
San Diego, CA 92126  
Tel. (858) 663-0162

e-mail: [drnageswar@yahoo.com](mailto:drnageswar@yahoo.com)

R.Nageswar, Ph.D.  
RR Labs Inc., 8013 Los Sabalso St.  
San Diego, CA 92126

Editors Emeritus: Bruce Gelin, Ph.D.  
David Busath, M.D.

Dr. Gelin was founder of MMCC Results and edited volumes 1-6.  
Dr. David Busath edited volumes 7-14

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Assistant Editors:

Anston Feenstra  
Vrije Univ., Amsterdam,  
Netherlands

Naresh Aerra  
Rational Labs, Hyderabad., India

R.Mutyala  
RR Labs Inc., San Diego, CA.

## **List of the Journals covered in MMCC Results Newsletter**

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Editorial and News

## **1. APPLICATIONS**

### **Small Molecules**

**General and Model Systems**

**Water and Solvation**

**Organic Solvents**

**Medicinal Chemistry and Drug Design**

**Quantitative Structure-Activity Relations**

**Host-Guest Systems**

**Inclusion Compounds**

**Carbon Adsorbent**

**Zeolites**

**Carbon Nanoparticles**

**Supramolecular Chemistry**

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**Explosives**

**Crystal Growth**

**Liquid Crystals**

**Lubricants**

### **Biopolymers**

**Bioinformatics**

**Protein Sequence Analysis and Alignment**

**Protein Secondary Structure**

**Threading and Fold Recognition**

**Protein Structure Prediction**

**Comparative or Homology Modeling**

**Protein Side-Chain Conformation Prediction**

**Peptide Conformational Analysis**

**Protein Structure Analysis**

**Protein Engineering**

**Protein Hydration**

**Protein Electrostatics and Titration**

**Protein Dynamics**

**Free Energy Calculations**

**Ligand Binding**

**Enzyme Catalysis**

**Protein-Protein Interactions**

**Membrane Proteins and Lipid-Peptide Interactions**

**Proteins and Surfaces**

**Protein Design**

**Protein Folding**

**Protein-Nucleic Acid Interactions**

**Nucleic Acids**

**Lipids and Surfactants**

**Carbohydrates**

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## **Surfaces, Catalysts, and Materials Subjects**

### **2. METHODOLOGY**

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##### **Conformational Search and Analysis**

###### **Structure-Based Focussing**

###### **Potentials and Parameters**

###### **Solvation Energy**

##### **Global Energy Minimization**

###### **Normal Modes Analysis**

###### **Molecular Dynamics**

###### **Monte Carlo**

###### **Free Energy Perturbation**

###### **QM/MM**

##### **Secondary Structure Prediction**

#### **Comparative or Homology Modeling**

##### **Peptide Conformational Analysis**

##### **Side Chain Structure Prediction**

###### **Protein Structure Prediction**

###### **Threading or Fold Prediction**

#### **Surface and Volume Determination**

##### **Structural Similarity Analysis**

###### **Protein Folding**

**Ligand Docking**

**Structure Determination**

### **3. JOURNAL REVIEWS**

**Journal of Computational Chemistry 20(11), August, 1999**

**Journal of Computational Chemistry 20(12), September, 1999**

**Bioinformatics 15(5) May, 1999**

**Bioinformatics 15(7) July, 1999**

**Bioinformatics 15(8) August, 1999**

### **4. BOOK REVIEWS**

### **5. SOFTWARE/NETWARE REVIEWS**

### **6. EMPLOYMENT OPPORTUNITIES**

Ads for this section cost \$200 and run for three months unless withdrawn. There is no size limit, but each ad can fully describe only one specific position with additional positions described only in general terms and as a group. Alternative formats will be accommodated within reason. Email or mail your text to the editor by the third Saturday of the month prior to publication (*N.B.*: no issues in August and January).

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### **7. UPCOMING MEETINGS**

#### **8. ADDRESSES OF PRINCIPAL AUTHORS**

The production sites for the corresponding or principal authors are given in brackets in the citations. When not designated by the publisher, the first author is assumed to be the principal. Current addresses are listed here.

Temple F. Smith

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Editor:

R.Nageswar Ph.D.

MMCC Results

RR Labs Inc.,

8013 Los Sabalos Street

San Diego, CA 92126

Tel. (858) 663-0162

E-mail: [drnageswar@yahoo.com](mailto:drnageswar@yahoo.com)

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