

Applications of *SnB* to Proteins



Russ Miller & Charles M. Weeks

Hauptman-Woodward Med. Res. Inst.

Principal Contributors:

C.-S. Chang

G.T. DeTitta

S.M. Gallo

H.A. Hauptman

H.G. Khalak

D.A. Langs

R. Miller

S. Potter

C.M. Weeks

Partial funding from NIH and NSF.

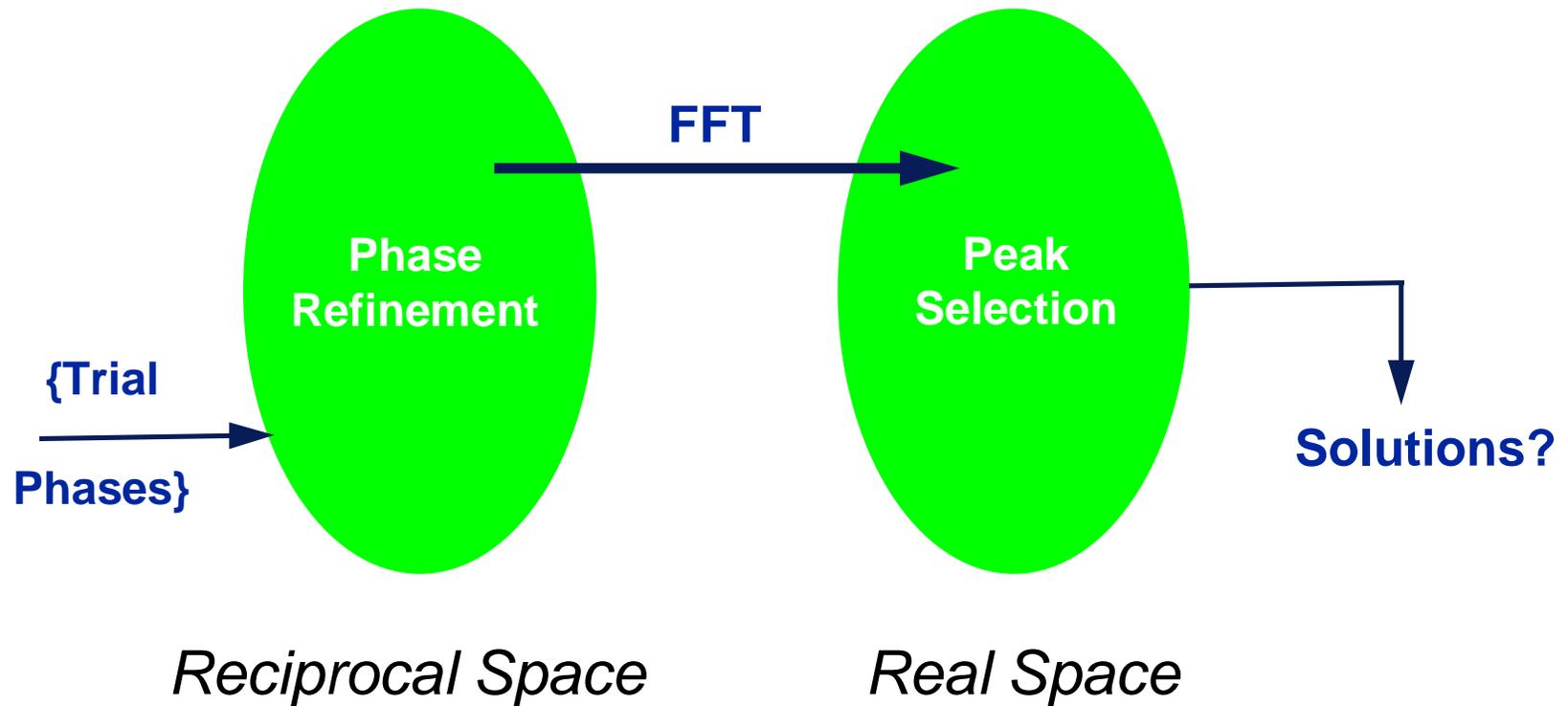
Outline of Talk

SnB

- ◆ *Shake-and-Bake: The Method*
 - *Cyclical Procedure*
 - *The Minimal Function*
- ◆ *SnB: The Program*
- ◆ *Applications of SnB*
- ◆ *SnB version 2.0*

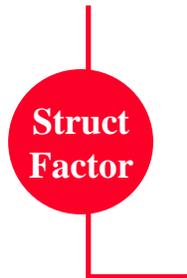
Conventional Direct Methods

SnB



Shake-and-Bake

{Trial
Structures}



FFT⁻¹



Shake-and-Bake

SnB

- ◆ **Direct Methods Optimization Technique**
- ◆ **Multiple Trial Structures**
- ◆ **Real Space \Leftrightarrow Reciprocal Space**
- ◆ **Phase Refinement Techniques**
 - **Parameter Shift (Minimal Function)**
 - **Tangent Formula**
- ◆ **Minimal Function as FOM**

The Minimal Function

SnB

$$R = \frac{\sum_T W_T (\cos \phi_T - est_T)^2}{\sum_T W_T}$$

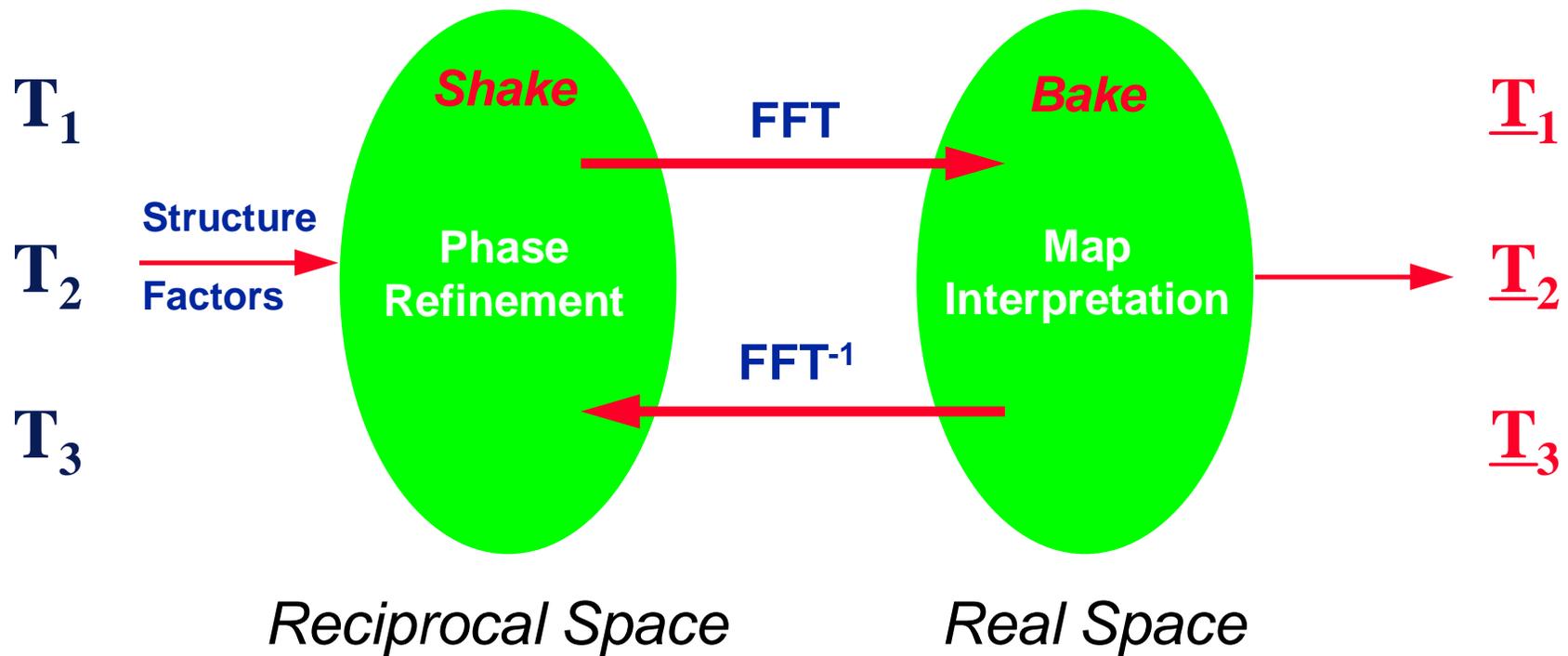
Triple: $\phi_T = \phi_h + \phi_k + \phi_{-h-k}$

$$W_T = \left(\frac{2}{N^{1/2}} \right) |E_h E_k E_{-h-k}|$$

est_T is the known expected value of $\cos \phi_T$

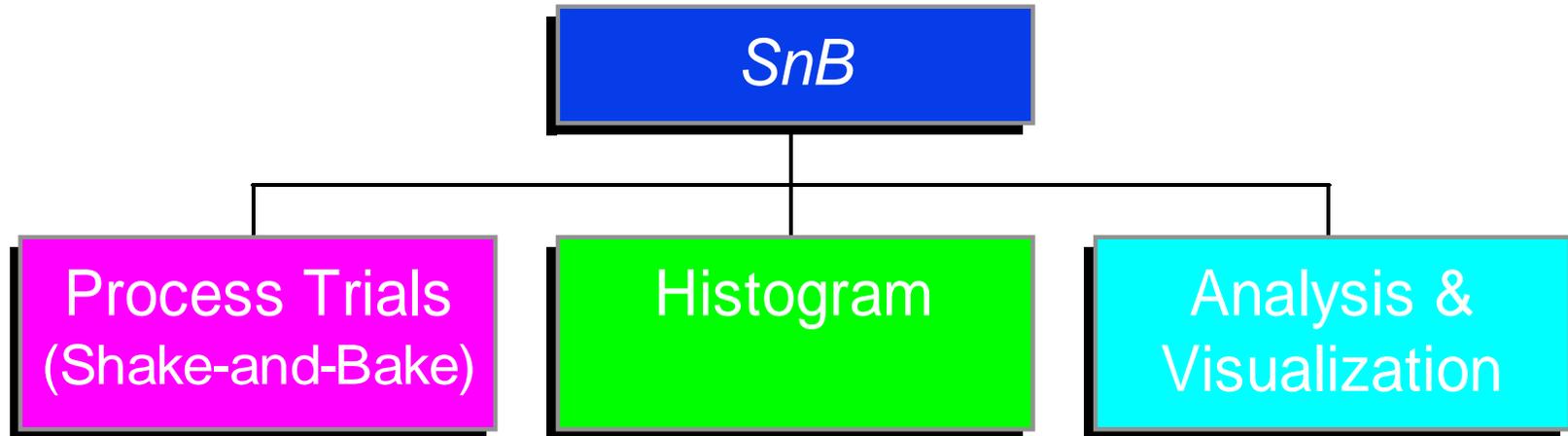
Shake-and-Bake

SnB



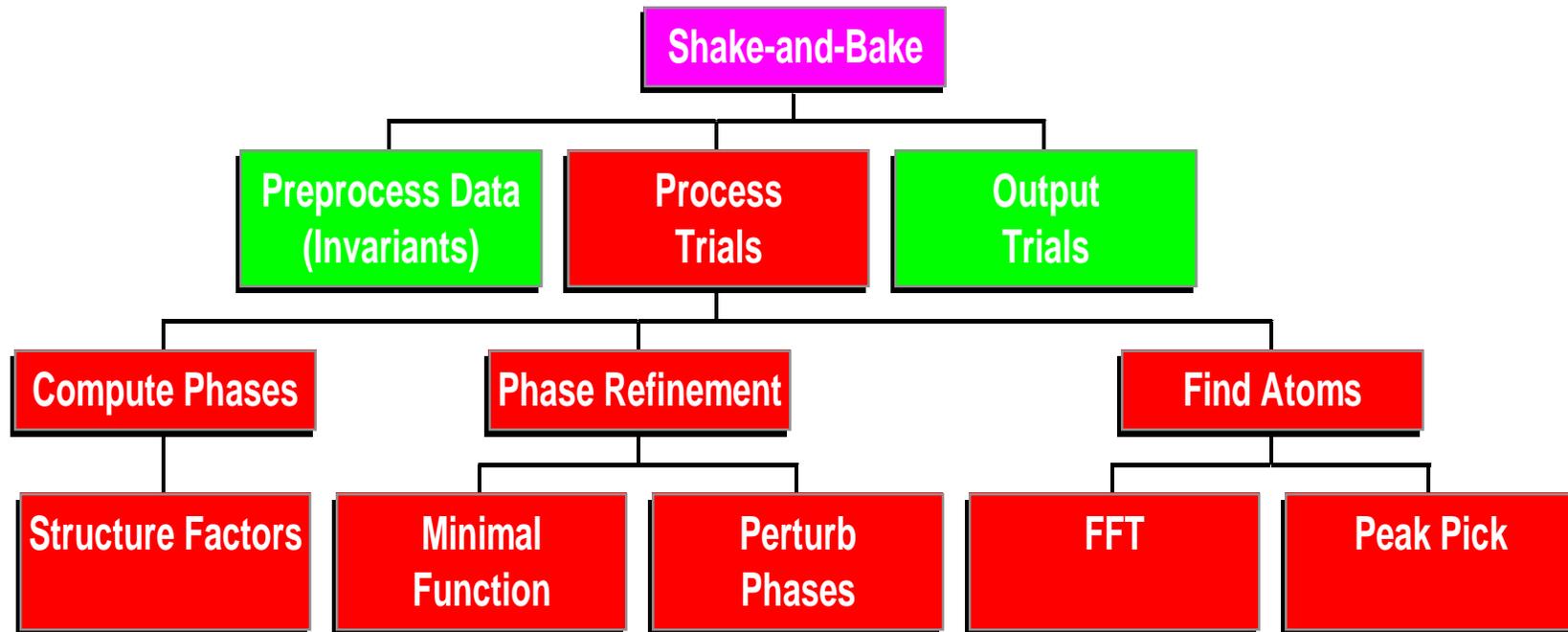
Structure of *SnB*

SnB



Shake-and-Bake

SnB



SnB Parameters

The logo for SnB, featuring the letters 'SnB' in a stylized, italicized font. The 'S' is red, the 'n' is grey, and the 'B' is red. The letters are slightly offset from each other, giving a 3D or layered appearance.

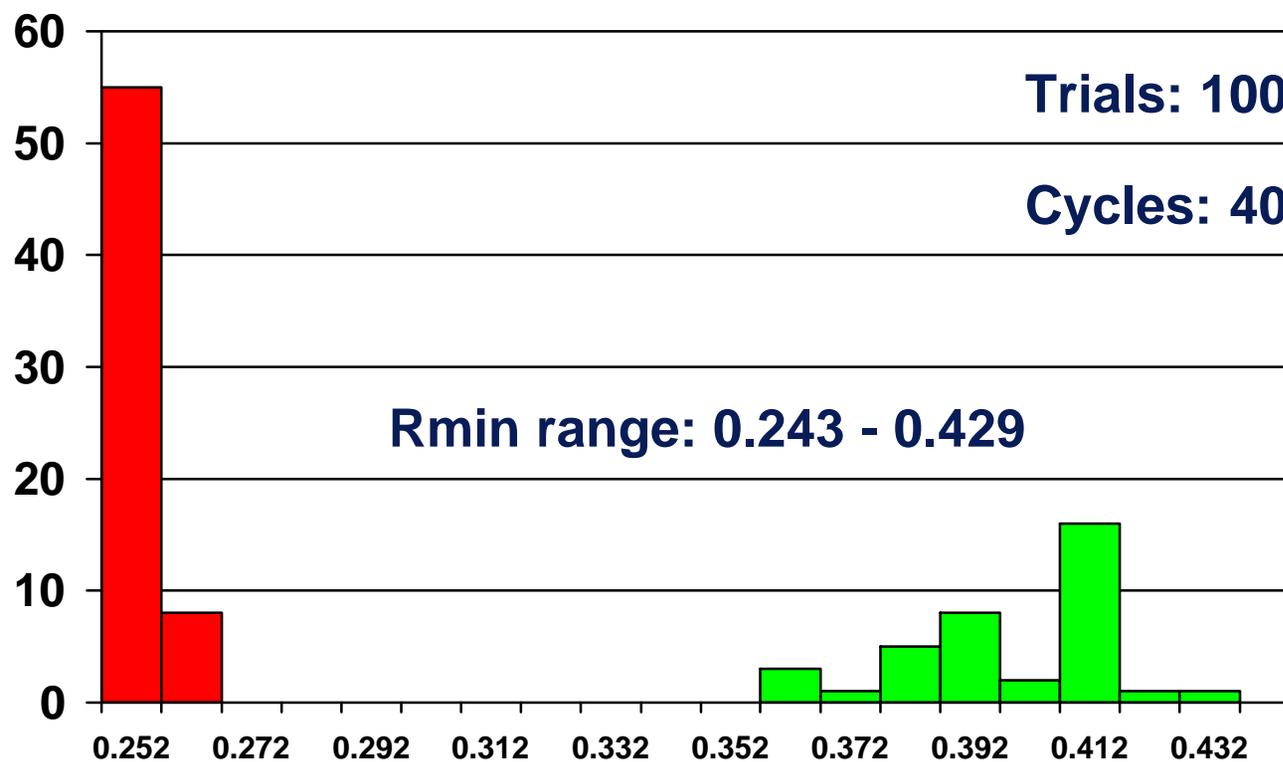
	Default	Ph8755	ToxII
Atoms (a.u.)	n	74	508
Phases	8n - 10n	740	5,000
Triples	70n - 100n	7,400	50,000
Cycles (PS)	n/2	40	255
Peaks recycled	0.8n - n	74	400
E-Fourier Steps	2	2	5

Sample Histogram (Ph8755)

SnB

Atoms: 74
Space Group: P1

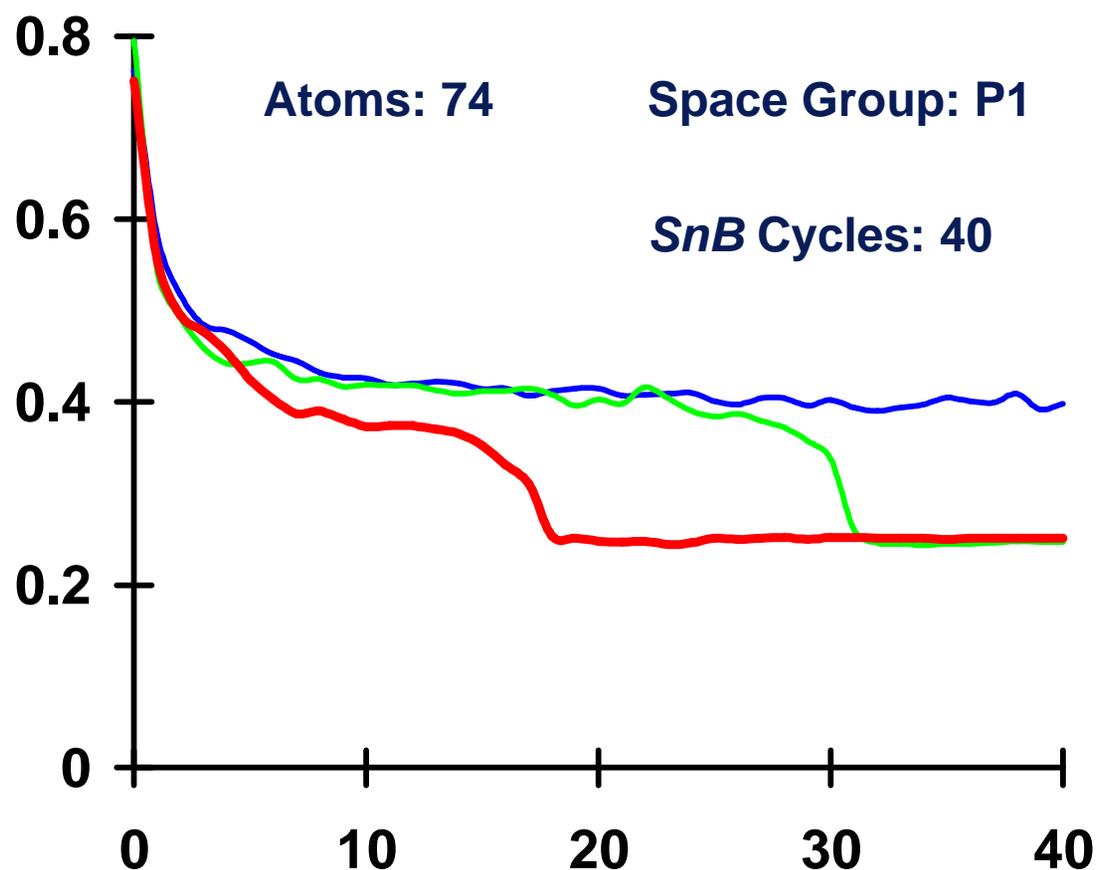
Phases: 740
Triples: 7,400



The Minimal Function is Diagnostic

Structure: Ph8755

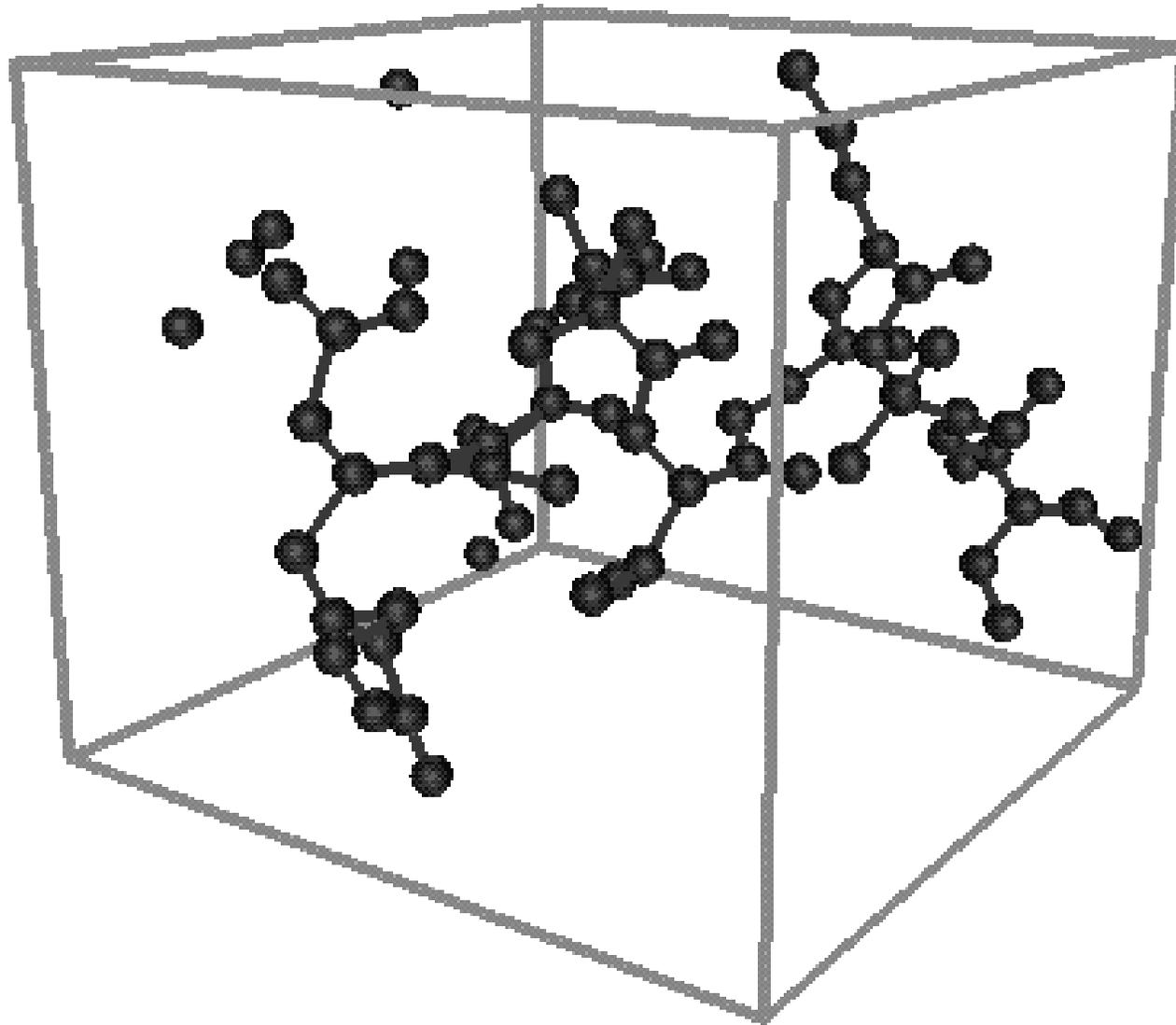
SnB



Visualization in *SnB* (Ph8755)

Geomview: Geometry Center, U. Minn.

SnB



ToxII: *SnB* Histogram

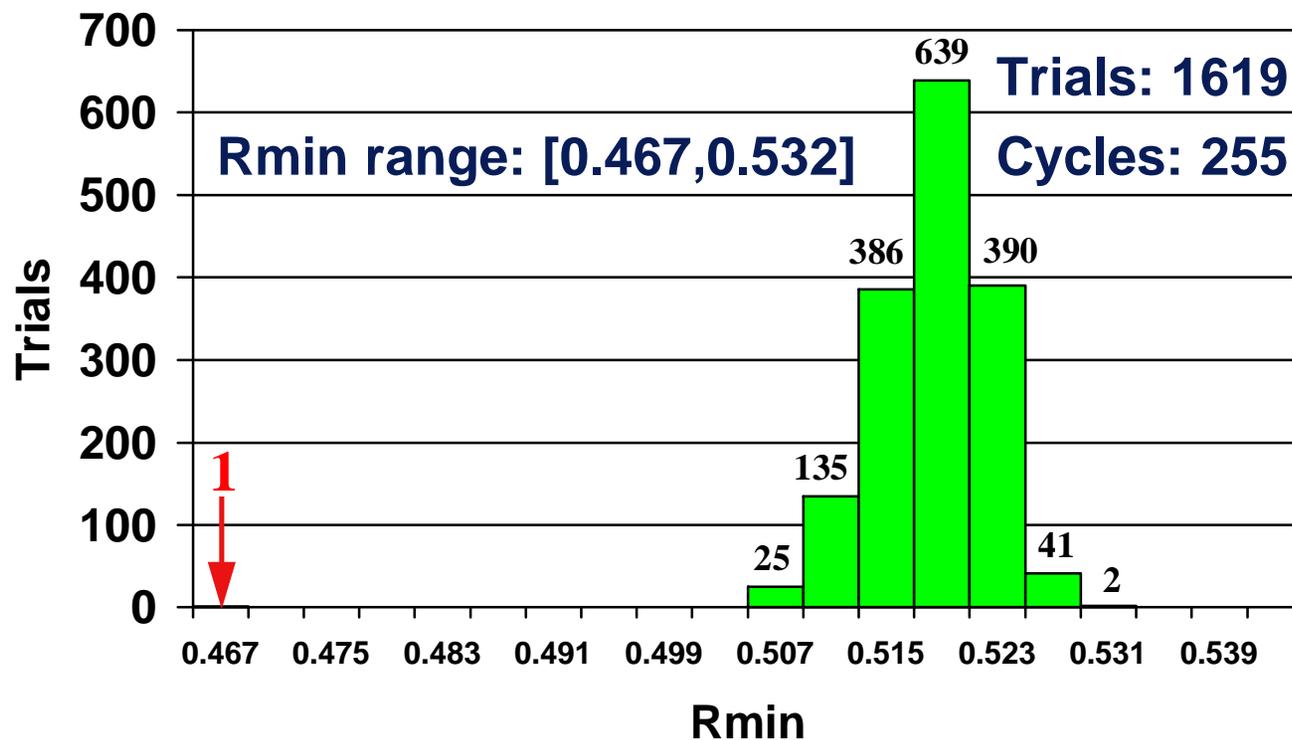
SnB

Atoms: 500

Phases: 5,000

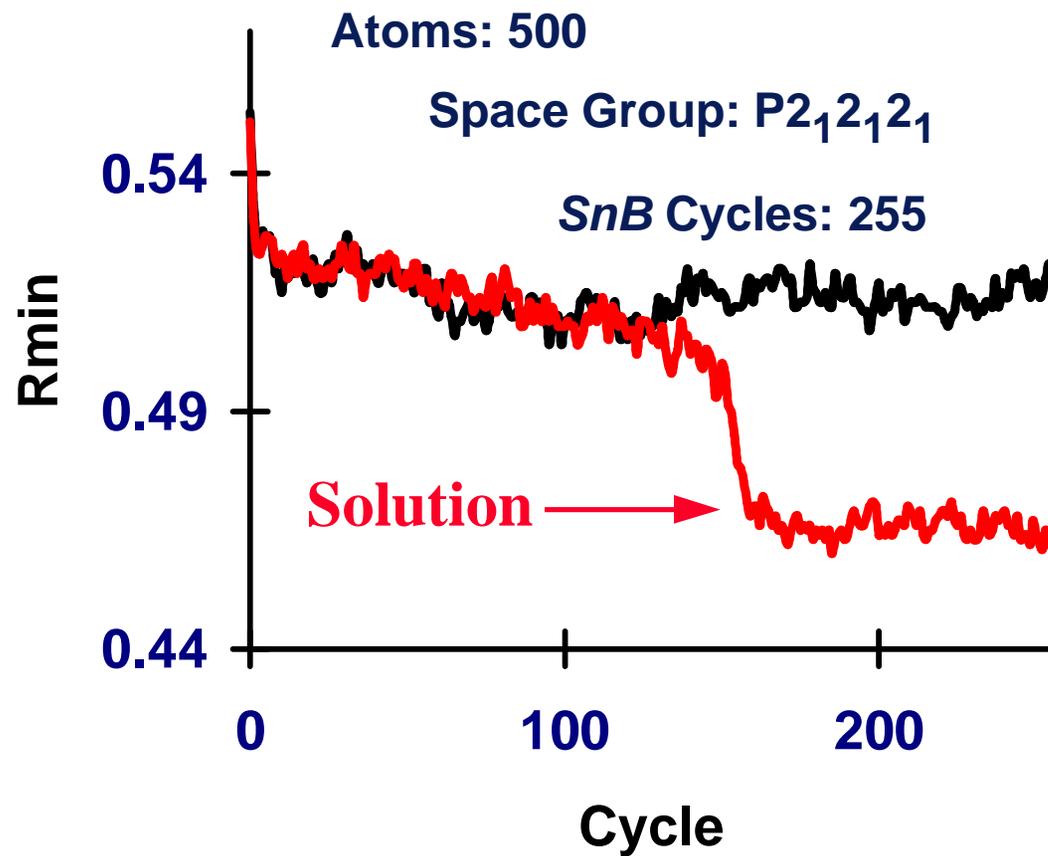
Space Group: $P2_12_12_1$

Triples: 50,000



Tox II: Trace of *SnB* Solution

SnB



Some *SnB* Solutions



STRUCTURE	LOCATION	ATOMS	SPACE GROUP
Scripps	Scripps	144	P1
Riboflavin tetrabutryate	MSC	180	P1
Vancomycin	Penn	255	P _{4,2,2}
Er-1 pheromone	UCLA	302	C2
Gramicidin A	HWI	317	P _{2,2,2} ₁
Crambin	HWI	400	P ₂ ₁
Alpha-1 peptide	OCI/U. of T.	450	P1
Rubredoxin	HWI	500	P ₂ ₁
ToxII	HWI	630	P _{2,2,2} ₁

Computing Platforms

SnB

◆ Unix Workstations

- ❑ SGI, Sun, DEC/Alpha
- ❑ IBM RS6000
- ❑ Wintel/Linux

◆ Parallel Computers

- ❑ Thinking Machines Corp. CM-5 (NCSA)
 - ❑ Cray T3D (PSC)
 - ❑ IBM SP2 (CTC)
- ## ◆ Cray C90 (PSC)

Looking Ahead: *SnB* vers. 2.0

- ◆ Includes Inverse FFT
- ◆ Additional Density Modification Options
- ◆ Improved Fourier Recycling: “Twice Baking”
- ◆ Input Improvements: $|E|$ Calculation
- ◆ Output Improvements: Compatibility
- ◆ (SIR/SAS/MAD Invariants with est. values)

PS02.03.16: Development of *SnB* Version 2

Related Presentations

SnB

- ◆ **MS01.03.05:** Direct Methods Phasing of a 450 Atom Structure (Prive and Eisenberg)
- ◆ **MS02.06.07:** The *Ab Initio* Structure and Refinement of a Scorpion Protein Toxin (Smith, et. al.)
- ◆ **NO.BL.01:** The SAS Maximal Principle (Hauptman)
- ◆ **PS02.03.16:** Development of *SnB* Version 2 (Weeks and Miller)
- ◆ **PS02.06.09:** Application of the SAS Tangent Formula to Multiple Site Problems (Chang, et. al.)
- ◆ **MS02.01.05:** The Application of Shake and Bake to Unsolved Structures (Haltiwanger and Eggleston)

Summary

SnB

- ◆ *Shake-and-Bake*: Enhanced Direct Methods
- ◆ Targeted at 100-800 atom structures
- ◆ Publicly available
- ◆ Recommend: LEVY / EVAL (Bob Blessing)
- ◆ Current Research: *SnB* version 2.0
- ◆ Addresses:
 - <http://www.hwi.buffalo.edu/SnB/>
 - snb-requests@hwi.buffalo.edu
 - snb-comments@hwi.buffalo.edu

SnB

