

# **Molecular Modeling Used as a Probe of Interactions to Study the Polymeric Glass Transition**

**Armand Soldera**

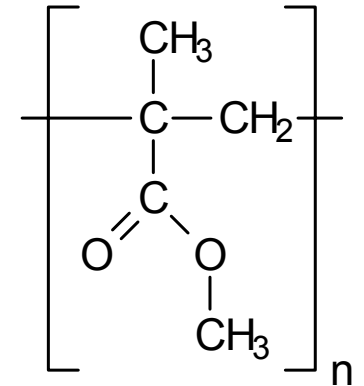
**Département de Chimie**



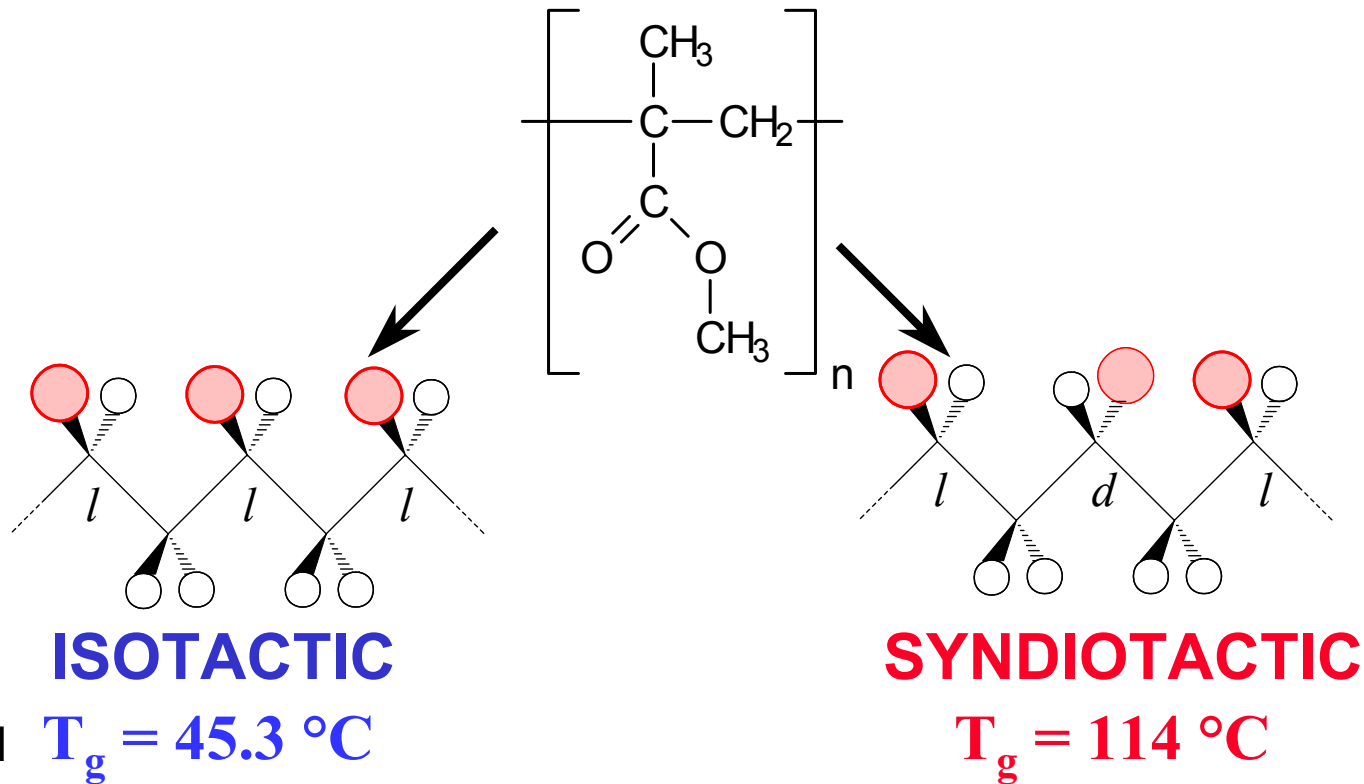
**HPCS May 13, 2003**

# Contents

- ✓ **Tacticity**
- ✓ **Simulation of the Amorphous Phase**
- ✓ **Dilatometric Simulation**
- ✓ **Energetic Analysis**
- ✓ **Local Dynamics**
- ✓ **Cooperativity**
- ✓ **Conclusions**



# Tacticity



Can we manage such a difference by the use of molecular modeling ?

If affirmative → better understanding of the difference → glass transition ...

# Simulation of the Amorphous Phase

## ✓ Design of a cubic box

- From the knowledge of the density and the mass of the polymer
- All the space is filled by replica of this box

## ✓ Chain design

- A propagation procedure (MC) is begun to design 1 polymer configuration
- The chain backbone is grown step by step looking for long range excluded volume

$$q'_{\xi\eta;i} = q_{\xi\eta;i} \frac{\exp\left[-\Delta U_{\eta;i}^{LR}/RT\right]}{\sum_{\eta'} q_{\xi\eta';i} \exp\left[-\Delta U_{\eta';i}^{LR}/RT\right]}$$

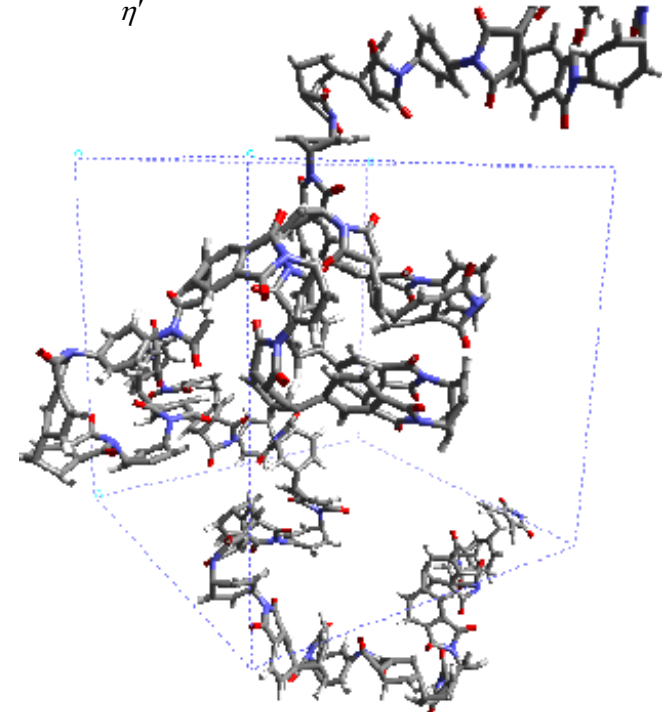
RIS

## ✓ Periodic Boundary Conditions

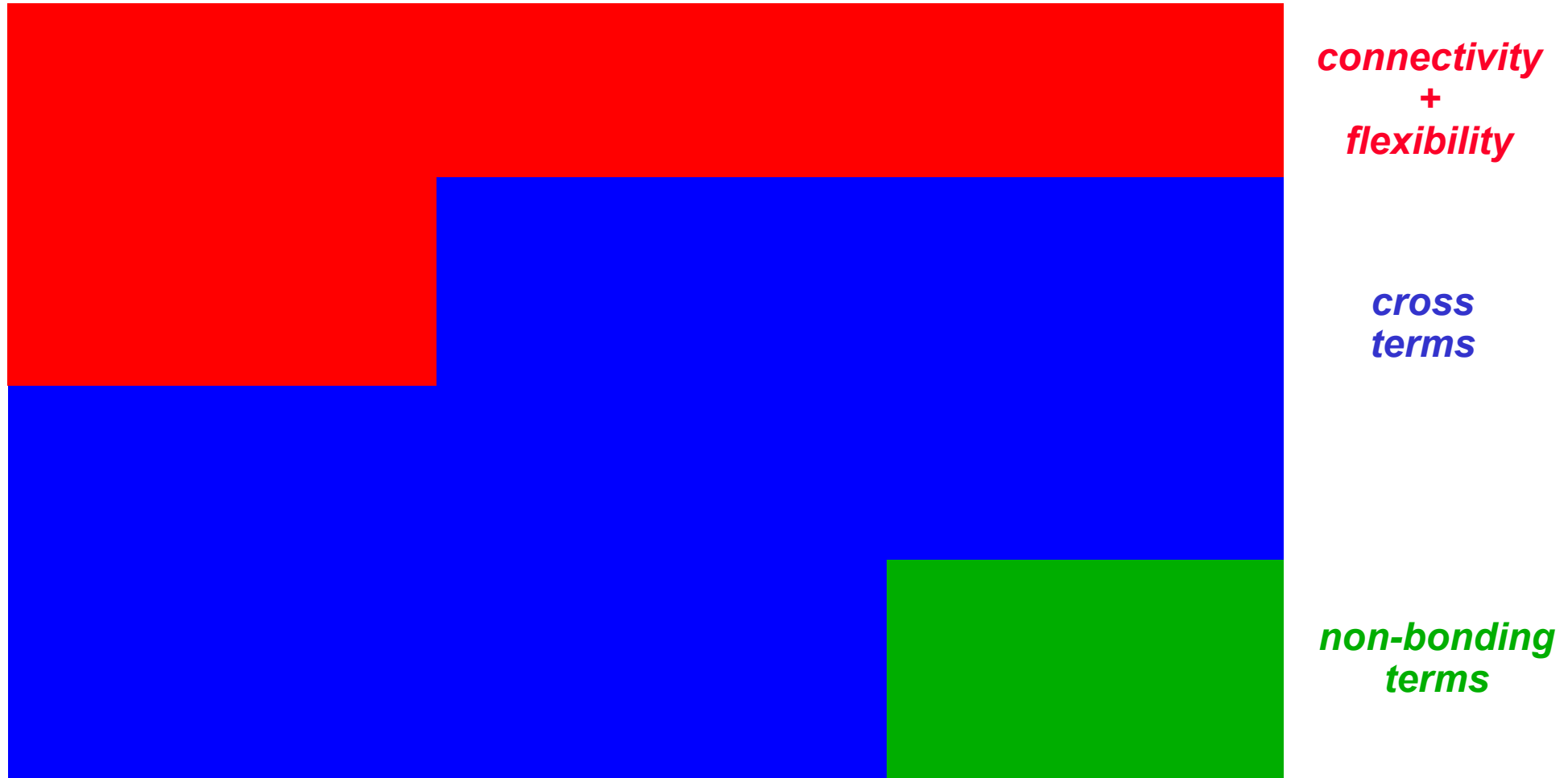
Each atom coming out from one face is automatically entering through the opposite face

## ✓ Relaxation Procedure

MD + minimization



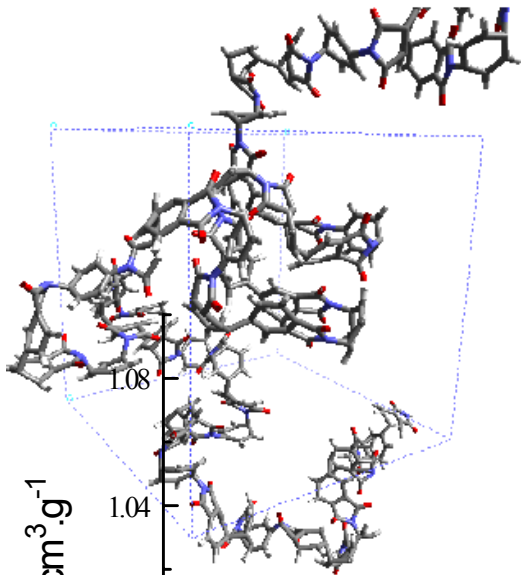
# *pcff* Force Field: schematic representation



# pcff Force Field: mathematical expression

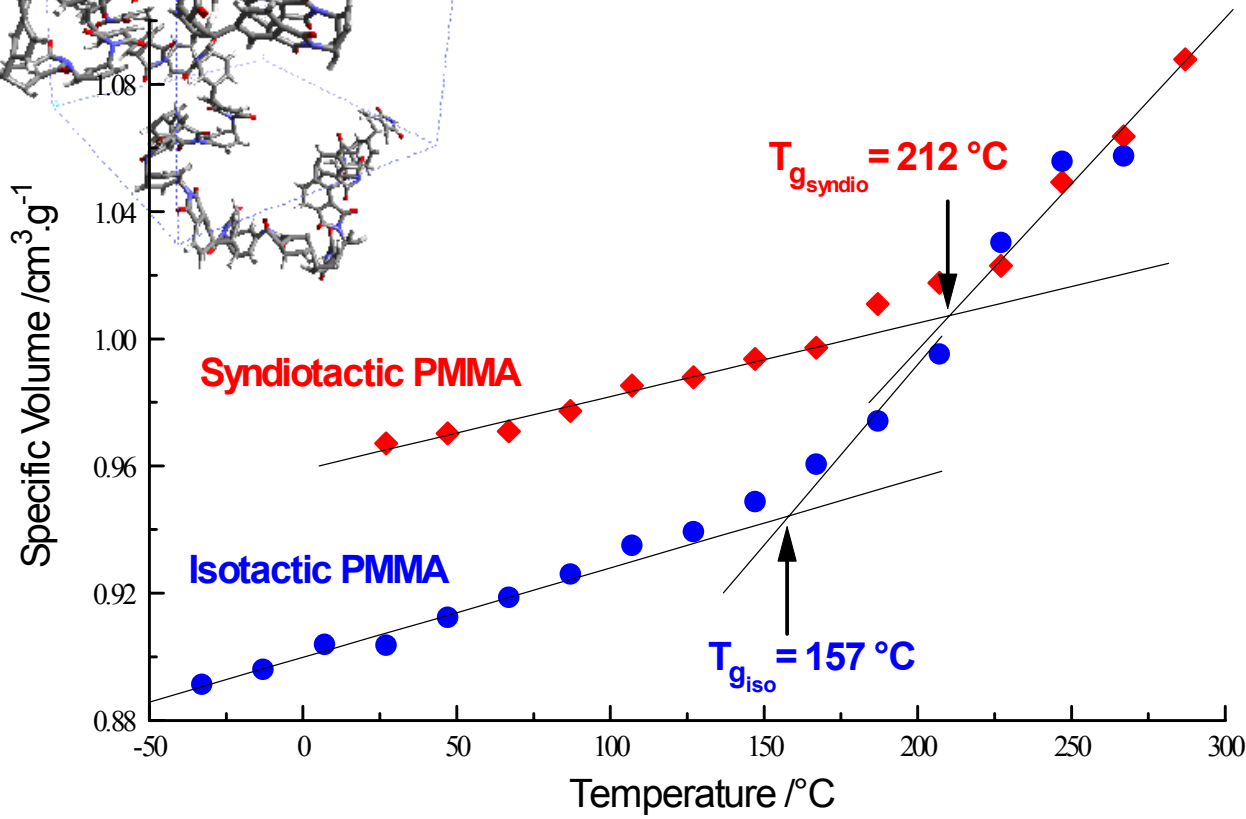
$$\begin{aligned}
 V = & \sum_b \left[ K_2 (b-b_0)^2 + K_3 (b-b_0)^3 + K_4 (b-b_0)^4 \right] + \sum_\theta \left[ H_2 (\theta-\theta_0)^2 + H_3 (\theta-\theta_0)^3 + H_4 (\theta-\theta_0)^4 \right] \\
 & + \sum_\phi \left[ V_1 \left[ 1 - \cos(\phi - \phi_1^0) \right] + V_2 \left[ 1 - \cos(2\phi - \phi_2^0) \right] + V_3 \left[ 1 - \cos(3\phi - \phi_3^0) \right] \right] + \sum_\chi K_\chi \chi^2 \\
 & + \sum_b \sum_{b'} F_{bb'} (b-b_0)(b'-b'_0) + \sum_\theta \sum_{\theta'} F_{\theta\theta'} (\theta-\theta_0)(\theta'-\theta'_0) + \sum_b \sum_\theta F_{b\theta} (b-b_0)(\theta-\theta_0) \\
 & + \sum_b \sum_\phi (b-b_0) [V_1 \cos \phi + V_2 \cos 2\phi + V_3 \cos 3\phi] + \sum_{b'} \sum_\phi (b'-b'_0) [V_1 \cos \phi + V_2 \cos 2\phi + V_3 \cos 3\phi] \\
 & + \sum_\theta \sum_\phi (\theta-\theta_0) [V_1 \cos \phi + V_2 \cos 2\phi + V_3 \cos 3\phi] + \sum_\phi \sum_\theta \sum_{\theta'} K_{\phi\theta\theta'} \cos \phi (\theta-\theta_0)(\theta'-\theta'_0) \\
 & + \sum_{i>j} \frac{q_i q_j}{\epsilon r_{ij}} + \sum_{i>j} \left[ \frac{A_{ij}}{r_{ij}^9} - \frac{B_{ij}}{r_{ij}^6} \right]
 \end{aligned}$$

# Dilatometric Simulation

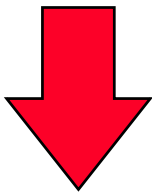


NPT ensemble  
Number of RU: 100  
Simulation time: 110 ps by data  
force field: *pcff*

12 hours  
in SGI O2000 /  
data



**Simulated  $\Delta T_g = 55 \text{ }^\circ\text{C}$**   
**Expected  $\Delta T_g = 69 \text{ }^\circ\text{C}$**



**Investigations to understand  
such a difference  
can be carried out**

# Energetic Analysis

## ✓ Principles

- The 2 PMMA configurations have the same force field parameters
- Changes in their molecular behavior will be directly linked to changes in their molecular characteristics



Energy differences

## ✓ Total Energy

$$E(\text{Iso}) - E(\text{Syndio}) = 10 \text{ kcal.mol}^{-1}$$



3 splits will be performed

1. Inter and intramolecular contributions
2. Inside the intramolecular part
3. Molecular contribution

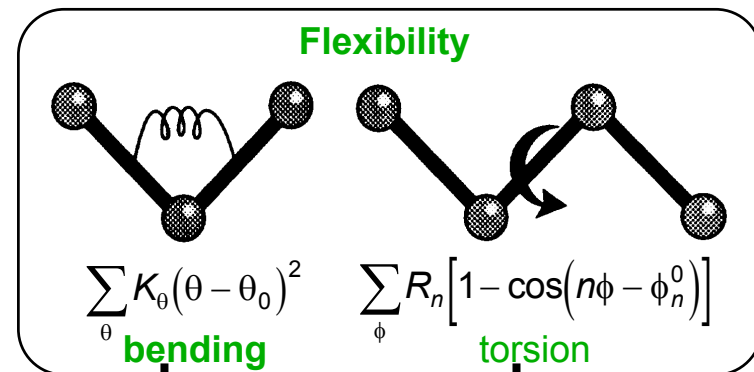
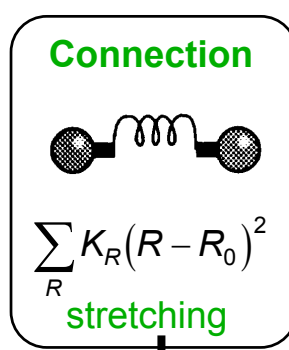
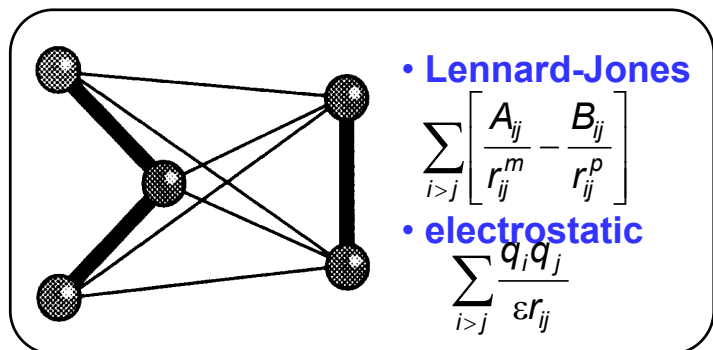


# Splits in the Energy Contributions

1) Total energy Intermolecular & Intramolecular

**-35** ( $\pm 8$ )

**45** ( $\pm 8$ )



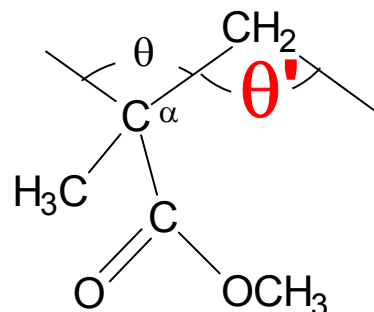
2) Intramolecular Energy

**15** ( $\pm 7$ )

**75** ( $\pm 10$ )

**15** ( $\pm 5$ )

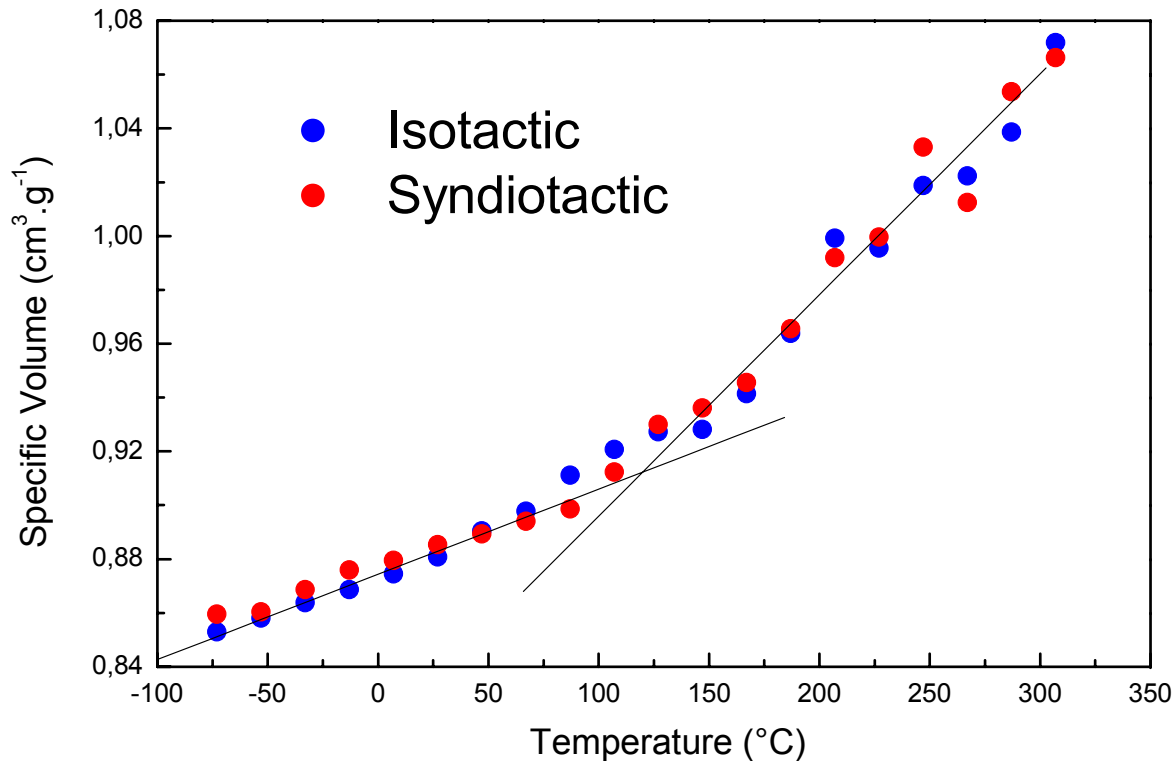
3) Bending Energy



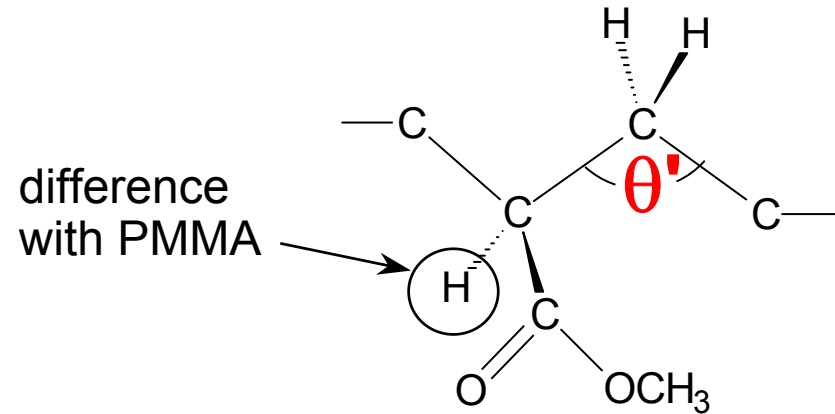
**Syndiotactic: 126.7°** ( $\pm 0.1$ )

**Isotactic: 127.8°** ( $\pm 0.1$ )

# T<sub>g</sub> Determination of PMA



$$T_g(PMA) < T_g(I - PMMA) < T_g(S - PMMA)$$



**No differences in T<sub>g</sub>s between the 2 PMA configurations, in agreement with experimental data**

# Energetic Analysis

## - Comparisons to PMA data -

### ✓ Intermolecular energy differences

$$E_{\text{inter}}(S - PMMA) > E_{\text{inter}}(I - PMMA) \gg E_{\text{inter}}(PMA)$$

$350 \text{ kcal. mol}^{-1} \qquad 258 \text{ kcal. mol}^{-1} \qquad 78 \text{ kcal. mol}^{-1}$

### ✓ Intramolecular energy differences

In the bending term associated with the intra-diad angle,  $\theta'$

$$\theta'(I - PMMA) > \theta'(S - PMMA) \gg \theta'(PMA)$$

$127.8^\circ \qquad 126.7^\circ \qquad 118.0^\circ$

### ✓ Conclusions

- Results are in agreement with the Free Volume Theory  
Higher interactions between neighboring polymer chains segments will give a higher  $T_g$
- Due to a greater aperture of  $\theta'$ , the isotactic chains should be more mobile



Study of the local dynamics

# Local Dynamics Analysis

## - Principles -

### ✓ Computation of the orientation function $P_2$

- From MD, acquisition of the bond autocorrelation function  $\langle (\mathbf{u}(t) \cdot \mathbf{u}(0)) \rangle$
- Computation of the 2nd Legendre polynomial term with respect to time,  $P_2(t)$

### ✓ Computation of the correlation time, $\tau_c$

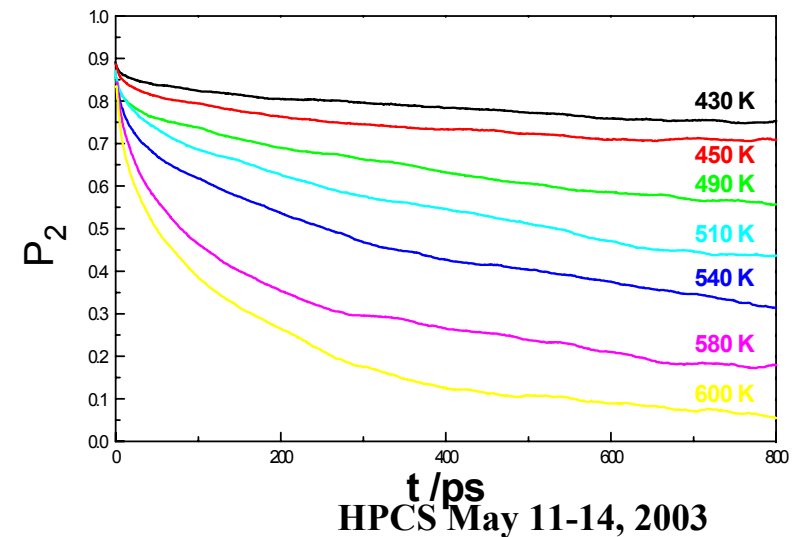
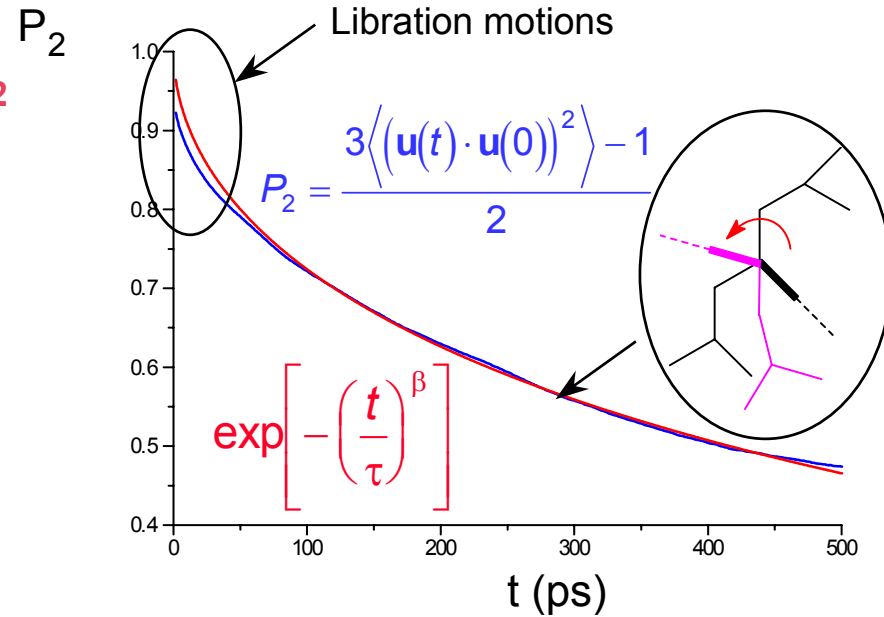
- Fit of  $P_2(t)$  with a stretching exponential, KWW

$$\tau_c = \int_0^{\infty} P_2(t) dt \quad \rightarrow \quad \tau_c = \frac{\tau}{\beta} \Gamma\left(\frac{1}{\beta}\right)$$

### ✓ Procedure is carried out at different temperatures

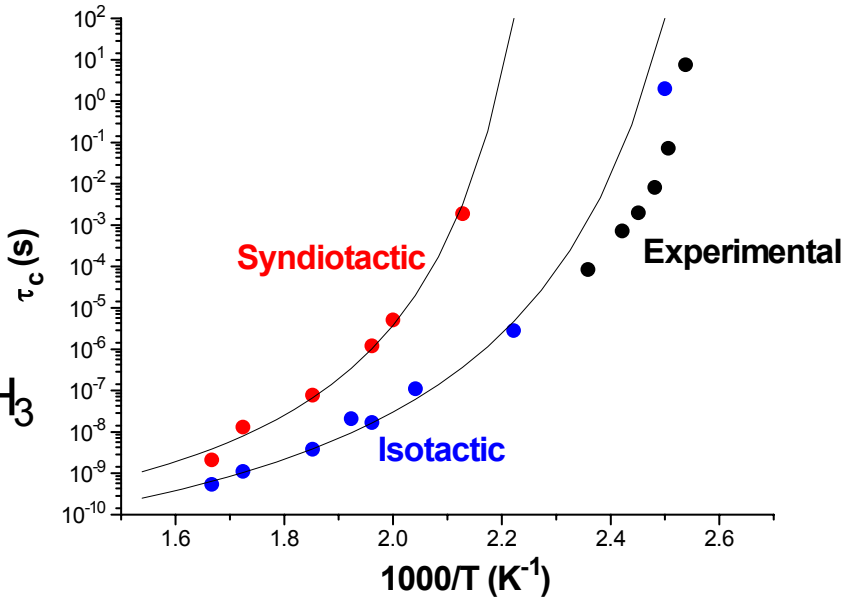
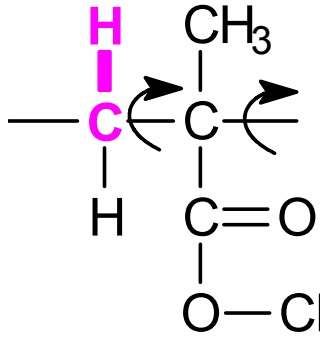
### ✓ Fit with a VFT equation (or WLF)

$$\tau(T) = A \exp\left(\frac{B}{T - T_0}\right)$$



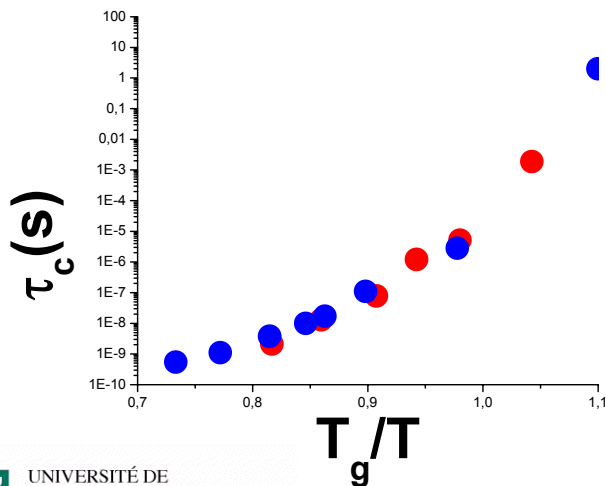
# Local Dynamics of the Backbone

✓ **Fit**



<b>B</b> (kJ.mol <sup>-1</sup> )
<b>11.9</b>
<b>12.8</b>

✓ **Results**



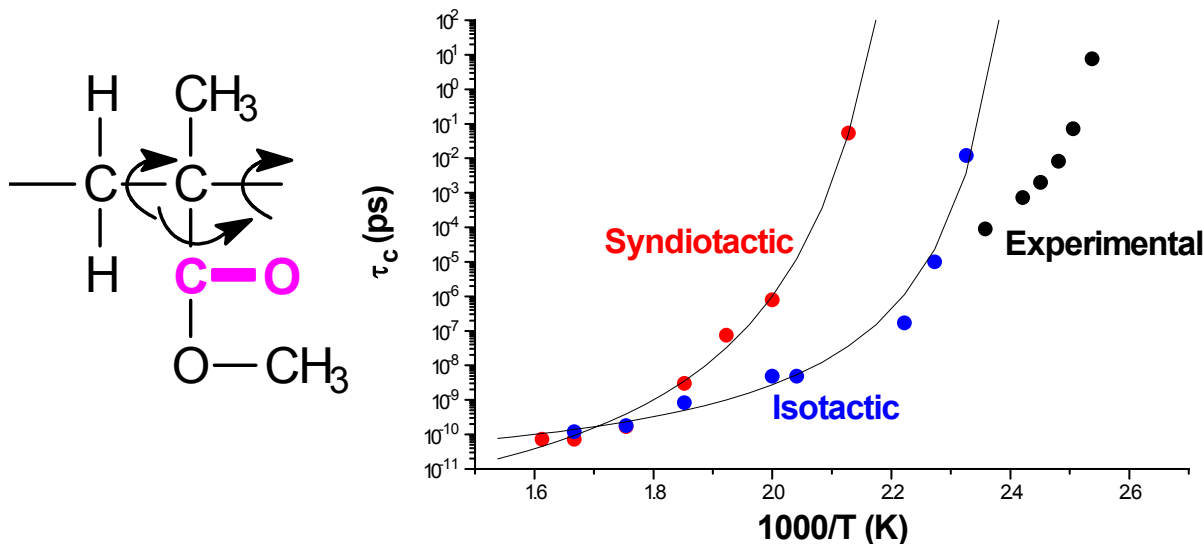
Behavior of the 2 isomers

- at  $T+T_g$ : **Comparable**
- at  $T$ : **Different**

**Study of the relaxation of the side chain**

# Local Dynamics of the Side-Chain

## ✓ Fit



<b>B</b> (kJ.mol <sup>-1</sup> )
<b>11.5</b>
<b>5</b>

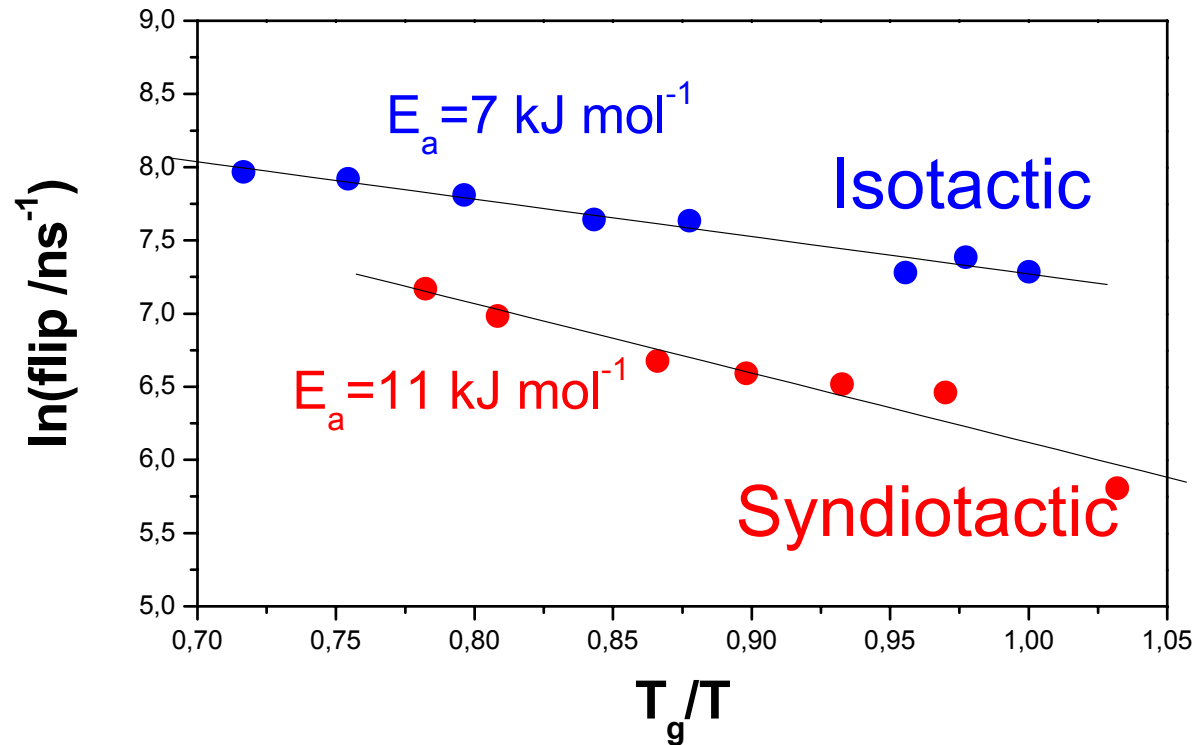
## ✓ Results

- Non-Arrhenian behavior, but such a relaxation corresponds to the  $\beta$  mode
- BUT, the simulation takes into account 3 motions:
  - » librational modes
  - » due to the side-chain (what we are interested in)
  - » due to the backbone

# Mobility of the side-chain

## ✓ Number of transitions of the side-chain

Computation of the number of transitions between the UP and DOWN states of C=O



The behavior is Arrhenian like !

# Compilation of the Results

## ✓ Correlation times

### - Backbone

- » Behavior of  $\tau_c(\text{C-H})$  is in agreement with published results:  
correlation times of iso PMMA are found inferior to the syndio PMMA ones
- » The backbones of the 2 configurations present the same behavior at  $T + T_g$ , therefore the difference in  $T_g$ s could not be explained

### - Side-Chain

- » The side-chains of Iso-PMMA show a greater mobility than the syndio ones
- » Behind this difference there lies a possible explanation of the difference in  $T_g$ s

## ✓ Comparison with experimental data

- From NMR experiments: compared with PEMA, PMMA showed that the greatest mobility of the side-chains induces a decrease of the lowest correlation time of the backbone
- Consequently, a higher side-chain rotation of iso PMMA generates a greater mobility of the backbone, and a greater mobility of the backbone explains a lower  $T_g$



$$T_g (\text{i-PMMA}) < T_g (\text{s-PMMA})$$



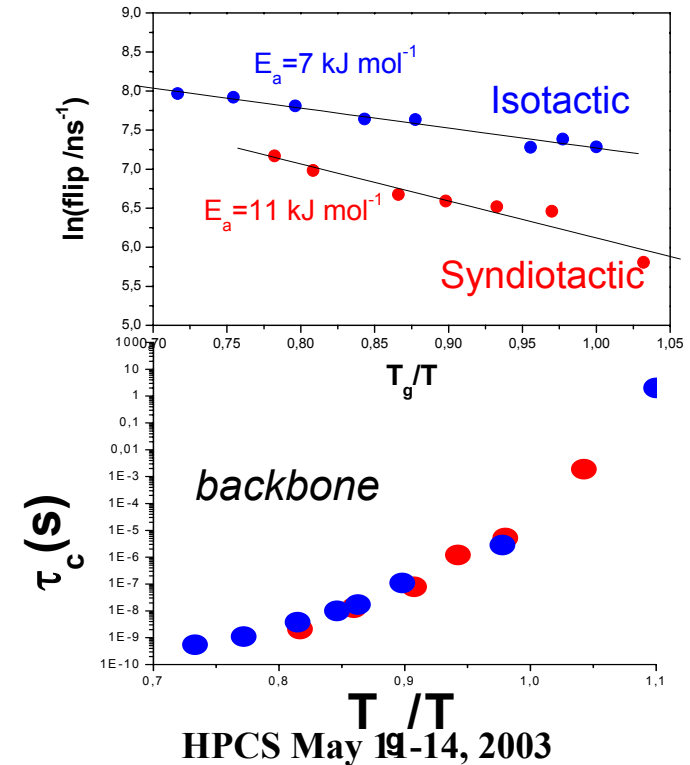
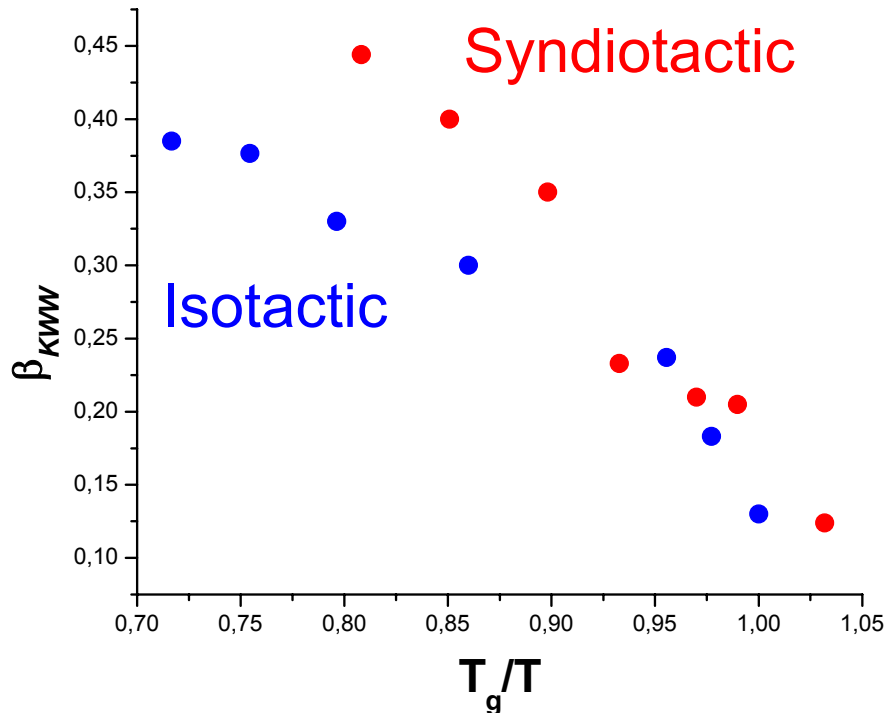
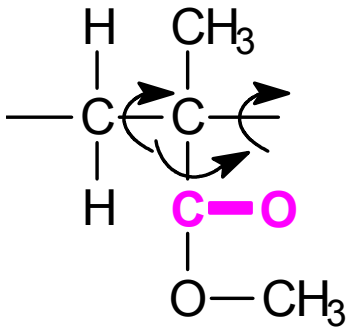
# Cooperativity ...

## ✓ Cooperativity observed by molecular simulation


The stretching function:  $\exp\left[-\left(\frac{t}{\tau}\right)^\beta\right]$

According to the Coupling Model:  $\beta = n - 1$  where  $n$  is the coupling parameter, it actually corresponds to a measure of the cooperativity

Consequently, the coupling between the side-chain and the backbone can be directly observed:



# Conclusions

- ✓ The **substitution of H by CH<sub>3</sub>** causes energetic variations
  - Augmentation of intermolecular interactions
  - Differentiation in the non-bonding interactions between the 2 stereomers
  - Aperture of the intra-diad angle to lessen side-chain interaction
  - Aperture of the intra-diad angle more important for the isotactic configuration
- ✓ The important mobility of the isotactic side-chain induces a greater mobility of the backbone, comparatively to the syndiotactic one: This cooperativity between the side-chain and the backbone was observed using molecular simulation
  -  Such a behavior tends to lower the  $T_g$  of the isotactic configuration
- ✓ **The comparison of simulated data to experimental results are in agreement with the free volume concept to explain the difference in  $T_g$  between the two stereomers**

# Questions

- ✓ **Is the local scale cooperativity observed here consistent with the CRR (Coupling Rearranging Region) theory ?**
- ✓ **Can the size or the shape of the CRR be affected by the PMMA tacticity ?**
- ✓ **The tacticity dependent size or anisotropic shape of the CRR may be a crucial point revealed by chain confinement:  $T_g$  (tacticity and film thickness)**
- ✓ **Further investigations will be performed on this topic in light of our molecular dynamics results.**

# Acknowledgement



**Pr. Yves Grohens**  
Université de Bretagne-Sud



UNIVERSITÉ DE  
**SHERBROOKE**



**InsightII:** Amorphous\_Cell and Discover3

**DL\_POLY**

