## Utility for Calculating Displacement Correlations for RMCProfile Configurations

Routine AtomCorr.exe calculates atomic correlation parameters for the .cfg files used by RMCProfile. These parameters are calculated as

$$C(\mathbf{n}_1, \mathbf{n}_2, \mathbf{R}) = \frac{\langle (\mathbf{u}_i \mathbf{n}_1) (\mathbf{u}_j \mathbf{n}_2) \rangle}{\sqrt{\langle (\mathbf{u}_i \mathbf{n}_1)^2 \rangle \langle (\mathbf{u}_j \mathbf{n}_2)^2 \rangle}}$$

where  $\mathbf{u}_i$  is the displacement of the *i*-th atom in the refined configuration relative to the position of this atom in the initial configuration (average crystal lattice),  $\mathbf{n}_1$  and  $\mathbf{n}_2$  are the unit vectors along the directions of interest, and  $\langle \rangle$  denote the average over configuration.

A set of atoms *i* is selected by specifying a range of their sequential numbers (first and last) in the \*.cfg file, whereas a set of atoms *j* is selected by specifying (1) a range of their sequential numbers (first and last) in the .cfg file and (2) a volume delimited by an intersection of a spherical layer around atoms *i* (radii  $r_{min}$  and  $r_{max}$ ) and a cone (axis **R** and angle  $\theta$ ) having its vertex at atoms *i*.

To use this utility, the \*.cfg files containing the starting and the refined configurations must be in the same directory with the **AtomCorr.exe** file. The input for **AtomCorr** can be entered either manually, following the on-screen instructions, or using a text file having an extension "**.cor**". This input file has the following structure:

Initial	! .cfg filename (no extension) for an initial configuration
Final	! .cfg filename (no extension) for a final configuration
1 4096	! range of atoms <i>i</i> in the .cfg file
4097 12288	! range of atoms <i>i</i> in the .cfg file
3.5 4.5	! <i>rmin</i> and <i>rmax</i> in Å
112	! component of vector <b>R</b> in Å
20	! Angle $\theta$ (degrees)
101	! components of vector $\mathbf{n}_1$
1 0 -1	! components of vector $\mathbf{n}_2$

Vectors  $\mathbf{R}$ ,  $\mathbf{n}_1$ , and  $\mathbf{n}_2$  are normalized by the program and, therefore, only relative values of their components are important. The program generates two output files: finalcomp.txt and final\_correl.txt.

The **finalcomp.txt** file contains the coordinates of all atoms in the **initial.cfg** file (RMC units) plus three components of the corresponding atomic displacements relative to the initial positions (Å) (the results are arranged in a six-column table). The file final\_correl.txt file contains calculated values of the correlation parameters and the *rms* displacements of atoms  $i (\sqrt{\langle (\mathbf{u}_i \mathbf{n}_1)^2 \rangle})$  and  $j (\sqrt{\langle (\mathbf{u}_i \mathbf{n}_1)^2 \rangle})$ .