This project has been done with utmost attention given to make use of the tool as easy as possible for the users with ample instructions and examples for further clarity before each input instance. Here are some more details in depth.

The default questionnaire is scid-5-cv. (Please add scid\_baseline.exe to the same directory as the scid\_generalisation.cpp in order to run the default.) Starting from questions to outputs and how to evaluate the outputs everything can be user inputted using this tool.

First the user has to enter space separated number of nodes and edges. Each node represented a class of questions like that of sleep or appetite and the edges represented which node to visit next.

Next the user has to input the set of vertices in pairs among whom there exists an edge. For example: If node 1 has connection to 2 and 3 and 2 has connection to 3, then the input can be -  
“1 2  
 1 3  
 2 3”

String responses are accepted although currently the tool works with 0 or 1 responses to the questions inputted by the clinician. The remaining input instances have been clearly explained in the code itself.