

/mnt/c/course\_3/CircuitGen  
\_Generator/src/generators  
\_Genetic/GenGenerator.h

```
graph BT; A["/mnt/c/course_3/CircuitGen_Generator/src/database_DataBaseGenerator.cpp"] --> C["/mnt/c/course_3/CircuitGen_Generator/src/generators_Genetic/GenGenerator.h"]; B["/mnt/c/course_3/CircuitGen_Generator/src/generators_Genetic/GenGenerator.cpp"] --> C;
```

This diagram illustrates the relationships between source files and a header file in a project. At the top is a gray box representing the header file: `/mnt/c/course_3/CircuitGen_Generator/src/generators_Genetic/GenGenerator.h`. Below it are two white boxes representing source files. The left box is `/mnt/c/course_3/CircuitGen_Generator/src/database_DataBaseGenerator.cpp` and the right box is `/mnt/c/course_3/CircuitGen_Generator/src/generators_Genetic/GenGenerator.cpp`. Blue arrows point from each source file box up to the header file box, indicating that both source files include or depend on this header file.

/mnt/c/course\_3/CircuitGen  
\_Generator/src/database  
\_DataBaseGenerator.cpp

/mnt/c/course\_3/CircuitGen  
\_Generator/src/generators  
\_Genetic/GenGenerator.cpp