N explorers are being read from a file (*explorers.dat*) each having a name, country of origin, number of discoveries and their relative distance from the *Holy Grail*. (See example of input)

The Holy Grail is a unique artefact which has a country of origin and an age, both read from a file (*holyGrail.dat)*.

- Read all the explorers from the input file
- Sort them based on the number of discoveries and print out (in file *discoveriesExplorers.dat*) the first ***m*** (is read from the keyboard at this step) explorers based on this *number of discoveries* field. If **m > n** simply print them all out. (first explorer printed → most discoveries)
- Read the details of the Holy Grail artefact from the file and re-sort the explorers based on their distance to the Holy Grail (first in array is closest to the Holy Grail)
- Print out the newly sorted array of explorers in a file called *grailExplorers.dat*

Example of input:

explorers.dat	holyGrail.dat
5 John,Romania,15,214 Miriam,USA,10,851 Arya,North,12,514 Leeroy,Russia,18,412 Jaque,France,13,721	Egypt,824

Example of output:

discoveriesExplorers.dat	grailExplorers.dat
Leeroy,Russia,18,412 John,Romania,15,214 Jaque,France,13,721 Arya,North,12,514 Miriam,USA,10,851	John,Romania,15,214 Leeroy,Russia,18,412 Arya,North,12,514 Jaque,France,13,721 Miriam,USA,10,851