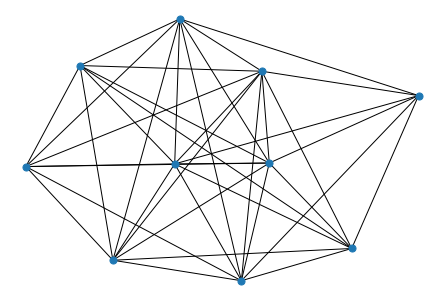
Biological Network Alignment Challenge Solution

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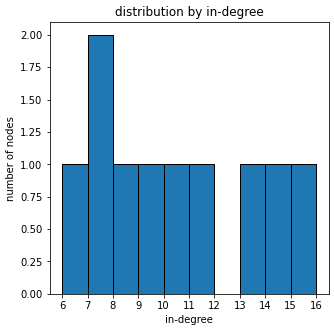
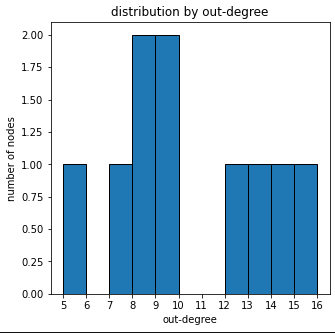
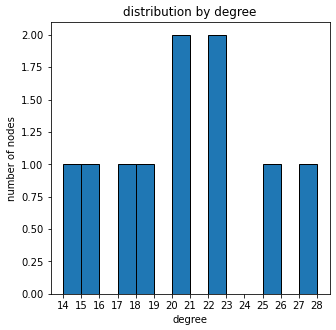
All the assumptions that I made include:

* The duplicate edges are counted as in/out-degree in histogram, since it is in the original input

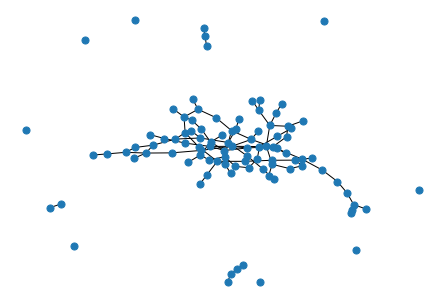
n10.txt



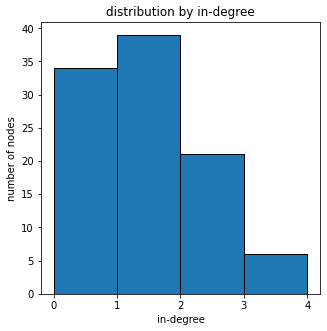
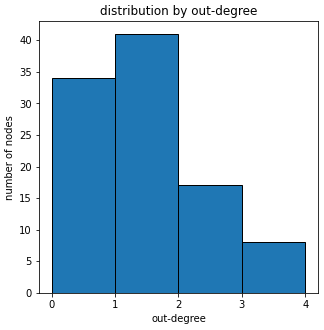
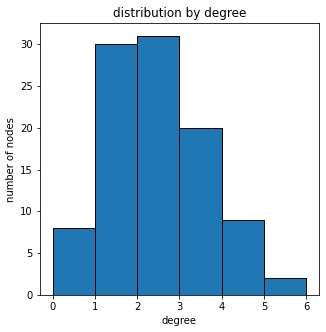
Connected Component: 1 Strongly Connected Component: 1

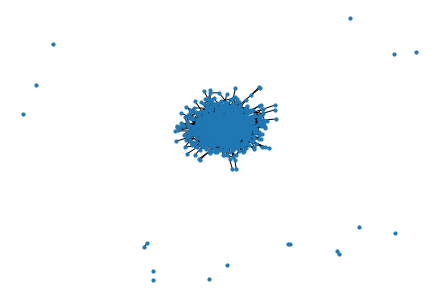
n100.txt



Connected Component: 12 Strongly Connected Component: 100

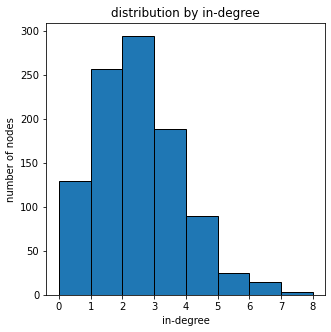
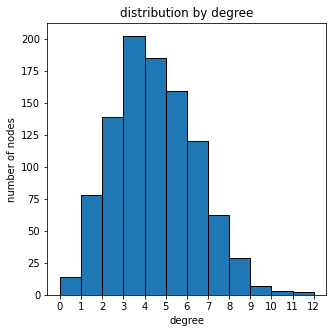
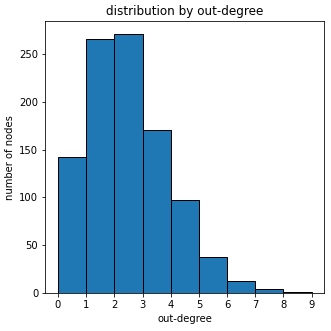
  

n1000.txt



Connected Component: 17

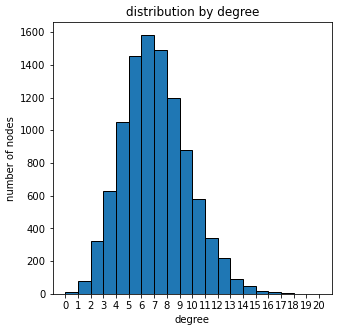
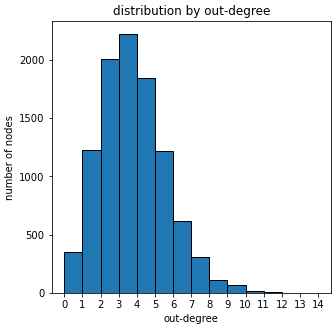
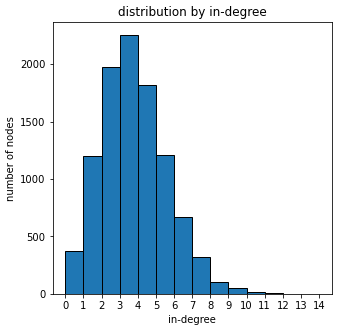
Strongly Connected Component: 383

n10000.txt

(graph not plotted)

Connected Component: 12 Strongly Connected Component: 814



s1.txt

(graph not plotted)

Connected Component: 6 Strongly Connected Component: 573

