Hw9

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Function

```
In[@]:= (*Power Law functions*)
     DegreeFreq[g_] := Map[{#[1], #[2] / VertexCount[g]} &, Tally[VertexDegree[g]]];
     CumDigDist[g_, min_, max_] :=
      Module[{df, cm, minpos, maxpos}, df = Sort[DegreeFreq[g], #1[1]] < #2[1] &];
       minpos = First@First@Position[df[All, 1], min];
       maxpos = First@First@Position[df[All, 1], max];
       cm = Reverse[Accumulate[Reverse[df[All, 2]]]]];
       Transpose[{df[minpos;; maxpos, 1], cm[minpos;; maxpos]]}]]
      (*Shift diagram functions*)
     RenameGraphN[g_, sl_] :=
      Module[{G = g, VrDg = s1, VrLs, VrLst, SVrLst, NewName, edges, NewEdges, ed},
       VrLs = VertexList[G];
       VrLst = Table[{VrDg[i], VrLs[i]}, {i, Length[VrDg]}];
       SVrLst = Sort[VrLst, #1[1] > #2[1] &];
       NewName = Table[SVrLst[i]][2], {i, Length[VrDg]}];
       edges = EdgeList[G];
       If[DirectedGraphQ[G],
        NewEdges = Table [Position [NewName, edges [i] [1] [1] [1] [1]
            Position[NewName, edges[i][2]][1][1], {i, Length[edges]}],
        NewEdges = Table [Position [NewName, edges [i] [1]] [1] [1] →
            Position[NewName, edges[i][2]][1][1], {i, Length[edges]}]];
       Graph[Range[Length[VrLst]], NewEdges]]
     SymmetryPoint[G_] :=
      Module[{s, EE = {}, EP = {}, PE = {}, PP = {}, ee = 0, ep = 0, pe = 0, pp = 0, tc, bc, tr, br},
       s = Length[VertexList[G]];
       ee = AdjacencyMatrix[G] [1] [1];
       pe = Total[Total[Take[AdjacencyMatrix[G], {2, s}, {1, 1}]]];
       ep = Total[Total[Take[AdjacencyMatrix[G], {1, 1}, {2, s}]]];
       pp = Total[Total[Take[AdjacencyMatrix[G], {2, s}, {2, s}]]];
       EE = \{0, ee / 2\};
       EP = \{0, ep / 2\};
       PE = \{0, pe / 2\};
       PP = {EdgeCount[G], pp / 2};
       For [i = 1, i < s - 1, i++;
        tc = Total[Total[Take[AdjacencyMatrix[G], {1, i-1}, {i, i}]]];
        bc = Total[Total[Take[AdjacencyMatrix[G], {i+1, s}, {i, i}]]];
        tr = Total[Total[Take[AdjacencyMatrix[G], {i, i}, {1, i - 1}]]];
        br = Total[Total[Take[AdjacencyMatrix[G], {i, i}, {i+1, s}]]];
        ee = ee + tc + tr + AdjacencyMatrix[G][i][i]];
        ep = ep + br - tc;
        pe = pe + bc - tr;
         pp = pp - bc - br - AdjacencyMatrix[G][[i]][i]];; EE = Join[EE, {ee / 2}];
```

```
EP = Join[EP, \{ep / 2\}];
   PE = Join[PE, {pe / 2}];
   PP = Join[PP, {pp / 2}];];
  {EE, EP + PE, PP}]
PlotShift[graph_, title_String] := Module[{a},
  a = ListLinePlot[SymmetryPoint[RenameGraphN[graph, VertexDegree[graph]]],
    PlotLegends \rightarrow Placed[{"I(\mathcal{E},\mathcal{E})", "I(\mathcal{P},\mathcal{E})", "I(\mathcal{P},\mathcal{P})"}, Bottom],
    PlotRange → All, AxesLabel → {"Core Size", "Number of Edges"},
    PlotStyle → {{Red, Thick}, {Blue, Thick}, {Green, Thick}},
    PlotLabel → Row[{"Shift diagram - ", title, " model"}]];
  Return[a];
 1
(*Community detection function, using the Louvaion method from HW7*)
(*Need to install this packet:*)
SetDirectory[NotebookDirectory[]];
PacletUninstall["IGraphM"];
Get["https://raw.githubusercontent.com/szhorvat/IGraphM/master/IGInstaller.m"];
<< "IGraphM`";
IGVersion[];
CommunityDetectionIGM[graph_, title_String] := Module[{cl, clcl, len, tot, plot},
  cl = IGCommunitiesMultilevel[graph];
clcl = cl["Communities"];
  len = Length[clcl];
tot = Table[Length[clcl[i]]], {i, 1, Length[clcl]}];
  plot = BarChart[tot, ChartLabels → Range[Length[clcl]],
    ChartStyle → "Pastel", LabelingFunction → (Placed[#, Center] &),
    PlotLabel → Row[{"Community detection - ", title, " model"}]];
  Return[plot];
The currently installed versions of IGraph/M are: {}
                                                          Name: IGraphM
Installing IGraph/M is complete: PacletObject
                                                           Version: 0.6.5
It can now be loaded using the command << IGraphM`
```

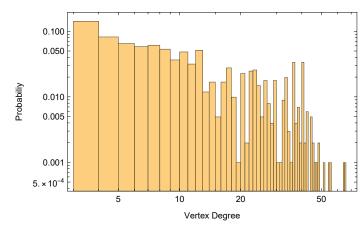
1. Find a network with power law degree distribution . Plot the degree distribution and try to find

```
in[o]:= net = ExampleData[{"NetworkGraph", "DiseaseGeneNetwork"}];
     Print["Vertex Count for the network: ", VertexCount[net]]
     maxD = Max[VertexDegree[net]];
     meanD = Mean[VertexDegree[net]] // N;
     Print["The mean of Vertex Count for the network: ", meanD]
     Print["The max Degree of the network: ", maxD]
     Vertex Count for the network: 1777
     The mean of Vertex Count for the network: 8.43106
     The max Degree of the network: 70
```

power exponent β .

```
In[@]:= Histogram[VertexDegree[net], {3, maxD, 1},
       "Probability", ScalingFunctions → {"Log", "Log"},
       FrameLabel \rightarrow {"Vertex Degree", "Probabiliy"}, Frame \rightarrow True]
      Powerfit = Fit[Log[CumDigDist[net, 5, maxD]], {1, x}, x]
      beta = -Powerfit[2][1];
      Print[Style["The power exponent \beta is: ", 20], beta, "\rightarrow 2\leq \beta \leq 3"]
      Show[ListLogLogPlot[CumDigDist[net, 5, maxD]],
       Plot[Powerfit, \{x, 1, 58\}, PlotStyle \rightarrow Green]]
```

Out[0]=

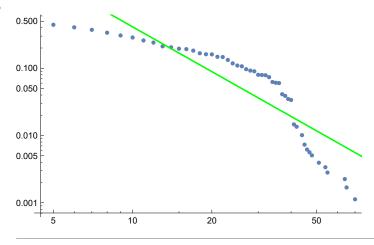


Out[0]=

4.20769 - 2.2119 x

The power exponent β is: $2.2119 \rightarrow 2 \le \beta \le 3$





2. Generate the symmetry point graph for PA with sevral sizes (on what fraction it is happends? what is the "core" fraction)

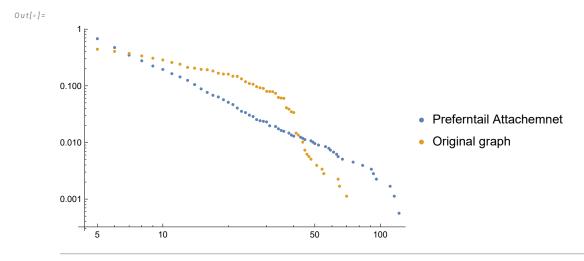
5

10

```
In[@]:= Pa = RandomGraph[BarabasiAlbertGraphDistribution[
            VertexCount[net], Round[EdgeCount[net] / VertexCount[net]]]];
       Print["Vertex Count for the PA: ", VertexCount[Pa]]
       Print["The mean of Vertex Count for the PA: ", Mean[VertexDegree[Pa]] // N]
       {Histogram[VertexDegree[net], {"Log", 15}, {"Log", "PDF"},
         PlotLabel → "Original Graph"], Histogram[VertexDegree[Pa],
          {"Log", 15}, {"Log", "PDF"}, PlotLabel → "Preferntail Attachment"]}
       PowerfitPa = Fit[Log[CumDigDist[Pa, 5, Max[VertexDegree[Pa]]]], {1, x}, x]
       Show[ListLogLogPlot[CumDigDist[Pa, 5, Max[VertexDegree[Pa]]]],
        Plot[PowerfitPa, \{x, 1, 58\}, PlotStyle \rightarrow Green]]
       ListLogLogPlot[{CumDigDist[Pa, 5, Max[VertexDegree[Pa]]],
         CumDigDist[net, 5, Max[VertexDegree[net]]]},
        PlotLegends → {"Preferntail Attachemnet", "Original graph"}]
       (*not work*)
       (*spPA=PlotShift[Pa,"PA"]*)
       Vertex Count for the PA: 1777
       The mean of Vertex Count for the PA: 7.98875
Out[0]=
                                              Preferntail Attachment
                    Original Graph
           0.50
                                      0.100
                                      0.010
           0.01
                                      0.001
        5. \times 10
                                       10^{-4}
                         10
                            20
Out[0]=
       2.80841 - 1.93525 x
Out[0]=
       0.100
       0.010
       0.001
```

50

100

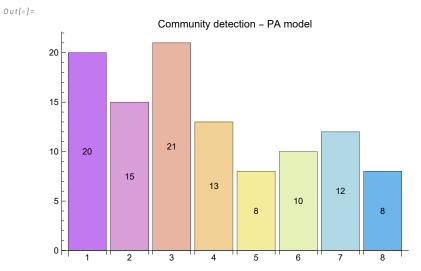


3. Add Preferential Attachment to the models you check "Network of Throne" with . Like we did in home work 8

```
In[*]:= (*Load the netork*)
     SetDirectory[NotebookDirectory[]];
     file = Rest[Import["stormofswords.csv"]];
     tribes = Import["tribes.csv"];
     nodes = Flatten[tribes[All, 1]];
     nodesTribe = Flatten[tribes[All, 2]];
     edges = \#[1] \leftrightarrow \#[2] \& /@ file[All, {1, 2}];
     G = Graph[nodes, edges];
```

```
In[*]:= (*Creat the PA - Graph*)
        PaG = RandomGraph[BarabasiAlbertGraphDistribution[
           VertexCount[G], Round[EdgeCount[G] / VertexCount[G]]]]
        (*Degree distribution*)
        degDisPa = Histogram[VertexDegree[PaG], {1},
          "Probability", PlotLabel → "Degree Distribution - PA model",
          \label{lem:chart-legends} \textbf{Chart-Legends} \rightarrow \textbf{Placed} \ [ \{ \texttt{StringForm} \ [ \texttt{"Mean degree: ``", Mean} \ [ \texttt{VertexDegree} \ [ \texttt{PaG} ] ] \ // \ N ] \ ,
              StringForm["Variance: ``", Variance[VertexDegree[PaG]] // N]}, Bottom]]
Out[0]=
Out[0]=
                    Degree Distribution - PA model
         0.35
         0.30
         0.25
         0.20
         0.15
         0.10
         0.05
        ■ Mean degree: 5.88785 ■ Variance: 25.5533
 In[*]:= (*Giant component*)
        GiantComponentG = Length[First[ConnectedComponents[PaG]]];
        Print["Giant component: ", GiantComponentG]
        (*Average distance, to avoide ∞ in un-connected graph,
        I took the largest connected components for that*)
        AverageDistancePa = Mean[Flatten[
              GraphDistanceMatrix[Subgraph[PaG, First[ConnectedComponents[PaG]]]]]] // N;
        Print["Average distance: ", AverageDistancePa]
        Giant component: 107
       Average distance: 2.5866
        (*Shift diagram, not work at all*)
        (*sdPa=PlotShift[PaG,"PA"]*)
```

In[0]:= (*Community detection, for Throne model I plot the true communities by tribes*) CommunityDetectionPa = CommunityDetectionIGM[PaG, "PA"]



Discuss our finding

For the graph DiseaseGeneNetwork network

for the discuss models, G (n, p) and G (n, m) models generate random graphs with varying edge configurations, while the block model and configuration model aim to simplify and analyze specific aspects of the original graph, such as community structure or degree distribution, respectively. The choice of model depends on the specific characteristics of the original graph that we want to preserve or explore.

The average distance and giant component are similarly present, and we can see that expected value (Degree) approximately the same for each model and the result is

N*P = 1777*0.0622465~110.612

and for the Diameter for all model is the same again and the calculate D~log (1777) = 3.249,log (110.612) = 2.043. We can see that is less then $\log (107) = 4.672$ as we expected. In summary, the differences between the original network and the graph models arise due to the specific generation methods employed by each model. These differences are purposeful and serve different analytical or modeling objectives, such as studying random graphs, understanding community structures, or analyzing degree distributions. The choice of model depends on the research. The average distance is around $\log (n)/\log(z) = (\log (1777)/\log(110.612) \sim 1.59)$.

1.59002

For the graph Throne network

N*P = 107*0.0622465~6.6

different analytical or modeling objectives, such as studying random graphs, understanding community structures, or analyzing degree distributions . The choice of model depends on the research . The average distance is around $log(n)/log(z) = log(107)/log(6.6) \sim 2.47624$

Log2[107] In[@]:= Log2[6.6] Out[0]=

2.47624