

## Distribution of number of neighbors on semi-directed Barab'asi-Albert networks with many initial neighbors

Dr. M. A. Sumour<sup>\*</sup>  
Dr. F. W. S. Lima<sup>\*\*</sup>  
Dr. M. A. Radwan<sup>\*</sup>  
Dr. M. M. Shabat<sup>\*\*\*</sup>

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m

m

m

m

.SDBA

SDBA2 SDBA1

m SDBA2

SDBA1 m

. m m.

### Abstract

On directed and undirected Barab'asi-Albert networks, when a new node has selected  $m$  old nodes as neighbors, then the  $m$  old nodes are added to the Kert'esz list, and the new node is also added  $m$  times to that list. These connections are made with  $m$  randomly selected elements of that list. If one adds to the list the  $m$  old nodes, plus only once and not  $m$  times the new node, one gets a semi-directed network (SDBA). Now we check the number of neighbors on two versions (SDBA1 and SDBA2) for semi-directed Barab'asi-Albert networks.

We found that SDBA2 does not give proper power laws for large  $m$  but it does so for small  $m$ , whereas SDBA1 gives proper power laws for large and small  $m$ . The resulting exponents vary with  $m$ .

**Keywords:** Directed BA networks , Undirected BA network , Semi-directed BA network.

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\* Physics Department, Al-Aqsa University, Gaza, Gaza Strip, Palestinian Authority.

\*\* Dietrich Stauffer Computational Physics Lab, Departamento de F'isica, Universidade Federal do Piaui, 64049-550, Teresina - PI, Brazil.

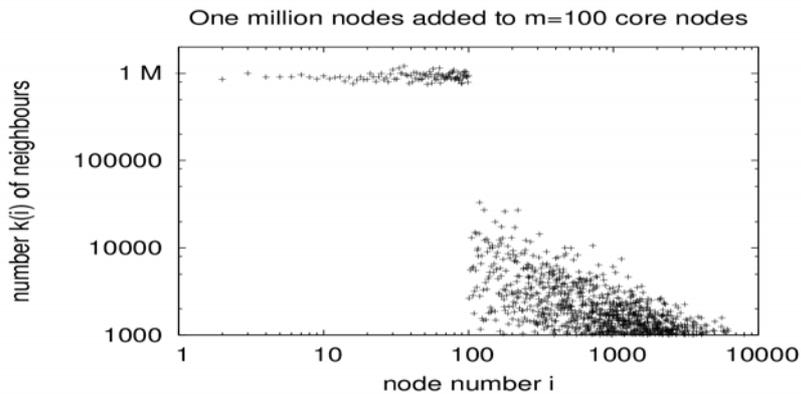
\*\*\*Physics Department, Islamic University P.o.Box 108, Gaza, Gaza Strip, Palestinian Authority.

## Distribution of number of neighbors ....

### 1. Introduction:

The Barba'si-Albert network [1,2] is growing such that the probability of a new site to be connected to one of the already existing sites, is proportional to the number of previous connections to this already existing site: The rich gets richer. In this way, each new site selects exactly  $m$  old sites as neighbors. In directed (DBA) and undirected (UBA) Barba'si-Albert networks, the network itself was built in the standard way, but when agents (spins) were put on the network nodes [2,3,4] the neighbor relations were such that if A has B as a neighbor, B in general does not have A as a neighbor for DBA while it does have for UBA.

The present work continues the study of two semi-directed BA networks, SDBA1 [5] and SDBA2 [6] for much larger  $m$  than before. In these semi-directed networks the exponent  $\gamma$  for the power law governing the decay of the number  $n(k)$  of nodes having  $k$  neighbors,  $n(k) \propto 1/k^\gamma$ , depends continuously on the parameter  $m$ , and we want to know its behavior for  $m \rightarrow \infty$ .



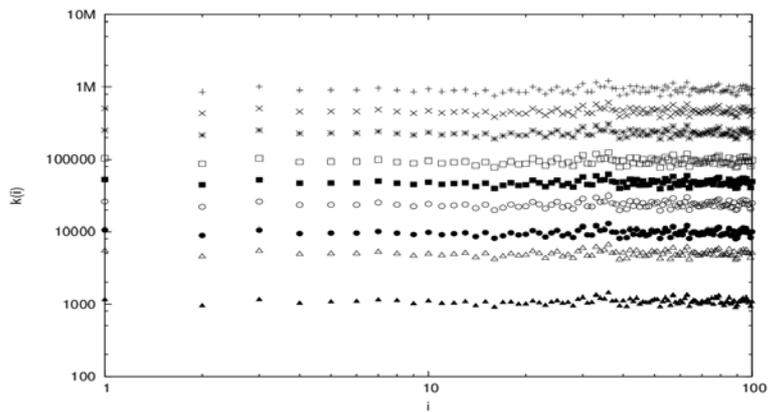
Figure(1): Number of neighbors  $k(i)$  influencing node  $(i)$  in SDBA2 networks with  $m=100$  and  $N = 1$  million.

### 2. Models and Simulations:

#### 2.1 Directed (DBA) and undirected (UBA) Barba'si-Albert network:

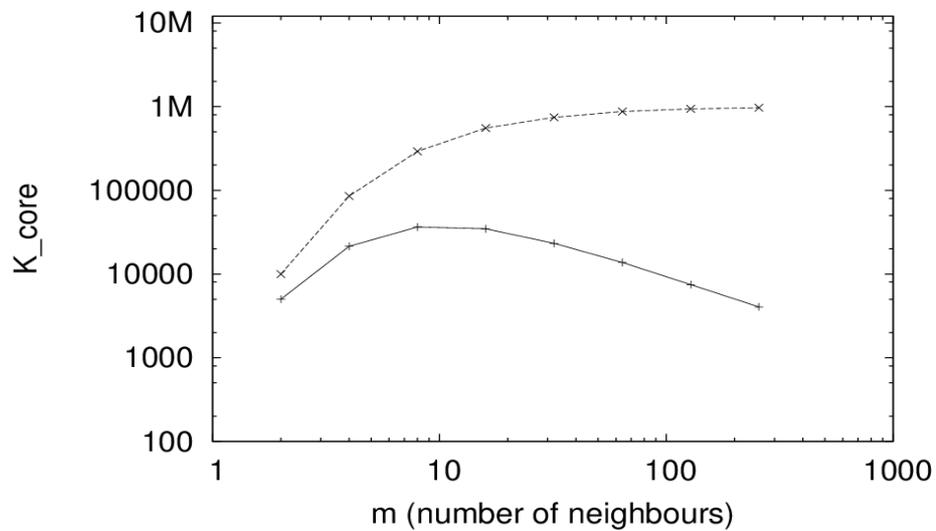
In the Barabasi-Albert network [1,2], if a new node selects  $m$  old nodes as neighbor, then the  $m$  old nodes are added to the Kert'esz list, and the new

node is also added  $m$  times to that list. Connections are made with  $m$  randomly selected elements of that list.



Figure(2): Number of neighbors  $k(i)$  versus the index of core nodes, at constant  $m=100$  and different  $N$

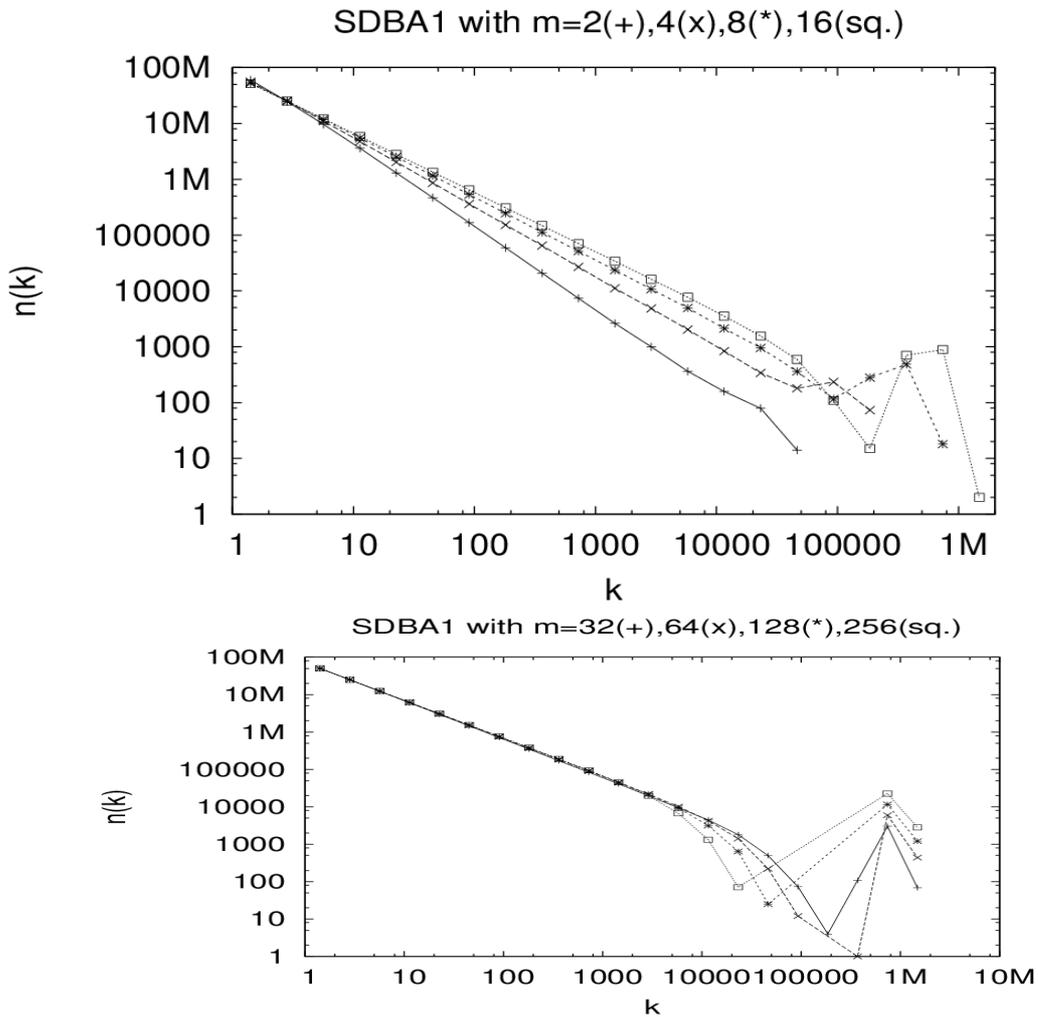
Core neighbours versus  $m$ , SDBA1 above and SDBA2 below



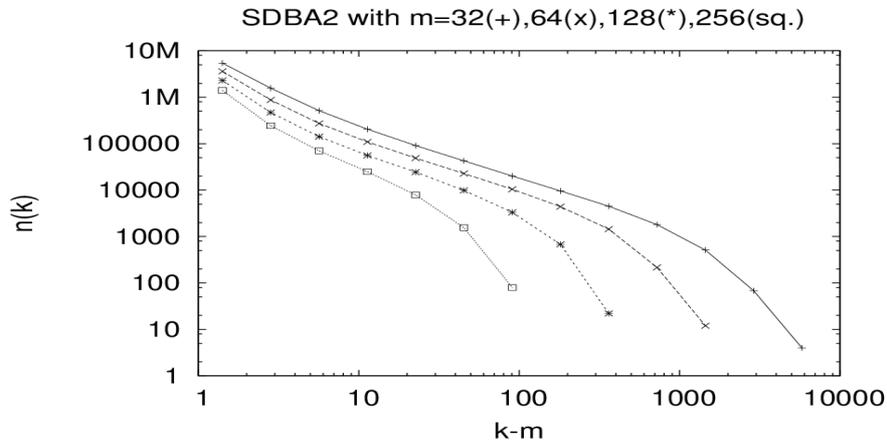
Figure(3): Average  $K_{core}$  versus  $m=2,4,8,16,32,64,128,$  and  $256$  with constant  $N=1,000,000$  for SDBA1 and SDBA2.

### Distribution of number of neighbors ....

In directed Barba'si-Albert networks DBA, the network itself was produced in the standard way, but then the neighbor relations were such that if A has B as a neighbor, B in general does not have A as a neighbor. The undirected Barabasi-Albert network UBA usually is grown in the same way, but then the neighbor relations were such that if A has B as a neighbor, B has A as a neighbor. The growth process always starts with a fully connected core of  $m$  nodes.

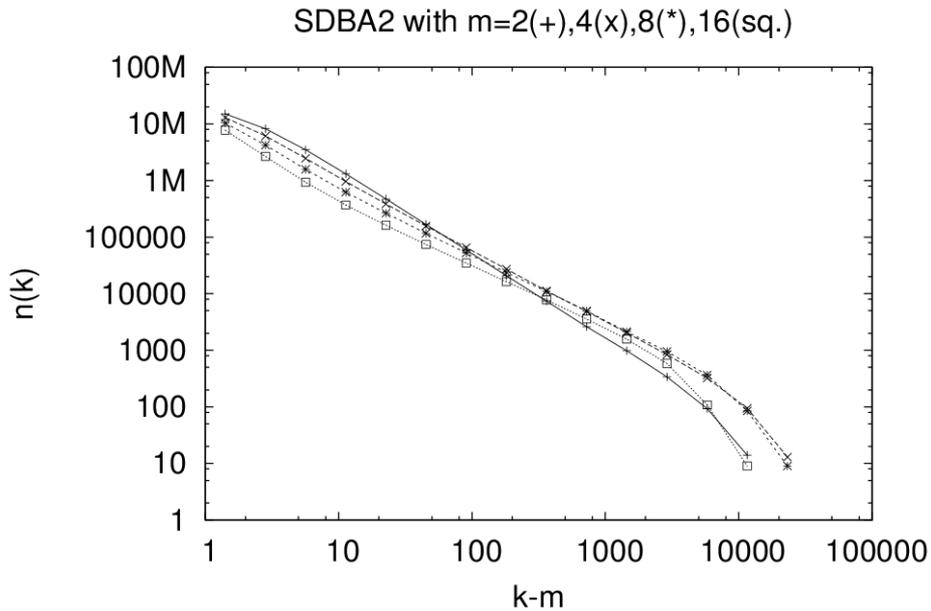


Figure(4): Number of nodes  $n(k)$  versus  $k$  with different  $m=2$  to  $256$  and  $N=1,000,000$  for SDBA1.



### 2.2 Semi-Directed Barba'asi-Albert network (SDBA):

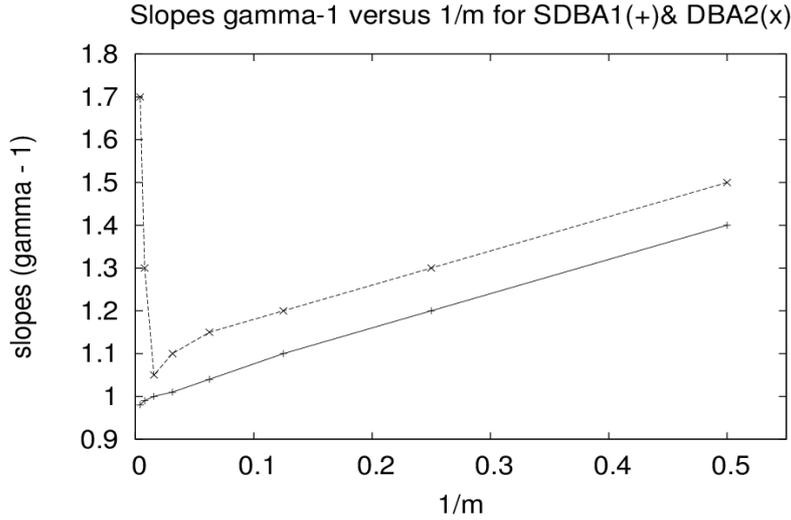
If in building a BA network one would only add the old nodes to the Kert'esz list after a new node was added, then only the initial core can be selected as neighbors, which is not interesting. But if one adds to the list the  $m$  old nodes, plus only once and not  $m$  times the new node, one has a semi-directed network (SDBA).



Figure(5): Number of nodes  $n(k)$  versus  $k-m$  with different  $m=2$  to  $256$  and  $N=1,000,000$  for SDBA1.

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The first version SDBA1 builds the network in the way of [5]. The new node selects  $m$  sites  $j$  which  $n$  will all influence, while  $n$  will be influenced only by the first selected  $j$ . Our second version SDBA2 inverts the direction of the spin interaction[6]:The new node  $n$  selects  $m$  sites  $j$  which will all influence  $n$ , while  $n$  will influence only the first selected  $j$ . Now we look at both SDBA1 and SDBA2 in the hope of finding the limit  $m \rightarrow \infty$ , more precisely for  $1 \ll m \ll N$ .



Figure(6): Plot of exponents  $\gamma - 1$  (slopes) versus  $1/m$  for the SDBA1 and SDBA2 with  $m=2,4,8,16,32,64,128$ , and  $256$ .

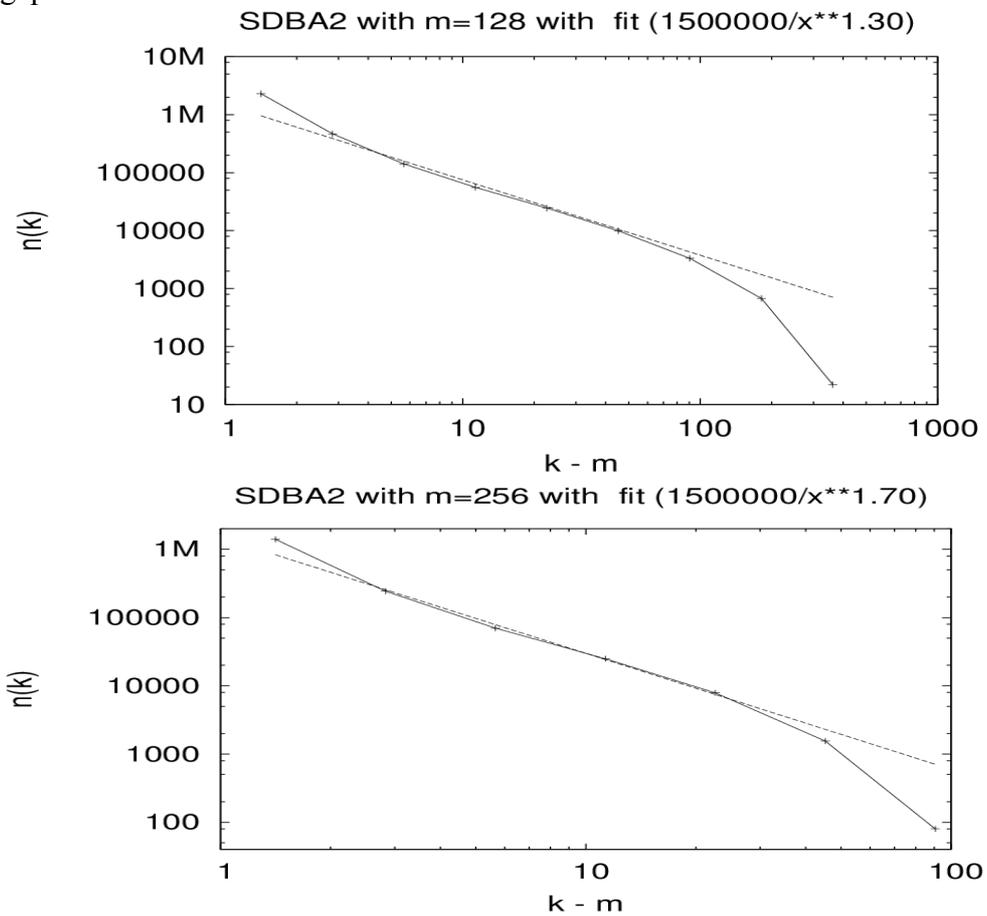
### 3. Results and discussion:

We used a simulation program that include two different programs Called SDBA1 and SDBA2 networks as shown in appendix (1-2). We determined powers of two and store in variable  $m=2,4,8,16,32,64,128$ , and  $256$ , and we fixed the network size (called maxtime in the programs)  $N=1,000,000$  and fixed an upper bond  $k_b=9$  million for  $k$  variable. First we measured the distribution nodes of  $n(k)$  number influenced by  $k$  neighbors in both programs SDBA2 and SDBA1 we concluded that both programs are support the result in [5]. The program SDBA2 shown a clear gap in  $k(i)$  number of neighbors, as shown in figure(1).

The new nodes  $n$  added to the initial core of  $m$  sites have at least  $k(n)=m$  (here  $m=100$ ) neighbors (see appendix 2 at label 1) most of result from the  $m$  core nodes, shown an important difference gap between SDBA1 and

SDBA2. Based on our simulation study for different sizes of lattice, by varying lattice size  $N$  (1,000,000, 500,000, 250,000, 100,000, 50,000, 10,000, 5,000, and 1,000) fixed  $m=100$ , the number of neighbors  $k(i)$  for the core nodes is shown in Figure(2).

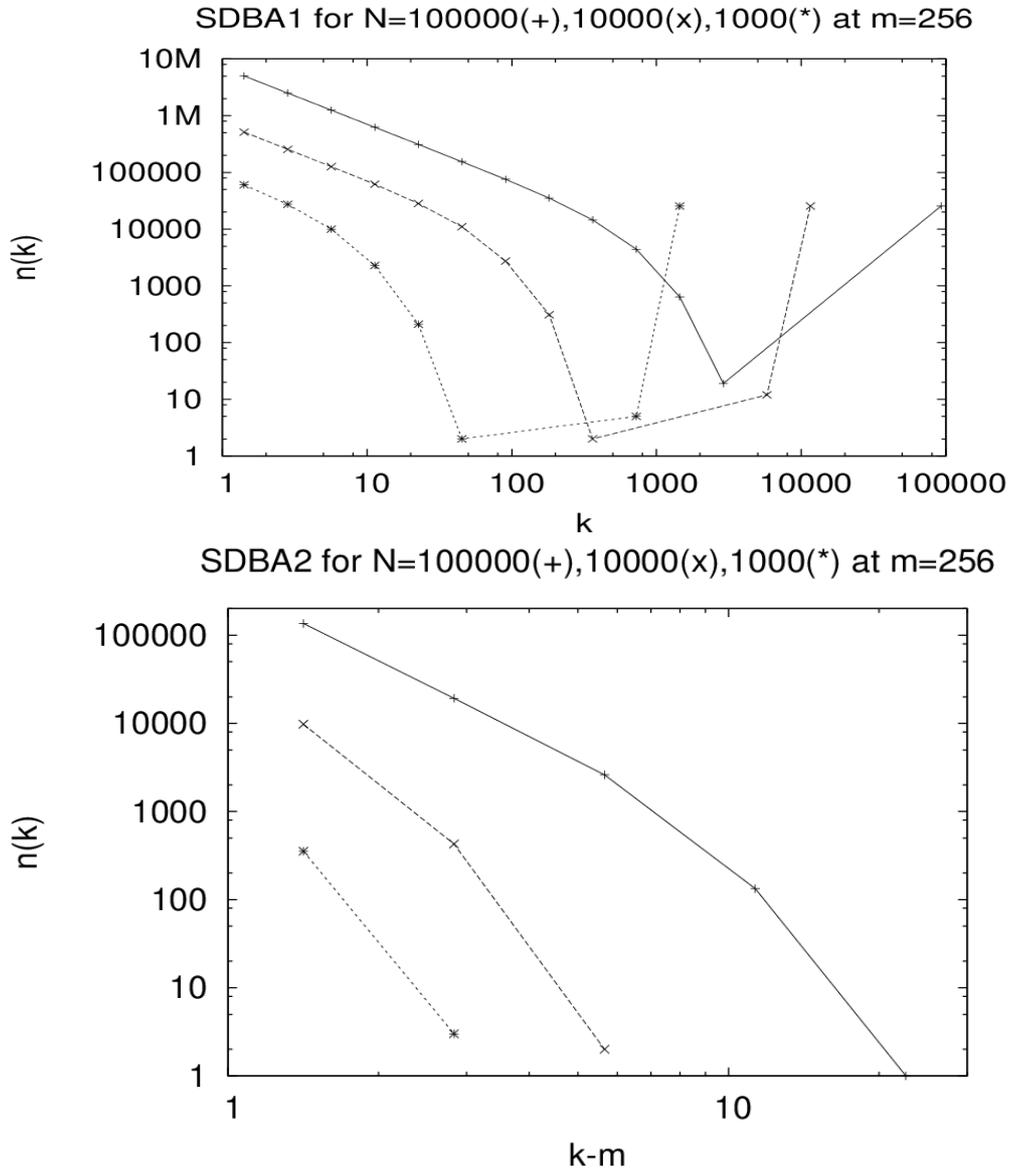
To plot the histograms  $n(k)$  of the numbers  $k$  of neighbors influencing a node, we bin them in powers of two. For SDBA2, because of the above gap, we count instead of  $k$  the number  $k-m$  of additional neighbors gained after the addition of a node to the network, in order to get smoother histograms. This means for SDBA2 we use only the right part of Figure(2) beyond the gap at  $m$ .



Figure(7): Number of nodes  $n(k)$  versus  $k-m$  links with  $m=128$  and  $256$ , at  $N=1,000,000$  for SDBA2

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In contrast, for SDBA1 these straight lines correspond to power laws without such a gap problem, we counted all  $k$  as usual and not the difference  $k-m$ .



Figure(8): Histograms of  $n(k)$  at smaller network sizes  $N$  for SDBA1 (top) and SDBA2 (bottom).

The average number  $k$  of neighbors for the  $m$  core nodes:

$$K_{core} = \sum_{i=1}^m k(i)/m,$$

is plotted in Figure(3), averaged over 100 samples. All following figures are also averaged over 100 samples.

As shown in figures(4 and 5), we can determine a nice slopes in log-log plots of  $n(k)$  versus  $k$  for SDBA1, while for large  $m$ , bad power law is seen for SDBA2. These slopes are summarized in Figure(6), which is the essence of main result of the research. The steep increase for SDBA2 near  $1/m=0$  is unreliable and due to the fact that Figure(5 bottom) does not give good power laws.

To see the problems for SDBA2 at large  $m$  we plotted as examples  $m=128$  and  $256$  which resulted a bad straight line, thus we fit the curves by  $1500000/x^{1.3}$  for  $m=128$ , and by  $1500000/x^{1.7}$  for  $m=256$ , as listed in figure(7).

The last figure(8) shown the strong influence of the finite network size  $N$  on the tails of the histogram, in both SDBA1 and SDBA2. for both SDBA1 and SDBA2, only for large networks used in Figures (1,3,4,5) with  $N = 1$  million do we get good power laws.

#### 4. Conclusion:

We concluded that SDBA2 the initial  $m$  core nodes have anomalous neighborhoods and that SDBA2 does not given proper power laws for large  $m$  but does so for small  $m$ . In contrast, SDBA1 given a nice power laws and its exponent  $\gamma-1$  approaches unity linearly in  $1/m$ , meaning  $n(k) \propto 1/k^2$  for large  $m$ , as opposed to  $1/k^3$  for standard BA networks at all  $m$ . In both SDBA1 and SDBA2 the exponents  $\gamma$  are not universal and a theoretical explanation for this continuous variation of the exponent should be searched for. The lack of good power laws for SDBA2(not SDBA1) might vanish if much larger networks can be simulated.

#### Acknowledgments:

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### Appendix(1):

Fortran program for Ising model on SDBA1 given in [5], without spins.

```
parameter(nrun=100,maxtime=1000000,m=64,iseed=2,max=maxtime+m,
1 kb=9000000,length=1+(1+m)*maxtime+m*(m-1))
integer*8 ibm
real*8 factor
dimension nklog(0:30)
dimension k(max), nk(kb), list(length)
data nk/kb*0/,nklog/31*0/
OPEN (UNIT=60,FILE='SDBA1-1NmA64.DAT')
OPEN (UNIT=70,FILE='SDBA1-2mA64.DAT')
WRITE (60,*)'#(nrun, maxtime, m, iseed)',nrun, maxtime,
m,iseed
ibm=2*iseed-1
factor=(0.25d0/2147483648.0d0)/2147483648.0d0
fac=1.0/0.69314
do 5 irun=1,nrun
do 3 i=1,m
do 7 j=(i-1)*(m-1)+1,(i-1)*(m-1)+m-1
7 list(j)=i
3 k(i)=m-1
L=m*(m-1)
if(m.eq.1) then
L=1
List(1)=1
k(1)=1
endif
C All m initial sites are connected with each other
do 1 n=m+1,max
```

```

do 2 new=1,m
4  ibm=ibm*16807
  j=1.d0+(ibm*factor+0.5d0)*L
  if(j.le.0.or.j.gt.L) goto 4
  j=list(j)
  list(L+new)=j
2  k(j)=k(j)+1
  L=L+m+1
  list(L)= n
1  k(n)=1
  write(60,*),'#(irun)', irun
  do 5 i= 1,max
5  nk(k(i))=nk(k(i))+1
  SUM=0
  do 6 i=1,m
  SUM=SUM+K(i)
6  write(60,*) i,k(i)
  do 9 i=1,kb
  j=alog(float(i))*fac
9  nklog(j)=nklog(j)+nk(i)
  AVERGESUM=SUM/m
  jmax=(1.0 + alog(float(kb))*fac)
  do 10 j=0,jmax
  if(nklog(j).ne.0)
    write(70,*)sqrt(2.0)*2**j,nklog(j),j,AVERGESUM
10 if(nklog(j).ne.0)print*,sqrt(2.0)*2**j,nklog(j),
  1 j,AVERGESUM
  stop
  end

```

### Appendix(2):

Fortran program for Ising model on SDBA2 given in[6], without spins.

```

parameter(nrun=100,maxtime=1000000,m=64,iseed=2,
1  max=maxtime+m kb=9000000,length=1+
1  (1+m)*maxtime+m*(m-1))
  integer*8 ibm
  real*8 factor
  dimension nklog(0:30)
  dimension k(max), nk(kb), list(length)
  data nk/kb*0/,nklog/31*0/,k/max*0/

```

### Distribution of number of neighbors ....

```
OPEN (UNIT=60,FILE='SDBA2-1M64.DAT')
OPEN (UNIT=70,FILE='SDBA2-2M64.DAT')
WRITE (60,*)'#(nrun, maxtime, m, iseed)',nrun,
1 maxtime, m,iseed
ibm=2*iseed-1
factor=(0.25d0/2147483648.0d0)/2147483648.0d0
fac=1.0/0.69314
do 5 irun=1,nrun
do 3 i=1,m
do 7 j=(i-1)*(m-1)+1,(i-1)*(m-1)+m-1
7 list(j)=i
3 k(i)= m-1
L= m*(m-1)
if(m.eq.1) then
L=1
List(1)=1
k(1)=1
endif
C All m initial sites are connected with each other
do 1 n=m+1,max
do 2 new=1,m
4 ibm=ibm*16807
j=1.d0+(ibm*factor+0.5d0)*L
if(j.le.0.or.j.gt.L) goto 4
j=list(j)
k(n)=k(n)+1
if(new.eq.1) k(j)=k(j)+1
if(k(j).gt.kb .or. k(n).gt.kb) stop 9
list(L+new)=j
2 continue
L=L+m+1
List(L)=n
1 k(n)=m
write(60,*) '# (irun )', irun
do 5 i= m+1,max
5 if(k(i).gt.m) nk(k(i)-m)=nk(k(i)-m)+1
SUM=0
```

```
do 6 i=1,m
SUM=SUM+K(i)
6  write(60,*) i,k(i)
do 9 i=1,kb
j=log(float(i))*fac
9  nklog(j)=nklog(j)+nk(i)
AVERGESUM=SUM/m
jmax=(1.0 + log(float(kb))*fac)
do 10 j=0,jmax
if(nklog(j).ne.0)
write(70,*)sqrt(2.0)*2**j,nklog(j),j,AVERGESUM
10 if(nklog(j).ne.0)print*,sqrt(2.0)*2**j,
1 nklog(j),j,AVERGESUM
stop
end
```