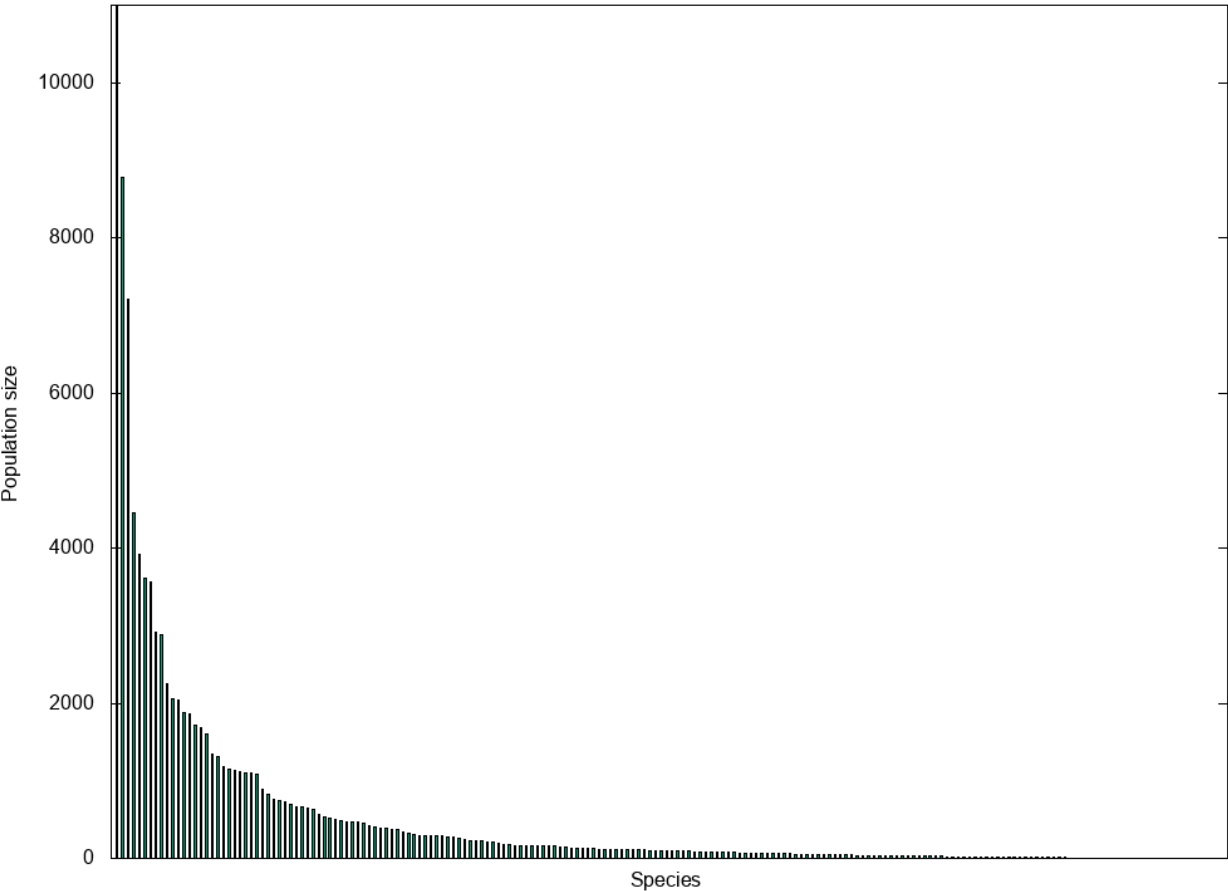


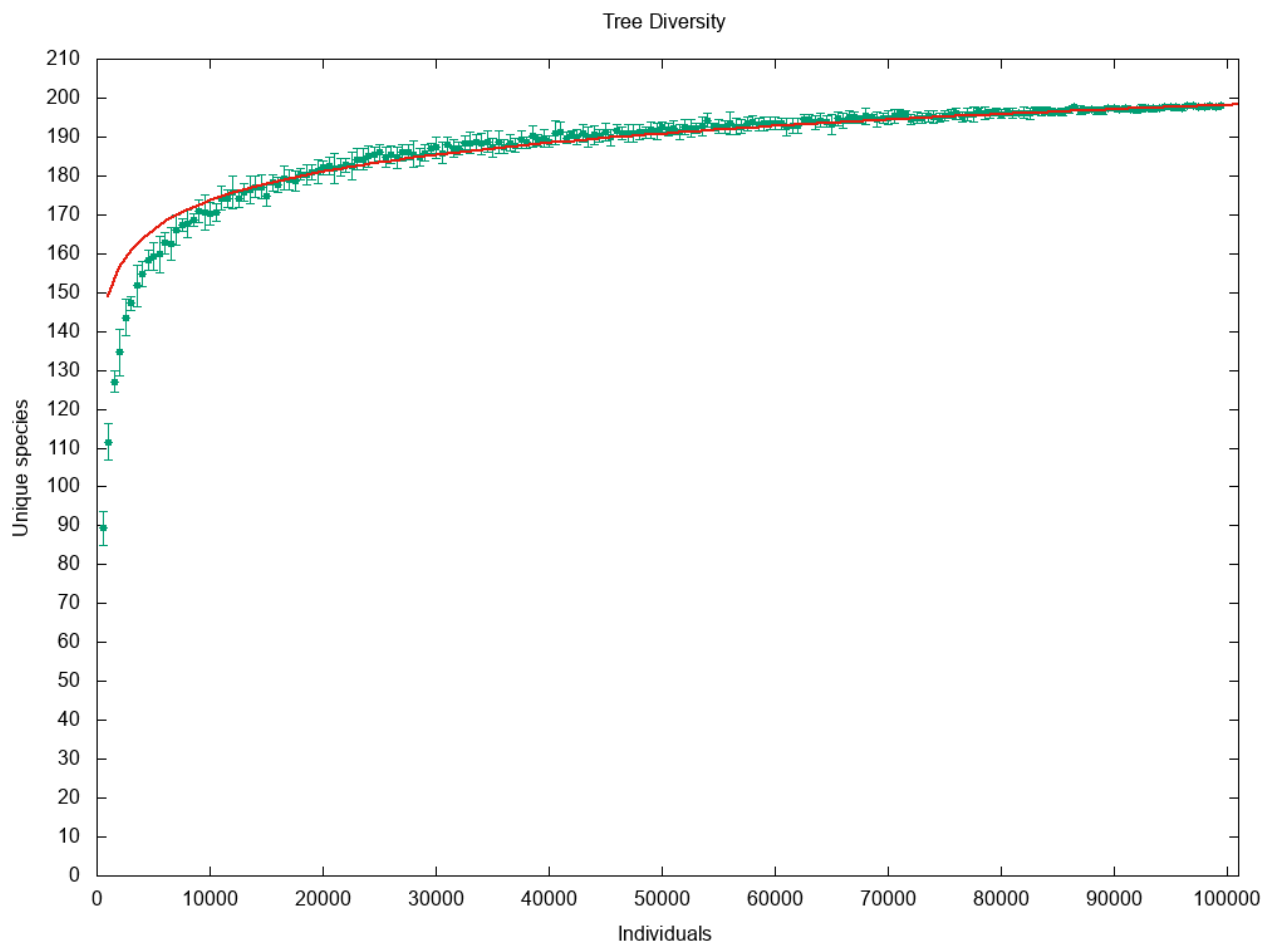
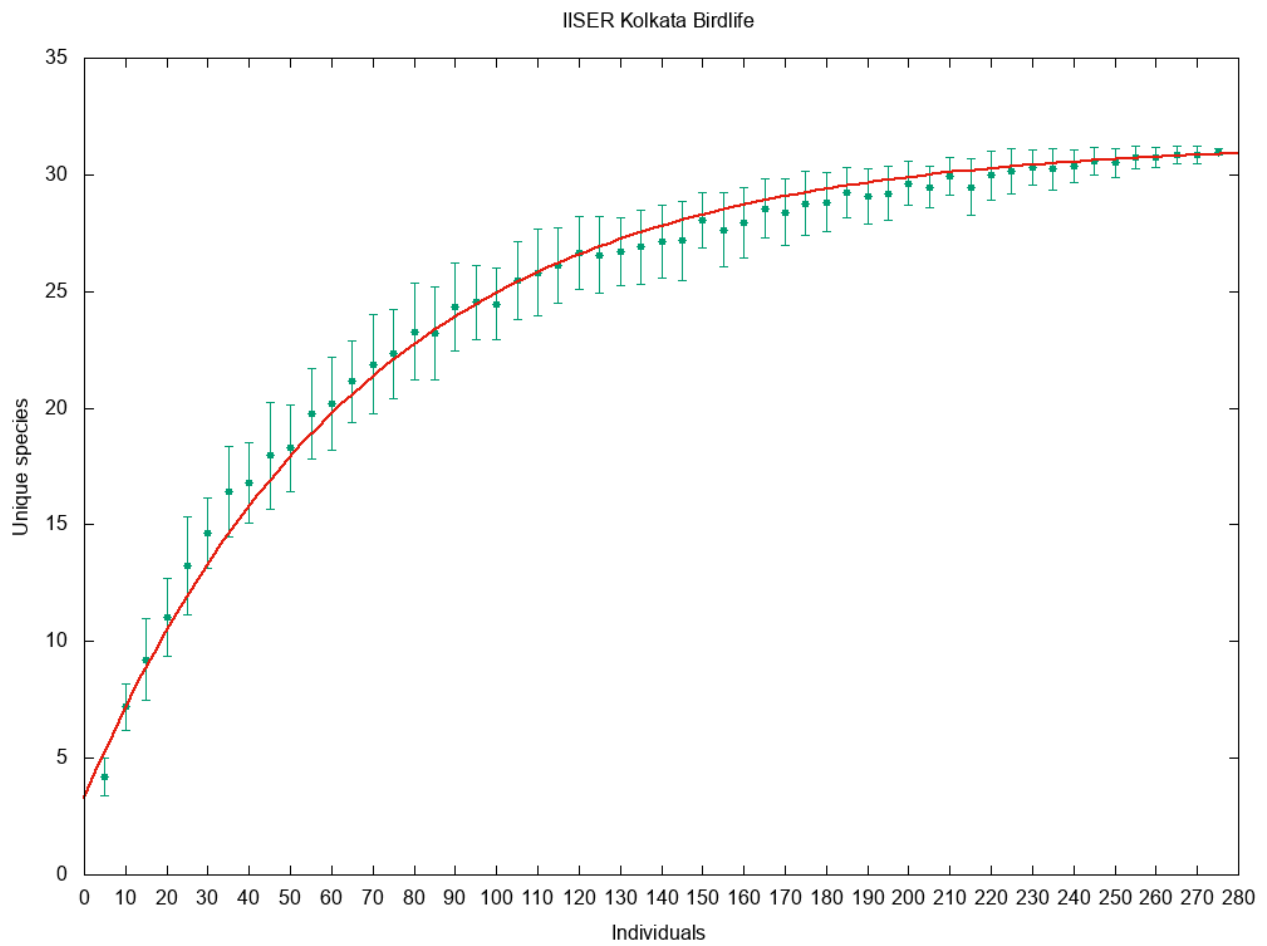
Species

Tree Diversity



Species

50 "common mynah"
40 "asian pied starling"
20 "black drongo"



```

18     "spotted dove"
16     "asian openbill"
14     "blyth's reed warbler"
11     "common iora"
10     "red-vented bulbul"
9      "scaly-breasted munia"
7      "yellow-bellied sunbird"
6      "white-throated kingfisher"
6      "rose-ringed parakeet"
6      "jungle babbler"
6      "great egret"
5      "plain prinia"
5      "grey sunbird"
5      "*(indian swift)"
4      "spotted owlet"
4      "oriental magpie robin"
4      "house crow"
3      "long-tailed shrike"
3      "brown-capped pygmy woodpecker"
2      "streaked-fantail warbler"
2      "rufous treepie"
2      "jungle mynah"
2      "cotton pygmy goose"
2      "cattle egret"
1      "green bee-eater"
1      "common flameback"
1      "*(scaly munia-like black crow)"
1      "*(black bird)"

```

```
#!/usr/bin/env python
```

```
# Calculates the simpson index of diversity for given data.
```

```
# Usage:
```

```
# ./simpson.py [frequencies]
```

```
#
```

```
# frequencies    list (space separated) of frequencies of species
```

```
from os import sys
```

```
import math
```

```
data = map(int, sys.argv[1:])
```

```
n = sum(data)
```

```
d = 0
```

```
for x in data:
```

```
    p = x * 1.0 / n
```

```
    d += p*p
```

```
print d
```

10976	"HUMBLA"	281	"GOMPSE"	60	"WALSPI"	4	"MEMEA_"
8788	"AGROIN"	261	"TRICER"	56	"NOTHFO"	3	"SYMPEL"
7203	"MESUNA"	237	"EUPHLO"	55	"SYZYAM"	3	"CASEZE"
4460	"UNIDEN"	232	"SYMPCO"	53	"EUGERI"	3	"BOMBCE"
3926	"GARCHE"	224	"PODATH"	52	"DYSOBI"	2	"SYZYMI"
3614	"SHORDI"	222	"SEMESU"	51	"SCOLAC"	2	"SYMPCC"
3567	"AGROHO"	216	"DILLRE"	50	"LIJNGA"	2	"PAGIDI"
2912	"PSYCNI"	215	"CALOBR"	49	"CHIOAL"	2	"EUONWA"
2886	"MESUFE"	197	"SYZYMA"	48	"CAMPZE"	2	"CLIDHI"
2249	"SHORWO"	179	"ACTIAL"	45	"MELAMA"	2	"CANTDI"
2062	"SHORAF"	174	"TIMOJA"	41	"PALAHI"	1	"WENDBI"
2037	"UROPEL"	167	"ELAESU"	41	"CALOTR"	1	"STRYMI"
1885	"SHORCR"	165	"GYRIWA"	40	"BYRSEL"	1	"PALAGR"
1859	"CULLCE"	163	"URANAP"	37	"SYZYOP"	1	"MASTNI"
1724	"SCHUCA"	163	"SYMPCU"	37	"DESMZE"	1	"KOKOZE"
1680	"PALACA"	162	"STRONA"	37	"ARTONO"	1	"FRAGZE"
1596	"SHORME"	161	"PUTRRO"	36	"MEMEB_"	1	"CARYUR"
1340	"MYRIDA"	159	"CINNVE"	35	"MEMESY"	1	"BRIDRE"
1320	"SHORTR"	154	"PAVEIN"	35	"MEMED_"		
1190	"GAERVA"	141	"SYZYLI"	33	"PRUNWA"		
1148	"CULLRO"	139	"MALLRH"	33	"PALALA"		
1126	"XYLOCH"	137	"GOMPTE"	30	"CLEIPA"		
1123	"SEMEWA"	132	"SYZYSY"	30	"CINNRI"		
1108	"PSYCDU"	131	"PRISTE"	29	"GARCTE"		
1098	"PALATH"	124	"SEMEAC"	29	"ARDIGA"		
1080	"PALAPE"	123	"MADHFU"	25	"MEMERY"		
896	"GAERRO"	121	"NEOLCA"	25	"MEMEC_"		
834	"ANISCI"	120	"DIOSQU"	25	"ABARBI"		
768	"DIOSAC"	115	"EURYAC"	23	"STEMCA"		
748	"GLOCAC"	113	"DIOSIN"	23	"DIOSRA"		
735	"SYZYNE"	112	"GLOCZE"	19	"OSMEGA"		
703	"MEMEAR"	110	"MEMEPR"	19	"GLOCNE"		
663	"MEMERO"	110	"DYSOPE"	18	"GARCTH"		
659	"CALOTH"	106	"LASIOB"	16	"HORSIR"		
646	"SEMEGA"	106	"ANTIPY"	16	"CLERIN"		
624	"MASTTE"	99	"MALLFU"	15	"MACAIN"		
562	"MANGZE"	97	"SYZYCY"	15	"LASIOL"		
539	"CRYPWI"	97	"MEMEVA"	15	"EUGEIN"		
512	"LITSLO"	97	"AXINZE"	15	"CHAEFE"		
500	"LEEAIN"	95	"SYMPHI"	14	"ADENBI"		
478	"ALLOZE"	94	"ISONLA"	13	"HOMACE"		
472	"SHORST"	91	"PSYCPL"	12	"SYZYCA"		
464	"BHESCE"	90	"GLYCPE"	12	"PERSMA"		
463	"HYDNOC"	88	"POMETO"	12	"CARABR"		
453	"SHORCN"	87	"CARACA"	11	"RANDGA"		
429	"DIOSHI"	86	"VITEAL"	11	"MEMEGR"		
412	"CHAECO"	86	"GONIHO"	11	"CANAZE"		
394	"NARGMA"	80	"SEMENI"	10	"DESMEL"		
392	"HOPEJU"	80	"GARCSP"	9	"MICRWA"		
374	"SYZYWI"	77	"ELAREGL"	9	"FICUHI"		
373	"SYZYSP"	75	"LITSGA"	7	"MACAPE"		
345	"IXORJU"	72	"DIPTHI"	7	"GLOCB_"		
328	"CHAECA"	71	"ERYTOB"	7	"DIOSCH"		
311	"PSEUCH"	70	"SYZYAQ"	6	"ARTOGO"		
297	"APAMSI"	68	"MEMECL"	5	"SWIEMA"		
295	"GARCMO"	68	"AGLAAP"	5	"DIOSSY"		
294	"CINNCA"	67	"CINNDU"	5	"DILLTR"		
287	"NOTHBE"	66	"DIOSAL"	5	"CHRYRO"		
287	"GLENUN"	63	"PSYCGL"	5	"BRIDMO"		
282	"LITSIT"	63	"HOPEDI"	5	"APORCA"		

```
#!/usr/bin/env python

# Calculates species-individual curve data by repeated, random sampling.

# Usage:
# ./species_individual.py trials step [frequencies]
#
# trials          number of trials for a given sample size
# step            increment of sample size
# frequencies     list (space separated) of frequencies of species

from os import sys
from statistics import stdev
import random
import math

trials = int(sys.argv[1])
step = int(sys.argv[2])
data = map(int, sys.argv[3:])
n = sum(data)

individuals = []
for i, frequency in enumerate(data):
    individuals += [i] * frequency

def count_unique(l):
    return len(set(l))

for i in range(step, n, step):
    counts = []
    for j in range(trials):
        counts.append(count_unique(random.sample(individuals, i)))
    s = stdev(counts)
    if s == 0:
        s = 1e-12
    print '{}\t{}\t{}'.format(i, 1.0 * sum(counts) / trials, s)
```

```
#!/usr/bin/env python

# Calculates the shannon index of diversity for given data.

# Usage:
# ./shannon.py [frequencies]
#
# frequencies     list (space separated) of frequencies of species

from os import sys
import math

data = map(int, sys.argv[1:])
n = sum(data)
h = 0

for x in data:
    p = x * 1.0 / n
    h += p * math.log(p)

print -h
```