Module "4": Assignment 1 Answer Key

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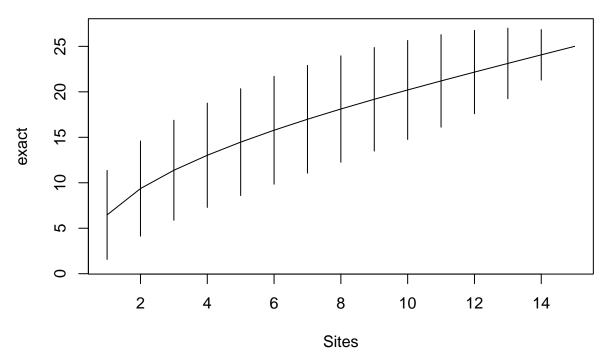
3.

##		homo_sapien canis_:	familiaris	felis_catus	canis_latrans		
##	1	3	0	0	18		
##	2	10	5	1	103		
##	3	47	2	1	21		
##	4	28	10	0	41		
##	5	114	16	0	40		
##	6	14	0	2	2		
##		<pre>geococcyx_californianus pecari_tajacu spermophilus_tereticaudus</pre>					
##	1		0	0		0	
##	2		3	2		0	
##	3		1	2		0	
##	4		0	0		0	
##	5		8	4		0	
##	6		0	10		0	
##		sylvilagus_audubon	_		naemorhous_mexication	anus lynx_r	ufus
##	1		47	3		0	0
##	2	•	77	0		0	0
##	3	•	48	0		0	0
##	4		6	0		0	0
##	5		55	0		2	2
##	6		67	0		0	4
##		<pre>procyon_lotor equs</pre>	_caballus p	ipilo_aberti	i otospermophilu	s_variegatu	IS
##		0	0	()		0
##		0	0	()		0
##		0	1	()		0
##		0	3	()		0
##	5	2	0)		0
##	6	0	0		3		52
##		spilogale_putorius	butorides_		onotrichia_leuco		_
##	_	0		0		0	0
##		0		0		0	0
##		0		0		0	0
##		0		0		0	0
##		0		1		0	0
##	6	0	_	0		2	0
##		anas_platyrhynchos	sayornis_n	_	= =	-	
	1	0		0	0	0	
##		^		^			
##	2	0		0	0	0	
	2	0 0 0		0 0 0	0	0)

```
## 5
                                            0
                        0
                                                                             0
## 6
                        0
                                                                 0
                                             1
                                                                             1
##
     junco_hyemalis campylorhynchus_brunneicapillus
## 1
                   0
## 2
                   0
                                                      0
## 3
                   0
                                                      0
## 4
                   0
                                                      0
                   0
                                                      0
## 5
## 6
                   1
                                                      0
  5.
## Warning in cor(x > 0): the standard deviation is zero
```

```
## Species Accumulation Curve
## Accumulation method: exact
## Call: specaccum(comm = scr)
##
##
## Sites
            1.000000 2.000000 3.000000 4.000000 5.000000 6.000000 7.000000
## Richness 6.466667 9.361905 11.375824 13.024176 14.470529 15.778821 16.984149
## sd
            2.445858 2.615566 2.749919 2.866508 2.936565 2.966045 2.962647
##
            8.000000 9.000000 10.000000 11.000000 12.000000 13.000000 14.000000
## Sites
## Richness 18.111422 19.180020 20.205128 21.198535 22.169231 23.123810 24.066667
            2.923536 2.845109 2.719706 2.542279 2.290225 1.937607 1.388844
## sd
##
## Sites
            15
## Richness 25
            0
## sd
```

6.



7.

Species chao chao.se jack1 jack1.se jack2 boot boot.se n ## All 25 116.4667 104.1535 38.06667 6.191033 49.39524 30.15487 2.9969 15

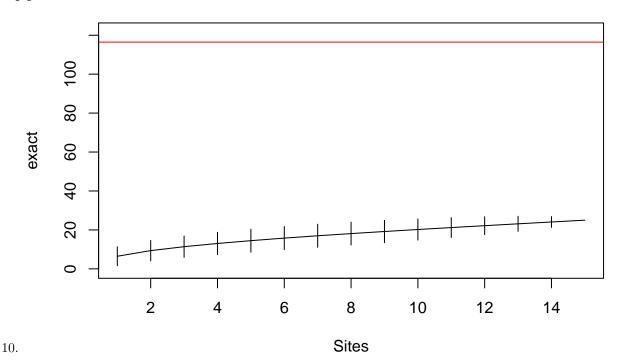
8.

[1] 25

[1] 116.4667

9.

[1] 91.46667



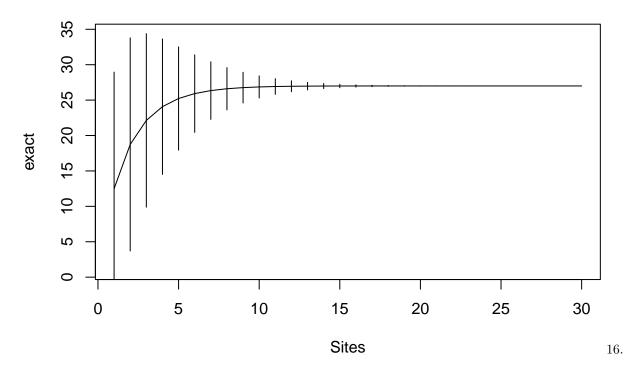
11.

CHA TRU VAI LOC OMB BLA HOT TOX VAN CHE BAR SPI GOU BRO PER BOU PSO ROT CAR ## 1 ## 2 ## 3 ## 4 ## 5 ## 6 ## TAN BCO PCH GRE GAR BBO ABL ANG ## 1 ## 2 ## 3 ## 4 ## 5 ## 6

14.

```
## Species Accumulation Curve
## Accumulation method: exact
## Call: specaccum(comm = fish)
##
##
             1.000000 2.000000 3.000000 4.000000 5.000000 6.000000 7.000000
## Sites
## Richness 12.500000 18.742529 22.133498 24.076665 25.226910 25.921113 26.344608
             8.224962 7.521837 6.125741 4.780134 3.645886 2.738364 2.032045
## sd
##
## Sites
             8.000000 9.000000 10.000000 11.000000 12.000000 13.000000 14.000000
## Richness 26.604116 26.763081 26.860035 26.918697 26.953776 26.974418 26.986307
             1.491678 1.083656 0.778985 0.553784 0.388989 0.269569 0.183966
## sd
##
            15.000000 16.00000 17.000000 18.000000 19.000000 20.000000 21.000000
## Sites
## Richness 26.992969 26.99657 26.998432 26.999339 26.999749 26.999917 26.999978
             0.123247 \quad 0.08070 \quad 0.051322 \quad 0.031424 \quad 0.018253 \quad 0.009885 \quad 0.004868
##
## Sites
            22.000000 2.30e+01 24 25 26 27 28 29 30
## Richness 26.999996 2.70e+01 27 27 27 27 27 27 27
            0.002031 7.01e-04 0 0 0 0 0 0 0
## sd
```

15.



Species chao chao.se jack1 jack1.se jack2 boot boot.se n ## All 27 27 0 27 0 27 27.00065 0.02849795 30

17.

[1] 27

[1] 27

18.

[1] 0

19.

