

STATS 205: Homework Assignment 5

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Solution to Problem 1

We say that two observations X_1 and X_2 are *independent* of one another with respect to a collection of events \mathcal{A} if

$$Pr \{X_1 \in A \text{ and } X_2 \in B\} = Pr \{X_1 \in A\} Pr \{X_2 \in B\}$$

where A and B are any two not necessarily distinct sets of outcomes belonging to \mathcal{A}^3 .

– 2.2.1 Independent Observations; Permutation, Parametric, and Bootstrap Tests of Hypotheses; Good, Phillip I

In deciding whether your own observations are exchangeable and a permutation test applicable, the key question is the one we posed in the very first chapter: Under the null hypothesis of no differences among the various experimental or survey groups, can we exchange the labels on the observations without significantly affecting the results?

– 2.2.2 Exchangeable Observations; Permutation, Parametric, and Bootstrap Tests of Hypotheses; Good, Phillip I

Solution to Problem 2

```
cysticerici <- c(28.9, 32.8, 12.0, 9.9, 15.0, 38.0, 12.5, 36.5, 8.6, 26.8);cysticerici
```

```
## [1] 28.9 32.8 12.0 9.9 15.0 38.0 12.5 36.5 8.6 26.8
```

```
worms_reco <- c(1.0, 7.7, 7.3, 7.9, 1.1, 3.5, 18.9, 33.9, 28.6, 25.0); worms_reco
```

```
## [1] 1.0 7.7 7.3 7.9 1.1 3.5 18.9 33.9 28.6 25.0
```

The null hypothesis is that the mean weight of introduced cysticerici *has no correlation with* the mean weight of worms recovered. That is,

$$H_0 : \tau = 0$$

The alternative hypothesis is that the mean weight of introduced cysticerici is *positively correlated with* the mean weight of worms recovered. That is,

$$H_A : \tau > 0$$

To test the null hypothesis against the alternative hypothesis, we will use the Kendall test, a distribution-free test for independence based on signs.

```
cor.test(x = cysticerici, y = worms_reco, method = "kendall", alt = "greater")
```

```
##
```

```
## Kendall's rank correlation tau
```

```
##
```

```
## data: cysticerci and worms_reco
## T = 19, p-value = 0.7578
## alternative hypothesis: true tau is greater than 0
## sample estimates:
##      tau
## -0.1555556
```

The p -value is 0.7578, which is *not* significant at the $\alpha = 0.05$ level. There is *not enough* evidence that the mean weight of introduced cysticerci is *positively correlated with* the mean weight of worms recovered.

Solution to Problem 3

```
cysticerci <- c(28.9, 32.8, 12.0, 9.9, 15.0, 38.0, 12.5, 36.5, 8.6, 26.8)
worms_reco <- c(1.0, 7.7, 7.3, 7.9, 1.1, 3.5, 18.9, 33.9, 28.6, 25.0)
cor.test(x = cysticerci, y = worms_reco, method = "kendall", alt = "greater")
```

```
##
## Kendall's rank correlation tau
##
## data: cysticerci and worms_reco
## T = 19, p-value = 0.7578
## alternative hypothesis: true tau is greater than 0
## sample estimates:
##      tau
## -0.1555556
```

The estimate for $\tau = -0.1555556$.

Solution to Problem 4

```
brain_weight = c(515, 286, 469, 410, 461, 436, 479, 198, 389, 262, 536); length(brain_weight)

## [1] 11

fiber_count = c(32500, 26800, 11410, 14850, 23640, 23820, 29840, 21830, 24650, 22500, 26000); length(fiber_count)

## [1] 11

library(bootstrap)
theta.hat = cor(brain_weight, fiber_count); theta.hat

## [1] 0.1604644

library(partitions)
n = length(brain_weight)
allCompositions = compositions(n, n); allCompositions[,1:length(brain_weight)]

##      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10] [,11]
## [1,]   11   10    9    8    7    6    5    4    3    2    1
## [2,]    0    1    2    3    4    5    6    7    8    9   10
## [3,]    0    0    0    0    0    0    0    0    0    0    0
## [4,]    0    0    0    0    0    0    0    0    0    0    0
## [5,]    0    0    0    0    0    0    0    0    0    0    0
## [6,]    0    0    0    0    0    0    0    0    0    0    0
## [7,]    0    0    0    0    0    0    0    0    0    0    0
```

```
## [8,] 0 0 0 0 0 0 0 0 0 0 0
## [9,] 0 0 0 0 0 0 0 0 0 0 0
## [10,] 0 0 0 0 0 0 0 0 0 0 0
## [11,] 0 0 0 0 0 0 0 0 0 0 0

allCompositions.sub = allCompositions[, sample(1:dim(allCompositions)[2], size=1000, replace=FALSE)]

draw.bootstrap.samples = function(df){
  n = dim(df)[1]
  ind = sample(n, replace = TRUE)
  cor.bootstrap.replicate = cor(df[ind, "LSAT"], df[ind, "GPA"])
  return(cor.bootstrap.replicate)
}

R = 1000
theta.hat.star = replicate(R, draw.bootstrap.samples(law))
# make a ggplot
library(ggplot2)

## Registered S3 methods overwritten by 'ggplot2':
## method from
## [.quosures rlang
## c.quosures rlang
## print.quosures rlang

theta.hat.star.df = data.frame(theta.hat.star = theta.hat.star)
theta.hat.star

## [1] 0.8484063 0.6316826 0.9825900 0.8255229 0.8495999 0.8261002
## [7] 0.8661347 0.8834630 0.9006001 0.5907339 0.7830618 0.8956721
## [13] 0.8390269 0.6767403 0.6655151 0.7880434 0.7669794 0.8333284
## [19] 0.4788920 0.7714328 0.6888578 0.6085110 0.8870055 0.8891343
## [25] 0.9293760 0.8941580 0.7842890 0.9759656 0.6580838 0.8803008
## [31] 0.7278119 0.9764674 0.6583863 0.6168595 0.6598359 0.8303769
## [37] 0.7441387 0.9125698 0.7720787 0.8052933 0.7456710 0.9079140
## [43] 0.7810588 0.7018577 0.7022426 0.9012286 0.9088847 0.9309482
## [49] 0.3903147 0.9605292 0.8423109 0.9381429 0.7806078 0.9464569
## [55] 0.7778007 0.6670899 0.8605475 0.7628063 0.8569390 0.6204809
## [61] 0.7548218 0.4333586 0.9514910 0.2215294 0.9487501 0.8885582
## [67] 0.6819842 0.6990087 0.8783708 0.6023873 0.8013960 0.3813482
## [73] 0.7119660 0.8667703 0.7203806 0.6057199 0.8547396 0.9245165
## [79] 0.8545298 0.6373720 0.7845749 0.9768518 0.8438509 0.9895442
## [85] 0.8665138 0.9117396 0.7704158 0.8589641 0.7834545 0.8452853
## [91] 0.9354389 0.7198367 0.8875415 0.9335897 0.7231350 0.8398740
## [97] 0.8276306 0.8507689 0.8705201 0.6667573 0.6629954 0.7455497
## [103] 0.8135639 0.7826830 0.9558701 0.8636942 0.6215780 0.5386313
## [109] 0.7424817 0.6646911 0.8607529 0.9353608 0.7096247 0.3969845
## [115] 0.8585453 0.8808895 0.8671265 0.9466575 0.7951984 0.8093433
## [121] 0.6461451 0.7237113 0.6086657 0.6420466 0.8083532 0.8101672
## [127] 0.7055363 0.8455851 0.8672329 0.9316990 0.8591594 0.7003659
## [133] 0.8052631 0.7597974 0.8761399 0.7878997 0.9123433 0.6228996
## [139] 0.5804809 0.7679912 0.9705725 0.7407414 0.8154191 0.8546031
## [145] 0.9173857 0.8029028 0.8388353 0.6528412 0.5988332 0.7513465
## [151] 0.8022712 0.7863913 0.6940034 0.7627176 0.8931142 0.8630833
## [157] 0.7928282 0.6684376 0.9324184 0.6394841 0.8584064 0.9252122
## [163] 0.8969546 0.8636759 0.6429309 0.7967141 0.8145145 0.8791452
## [169] 0.8398039 0.9399158 0.9410929 0.9412756 0.6638791 0.8583365
```

```

## [175] 0.6737819 0.7699293 0.8347644 0.7492324 0.5921786 0.8119755
## [181] 0.6229308 0.4989414 0.8746569 0.7255861 0.8609801 0.8166173
## [187] 0.5334794 0.9104799 0.7811890 0.7510102 0.7829000 0.8339260
## [193] 0.8063302 0.8625832 0.7044331 0.9388681 0.9334327 0.7058491
## [199] 0.7693600 0.9093276 0.9033053 0.7995914 0.7252945 0.8471418
## [205] 0.6337806 0.7837673 0.9531264 0.8907437 0.8434617 0.7711600
## [211] 0.8188653 0.7063732 0.8508669 0.7798309 0.8505594 0.5847226
## [217] 0.7568213 0.9045626 0.9496660 0.6024775 0.8918807 0.7941885
## [223] 0.8873180 0.9107763 0.7447879 0.7336387 0.7119080 0.7073913
## [229] 0.7109964 0.8043846 0.5885016 0.4996746 0.7431657 0.4651076
## [235] 0.8195815 0.8087968 0.7689289 0.8219518 0.8281651 0.8376250
## [241] 0.8574205 0.7711048 0.8520056 0.8012227 0.6669427 0.6382964
## [247] 0.8190446 0.7861443 0.8600243 0.9412712 0.8640642 0.8747762
## [253] 0.8222612 0.6205937 0.6600079 0.7727376 0.7299584 0.9640808
## [259] 0.9863439 0.8388737 0.9288310 0.8073237 0.6721381 0.7015188
## [265] 0.5752959 0.6181041 0.7243230 0.7778983 0.7793896 0.9727994
## [271] 0.8436386 0.8879080 0.7565744 0.8116215 0.6835844 0.7701193
## [277] 0.7333859 0.5970968 0.8611212 0.6305111 0.8013348 0.4775221
## [283] 0.9328511 0.7057912 0.7118070 0.7650197 0.7758004 0.8572204
## [289] 0.8284686 0.7688830 0.7862307 0.9823026 0.8135034 0.9262331
## [295] 0.2935749 0.5046892 0.7306630 0.8373255 0.8605431 0.8600511
## [301] 0.5796348 0.8811059 0.8190785 0.8212030 0.8756490 0.8010216
## [307] 0.9301198 0.7029825 0.8037159 0.8432400 0.5073820 0.7282265
## [313] 0.8187329 0.7305424 0.7428246 0.9248807 0.6800046 0.7205126
## [319] 0.8390958 0.6623474 0.9082076 0.9042436 0.7754951 0.4234463
## [325] 0.4791721 0.5839406 0.6449879 0.8360886 0.8220325 0.8682514
## [331] 0.9462763 0.4927805 0.8702052 0.8297424 0.5743866 0.8638268
## [337] 0.6200188 0.8575615 0.8827770 0.8560487 0.7048624 0.8223029
## [343] 0.8107322 0.8474215 0.6446622 0.7949617 0.8398069 0.8037369
## [349] 0.5738176 0.9621100 0.9510679 0.8073415 0.9045813 0.8311709
## [355] 0.7908642 0.7723811 0.8965577 0.9379014 0.9125163 0.7155668
## [361] 0.9177321 0.9032003 0.7453182 0.8475961 0.6763958 0.9111765
## [367] 0.7215191 0.8975345 0.7182437 0.9327780 0.7749827 0.4613007
## [373] 0.7699416 0.6244720 0.9435499 0.7912792 0.9179509 0.9120472
## [379] 0.6592434 0.7020889 0.7949157 0.7660971 0.7632987 0.8142261
## [385] 0.6814763 0.7778572 0.8305780 0.8733795 0.8894003 0.4779211
## [391] 0.9264889 0.7010274 0.7922144 0.7724571 0.4337690 0.3142486
## [397] 0.8134615 0.7991898 0.3445492 0.7112154 0.8405469 0.8301388
## [403] 0.7269186 0.6638092 0.9012427 0.7914690 0.6230790 0.8195173
## [409] 0.8708369 0.6038329 0.8005954 0.9309686 0.8932810 0.9338088
## [415] 0.6814781 0.9188700 0.6429278 0.9313080 0.6516503 0.8313922
## [421] 0.7807945 0.8019565 0.6132260 0.6138717 0.8568591 0.6699759
## [427] 0.8126850 0.5933424 0.8600165 0.4065893 0.6663190 0.3710635
## [433] 0.7437318 0.7935980 0.4528464 0.8801452 0.9175498 0.6685032
## [439] 0.7230661 0.7272831 0.5718904 0.8832916 0.7332020 0.6743675
## [445] 0.3444414 0.8628954 0.8689840 0.9057621 0.9358893 0.6345076
## [451] 0.9175508 0.7385811 0.9232594 0.9593707 0.7216116 0.6891631
## [457] 0.6704509 0.4139246 0.9242060 0.9311526 0.7402539 0.6674717
## [463] 0.8841071 0.9151263 0.8195928 0.9084508 0.8873319 0.9204116
## [469] 0.8544382 0.7491610 0.9467864 0.6285715 0.6559344 0.6606244
## [475] 0.7933110 0.6867611 0.7557185 0.7935863 0.8332683 0.8130354
## [481] 0.6002205 0.7569397 0.7491856 0.9443432 0.7715007 0.8711288
## [487] 0.9059689 0.6856878 0.7493826 0.7578152 0.9789478 0.7794192
## [493] 0.8706300 0.9263026 0.8759279 0.4862369 0.6639006 0.5864141

```

```

## [499] 0.9066888 0.7381651 0.7182064 0.9228661 0.6913695 0.6998998
## [505] 0.7258716 0.9704164 0.8368530 0.8511458 0.6464158 0.4453887
## [511] 0.8752147 0.7443561 0.7616496 0.9006988 0.8803353 0.4960341
## [517] 0.9145048 0.6525035 0.8582500 0.9303695 0.9000041 0.6161071
## [523] 0.7345535 0.4711859 0.8747066 0.5995823 0.4896198 0.9320428
## [529] 0.8153481 0.6573873 0.8389320 0.7346586 0.8140948 0.7203778
## [535] 0.5798306 0.8660962 0.7715967 0.8515778 0.6739306 0.6711454
## [541] 0.7647722 0.6650198 0.9511903 0.9523745 0.6669188 0.8709629
## [547] 0.7654281 0.9348194 0.8651852 0.6298582 0.9241811 0.8866581
## [553] 0.8212606 0.9073364 0.8106002 0.4884523 0.8923018 0.8773432
## [559] 0.7272449 0.7785269 0.8209021 0.8376321 0.7624946 0.9607840
## [565] 0.9524087 0.8614988 0.6729730 0.8311742 0.7773052 0.5794988
## [571] 0.7323972 0.7686293 0.8580465 0.8619659 0.9010520 0.4769893
## [577] 0.7942936 0.5379205 0.9109302 0.5860928 0.7317986 0.9508078
## [583] 0.9096985 0.8032535 0.9052325 0.7539075 0.8297247 0.7278393
## [589] 0.8649035 0.5617298 0.7068490 0.6855866 0.6978611 0.7283221
## [595] 0.8676930 0.9195045 0.5448878 0.8920227 0.7750389 0.8268857
## [601] 0.7780530 0.9850521 0.8686554 0.7108613 0.6853134 0.5752779
## [607] 0.9046527 0.7171031 0.7826220 0.5600183 0.7368691 0.8900705
## [613] 0.8676643 0.4286164 0.6754064 0.5110668 0.9446405 0.9151668
## [619] 0.9158021 0.6438445 0.6674373 0.7676561 0.7805144 0.5157234
## [625] 0.7919814 0.8960291 0.7346113 0.7220085 0.8223600 0.6798229
## [631] 0.5823128 0.6697052 0.5936597 0.6532075 0.7010886 0.8505122
## [637] 0.6500789 0.6746217 0.8164387 0.8812137 0.5917140 0.8547538
## [643] 0.6146122 0.9554979 0.9550466 0.7423891 0.8707321 0.7445611
## [649] 0.7862246 0.7411131 0.7561633 0.5966908 0.8857497 0.7916467
## [655] 0.8607571 0.8229016 0.6133203 0.6422552 0.8299774 0.6258596
## [661] 0.9783156 0.8823634 0.9052889 0.9465145 0.8553889 0.9196763
## [667] 0.7281712 0.8185132 0.7976887 0.8373100 0.9329585 0.9085770
## [673] 0.7151374 0.5774354 0.8594549 0.9422850 0.8124203 0.7304725
## [679] 0.9195452 0.5686190 0.7741607 0.8244837 0.5209978 0.9634176
## [685] 0.6711526 0.6299782 0.9918361 0.7240248 0.9125133 0.9153378
## [691] 0.6313603 0.7279531 0.8245974 0.7958124 0.6075838 0.5402864
## [697] 0.5136750 0.9184433 0.6808469 0.8992837 0.7501455 0.8097748
## [703] 0.8573210 0.6143540 0.8232470 0.8872820 0.8845771 0.8229001
## [709] 0.7578870 0.9253479 0.5711217 0.8990287 0.4580262 0.8207627
## [715] 0.8425417 0.7745900 0.7072784 0.7937883 0.8725666 0.7029809
## [721] 0.6661814 0.7327877 0.6924548 0.9177936 0.3552199 0.7875509
## [727] 0.8985668 0.7727130 0.9532091 0.6890796 0.5399657 0.6172611
## [733] 0.7321934 0.8123198 0.8731853 0.5422638 0.8813413 0.5839453
## [739] 0.7234418 0.8506626 0.8863809 0.4429041 0.9468900 0.9132313
## [745] 0.7793572 0.8386513 0.7363930 0.9224509 0.8211493 0.8318534
## [751] 0.9054196 0.8238898 0.6307103 0.7839515 0.8506296 0.8085137
## [757] 0.7629900 0.8463466 0.8196959 0.7299904 0.7350514 0.8615653
## [763] 0.8864065 0.9231701 0.6505283 0.9351291 0.6261416 0.9401788
## [769] 0.9548218 0.5200688 0.9386668 0.5103738 0.7594599 0.5858365
## [775] 0.7973548 0.5150957 0.7861382 0.8141736 0.5875679 0.9021349
## [781] 0.6972597 0.9148199 0.7459760 0.8521829 0.9274202 0.7274581
## [787] 0.6557685 0.9848333 0.5815401 0.7764410 0.7728602 0.9132548
## [793] 0.5109258 0.7214993 0.7728011 0.5422207 0.7738965 0.6099208
## [799] 0.6224529 0.6362466 0.7730588 0.7149808 0.6859459 0.8378129
## [805] 0.8085028 0.8538206 0.9799279 0.6191026 0.9292810 0.6483710
## [811] 0.9150360 0.8773784 0.9632035 0.9179394 0.9416594 0.6888332
## [817] 0.9194930 0.7937190 0.7557179 0.9382845 0.7731015 0.7591713

```

```
## [823] 0.7718074 0.7982163 0.5895211 0.8682406 0.8105870 0.7836722
## [829] 0.8130234 0.6664242 0.9094402 0.9076426 0.7567746 0.5517168
## [835] 0.7234137 0.8803638 0.5812526 0.8841997 0.7971025 0.9285724
## [841] 0.6777402 0.4820797 0.8526137 0.9656699 0.8018406 0.9133694
## [847] 0.8066360 0.6796558 0.2821170 0.5500308 0.9659223 0.8080355
## [853] 0.7529393 0.9305615 0.8468762 0.7848164 0.6876661 0.9172720
## [859] 0.8940477 0.9411414 0.7188100 0.8267012 0.4774145 0.4832236
## [865] 0.5790943 0.7640526 0.7148584 0.4473728 0.6422627 0.7748095
## [871] 0.8609262 0.6587911 0.5590200 0.7680032 0.7388040 0.7033241
## [877] 0.4765397 0.9485446 0.9076330 0.7955476 0.9030230 0.8491559
## [883] 0.9028816 0.9570602 0.8388348 0.4999516 0.9084215 0.7718385
## [889] 0.9452761 0.7729639 0.4825567 0.8914957 0.8902164 0.6285405
## [895] 0.8772998 0.7694990 0.8855217 0.7115879 0.9256531 0.7271115
## [901] 0.6332688 0.8980526 0.7597185 0.9587484 0.3277952 0.6957912
## [907] 0.9436030 0.5329854 0.8547113 0.7705394 0.5850841 0.9441016
## [913] 0.9380508 0.9312880 0.9250382 0.8290341 0.4513537 0.8027011
## [919] 0.8796030 0.8941213 0.7429799 0.7718867 0.7303820 0.8976351
## [925] 0.7518645 0.9142553 0.9741537 0.7224712 0.8102082 0.8440600
## [931] 0.8740437 0.8203747 0.7776758 0.6225317 0.6047948 0.9554754
## [937] 0.6744197 0.8601355 0.8238910 0.6497458 0.7407098 0.7217674
## [943] 0.9189859 0.8172124 0.7787066 0.7413331 0.6520937 0.9774700
## [949] 0.8218603 0.7213974 0.7551289 0.8642981 0.8278792 0.7640785
## [955] 0.8054259 0.8014907 0.7326306 0.8021894 0.9288738 0.8657932
## [961] 0.8309316 0.8215137 0.8636100 0.7018080 0.6164522 0.7789584
## [967] 0.7206287 0.7261358 0.8920904 0.7588431 0.7755221 0.8870612
## [973] 0.8663629 0.7928683 0.6318836 0.7115789 0.6551894 0.9253604
## [979] 0.7122619 0.6910876 0.5971311 0.5430247 0.5675155 0.7052825
## [985] 0.8187125 0.6553523 0.5903269 0.9336316 0.6538122 0.8342932
## [991] 0.5852040 0.6441082 0.8765090 0.7962291 0.9175517 0.5701648
## [997] 0.9694874 0.8415724 0.8928194 0.5996846
```

```
theta.hat.star.df
```

```
##      theta.hat.star
## 1      0.8484063
## 2      0.6316826
## 3      0.9825900
## 4      0.8255229
## 5      0.8495999
## 6      0.8261002
## 7      0.8661347
## 8      0.8834630
## 9      0.9006001
## 10     0.5907339
## 11     0.7830618
## 12     0.8956721
## 13     0.8390269
## 14     0.6767403
## 15     0.6655151
## 16     0.7880434
## 17     0.7669794
## 18     0.8333284
## 19     0.4788920
## 20     0.7714328
## 21     0.6888578
```

## 22	0.6085110
## 23	0.8870055
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## 32	0.9764674
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## 37	0.7441387
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## 40	0.8052933
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## 45	0.7022426
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## 87	0.7704158
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## 812	0.8773784
## 813	0.9632035
## 814	0.9179394
## 815	0.9416594
## 816	0.6888332
## 817	0.9194930
## 818	0.7937190
## 819	0.7557179
## 820	0.9382845
## 821	0.7731015
## 822	0.7591713
## 823	0.7718074
## 824	0.7982163
## 825	0.5895211
## 826	0.8682406
## 827	0.8105870
## 828	0.7836722
## 829	0.8130234
## 830	0.6664242
## 831	0.9094402

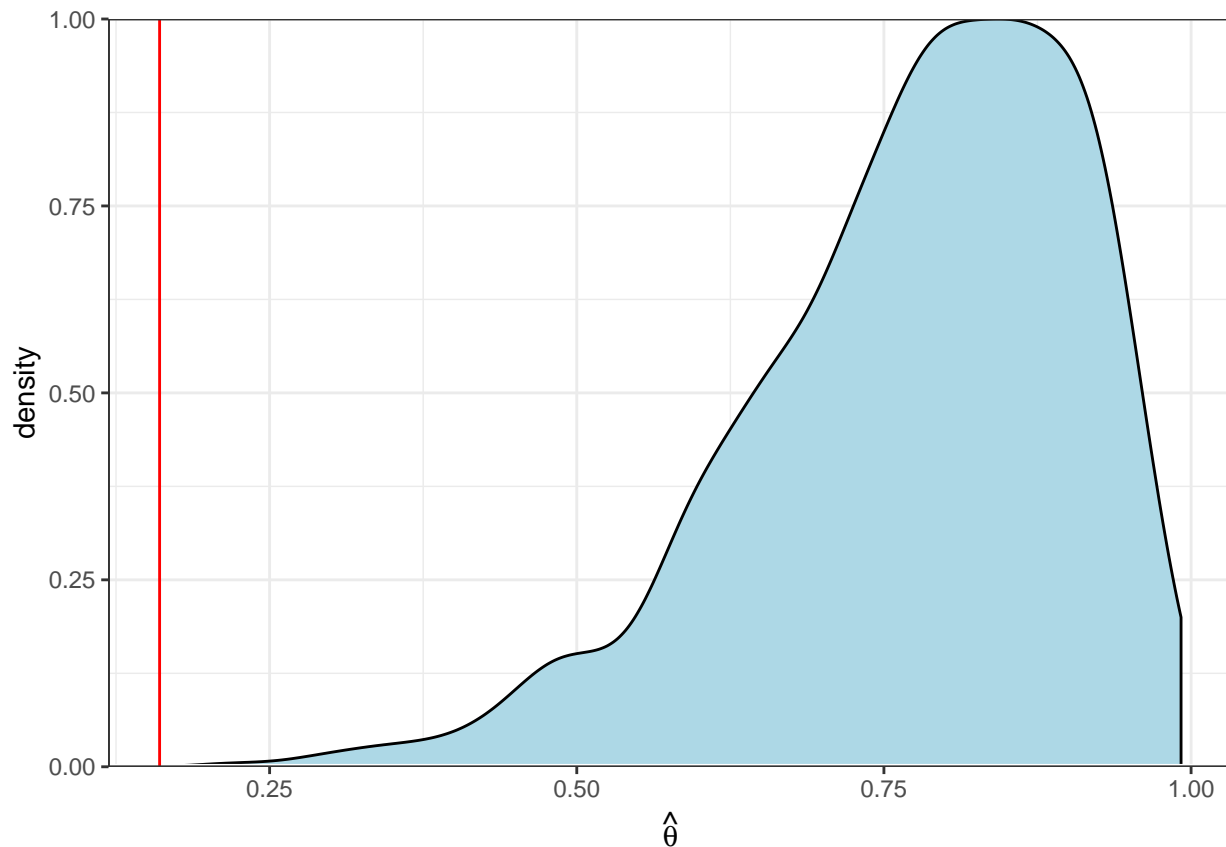
## 832	0.9076426
## 833	0.7567746
## 834	0.5517168
## 835	0.7234137
## 836	0.8803638
## 837	0.5812526
## 838	0.8841997
## 839	0.7971025
## 840	0.9285724
## 841	0.6777402
## 842	0.4820797
## 843	0.8526137
## 844	0.9656699
## 845	0.8018406
## 846	0.9133694
## 847	0.8066360
## 848	0.6796558
## 849	0.2821170
## 850	0.5500308
## 851	0.9659223
## 852	0.8080355
## 853	0.7529393
## 854	0.9305615
## 855	0.8468762
## 856	0.7848164
## 857	0.6876661
## 858	0.9172720
## 859	0.8940477
## 860	0.9411414
## 861	0.7188100
## 862	0.8267012
## 863	0.4774145
## 864	0.4832236
## 865	0.5790943
## 866	0.7640526
## 867	0.7148584
## 868	0.4473728
## 869	0.6422627
## 870	0.7748095
## 871	0.8609262
## 872	0.6587911
## 873	0.5590200
## 874	0.7680032
## 875	0.7388040
## 876	0.7033241
## 877	0.4765397
## 878	0.9485446
## 879	0.9076330
## 880	0.7955476
## 881	0.9030230
## 882	0.8491559
## 883	0.9028816
## 884	0.9570602
## 885	0.8388348

## 886	0.4999516
## 887	0.9084215
## 888	0.7718385
## 889	0.9452761
## 890	0.7729639
## 891	0.4825567
## 892	0.8914957
## 893	0.8902164
## 894	0.6285405
## 895	0.8772998
## 896	0.7694990
## 897	0.8855217
## 898	0.7115879
## 899	0.9256531
## 900	0.7271115
## 901	0.6332688
## 902	0.8980526
## 903	0.7597185
## 904	0.9587484
## 905	0.3277952
## 906	0.6957912
## 907	0.9436030
## 908	0.5329854
## 909	0.8547113
## 910	0.7705394
## 911	0.5850841
## 912	0.9441016
## 913	0.9380508
## 914	0.9312880
## 915	0.9250382
## 916	0.8290341
## 917	0.4513537
## 918	0.8027011
## 919	0.8796030
## 920	0.8941213
## 921	0.7429799
## 922	0.7718867
## 923	0.7303820
## 924	0.8976351
## 925	0.7518645
## 926	0.9142553
## 927	0.9741537
## 928	0.7224712
## 929	0.8102082
## 930	0.8440600
## 931	0.8740437
## 932	0.8203747
## 933	0.7776758
## 934	0.6225317
## 935	0.6047948
## 936	0.9554754
## 937	0.6744197
## 938	0.8601355
## 939	0.8238910

## 940	0.6497458
## 941	0.7407098
## 942	0.7217674
## 943	0.9189859
## 944	0.8172124
## 945	0.7787066
## 946	0.7413331
## 947	0.6520937
## 948	0.9774700
## 949	0.8218603
## 950	0.7213974
## 951	0.7551289
## 952	0.8642981
## 953	0.8278792
## 954	0.7640785
## 955	0.8054259
## 956	0.8014907
## 957	0.7326306
## 958	0.8021894
## 959	0.9288738
## 960	0.8657932
## 961	0.8309316
## 962	0.8215137
## 963	0.8636100
## 964	0.7018080
## 965	0.6164522
## 966	0.7789584
## 967	0.7206287
## 968	0.7261358
## 969	0.8920904
## 970	0.7588431
## 971	0.7755221
## 972	0.8870612
## 973	0.8663629
## 974	0.7928683
## 975	0.6318836
## 976	0.7115789
## 977	0.6551894
## 978	0.9253604
## 979	0.7122619
## 980	0.6910876
## 981	0.5971311
## 982	0.5430247
## 983	0.5675155
## 984	0.7052825
## 985	0.8187125
## 986	0.6553523
## 987	0.5903269
## 988	0.9336316
## 989	0.6538122
## 990	0.8342932
## 991	0.5852040
## 992	0.6441082
## 993	0.8765090


```
## 994      0.7962291
## 995      0.9175517
## 996      0.5701648
## 997      0.9694874
## 998      0.8415724
## 999      0.8928194
## 1000     0.5996846
```

```
ggplot(theta.hat.star.df) +
  geom_density(aes(x = theta.hat.star, y = ..scaled..),
    fill = "lightblue") +
  geom_hline(yintercept=0, colour="white", size=1) +
  theme_bw() +
  ylab("density") +
  xlab(bquote(hat(theta))) +
  geom_vline(xintercept = theta.hat, col = "red")+
  scale_y_continuous(expand = c(0,0))
```



Solution to Problem 5

```
cysticerici <- c(28.9, 32.8, 12.0, 9.9, 15.0, 38.0, 12.5, 36.5, 8.6, 26.8)
worms_reco <- c(1.0, 7.7, 7.3, 7.9, 1.1, 3.5, 18.9, 33.9, 28.6, 25.0)
```

The null hypothesis is that the mean weight of introduced cysticerici *has no correlation with* the mean weight of worms recovered. That is,

$$H_0 : r_s < r_{s,\alpha}$$

The alternative hypothesis is that the mean weight of introduced cysticerci is *positively correlated with* the mean weight of worms recovered. That is,

$$H_A : r_s \geq r_{s,\alpha}$$

Otherwise, do not reject.

To test the null hypothesis against the alternative hypothesis, we will use the Spearman test, a distribution-free test for independence based on ranks.

```
# this method of performing the test was given in the textbook
library(SuppDists)
qSpearman(p = 0.05, r = 10)
```

```
## [1] -0.5393939
```

Since $r_{s,\alpha} = -0.5393939$, we will reject the null hypothesis only if $r_s \geq -0.5393939$.

Calculating r_s ,

```
cor(x = cysticerci, y = worms_reco, method = "spearman")
```

```
## [1] -0.2
```

Since $r_s = -0.2$ and $r_{s,\alpha} = -0.5393939$, the statement $r_s \geq r_{s,\alpha}$ is *true*. Thus, we *reject* the null hypothesis. There is *sufficient* evidence that the mean weight of introduced cysticerci is *positively correlated with* the mean weight of worms recovered.

NOTE: At this point, I tried to use `cor.test()` with `method = "spearman"` but I got a different result than I expected, and I'm not sure why. Maybe I'm interpreting the output incorrectly?

```
cor.test(x = cysticerci, y = worms_reco, method = "spearman", alternative = "greater")
```

```
##
## Spearman's rank correlation rho
##
## data: cysticerci and worms_reco
## S = 198, p-value = 0.72
## alternative hypothesis: true rho is greater than 0
## sample estimates:
## rho
## -0.2
```

The p -value is 0.72, which is *not* significant at the $\alpha = 0.05$ level. There is *not enough* evidence that the mean weight of introduced cysticerci is *positively correlated with* the mean weight of worms recovered.

Solution to Problem 6

```
x = c(0, 5000, 10000, 15000, 20000, 25000, 30000, 100000)
y = c(0.924, 0.988, 0.992, 1.118, 1.133, 1.145, 1.157, 1.357)
```

The null hypothesis is that the mean weight of introduced cysticerici *has no correlation with* the mean weight of worms recovered. That is,

$$H_0 : \beta = \beta_0$$

$$H_0 : \beta = 0$$

The alternative hypothesis is that the mean weight of introduced cysticerici is *positively correlated with* the mean weight of worms recovered. That is,

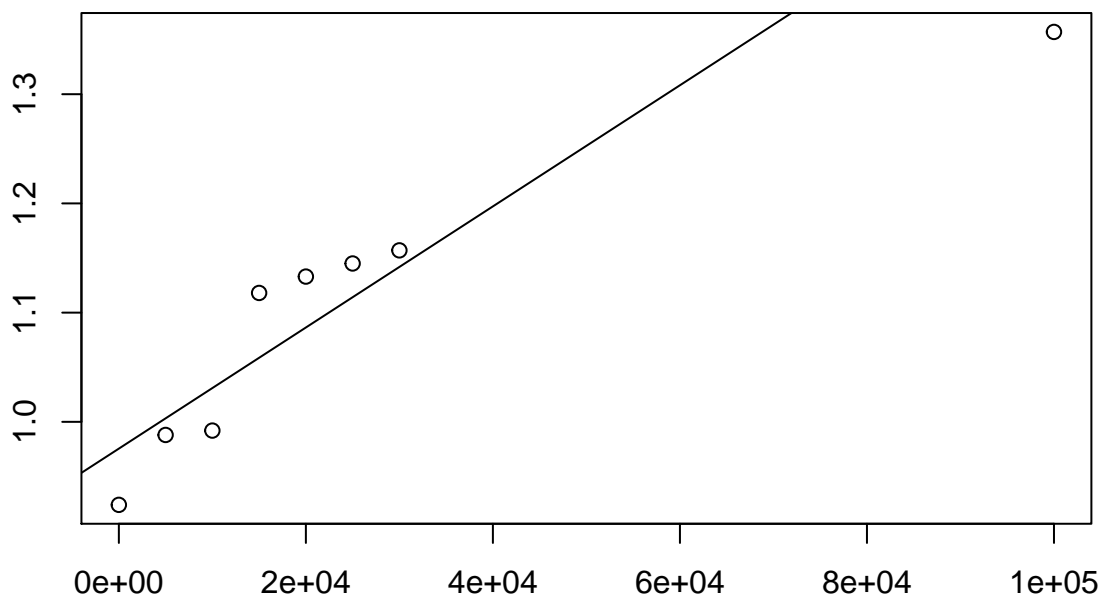
$$H_A : \beta > \beta_0$$

$$H_A : \beta > 0$$

To test the null hypothesis against the alternative hypothesis, we will use the Theil test, a distribution-free test for the slope of the regression line.

```
library(NSM3)
```

```
## Loading required package: combinat
##
## Attaching package: 'combinat'
## The following object is masked from 'package:utils':
##
##     combn
## Loading required package: MASS
## Loading required package: survival
## fANCOVA 0.5-1 loaded
theil(x, y, alpha=0.05, beta.0=0, type = "u")
```



```
## Alternative: beta greater than 0
## C = 28, C.bar = 1, P = 0
## beta.hat = 0
```

```
## alpha.hat = 0.975
##
## 1 - alpha = 0.95 upper bound for beta:
## -Inf, 0
```

```
theil.fit = theil(x,
  y,
  beta.0 = 0,
  slopes=TRUE,
  type = "u",
  doplot = FALSE)
theil.fit
```

```
## Alternative: beta greater than 0
## C = 28, C.bar = 1, P = 0
## beta.hat = 0
## alpha.hat = 0.975
##
```

```
## All slopes:
## i j S.ij
## 1 2 1.280000e-05
## 1 3 6.800000e-06
## 1 4 1.293333e-05
## 1 5 1.045000e-05
## 1 6 8.840000e-06
## 1 7 7.766667e-06
## 1 8 4.330000e-06
## 2 3 8.000000e-07
## 2 4 1.300000e-05
## 2 5 9.666667e-06
## 2 6 7.850000e-06
## 2 7 6.760000e-06
## 2 8 3.884211e-06
## 3 4 2.520000e-05
## 3 5 1.410000e-05
## 3 6 1.020000e-05
## 3 7 8.250000e-06
## 3 8 4.055556e-06
## 4 5 3.000000e-06
## 4 6 2.700000e-06
## 4 7 2.600000e-06
## 4 8 2.811765e-06
## 5 6 2.400000e-06
## 5 7 2.400000e-06
## 5 8 2.800000e-06
## 6 7 2.400000e-06
## 6 8 2.826667e-06
## 7 8 2.857143e-06
##
```

```
##
##
## 1 - alpha = 0.95 upper bound for beta:
## -Inf, 0
```

```
theil.output = theil(x,
  y,
```

```

beta.0 = 0,
slopes=TRUE,
type = "u", doplot = FALSE, alpha = .05)
c(theil.output$L, theil.output$U)

```

```
## [1] -Inf      0
```

TODO: Interpret these results correctly.

Solution to Problem 7

```

height = c(42.8, 63.5, 37.5, 39.5, 45.5, 38.5, 43.0, 22.5, 37.0, 23.5, 33.0, 58.0)
weight = c(40.0, 93.5, 35.5, 30.0, 52.0, 17.0, 38.5, 8.5, 33.0, 9.5, 21.0, 79.0)
heart_catheter_length = c(37.0, 49.5, 34.5, 36.0, 43.0, 28.0, 37.0, 20.0, 33.5, 30.5, 38.5, 47.0)

cor.test(x = height, y = heart_catheter_length, method = "pearson")

```

```

##
## Pearson's product-moment correlation
##
## data: height and heart_catheter_length
## t = 5.8936, df = 10, p-value = 0.0001524
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.6216270 0.9663721
## sample estimates:
##      cor
## 0.8811691

```

```
cor.test(x = weight, y = heart_catheter_length, method = "pearson")
```

```

##
## Pearson's product-moment correlation
##
## data: weight and heart_catheter_length
## t = 6.3033, df = 10, p-value = 8.871e-05
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
##  0.6568763 0.9700971
## sample estimates:
##      cor
## 0.8938226

```

From the Pearson correlation tests, there is *strong evidence* that, individually, height and weight contribute to the determination of heart catheter length.

```

library(Rfit)

r.01 <- rfit(heart_catheter_length ~ height)
f.01 <- rfit(heart_catheter_length ~ height + weight)
first_drop_test <- drop.test(f.01, r.01)
first_drop_test

```

```

##
## Drop in Dispersion Test

```

```
## F-Statistic      p-value
##      1.55202      0.24429

r.02 <- rfit(heart_catheter_length ~ weight)
second_drop_test <- drop.test(f.01, r.02)
second_drop_test
```

```
##
## Drop in Dispersion Test
## F-Statistic      p-value
##      0.014435      0.907007
```

However, based on the large p-values from the Drop in Dispersion tests, there is *not enough evidence* to suggest that height or weight contribute significantly over each other to the determination of heart catheter length.

Note

Treating length of heart catheter as the *independent* variable, test for the importance of height and weight in *determining* the required catheter length.

If height and weight are the *determiners* of length of heart catheter, length of heart catheter must be the *dependent* variable.