

StateByState

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This document gives an overall analyst to the b/g ratio over States

Load packages

```
require(data.table)
```

```
## Loading required package: data.table
```

```
## Warning: package 'data.table' was built under R version 3.1.3
```

```
require(agricolae)
```

```
## Loading required package: agricolae
```

```
## Warning: package 'agricolae' was built under R version 3.1.3
```

```
require(plyr)
```

```
## Loading required package: plyr
```

```
## Warning: package 'plyr' was built under R version 3.1.3
```

```
require(dplyr)
```

```
## Loading required package: dplyr
```

```
## Warning: package 'dplyr' was built under R version 3.1.3
```

```
##
```

```
## Attaching package: 'dplyr'
```

```
##
```

```
## The following objects are masked from 'package:plyr':
```

```
##
```

```
##      arrange, count, desc, failwith, id, mutate, rename, summarise,
```

```
##      summarize
```

```
##
```

```
## The following objects are masked from 'package:data.table':
```

```
##
```

```
##      between, last
```

```
##
```

```
## The following objects are masked from 'package:stats':
```

```
##
##      filter, lag
##
## The following objects are masked from 'package:base':
##
##      intersect, setdiff, setequal, union
```

Load data

```
partD_npi <- read.csv("F:/Academic/Stat 992/group project/prescription_fraud/analysis.py/NPI_bg.csv")
names(partD_npi)
```

```
## [1] "b.g" "b.g.with.claim"
## [3] "NPI" "NPPES_PROVIDER_LAST_ORG_NAME"
## [5] "NPPES_PROVIDER_FIRST_NAME" "NPPES_PROVIDER_CITY"
## [7] "NPPES_PROVIDER_STATE" "SPECIALTY_DESC"
## [9] "DESCRIPTION_FLAG" "BENE_COUNT"
## [11] "TOTAL_CLAIM_COUNT" "TOTAL_DAY_SUPPLY"
## [13] "TOTAL_DRUG_COST" "BENE_COUNT_GE65"
## [15] "BENE_COUNT_GE65_REDACT_FLAG" "TOTAL_CLAIM_COUNT_GE65"
## [17] "GE65_REDACT_FLAG" "TOTAL_DAY_SUPPLY_GE65"
## [19] "TOTAL_DRUG_COST_GE65"
```

ANOVA without counting claim

```
data <- cbind(partD_npi, trt=factor(partD_npi$NPPES_PROVIDER_STATE, labels=1:length(unique(partD_npi$NPPES_PROVIDER_STATE))))
model <- aov(b.g ~ trt, data = data)
summary(model)
```

```
##              Df Sum Sq Mean Sq F value Pr(>F)
## trt           60    234   3.903   55.26 <2e-16 ***
## Residuals    807959  57067   0.071
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

The model shows significant result to States ## Doing LSD and HSD test

```
lsd.out <- LSD.test(model, "trt", p.adj = "bonferroni")
hsd.out <- HSD.test(model, "trt", group = T)

#bar.group(lsd.out$groups, ylim=c(0,45), density=4, border="blue")
#bar.group(hsd.out$groups, ylim=c(0,45), density=4, border="blue")

#lsd.out$groups
#hsd.out$groups
```

match the group number with States with the data we use

```
id.group <- data %>%
  dplyr::select(NPPES_PROVIDER_STATE, trt) %>%
  unique() %>%
  arrange(trt)
groups <- as.data.frame(hsd.out$groups)
groups$trt <- as.integer(groups$trt)
id.group$trt <- as.integer(id.group$trt)
result <- plyr::join(groups, id.group)
```

Joining by: trt

result

##	trt	means	M	NPPES_PROVIDER_STATE
## 1	57	0.8296296	a	WI
## 2	1	0.7536630	a	AA
## 3	59	0.7460317	a	WY
## 4	8	0.7435175	a	AZ
## 5	41	0.7099463	a	NY
## 6	58	0.7042467	ab	WV
## 7	18	0.6931382	ab	IA
## 8	50	0.6802200	ab	TN
## 9	25	0.6795626	ab	MA
## 10	22	0.6737069	ab	KS
## 11	51	0.6691765	ab	TX
## 12	20	0.6686648	ab	IL
## 13	31	0.6632435	ab	MP
## 14	10	0.6620653	ab	CO
## 15	45	0.6607997	ab	PA
## 16	19	0.6589100	ab	ID
## 17	42	0.6584061	ab	OH
## 18	33	0.6558891	ab	MT
## 19	4	0.6534401	ab	AL
## 20	53	0.6524948	ab	VA
## 21	44	0.6508540	ab	OR
## 22	30	0.6506284	ab	MO
## 23	3	0.6492230	ab	AK
## 24	52	0.6489724	ab	UT
## 25	61	0.6479112	ab	ZZ
## 26	36	0.6475830	ab	NE
## 27	29	0.6462845	ab	MN
## 28	40	0.6455697	ab	NV
## 29	6	0.6453119	ab	AR
## 30	35	0.6446090	ab	ND
## 31	48	0.6442566	ab	SC
## 32	21	0.6441525	ab	IN
## 33	9	0.6422798	ab	CA
## 34	13	0.6404267	ab	DE
## 35	49	0.6398098	ab	SD
## 36	37	0.6395605	ab	NH

```
## 37 24 0.6385870 ab LA
## 38 16 0.6380884 ab GU
## 39 15 0.6362533 ab GA
## 40 27 0.6347261 ab ME
## 41 2 0.6345081 ab AE
## 42 39 0.6330565 ab NM
## 43 28 0.6327925 ab MI
## 44 60 0.6299534 ab XX
## 45 34 0.6294399 ab NC
## 46 43 0.6276517 ab OK
## 47 56 0.6271792 ab WA
## 48 11 0.6267211 ab CT
## 49 46 0.6261382 ab PR
## 50 5 0.6253142 ab AP
## 51 54 0.6246399 ab VI
## 52 32 0.6230888 ab MS
## 53 7 0.6223479 ab AS
## 54 14 0.6215336 ab FL
## 55 26 0.6209656 ab MD
## 56 55 0.6207932 ab VT
## 57 47 0.6204590 ab RI
## 58 38 0.6169197 ab NJ
## 59 23 0.6104840 ab KY
## 60 17 0.6103561 ab HI
## 61 12 0.5715370 ab DC
```

ANOVA with counting claim

```
data <- cbind(partD_npi, trt=factor(partD_npi$NPPE_PROVIDER_STATE, labels=1:length(unique(partD_npi$NPPE_PROVIDER_STATE))))
model <- aov(b.g.with.claim ~ trt, data = data)
summary(model)
```

```
##              Df      Sum Sq Mean Sq F value Pr(>F)
## trt          60 1.019e+08 1698278    134 <2e-16 ***
## Residuals   807959 1.024e+10   12675
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
```

The model shows significant result to States ## Doing LSD and HSD test

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lsd.out <- LSD.test(model, "trt", p.adj = "bonferroni")
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#bar.group(lsd.out$groups, ylim=c(0,45), density=4, border="blue")
#bar.group(hsd.out$groups, ylim=c(0,45), density=4, border="blue")

#lsd.out$groups
#hsd.out$groups
```

match the group number with States with the data we use

```
id.group <- data %>%
  dplyr::select(NPPES_PROVIDER_STATE, trt) %>%
  unique() %>%
  arrange(trt)
groups <- as.data.frame(hsd.out$groups)
groups$trt <- as.integer(groups$trt)
id.group$trt <- as.integer(id.group$trt)
result <- plyr::join(groups, id.group)
```

Joining by: trt

result

##	trt	means	M	NPPES_PROVIDER_STATE
## 1	41	94.588324	a	NY
## 2	34	79.634906	b	NC
## 3	56	72.097695	c	WA
## 4	26	67.739304	cd	MD
## 5	16	64.588452	de	GU
## 6	10	61.551558	def	CO
## 7	46	61.181660	ef	PR
## 8	54	60.247493	efg	VI
## 9	6	59.412172	fg	AR
## 10	7	58.030295	fg	AS
## 11	24	56.585151	fgh	LA
## 12	28	54.672548	ghi	MI
## 13	43	54.398429	ghij	OK
## 14	17	54.165338	ghij	HI
## 15	15	53.394622	ghij	GA
## 16	14	51.431418	hijk	FL
## 17	30	50.739024	hijkl	MO
## 18	38	49.895729	ijkl	NJ
## 19	37	49.478959	jkl	NH
## 20	35	48.827277	jklm	ND
## 21	29	48.100536	jklmn	MN
## 22	44	47.783472	jklmn	OR
## 23	40	47.585188	klmn	NV
## 24	42	46.637912	klmn	OH
## 25	47	45.725775	lmn	RI
## 26	49	44.721457	lmn	SD
## 27	11	43.562404	lmno	CT
## 28	61	43.001584	mno	ZZ
## 29	27	42.754208	mnop	ME
## 30	53	42.411970	mnop	VA
## 31	39	42.214424	mnop	NM
## 32	60	41.161856	nop	XX
## 33	9	40.064134	nopq	CA
## 34	13	39.604225	nopq	DE
## 35	33	39.155338	nopq	MT
## 36	21	38.944621	nopq	IN

## 37	22	38.857150	nopq	KS
## 38	20	38.226426	nopq	IL
## 39	36	37.668456	opq	NE
## 40	51	37.251280	opqr	TX
## 41	52	37.223366	opqr	UT
## 42	32	36.700499	opqr	MS
## 43	5	34.830763	opqrs	AP
## 44	48	34.106327	opqrs	SC
## 45	18	33.723717	pqrs	IA
## 46	55	32.463806	pqrs	VT
## 47	2	31.966822	qrs	AE
## 48	3	31.129564	qrs	AK
## 49	8	30.883649	qrst	AZ
## 50	31	30.392160	qrst	MP
## 51	19	27.389842	rst	ID
## 52	50	27.241004	rst	TN
## 53	59	25.824322	rst	WY
## 54	25	22.888835	rst	MA
## 55	23	21.081185	st	KY
## 56	58	17.593594	st	WV
## 57	4	17.087016	t	AL
## 58	12	8.538531	t	DC
## 59	57	6.279915	t	WI
## 60	1	3.832884	t	AA
## 61	45	2.962611	t	PA