

BIMI-6100 Elements in Biomedical Informatics

Fall 2023

David Crosslin

Tulane University School of Medicine
John W. Deming Department of Medicine
Division of Biomedical Informatics and Genomics New Orleans, LA

March 11, 2024

Contents

1	Hmisc describe function - rhc data	2
2	Hmisc summary function - rhc data	10
3	ols function output	11

List of Figures

1	plot(summary(m))	13
2	plot(anova(m))	13

List of Tables

1	Descriptive Statistics by sex	10
2	Effects Response : wblc1	12
3	Analysis of Variance for wblc1	12

1 Hmisc describe function - rhc data

rhc[, -1]
62 Variables 5735 Observations

cat1													
	n	missing	distinct										
	5735	0	9										
ARF (2490, 0.434), CHF (456, 0.080), Cirrhosis (224, 0.039), Colon Cancer (7, 0.001), Coma (436, 0.076), COPD (457, 0.080), Lung Cancer (39, 0.007), MOSF w/Malignancy (399, 0.070), MOSF w/Sepsis (1227, 0.214)													
cat2													
	n	missing	distinct										
	1200	4535	6										
Value				Cirrhosis	Colon Cancer			Coma			Lung Cancer		
Frequency				38	2			90			15		
Proportion				0.032	0.002			0.075			0.013		
Value				MOSF w/Malignancy	MOSF w/Sepsis								
Frequency				229	826								
Proportion				0.191	0.688								
ca													
	n	missing	distinct										
	5735	0	3										
Value				Metastatic	No	Yes							
Frequency				384	4379	972							
Proportion				0.067	0.764	0.169							
sadmte													
	n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
	5735	0	1401	1	11639	589.7	10831	10912	11164	11759	12097	12298	12363
lowest : 10754 10755 10756 10757 10758, highest: 12437 12438 12439 12440 12441													
dschdte													
	n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
	5734	1	1486	1	11660	589.2	10854	10939	11184	11777	12120	12319	12386
lowest : 10757 10758 10759 10763 10765, highest: 12492 12493 12505 12524 12560													
dthdte													
	n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
	3722	2013	1592	1	11754	619.4	10881	10991	11267	11832	12208	12418	12533
lowest : 10757 10758 10759 10763 10766, highest: 12772 12773 12778 12780 12783													
lstctdte													
	n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
	5735	0	932	1	11781	602.6	10942	11038	11316	11868	12244	12440	12522
lowest : 10756 10758 10762 10766 10768, highest: 12634 12638 12640 12642 12644													

death

n missing distinct
5735 0 2

Value No Yes
Frequency 2013 3722
Proportion 0.351 0.649

cardiohx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.436 1013 0.1766 0.2909

chfhx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.439 1021 0.178 0.2927

dementhx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.266 564 0.09834 0.1774

psychhx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.188 386 0.06731 0.1256

chrpulhx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.461 1089 0.1899 0.3077

renalhx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.127 255 0.04446 0.08499

liverhx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.195 401 0.06992 0.1301

gibledhx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.094 185 0.03226 0.06245

malighx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.53 1316 0.2295 0.3537

immunhx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.59 1543 0.269 0.3934

transhx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.306 662 0.1154 0.2042

amihx

n missing distinct Info Sum Mean Gmd
5735 0 2 0.101 200 0.03487 0.06733

age

n missing distinct Info Mean Gmd .05 .10 .25 .50 .75 .90 .95
5735 0 5036 1 61.38 18.86 30.02 36.97 50.15 64.05 73.93 80.88 85.00

lowest : 18.042 18.119 18.146 18.163 18.19 , highest: 97.84 98.089 100.246 100.849 101.848

sex

n missing distinct
5735 0 2

Value Female Male
Frequency 2543 3192
Proportion 0.443 0.557

edu

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	42	0.977	11.68	3.344	7	8	10	12	13	16	17

lowest : 0 1 2 3 4, highest: 24 25 26 27 30

surv2mdl

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	1522	1	0.5925	0.2177	0.2080	0.3100	0.4709	0.6280	0.7430	0.8165	0.8470

lowest : 0
highest: 0.938965 0.000999928 0.00199986 0.002 0.00299978
0.939 0.94 0.941 0.962

das2d3pc

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	1023	0.999	20.5	5.964	13.50	14.75	16.06	19.75	23.43	29.14	29.15

lowest : 11 12 12.5 13 13.5, highest: 30.5 31 31.5 32 33

t3d30

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	29	0.699	23.61	9.454	3	5	16	30	30	30	30

lowest : 2 3 4 5 6, highest: 26 27 28 29 30

dth30

n	missing	distinct
5735	0	2

Value	No	Yes
Frequency	3817	1918
Proportion	0.666	0.334

aps1

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	123	1	54.67	22.45	24	30	41	54	67	81	90

lowest : 3 4 6 7 8, highest: 127 129 130 135 147

scomal

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	11	0.839	21	29.49	0	0	0	0	41	61	100

Value	0	9	26	37	41	44	55	61	89	94	100
Frequency	3105	497	383	259	214	466	132	175	62	81	361
Proportion	0.541	0.087	0.067	0.045	0.037	0.081	0.023	0.031	0.011	0.014	0.063

For the frequency table, variable is rounded to the nearest 1

meanbp1

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	178	1	78.52	41.61	38	42	50	63	115	133	145

lowest : 0 10 13 15 17, highest: 210 212 222 242 259

wblc1

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	520	1	15.65	11.3	1.700	4.300	8.398	14.100	20.049	27.500	34.000

lowest : 0
highest: 100.391 0.0499954 0.0999908 0.109985 0.119995
112.188 171 178.594 192

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	189	1	115.2	45.1	45	58	97	124	141	160	171

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	72	0.997	28.09	15.87	8.0	8.4	14.0	30.0	38.0	44.0	50.0

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	118	0.999	37.62	1.984	34.90	35.40	36.09	38.09	39.00	39.59	40.00

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	1342	0.999	222.3	127.7	68.00	86.66	133.31	202.50	316.62	360.00	423.44

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	57	0.916	3.093	0.7435	1.8	2.0	2.6	3.5	3.5	3.5	3.9

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	450	1	31.87	9.214	21.00	23.00	26.10	30.00	36.30	44.59	47.03

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.9
5735	0	266	0.954	2.267	2.721	0.4000	0.5000	0.7999	1.0098	1.3999	4.0996	8.898

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	148	0.999	2.133	1.786	0.600	0.700	1.000	1.500	2.400	4.300	6.199

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	73	0.998	136.8	8.352	125	128	132	136	142	146	150

pot1

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	81	0.998	4.067	1.09	2.8	3.0	3.4	3.8	4.6	5.5	6.0

lowest : 1.09985 1.19995 1.3999 1.69995 1.7998 , highest: 9 9.29883 9.39844 10.3984 11.8984

paco21

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	266	0.998	38.75	12.98	23	26	31	37	42	53	63

lowest : 1 2 6 8 9 , highest: 133 148 151 155.281 156

ph1

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	96	0.999	7.388	0.1162	7.189	7.250	7.340	7.400	7.460	7.500	7.529

lowest : 6.5791 6.70996 6.75 6.76953 6.80957, highest: 7.69922 7.70996 7.71973 7.75 7.76953

swang1

n	missing	distinct
5735	0	2

Value	No	RHC
Frequency	3551	2184
Proportion	0.619	0.381

wtkilo1

n	missing	distinct	Info	Mean	Gmd	.05	.10	.25	.50	.75	.90	.95
5735	0	922	0.999	67.83	30.6	0.0	38.5	56.3	70.0	83.7	98.5	109.1

lowest : 0 19.5 24.1 24.3 26.2 , highest: 200 200.8 204.5 209.2 244

dnr1

n	missing	distinct
5735	0	2

Value	No	Yes
Frequency	5081	654
Proportion	0.886	0.114

ninsclas

n	missing	distinct
5735	0	6

Value	Medicaid	Medicare	Medicare & Medicaid	No insurance
Frequency	647	1458	374	322
Proportion	0.113	0.254	0.065	0.056

Value	Private	Private & Medicare
Frequency	1698	1236
Proportion	0.296	0.216

resp

n	missing	distinct
5735	0	2

Value	No	Yes
Frequency	3622	2113
Proportion	0.632	0.368

card

	n	missing	distinct
	5735	0	2

Value	No	Yes
Frequency	3804	1931
Proportion	0.663	0.337

neuro

	n	missing	distinct
	5735	0	2

Value	No	Yes
Frequency	5042	693
Proportion	0.879	0.121

gastr

	n	missing	distinct
	5735	0	2

Value	No	Yes
Frequency	4793	942
Proportion	0.836	0.164

renal

	n	missing	distinct
	5735	0	2

Value	No	Yes
Frequency	5440	295
Proportion	0.949	0.051

meta

	n	missing	distinct
	5735	0	2

Value	No	Yes
Frequency	5470	265
Proportion	0.954	0.046

hema

	n	missing	distinct
	5735	0	2

Value	No	Yes
Frequency	5381	354
Proportion	0.938	0.062

seps

	n	missing	distinct
	5735	0	2

Value	No	Yes
Frequency	4704	1031
Proportion	0.82	0.18

trauma
n missing distinct
5735 0 2

Value No Yes
Frequency 5683 52
Proportion 0.991 0.009

ortho
n missing distinct
5735 0 2

Value No Yes
Frequency 5728 7
Proportion 0.999 0.001

addl3p
n missing distinct Info Mean Gmd
1439 4296 8 0.832 1.182 1.699

Value 0.00 0.98 1.96 2.94 3.99 4.97 5.95 7.00
Frequency 778 296 130 43 49 64 53 26
Proportion 0.541 0.206 0.090 0.030 0.034 0.044 0.037 0.018

For the frequency table, variable is rounded to the nearest 0.07

urin1
n missing distinct Info Mean Gmd .05 .10 .25 .50 .75 .90 .95
2707 3028 1440 1 2192 1655 120.0 473.8 1110.0 1927.0 2955.0 4230.0 5078.5

lowest : 0 2.11 3 3.52 3.69, highest: 8340 8370 8392 8580 9000

race
n missing distinct
5735 0 3

Value black other white
Frequency 920 355 4460
Proportion 0.160 0.062 0.778

income
n missing distinct
5735 0 4

Value > \$50k \$11-\$25k \$25-\$50k Under \$11k
Frequency 451 1165 893 3226
Proportion 0.079 0.203 0.156 0.563

ptid
n missing distinct Info Mean Gmd .05 .10 .25 .50 .75 .90 .95
5735 0 5735 1 5134 3432 516.7 997.4 2561.5 5131.0 7689.0 9233.6 9786.9

lowest : 5 7 9 10 11, highest: 10270 10272 10273 10277 10278

Table 1: Descriptive Statistics by sex

	N	Female <i>N</i> = 2543			Male <i>N</i> = 3192			Combined <i>N</i> = 5735			Test Statistic
death : Yes	5735	64%	(1624)		66%	(2098)		65%	(3722)		$\chi_1^2=2.16$, P=0.141 ¹
race : black	5735	18%	(465)		14%	(455)		16%	(920)		$\chi_2^2=17.2$, P<0.001 ¹
other		6%	(157)		6%	(198)		6%	(355)		
white		76%	(1921)		80%	(2539)		78%	(4460)		
age	5735	50.4	64.8	74.9	49.9	63.4	73.0	50.1	64.0	73.9	$F_{1,5733}=9.63$, P=0.002 ²
bili1	5735	0.70	1.01	1.10	0.90	1.01	1.60	0.80	1.01	1.40	$F_{1,5733}=54.1$, P<0.001 ²
crea1	5735	0.8	1.3	2.3	1.1	1.6	2.5	1.0	1.5	2.4	$F_{1,5733}=101$, P<0.001 ²

a b c represent the lower quartile *a*, the median *b*, and the upper quartile *c* for continuous variables. *N* is the number of non-missing values. Numbers after percents are frequencies. Tests used: ¹Pearson test; ²Wilcoxon test

2 Hmisc summary function - rhc data

3 ols function output

Table 2: Effects Response : wblc1

	Low	High	Δ	Effect	S.E.	Lower 0.95	Upper 0.95
cardiohx	0	1	1	-0.934	0.412	-1.7420	-0.127
race — black:white	3	1		0.203	0.436	-0.6507	1.057
race — other:white	3	2		-2.077	0.654	-3.3590	-0.796
sex — Female:Male	2	1		0.521	0.316	-0.0998	1.141
income — > \$50k:Under \$11k	4	1		-1.059	0.600	-2.2349	0.117
income — \$11—\$25k:Under \$11k	4	2		0.435	0.408	-0.3644	1.234
income — \$25—\$50k:Under \$11k	4	3		-1.155	0.451	-2.0397	-0.271

Table 3: Analysis of Variance for wblc1

	d.f.	Partial SS	MS	F	P
race	2	1515	758	5.40	0.0045
sex	1	379	379	2.71	0.1000
income	3	1726	575	4.10	0.0064
cardiohx	1	721	721	5.14	0.0234
REGRESSION	7	4413	630	4.50	<0.0001
ERROR	5727	803152	140		

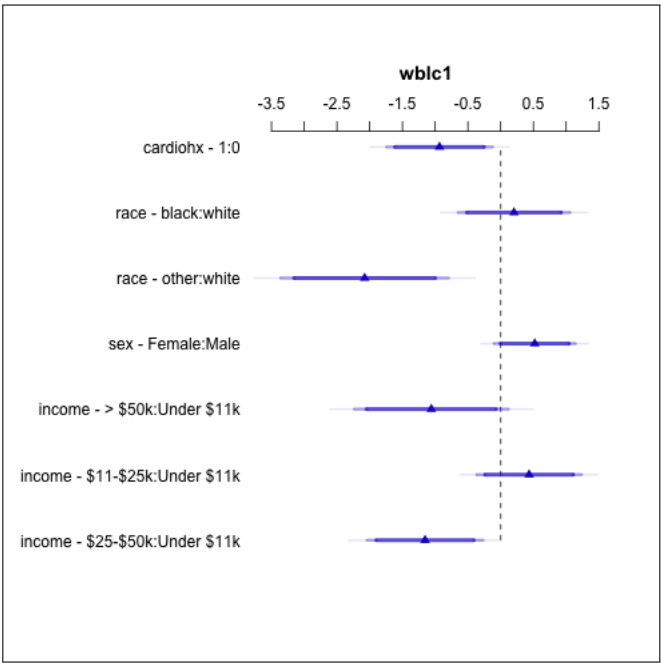


Figure 1: `plot(summary(m))`

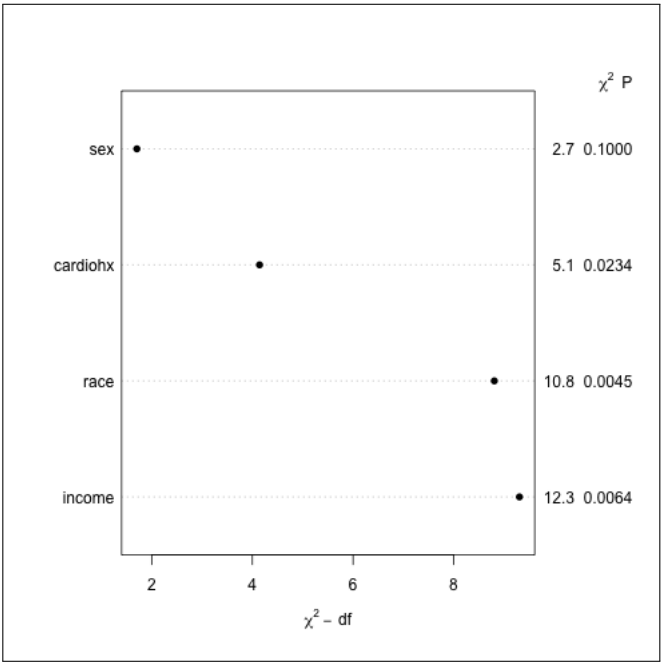


Figure 2: `plot(anova(m))`