ParaTest

Guillem

6 de febrero de 2020

Contraste de μ de normal con σ desconocida con t.test

```
# el parámetro alternative puede tomar tres valores: "two.sided" (por defecto, bilateral), "less" (mu <
set.seed(2020)
flores_ele <- sample(1:150, 40, replace = TRUE)
long_sep_mue <- iris[flores_ele,]$Sepal.Length</pre>
t.test(long_sep_mue, mu = 5.7, conf.level = 0.95, alternative = "greater")
##
##
  One Sample t-test
##
## data: long_sep_mue
## t = 2.5864, df = 39, p-value = 0.006773
## alternative hypothesis: true mean is greater than 5.7
## 95 percent confidence interval:
## 5.816769
                  Inf
## sample estimates:
## mean of x
##
       6.035
test_t <- t.test(long_sep_mue, mu = 5.7, conf.level = 0.95, alternative = "greater")
test_t$conf.int
## [1] 5.816769
## attr(,"conf.level")
## [1] 0.95
test_t$p.value
## [1] 0.006772975
```

Contrates de proporciones p con binom.test

```
binom.test(21, 25, p=0.8,alternative = "greater", conf.leve = 0.95)

##

## Exact binomial test

##

## data: 21 and 25

## number of successes = 21, number of trials = 25, p-value = 0.4207

## alternative hypothesis: true probability of success is greater than 0.8

## 95 percent confidence interval:

## 0.6703917 1.0000000
```

```
## sample estimates:
## probability of success
## 0.84
```

library(MASS)

birthwt

##		low	age	lwt	race	smoke	ptl	ht	ui	ftv	bwt
##	85	0	19	182	2	0	0	0	1	0	2523
##	86	0	33	155	3	0	0	0	0	3	2551
##	87	0	20	105	1	1	0	0	0	1	2557
##	88	0	21	108	1	1	0	0	1	2	2594
##	89	0	18	107	1	1	0	0	1	0	2600
##	91	0	21	124	3	0	0	0	0	0	2622
##	92	0	22	118	1	0	0	0	0	1	2637
##	93	0	17	103	3	0	0	0	0	1	2637
##	94	0	29	123	1	1	0	0	0	1	2663
##	95	0	26	113	1	1	0	0	0	0	2665
##	96	0	19	95	3	0	0	0	0	0	2722
##	97	0	19	150	3	0	0	0	0	1	2733
##	98	0	22	95	3	0	0	1	0	0	2751
##	99	0	30	107	3	0	1	0	1	2	2750
##	100	0	18	100	1	1	0	0	0	0	2769
##	101	0	18	100	1	1	0	0	0	0	2769
##	102	0	15	98	2	0	0	0	0	0	2778
##	103	0	25	118	1	1	0	0	0	3	2782
##	104	0	20	120	3	0	0	0	1	0	2807
##	105	0	28	120	1	1	0	0	0	1	2821
##	106	0	32	121	3	0	0	0	0	2	2835
##	107	0	31	100	1	0	0	0	1	3	2835
##	108	0	36	202	1	0	0	0	0	1	2836
##	109	0	28	120	3	0	0	0	0	0	2863
##	111	0	25	120	3	0	0	0	1	2	2877
##	112	0	28	167	1	0	0	0	0	0	2877
##	113	0	17	122	1	1	0	0	0	0	2906
##	114	0	29	150	1	0	0	0	0	2	2920
##	115	0	26	168	2	1	0	0	0	0	2920
## ##	116 117	0	17 17	113	2 2	0	0	0	0	1 1	2920 2920
##	117	0	24	113 90	1	1	1	0	0	1	2948
##	119	0	35	121	2	1	1	0	0	1	2948
##	120	0	25	155	1	0	0	0	0	1	2977
##	121	0	25	125	2	0	0	0	0	0	2977
##	123	0	29	140	1	1	0	0	0	2	2977
##	124	0	19	138	1	1	0	0	0	2	2977
##	125	0	27	124	1	1	0	0	0	0	2922
##	126	0	31	215	1	1	0	0	0	2	3005
##	127	0	33	109	1	1	0	0	0	1	3033
##	128	0	21	185	2	1	0	0	0	2	3042
##	129	0	19	189	1	0	0	0	0	2	3062
##	130	0	23	130	2	0	0	0	0	1	3062
##	131	0	21	160	1	0	0	0	0	0	3062
##	132	0	18	90	1	1	0	0	1	0	3062
##	133	0	18	90	1	1	0	0	1	0	3062
##	134	0	32	132	1	0	0	0	0	4	3080

##	135	0	19	132	3	0	0	0	0	0 3090
##	136	0	24	115	1	0	0	0	0	2 3090
##	137	0	22	85	3	1	0	0	0	0 3090
##	138	0	22	120	1	0	0	1	0	1 3100
##	139	0	23	128	3	0	0	0	0	0 3104
##	140	0	22	130	1	1	0	0	0	0 3132
##	141	0	30	95	1	1	0	0	0	2 3147
##	142	0	19	115	3	0	0	0	0	0 3175
##	143	0	16	110	3	0	0	0	0	0 3175
##	144	0	21	110	3	1	0	0	1	0 3203
##	145	0	30	153	3	0	0	0	0	0 3203
##	146	0	20	103	3	0	0	0	0	0 3203
##	147	0	17	119	3	0	0	0	0	0 3225
##	148	0	17	119	3	0	0	0	0	0 3225
##	149	0	23	119	3	0	0	0	0	2 3232
##	150	0	24	110	3	0	0	0	0	0 3232
##	151	0	28	140	1	0	0	0	0	0 3234
##	154	0	26	133	3	1	2	0	0	0 3260
##	155	0	20	169	3	0	1	0	1	1 3274
##	156	0	24	115	3	0	0	0	0	2 3274
##	159	0	28	250	3	1	0	0	0	6 3303
##	160	0	20	141	1	0	2	0	1	1 3317
##	161	0	22	158	2	0	1	0	0	2 3317
##	162	0	22	112	1	1	2	0	0	0 3317
##	163	0	31	150	3	1	0	0	0	2 3321
##	164	0	23	115	3	1	0	0	0	1 3331
##	166	0	16	112	2	0	0	0	0	0 3374
##	167	0	16	135	1	1	0	0	0	0 3374
##	168	0	18	229	2	0	0	0	0	0 3402
##	169	0	25	140	1	0	0	0	0	1 3416
##	170	0	32	134	1	1	1	0	0	4 3430
##	172	0	20	121	2	1	0	0	0	0 3444
##	173	0	23	190	1	0	0	0	0	0 3459
##	174	0	22	131	1	0	0	0	0	1 3460
##	175	0	32	170	1	0	0	0	0	0 3473
##	176	0	30	110	3	0	0	0	0	0 3544
##	177	0	20	127	3	0	0	0	0	0 3487
##	179	0	23	123	3	0	0	0	0	0 3544
##	180	0	23 17	120	3	1		0		0 3544
##	181	0			3		0	0	0	
			19	105		0			0	
##	182	0	23	130	1	0	0	0	0	0 3586
##	183	0	36	175	1	0	0	0	0	0 3600
##	184	0	22	125	1	0	0	0	0	1 3614
##	185	0	24	133	1	0	0	0	0	0 3614
##	186	0	21	134	3	0	0	0	0	2 3629
##	187	0	19	235	1	1	0	1	0	0 3629
##	188	0	25	95	1	1	3	0	1	0 3637
##	189	0	16	135	1	1	0	0	0	0 3643
##	190	0	29	135	1	0	0	0	0	1 3651
##	191	0	29	154	1	0	0	0	0	1 3651
##	192	0	19	147	1	1	0	0	0	0 3651
##	193	0	19	147	1	1	0	0	0	0 3651
##	195	0	30	137	1	0	0	0	0	1 3699
##	196	0	24	110	1	0	0	0	0	1 3728

## 197	0	19 184	1	1	0	1	0	0 3756
## 199	0	24 110	3	0	1	0	0	0 3770
## 200	0	23 110	1	0	0	0	0	1 3770
## 201	0	20 120	3	0	0	0	0	0 3770
## 202	0	25 241	2	0	0	1	0	0 3790
## 203	0	30 112	1	0	0	0	0	1 3799
## 204	0	22 169	1	0	0	0	0	0 3827
## 205	0	18 120	1	1	0	0	0	2 3856
## 206	0	16 170	2	0	0	0	0	4 3860
## 207	0	32 186	1	0	0	0	0	2 3860
## 208	0	18 120	3	0	0	0	0	1 3884
## 209	0	29 130	1	1	0	0	0	2 3884
## 203	0	33 117	1	0	0	0	1	1 3912
## 210 ## 211	0		1	1	0	0	0	0 3940
## 212	0	28 134	3	0	0	0	0	1 3941
## 213	0	14 135	1	0	0	0	0	0 3941
## 214	0	28 130	3	0	0	0	0	0 3969
## 215	0	25 120	1	0	0	0	0	2 3983
## 216	0	16 95	3	0	0	0	0	1 3997
## 217	0	20 158	1	0	0	0	0	1 3997
## 218	0	26 160	3	0	0	0	0	0 4054
## 219	0	21 115	1	0	0	0	0	1 4054
## 220	0	22 129	1	0	0	0	0	0 4111
## 221	0	25 130	1	0	0	0	0	2 4153
## 222	0	31 120	1	0	0	0	0	2 4167
## 223	0	35 170	1	0	1	0	0	1 4174
## 224	0	19 120	1	1	0	0	0	0 4238
## 225	0	24 116	1	0	0	0	0	1 4593
## 226	0	45 123	1	0	0	0	0	1 4990
## 4	1	28 120	3	1	1	0	1	0 709
## 10	1	29 130	1	0	0	0	1	2 1021
## 11	1	34 187	2	1	0	1	0	0 1135
## 13	1	25 105	3	0	1	1	0	0 1330
## 15	1	25 85	3	0	0	0	1	0 1474
## 16	1	27 150	3	0	0	0	0	0 1588
## 17	1	23 97	3	0	0	0	1	1 1588
## 18	1	24 128	2	0	1	0	0	1 1701
## 19	1	24 132	3	0	0	1	0	0 1729
## 20	1	21 165	1	1	0	1	0	1 1790
## 22	1	32 105	1	1	0	0	0	0 1818
## 23	1	19 91	1	1	2	0	1	0 1885
## 24	1	25 115	3	0	0	0	0	0 1893
## 2 4 ## 25	1	16 130	3	0	0	0	0	1 1899
## 26	1	25 92	1	1	0	0	0	0 1928
## 20 ## 27	1	20 150	1	1	0	0	0	2 1928
## 21 ## 28	1	20 150	2	0	0	0	1	2 1928
## 28 ## 29						0		
	1	24 155	1	1	1		0	
## 30 ## 31	1	21 103	3	0	0	0	0	0 1970
## 31	1	20 125	3	0	0	0	1	0 2055
## 32	1	25 89	3	0	2	0	0	1 2055
## 33	1	19 102	1	0	0	0	0	2 2082
## 34	1	19 112	1	1	0	0	1	0 2084
## 35	1	26 117	1	1	1	0	0	0 2084
## 36	1	24 138	1	0	0	0	0	0 2100

```
## 37
         1 17 130
                                             0 2125
## 40
            20 120
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## 42
            22 130
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            27 130
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## 49
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            18 148
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                                             0 2282
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         1
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## 51
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## 52
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## 54
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## 56
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            31 102
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## 57
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## 59
         1
            23 187
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## 60
            20 122
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## 61
         1 24 105
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                                             0 2381
## 62
         1 15 115
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## 63
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            23 120
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## 65
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            30 142
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## 67
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                                             1 2410
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## 68
         1 17 120
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            23 110
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                                             0 2424
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         1 17 120
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## 71
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         1 26 154
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## 76
            20 105
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## 79
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            28 95
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                                             2 2466
                       1
## 81
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            14 100
                       3
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## 82
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            23 94
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## 83
            17 142
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                                             0 2495
## 84
            21 130
                                  0
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                                     1
                                        0
         1
                       1
                              1
set.seed(1001)
madres_elegidas <- sample(1:189,40, replace = TRUE)</pre>
muestra_madres <- birthwt[madres_elegidas,]</pre>
x <- table(muestra_madres$smoke)[2]
binom.test(x,length(madres_elegidas), p=0.3, alternative = "greater")
##
##
    Exact binomial test
##
## data: x and length(madres_elegidas)
## number of successes = 17, number of trials = 40, p-value = 0.06331
## alternative hypothesis: true probability of success is greater than 0.3
  95 percent confidence interval:
    0.2918466 1.0000000
## sample estimates:
   probability of success
##
                     0.425
```

Rechazaremos la	H_0 ya	a que el	p_valor	es r	nenor	a 0.05	que	es el	nivel	de s	signific	ación	que	hemos	elegido