

R version 4.1.2 (2021-11-01) -- "Bird Hippie"
Copyright (C) 2021 The R Foundation for Statistical Computing
Platform: x86_64-w64-mingw32/x64 (64-bit)

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Natural language support but running in an English locale

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Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.

Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

[Previously saved workspace restored]

```
> > source("D:\\Dropbox\\Documents\\TAGS.R")
> TAGS()
```

TAGS V.2.0 is a R program developed by
R. Pouillot and G. Gerbier, Agence Française de Sécurité Sanitaire des Aliments, France.
r.pouillot@afssa.fr

Its purpose is to evaluate diagnostic tests in the absence of a gold standard,
using Maximum Likelihood Estimation (Newton Raphson and Expectation Maximisation
algorithms).

For further details: see
Pouillot R., Gerbier G. (2001)
'Tags' a program for validation of the diagnostic values of tests in the absence of a gold
standard.
Proceedings of the Society for Veterinary Epidemiology and Preventive Medicine,
Noordwijkerhout, The Netherlands, 28th - 30th March 2001: 37-48

Reference(s) population(s) data may be used (population(s) with a known infection status)

Evaluation may be used as soon as df>=parameters
A goodness-of-fit test and residual correlations are then provided

Three sets of data may be used as examples :

Hui and Walter (Biometrics, 1980, 36:167-171): Enter hui

Saegeaman et al (Vet Record, 1999, 145:214-8): Enter sae

Handelman's dentistry data (JADA, 1986, 113:751-754): Enter qu

If you want to enter new data: Enter new

If you want to use loaded data: Enter the name the program has already given to you

data Set ? new

Enter a name for your data : new_8_double

ENTER YOUR data

Number of tests (between 1 and 10) ? 8

Number of population without unknown status (between 1 and 10) ? 2

Do you have Reference Population(s) ?

No : Enter 0

Yes, Disease free : Enter 1

Yes, Infected : Enter 2

Yes, one Disease-free and one Infected : Enter 3

0

population with unknown status 1

Number of results 0 0 0 0 0 0 0 in population with unknown status 1 : 254

Number of results 1 0 0 0 0 0 0 in population with unknown status 1 : 0

Number of results 0 1 0 0 0 0 0 in population with unknown status 1 : 0

Number of results 1 1 0 0 0 0 0 in population with unknown status 1 : 0

Number of results 0 0 1 0 0 0 0 in population with unknown status 1 : 0

Number of results 1 0 1 0 0 0 0 in population with unknown status 1 : 0

Number of results 0 1 1 0 0 0 0 in population with unknown status 1 : 0

Number of results 1 1 1 0 0 0 0 in population with unknown status 1 : 0

Number of results 0 0 0 1 0 0 0 in population with unknown status 1 : 0

Number of results 1 0 0 1 0 0 0 in population with unknown status 1 : 0

Number of results 0 1 0 1 0 0 0 in population with unknown status 1 : 0

Number of results 1 1 0 1 0 0 0 in population with unknown status 1 : 0

Number of results 0 0 1 1 0 0 0 in population with unknown status 1 : 0

Number of results 1 0 1 1 0 0 0 in population with unknown status 1 : 0

Number of results 0 1 1 1 0 0 0 in population with unknown status 1 : 0

Number of results 1 1 1 1 0 0 0 in population with unknown status 1 : 0

Number of results 0 0 0 0 1 0 0 0 in population with unknown status 1 : 0

Number of results 1 0 0 0 1 0 0 0 in population with unknown status 1 : 0

Number of results 0 1 0 0 1 0 0 0 in population with unknown status 1 : 0

Number of results 1 1 0 0 1 0 0 0 in population with unknown status 1 : 0

Number of results 0 0 1 0 1 0 0 0 in population with unknown status 1 : 0

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254	1	0	1	1	1	1	1	1	0	0	0	0
255	0	1	1	1	1	1	1	1	0	0	0	0
256	1	1	1	1	1	1	1	1	44	31	0	0

Is it OK ?

No : Enter 0

Yes : Enter 1

1

Default Best Guess:

pre1 pre2 Sp1 Sp2 Sp3 Sp4 Sp5 Sp6 Sp7 Sp8 Se1 Se2 Se3 Se4

Best Guess 0.2 0.2 0.95 0.95 0.95 0.95 0.95 0.95 0.95 0.95 0.8 0.8 0.8 0.8

Se5 Se6 Se7 Se8

Best Guess 0.8 0.8 0.8 0.8

Do you have Best Guess?

No : Enter 0

Yes : Enter 1

0

Would you like Bootstrap Confidence intervals

(CAUTION: this may be time consuming. It is recommended to ask for it in a second step!)

No : Enter 0

Yes : Enter 1

1

How many iterations? (between 50 and 5000)

5000

DATA SUMMARY

2 Population(s); 8 Tests; 0 Reference Population(s)

df: 510 ; parameters: 18

3	0	1	0	0	0	0	0	0	0	0	0	0
4	1	1	0	0	0	0	0	0	0	0	0	0
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6	1	0	1	0	0	0	0	0	0	0	0	0
7	0	1	1	0	0	0	0	0	0	0	0	0
8	1	1	1	0	0	0	0	0	0	0	0	0
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23	0	1	1	0	1	0	0	0	0	0	0	0
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25	0	0	0	1	1	0	0	0	0	0	0	0
26	1	0	0	1	1	0	0	0	0	0	0	0
27	0	1	0	1	1	0	0	0	0	0	0	0
28	1	1	0	1	1	0	0	0	0	0	0	0
29	0	0	1	1	1	0	0	0	0	0	0	0
30	1	0	1	1	1	0	0	0	0	0	0	0
31	0	1	1	1	1	0	0	0	0	0	0	0
32	1	1	1	1	1	0	0	0	0	0	0	0
33	0	0	0	0	0	1	0	0	0	0	0	0
34	1	0	0	0	0	1	0	0	0	0	0	0
35	0	1	0	0	0	1	0	0	0	0	0	0
36	1	1	0	0	0	1	0	0	0	0	0	0
37	0	0	1	0	0	1	0	0	0	0	0	0
38	1	0	1	0	0	1	0	0	0	0	0	0
39	0	1	1	0	0	1	0	0	0	0	0	0
40	1	1	1	0	0	1	0	0	0	0	0	0
41	0	0	0	1	0	1	0	0	0	0	0	0
42	1	0	0	1	0	1	0	0	0	0	0	0
43	0	1	0	1	0	1	0	0	0	0	0	0
44	1	1	0	1	0	1	0	0	0	0	0	0
45	0	0	1	1	0	1	0	0	0	0	0	0
46	1	0	1	1	0	1	0	0	0	0	0	0

47	0	1	1	1	0	1	0	0	0	0	0	0
48	1	1	1	1	0	1	0	0	0	0	0	0
49	0	0	0	0	1	1	0	0	0	0	0	0
50	1	0	0	0	1	1	0	0	0	0	0	0
51	0	1	0	0	1	1	0	0	0	0	0	0
52	1	1	0	0	1	1	0	0	0	0	0	0
53	0	0	1	0	1	1	0	0	0	0	0	0
54	1	0	1	0	1	1	0	0	0	0	0	0
55	0	1	1	0	1	1	0	0	0	0	0	0
56	1	1	1	0	1	1	0	0	0	0	0	0
57	0	0	0	1	1	1	0	0	0	0	0	0
58	1	0	0	1	1	1	0	0	0	0	0	0
59	0	1	0	1	1	1	0	0	0	0	0	0
60	1	1	0	1	1	1	0	0	0	0	0	0
61	0	0	1	1	1	1	0	0	0	0	0	0
62	1	0	1	1	1	1	0	0	0	0	0	0
63	0	1	1	1	1	1	0	0	0	0	0	0
64	1	1	1	1	1	1	0	0	0	0	0	0
65	0	0	0	0	0	0	1	0	0	0	0	0
66	1	0	0	0	0	0	1	0	0	0	0	0
67	0	1	0	0	0	0	1	0	0	0	0	0
68	1	1	0	0	0	0	1	0	0	0	0	0
69	0	0	1	0	0	0	1	0	0	0	0	0
70	1	0	1	0	0	0	1	0	0	0	0	0
71	0	1	1	0	0	0	1	0	0	0	0	0
72	1	1	1	0	0	0	1	0	0	0	0	0
73	0	0	0	1	0	0	1	0	0	0	0	0
74	1	0	0	1	0	0	1	0	0	0	0	0
75	0	1	0	1	0	0	1	0	0	0	0	0
76	1	1	0	1	0	0	1	0	0	0	0	0
77	0	0	1	1	0	0	1	0	0	0	0	0
78	1	0	1	1	0	0	1	0	0	0	0	0
79	0	1	1	1	0	0	1	0	0	0	0	0
80	1	1	1	1	0	0	1	0	0	0	0	0
81	0	0	0	0	1	0	1	0	0	0	0	0
82	1	0	0	0	1	0	1	0	0	0	0	0
83	0	1	0	0	1	0	1	0	0	0	0	0
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86	1	0	1	0	1	0	1	0	0	0	0	0
87	0	1	1	0	1	0	1	0	0	0	0	0
88	1	1	1	0	1	0	1	0	0	0	0	0
89	0	0	0	1	1	0	1	0	0	0	0	0
90	1	0	0	1	1	0	1	0	0	0	0	0

91	0	1	0	1	1	0	1	0	0	0	0	0
92	1	1	0	1	1	0	1	0	0	0	0	0
93	0	0	1	1	1	0	1	0	0	0	0	0
94	1	0	1	1	1	0	1	0	0	0	0	0
95	0	1	1	1	1	0	1	0	0	0	0	0
96	1	1	1	1	1	0	1	0	0	0	0	0
97	0	0	0	0	0	1	1	0	0	0	0	0
98	1	0	0	0	0	1	1	0	0	0	0	0
99	0	1	0	0	0	1	1	0	0	0	0	0
100	1	1	0	0	0	1	1	0	0	0	0	0
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107	0	1	0	1	0	1	1	0	0	0	0	0
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109	0	0	1	1	0	1	1	0	0	2	0	0
110	1	0	1	1	0	1	1	0	0	0	0	0
111	0	1	1	1	0	1	1	0	1	0	0	0
112	1	1	1	1	0	1	1	0	0	0	0	0
113	0	0	0	0	1	1	1	0	0	0	0	0
114	1	0	0	0	1	1	1	0	0	0	0	0
115	0	1	0	0	1	1	1	0	0	0	0	0
116	1	1	0	0	1	1	1	0	0	0	0	0
117	0	0	1	0	1	1	1	0	0	0	0	0
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119	0	1	1	0	1	1	1	0	0	0	0	0
120	1	1	1	0	1	1	1	0	0	0	0	0
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122	1	0	0	1	1	1	1	0	0	0	0	0
123	0	1	0	1	1	1	1	0	0	0	0	0
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127	0	1	1	1	1	1	1	0	2	0	0	0
128	1	1	1	1	1	1	1	0	0	0	0	0
129	0	0	0	0	0	0	0	1	0	0	0	0
130	1	0	0	0	0	0	0	1	0	0	0	0
131	0	1	0	0	0	0	0	1	0	0	0	0
132	1	1	0	0	0	0	0	1	0	0	0	0
133	0	0	1	0	0	0	0	1	0	0	0	0
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136	1	1	1	0	0	0	0	1	0	0	0	0
137	0	0	0	1	0	0	0	1	0	0	0	0
138	1	0	0	1	0	0	0	1	0	0	0	0
139	0	1	0	1	0	0	0	1	0	0	0	0
140	1	1	0	1	0	0	0	1	0	0	0	0
141	0	0	1	1	0	0	0	1	0	0	0	0
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143	0	1	1	1	0	0	0	1	0	0	0	0
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147	0	1	0	0	1	0	0	1	0	0	0	0
148	1	1	0	0	1	0	0	1	0	0	0	0
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151	0	1	1	0	1	0	0	1	0	0	0	0
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156	1	1	0	1	1	0	0	1	0	0	0	0
157	0	0	1	1	1	0	0	1	0	0	0	0
158	1	0	1	1	1	0	0	1	0	0	0	0
159	0	1	1	1	1	0	0	1	0	0	0	0
160	1	1	1	1	1	0	0	1	0	0	0	0
161	0	0	0	0	0	1	0	1	0	0	0	0
162	1	0	0	0	0	0	1	0	1	0	0	0
163	0	1	0	0	0	1	0	1	6	4	0	0
164	1	1	0	0	0	1	0	1	0	0	0	0
165	0	0	1	0	0	1	0	1	0	0	0	0
166	1	0	1	0	0	1	0	1	0	0	0	0
167	0	1	1	0	0	1	0	1	0	0	0	0
168	1	1	1	0	0	1	0	1	0	0	0	0
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171	0	1	0	1	0	1	0	1	0	0	0	0
172	1	1	0	1	0	1	0	1	0	0	0	0
173	0	0	1	1	0	1	0	1	0	0	0	0
174	1	0	1	1	0	1	0	1	0	0	0	0
175	0	1	1	1	0	1	0	1	0	0	0	0
176	1	1	1	1	0	1	0	1	0	0	0	0
177	0	0	0	0	1	1	0	1	0	0	0	0
178	1	0	0	0	1	1	0	1	0	0	0	0

179	0	1	0	0	1	1	0	1	0	0	0	0
180	1	1	0	0	1	1	0	1	0	0	0	0
181	0	0	1	0	1	1	0	1	0	0	0	0
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184	1	1	1	0	1	1	0	1	0	0	0	0
185	0	0	0	1	1	1	0	1	0	0	0	0
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187	0	1	0	1	1	1	0	1	0	0	0	0
188	1	1	0	1	1	1	0	1	0	0	0	0
189	0	0	1	1	1	1	0	1	0	0	0	0
190	1	0	1	1	1	1	0	1	0	0	0	0
191	0	1	1	1	1	1	0	1	0	0	0	0
192	1	1	1	1	1	1	0	1	0	0	0	0
193	0	0	0	0	0	0	1	1	4	2	0	0
194	1	0	0	0	0	0	1	1	0	0	0	0
195	0	1	0	0	0	0	1	1	0	0	0	0
196	1	1	0	0	0	0	1	1	0	0	0	0
197	0	0	1	0	0	0	1	1	11	9	0	0
198	1	0	1	0	0	0	1	1	0	0	0	0
199	0	1	1	0	0	0	1	1	0	0	0	0
200	1	1	1	0	0	0	1	1	0	0	0	0
201	0	0	0	1	0	0	1	1	0	0	0	0
202	1	0	0	1	0	0	1	1	0	0	0	0
203	0	1	0	1	0	0	1	1	0	0	0	0
204	1	1	0	1	0	0	1	1	0	0	0	0
205	0	0	1	1	0	0	1	1	0	0	0	0
206	1	0	1	1	0	0	1	1	0	0	0	0
207	0	1	1	1	0	0	1	1	0	0	0	0
208	1	1	1	1	0	0	1	1	0	0	0	0
209	0	0	0	0	1	0	1	1	0	0	0	0
210	1	0	0	0	1	0	1	1	0	0	0	0
211	0	1	0	0	1	0	1	1	0	0	0	0
212	1	1	0	0	1	0	1	1	0	0	0	0
213	0	0	1	0	1	0	1	1	0	0	0	0
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215	0	1	1	0	1	0	1	1	0	0	0	0
216	1	1	1	0	1	0	1	1	0	0	0	0
217	0	0	0	1	1	0	1	1	0	0	0	0
218	1	0	0	1	1	0	1	1	0	0	0	0
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220	1	1	0	1	1	0	1	1	0	0	0	0
221	0	0	1	1	1	0	1	1	0	0	0	0
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223	0	1	1	1	1	0	1	1	0	0	0	0
224	1	1	1	1	1	0	1	1	0	0	0	0
225	0	0	0	0	0	1	1	1	0	0	0	0
226	1	0	0	0	0	1	1	1	0	0	0	0
227	0	1	0	0	0	1	1	1	0	0	0	0
228	1	1	0	0	0	1	1	1	0	0	0	0
229	0	0	1	0	0	1	1	1	0	0	0	0
230	1	0	1	0	0	1	1	1	0	0	0	0
231	0	1	1	0	0	1	1	1	0	0	0	0
232	1	1	1	0	0	1	1	1	7	1	0	0
233	0	0	0	1	0	1	1	1	0	0	0	0
234	1	0	0	1	0	1	1	1	0	1	0	0
235	0	1	0	1	0	1	1	1	0	0	0	0
236	1	1	0	1	0	1	1	1	0	0	0	0
237	0	0	1	1	0	1	1	1	0	0	0	0
238	1	0	1	1	0	1	1	1	0	0	0	0
239	0	1	1	1	0	1	1	1	0	0	0	0
240	1	1	1	1	0	1	1	1	7	10	0	0
241	0	0	0	0	1	1	1	1	0	0	0	0
242	1	0	0	0	1	1	1	1	0	0	0	0
243	0	1	0	0	1	1	1	1	0	0	0	0
244	1	1	0	0	1	1	1	1	0	0	0	0
245	0	0	1	0	1	1	1	1	0	0	0	0
246	1	0	1	0	1	1	1	1	0	0	0	0
247	0	1	1	0	1	1	1	1	0	0	0	0
248	1	1	1	0	1	1	1	1	0	0	0	0
249	0	0	0	1	1	1	1	1	0	0	0	0
250	1	0	0	1	1	1	1	1	0	0	0	0
251	0	1	0	1	1	1	1	1	0	0	0	0
252	1	1	0	1	1	1	1	1	0	0	0	0
253	0	0	1	1	1	1	1	1	0	0	0	0
254	1	0	1	1	1	1	1	1	0	0	0	0
255	0	1	1	1	1	1	1	1	0	0	0	0
256	1	1	1	1	1	1	1	1	44	31	0	0

pre1 pre2 Sp1 Sp2 Sp3 Sp4 Sp5 Sp6 Sp7 Sp8 Se1 Se2 Se3 Se4

Best Guess 0.2 0.2 0.95 0.95 0.95 0.95 0.95 0.95 0.95 0.95 0.8 0.8 0.8 0.8

Se5 Se6 Se7 Se8

Best Guess 0.8 0.8 0.8 0.8

EXPECTATION MAXIMISATION

\$Iterations

[1] 7

\$LogLikelihood
[1] -857.8069

\$Estimations
pre1 pre2 Sp1 Sp2 Sp3 Sp4 Sp5 Sp6 Sp7 Sp8 Se1 Se2
Est 0.1864 0.2196 1 0.977 0.954 1 0.9977 0.977 0.9402 0.9173 0.9443 0.9629
Se3 Se4 Se5 Se6 Se7 Se8
Est 0.9814 0.9165 0.7313 0.9907 0.9906 0.9352

NEWTON-RAPHSON
\$Iterations
[1] 30

\$LogLikelihood
[1] -857.8069

\$Estimations
pre1 pre2 Sp1 Sp2 Sp3 Sp4 Sp5 Sp6 Sp7 Sp8 Se1
Est 0.1864 0.2196 1 0.9770 0.9540 1 0.9977 0.9770 0.9402 0.9173 0.9443
Clinf 0.1484 0.1681 NaN 0.9578 0.9298 NaN 0.9839 0.9578 0.9137 0.8874 0.8814
Clsup 0.2316 0.2814 NaN 0.9876 0.9702 1 0.9997 0.9876 0.9590 0.9398 0.9748
Se2 Se3 Se4 Se5 Se6 Se7 Se8
Est 0.9629 0.9814 0.9165 0.7313 0.9907 0.9906 0.9352
Clinf 0.9052 0.9285 0.8472 0.6401 0.9366 0.9364 0.8702
Clsup 0.9860 0.9953 0.9560 0.8064 0.9987 0.9987 0.9688

WARNING 1: test results are assumed to be independent conditional on infection or disease status

WARNING 2: tests are supposed to have constant sensitivity and specificity in all populations

Expected Results (NR) and Goodness-of-fit test

\$Expected
test1 test2 test3 test4 test5 test6 test7 test8 pop1 pop2 RefInd RefInf
1 0 0 0 0 0 0 0 0 254 144 0 0
2 1 0 0 0 0 0 0 0 0 0 0 0
3 0 1 0 0 0 0 0 0 0 0 0 0
4 1 1 0 0 0 0 0 0 0 0 0 0
5 0 0 1 0 0 0 0 0 0 0 0 0
6 1 0 1 0 0 0 0 0 0 0 0 0
7 0 1 1 0 0 0 0 0 0 0 0 0
8 1 1 1 0 0 0 0 0 0 0 0 0
9 0 0 0 1 0 0 0 0 0 0 0 0

10	1	0	0	1	0	0	0	0	0	0	0
11	0	1	0	1	0	0	0	0	0	0	0
12	1	1	0	1	0	0	0	0	0	0	0
13	0	0	1	1	0	0	0	0	0	0	0
14	1	0	1	1	0	0	0	0	0	0	0
15	0	1	1	1	0	0	0	0	0	0	0
16	1	1	1	1	0	0	0	0	0	0	0
17	0	0	0	0	1	0	0	0	0	1	0
18	1	0	0	0	1	0	0	0	0	0	0
19	0	1	0	0	1	0	0	0	0	0	0
20	1	1	0	0	1	0	0	0	0	0	0
21	0	0	1	0	1	0	0	0	0	0	0
22	1	0	1	0	1	0	0	0	0	0	0
23	0	1	1	0	1	0	0	0	0	0	0
24	1	1	1	0	1	0	0	0	1	0	0
25	0	0	0	1	1	0	0	0	0	0	0
26	1	0	0	1	1	0	0	0	0	0	0
27	0	1	0	1	1	0	0	0	0	0	0
28	1	1	0	1	1	0	0	0	0	0	0
29	0	0	1	1	1	0	0	0	0	0	0
30	1	0	1	1	1	0	0	0	0	0	0
31	0	1	1	1	1	0	0	0	0	0	0
32	1	1	1	1	1	0	0	0	0	0	0
33	0	0	0	0	0	1	0	0	0	0	0
34	1	0	0	0	0	1	0	0	0	0	0
35	0	1	0	0	0	1	0	0	0	0	0
36	1	1	0	0	0	1	0	0	0	0	0
37	0	0	1	0	0	1	0	0	0	0	0
38	1	0	1	0	0	1	0	0	0	0	0
39	0	1	1	0	0	1	0	0	0	0	0
40	1	1	1	0	0	1	0	0	0	0	0
41	0	0	0	1	0	1	0	0	0	0	0
42	1	0	0	1	0	1	0	0	0	0	0
43	0	1	0	1	0	1	0	0	0	0	0
44	1	1	0	1	0	1	0	0	0	0	0
45	0	0	1	1	0	1	0	0	0	0	0
46	1	0	1	1	0	1	0	0	0	0	0
47	0	1	1	1	0	1	0	0	0	0	0
48	1	1	1	1	0	1	0	0	0	0	0
49	0	0	0	0	1	1	0	0	0	0	0
50	1	0	0	0	1	1	0	0	0	0	0
51	0	1	0	0	1	1	0	0	0	0	0
52	1	1	0	0	1	1	0	0	0	0	0
53	0	0	1	0	1	1	0	0	0	0	0

54	1	0	1	0	1	1	0	0	0	0	0	0
55	0	1	1	0	1	1	0	0	0	0	0	0
56	1	1	1	0	1	1	0	0	0	0	0	0
57	0	0	0	1	1	1	0	0	0	0	0	0
58	1	0	0	1	1	1	0	0	0	0	0	0
59	0	1	0	1	1	1	0	0	0	0	0	0
60	1	1	0	1	1	1	0	0	0	0	0	0
61	0	0	1	1	1	1	0	0	0	0	0	0
62	1	0	1	1	1	1	0	0	0	0	0	0
63	0	1	1	1	1	1	0	0	0	0	0	0
64	1	1	1	1	1	1	0	0	0	0	0	0
65	0	0	0	0	0	0	1	0	0	0	0	0
66	1	0	0	0	0	0	1	0	0	0	0	0
67	0	1	0	0	0	0	1	0	0	0	0	0
68	1	1	0	0	0	0	1	0	0	0	0	0
69	0	0	1	0	0	0	1	0	0	0	0	0
70	1	0	1	0	0	0	1	0	0	0	0	0
71	0	1	1	0	0	0	1	0	0	0	0	0
72	1	1	1	0	0	0	1	0	0	0	0	0
73	0	0	0	1	0	0	1	0	0	0	0	0
74	1	0	0	1	0	0	1	0	0	0	0	0
75	0	1	0	1	0	0	1	0	0	0	0	0
76	1	1	0	1	0	0	1	0	0	0	0	0
77	0	0	1	1	0	0	1	0	0	0	0	0
78	1	0	1	1	0	0	1	0	0	0	0	0
79	0	1	1	1	0	0	1	0	0	0	0	0
80	1	1	1	1	0	0	1	0	0	0	0	0
81	0	0	0	0	1	0	1	0	0	0	0	0
82	1	0	0	0	1	0	1	0	0	0	0	0
83	0	1	0	0	1	0	1	0	0	0	0	0
84	1	1	0	0	1	0	1	0	0	0	0	0
85	0	0	1	0	1	0	1	0	0	0	0	0
86	1	0	1	0	1	0	1	0	0	0	0	0
87	0	1	1	0	1	0	1	0	0	0	0	0
88	1	1	1	0	1	0	1	0	0	0	0	0
89	0	0	0	1	1	0	1	0	0	0	0	0
90	1	0	0	1	1	0	1	0	0	0	0	0
91	0	1	0	1	1	0	1	0	0	0	0	0
92	1	1	0	1	1	0	1	0	0	0	0	0
93	0	0	1	1	1	0	1	0	0	0	0	0
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95	0	1	1	1	1	0	1	0	0	0	0	0
96	1	1	1	1	1	0	1	0	0	0	0	0
97	0	0	0	0	0	0	1	1	0	0	0	0

98	1	0	0	0	0	1	1	0	0	0	0	0
99	0	1	0	0	0	1	1	0	0	0	0	0
100	1	1	0	0	0	1	1	0	0	0	0	0
101	0	0	1	0	0	1	1	0	0	0	0	0
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103	0	1	1	0	0	1	1	0	0	0	0	0
104	1	1	1	0	0	1	1	0	0	0	0	0
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106	1	0	0	1	0	1	1	0	0	0	0	0
107	0	1	0	1	0	1	1	0	0	0	0	0
108	1	1	0	1	0	1	1	0	0	0	0	0
109	0	0	1	1	0	1	1	0	0	2	0	0
110	1	0	1	1	0	1	1	0	0	0	0	0
111	0	1	1	1	0	1	1	0	1	0	0	0
112	1	1	1	1	0	1	1	0	0	0	0	0
113	0	0	0	0	1	1	1	0	0	0	0	0
114	1	0	0	0	1	1	1	0	0	0	0	0
115	0	1	0	0	1	1	1	0	0	0	0	0
116	1	1	0	0	1	1	1	0	0	0	0	0
117	0	0	1	0	1	1	1	0	0	0	0	0
118	1	0	1	0	1	1	1	0	0	0	0	0
119	0	1	1	0	1	1	1	0	0	0	0	0
120	1	1	1	0	1	1	1	0	0	0	0	0
121	0	0	0	1	1	1	1	0	1	0	0	0
122	1	0	0	1	1	1	1	0	0	0	0	0
123	0	1	0	1	1	1	1	0	0	0	0	0
124	1	1	0	1	1	1	1	0	0	0	0	0
125	0	0	1	1	1	1	1	0	0	0	0	0
126	1	0	1	1	1	1	1	0	0	0	0	0
127	0	1	1	1	1	1	1	0	2	0	0	0
128	1	1	1	1	1	1	1	0	0	0	0	0
129	0	0	0	0	0	0	0	1	0	0	0	0
130	1	0	0	0	0	0	0	1	0	0	0	0
131	0	1	0	0	0	0	0	1	0	0	0	0
132	1	1	0	0	0	0	0	1	0	0	0	0
133	0	0	1	0	0	0	0	1	0	0	0	0
134	1	0	1	0	0	0	0	1	0	0	0	0
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139	0	1	0	1	0	0	0	1	0	0	0	0
140	1	1	0	1	0	0	0	1	0	0	0	0
141	0	0	1	1	0	0	0	1	0	0	0	0

142	1	0	1	1	0	0	0	1	0	0	0	0
143	0	1	1	1	0	0	0	1	0	0	0	0
144	1	1	1	1	0	0	0	1	0	0	0	0
145	0	0	0	0	1	0	0	1	0	0	0	0
146	1	0	0	0	1	0	0	1	0	0	0	0
147	0	1	0	0	1	0	0	1	0	0	0	0
148	1	1	0	0	1	0	0	1	0	0	0	0
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150	1	0	1	0	1	0	0	1	0	0	0	0
151	0	1	1	0	1	0	0	1	0	0	0	0
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153	0	0	0	1	1	0	0	1	0	0	0	0
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155	0	1	0	1	1	0	0	1	0	0	0	0
156	1	1	0	1	1	0	0	1	0	0	0	0
157	0	0	1	1	1	0	0	1	0	0	0	0
158	1	0	1	1	1	0	0	1	0	0	0	0
159	0	1	1	1	1	0	0	1	0	0	0	0
160	1	1	1	1	1	0	0	1	0	0	0	0
161	0	0	0	0	0	1	0	1	0	0	0	0
162	1	0	0	0	0	1	0	1	0	0	0	0
163	0	1	0	0	0	1	0	1	6	4	0	0
164	1	1	0	0	0	1	0	1	0	0	0	0
165	0	0	1	0	0	1	0	1	0	0	0	0
166	1	0	1	0	0	1	0	1	0	0	0	0
167	0	1	1	0	0	1	0	1	0	0	0	0
168	1	1	1	0	0	1	0	1	0	0	0	0
169	0	0	0	1	0	1	0	1	0	0	0	0
170	1	0	0	1	0	1	0	1	0	0	0	0
171	0	1	0	1	0	1	0	1	0	0	0	0
172	1	1	0	1	0	1	0	1	0	0	0	0
173	0	0	1	1	0	1	0	1	0	0	0	0
174	1	0	1	1	0	1	0	1	0	0	0	0
175	0	1	1	1	0	1	0	1	0	0	0	0
176	1	1	1	1	0	1	0	1	0	0	0	0
177	0	0	0	0	1	1	0	1	0	0	0	0
178	1	0	0	0	1	1	0	1	0	0	0	0
179	0	1	0	0	1	1	0	1	0	0	0	0
180	1	1	0	0	1	1	0	1	0	0	0	0
181	0	0	1	0	1	1	0	1	0	0	0	0
182	1	0	1	0	1	1	0	1	0	0	0	0
183	0	1	1	0	1	1	0	1	0	0	0	0
184	1	1	1	0	1	1	0	1	0	0	0	0
185	0	0	0	1	1	1	0	1	0	0	0	0

186	1	0	0	1	1	1	0	1	0	0	0	0
187	0	1	0	1	1	1	0	1	0	0	0	0
188	1	1	0	1	1	1	0	1	0	0	0	0
189	0	0	1	1	1	1	0	1	0	0	0	0
190	1	0	1	1	1	1	0	1	0	0	0	0
191	0	1	1	1	1	1	0	1	0	0	0	0
192	1	1	1	1	1	1	0	1	0	0	0	0
193	0	0	0	0	0	0	1	1	4	2	0	0
194	1	0	0	0	0	0	1	1	0	0	0	0
195	0	1	0	0	0	0	1	1	0	0	0	0
196	1	1	0	0	0	0	1	1	0	0	0	0
197	0	0	1	0	0	0	1	1	11	9	0	0
198	1	0	1	0	0	0	1	1	0	0	0	0
199	0	1	1	0	0	0	1	1	0	0	0	0
200	1	1	1	0	0	0	1	1	0	0	0	0
201	0	0	0	1	0	0	1	1	0	0	0	0
202	1	0	0	1	0	0	1	1	0	0	0	0
203	0	1	0	1	0	0	1	1	0	0	0	0
204	1	1	0	1	0	0	1	1	0	0	0	0
205	0	0	1	1	0	0	1	1	0	0	0	0
206	1	0	1	1	0	0	1	1	0	0	0	0
207	0	1	1	1	0	0	1	1	0	0	0	0
208	1	1	1	1	0	0	1	1	0	0	0	0
209	0	0	0	0	1	0	1	1	0	0	0	0
210	1	0	0	0	1	0	1	1	0	0	0	0
211	0	1	0	0	1	0	1	1	0	0	0	0
212	1	1	0	0	1	0	1	1	0	0	0	0
213	0	0	1	0	1	0	1	1	0	0	0	0
214	1	0	1	0	1	0	1	1	0	0	0	0
215	0	1	1	0	1	0	1	1	0	0	0	0
216	1	1	1	0	1	0	1	1	0	0	0	0
217	0	0	0	1	1	0	1	1	0	0	0	0
218	1	0	0	1	1	0	1	1	0	0	0	0
219	0	1	0	1	1	0	1	1	0	0	0	0
220	1	1	0	1	1	0	1	1	0	0	0	0
221	0	0	1	1	1	0	1	1	0	0	0	0
222	1	0	1	1	1	0	1	1	0	0	0	0
223	0	1	1	1	1	0	1	1	0	0	0	0
224	1	1	1	1	1	0	1	1	0	0	0	0
225	0	0	0	0	0	1	1	1	0	0	0	0
226	1	0	0	0	0	1	1	1	0	0	0	0
227	0	1	0	0	0	1	1	1	0	0	0	0
228	1	1	0	0	0	1	1	1	0	0	0	0
229	0	0	1	0	0	1	1	1	0	0	0	0

230	1	0	1	0	0	1	1	1	0	0	0	0
231	0	1	1	0	0	1	1	1	0	0	0	0
232	1	1	1	0	0	1	1	1	7	1	0	0
233	0	0	0	1	0	1	1	1	0	0	0	0
234	1	0	0	1	0	1	1	1	0	1	0	0
235	0	1	0	1	0	1	1	1	0	0	0	0
236	1	1	0	1	0	1	1	1	0	0	0	0
237	0	0	1	1	0	1	1	1	0	0	0	0
238	1	0	1	1	0	1	1	1	0	0	0	0
239	0	1	1	1	0	1	1	1	0	0	0	0
240	1	1	1	1	0	1	1	1	7	10	0	0
241	0	0	0	0	1	1	1	1	0	0	0	0
242	1	0	0	0	1	1	1	1	0	0	0	0
243	0	1	0	0	1	1	1	1	0	0	0	0
244	1	1	0	0	1	1	1	1	0	0	0	0
245	0	0	1	0	1	1	1	1	0	0	0	0
246	1	0	1	0	1	1	1	1	0	0	0	0
247	0	1	1	0	1	1	1	1	0	0	0	0
248	1	1	1	0	1	1	1	1	0	0	0	0
249	0	0	0	1	1	1	1	1	0	0	0	0
250	1	0	0	1	1	1	1	1	0	0	0	0
251	0	1	0	1	1	1	1	1	0	0	0	0
252	1	1	0	1	1	1	1	1	0	0	0	0
253	0	0	1	1	1	1	1	1	0	0	0	0
254	1	0	1	1	1	1	1	1	0	0	0	0
255	0	1	1	1	1	1	1	1	0	0	0	0
256	1	1	1	1	1	1	1	1	44	31	0	0

ExpPop1 ExpPop2 ExpRefInd ExpRefInf

1	215.50	125.38	0	0
2	0.00	0.00	0	0
3	5.06	2.95	0	0
4	0.00	0.00	0	0
5	10.38	6.04	0	0
6	0.00	0.00	0	0
7	0.24	0.14	0	0
8	0.00	0.00	0	0
9	0.00	0.00	0	0
10	0.00	0.00	0	0
11	0.00	0.00	0	0
12	0.00	0.00	0	0
13	0.00	0.00	0	0
14	0.00	0.00	0	0
15	0.00	0.00	0	0
16	0.00	0.00	0	0

17	0.50	0.29	0	0
18	0.00	0.00	0	0
19	0.01	0.01	0	0
20	0.00	0.00	0	0
21	0.02	0.01	0	0
22	0.00	0.00	0	0
23	0.00	0.00	0	0
24	0.00	0.00	0	0
25	0.00	0.00	0	0
26	0.00	0.00	0	0
27	0.00	0.00	0	0
28	0.00	0.00	0	0
29	0.00	0.00	0	0
30	0.00	0.00	0	0
31	0.00	0.00	0	0
32	0.00	0.00	0	0
33	5.06	2.95	0	0
34	0.00	0.00	0	0
35	0.12	0.07	0	0
36	0.00	0.00	0	0
37	0.24	0.14	0	0
38	0.00	0.00	0	0
39	0.01	0.00	0	0
40	0.00	0.00	0	0
41	0.00	0.00	0	0
42	0.00	0.00	0	0
43	0.00	0.00	0	0
44	0.00	0.00	0	0
45	0.00	0.00	0	0
46	0.00	0.00	0	0
47	0.00	0.00	0	0
48	0.01	0.01	0	0
49	0.01	0.01	0	0
50	0.00	0.00	0	0
51	0.00	0.00	0	0
52	0.00	0.00	0	0
53	0.00	0.00	0	0
54	0.00	0.00	0	0
55	0.00	0.00	0	0
56	0.00	0.00	0	0
57	0.00	0.00	0	0
58	0.00	0.00	0	0
59	0.00	0.00	0	0
60	0.00	0.00	0	0

61	0.00	0.00	0	0
62	0.00	0.00	0	0
63	0.00	0.00	0	0
64	0.02	0.02	0	0
65	13.69	7.97	0	0
66	0.00	0.00	0	0
67	0.32	0.19	0	0
68	0.00	0.00	0	0
69	0.66	0.38	0	0
70	0.00	0.00	0	0
71	0.02	0.01	0	0
72	0.00	0.00	0	0
73	0.00	0.00	0	0
74	0.00	0.00	0	0
75	0.00	0.00	0	0
76	0.00	0.00	0	0
77	0.00	0.00	0	0
78	0.00	0.00	0	0
79	0.00	0.00	0	0
80	0.01	0.01	0	0
81	0.03	0.02	0	0
82	0.00	0.00	0	0
83	0.00	0.00	0	0
84	0.00	0.00	0	0
85	0.00	0.00	0	0
86	0.00	0.00	0	0
87	0.00	0.00	0	0
88	0.00	0.00	0	0
89	0.00	0.00	0	0
90	0.00	0.00	0	0
91	0.00	0.00	0	0
92	0.00	0.00	0	0
93	0.00	0.00	0	0
94	0.00	0.00	0	0
95	0.00	0.00	0	0
96	0.02	0.02	0	0
97	0.32	0.19	0	0
98	0.00	0.00	0	0
99	0.01	0.00	0	0
100	0.00	0.00	0	0
101	0.02	0.01	0	0
102	0.00	0.00	0	0
103	0.01	0.00	0	0
104	0.08	0.06	0	0

105	0.00	0.00	0	0
106	0.00	0.00	0	0
107	0.00	0.00	0	0
108	0.02	0.01	0	0
109	0.00	0.00	0	0
110	0.03	0.02	0	0
111	0.05	0.04	0	0
112	0.88	0.63	0	0
113	0.00	0.00	0	0
114	0.00	0.00	0	0
115	0.00	0.00	0	0
116	0.00	0.00	0	0
117	0.00	0.00	0	0
118	0.01	0.01	0	0
119	0.01	0.01	0	0
120	0.22	0.16	0	0
121	0.00	0.00	0	0
122	0.00	0.00	0	0
123	0.00	0.00	0	0
124	0.05	0.03	0	0
125	0.01	0.00	0	0
126	0.09	0.07	0	0
127	0.14	0.10	0	0
128	2.40	1.71	0	0
129	19.43	11.31	0	0
130	0.00	0.00	0	0
131	0.46	0.27	0	0
132	0.00	0.00	0	0
133	0.94	0.54	0	0
134	0.00	0.00	0	0
135	0.02	0.01	0	0
136	0.00	0.00	0	0
137	0.00	0.00	0	0
138	0.00	0.00	0	0
139	0.00	0.00	0	0
140	0.00	0.00	0	0
141	0.00	0.00	0	0
142	0.00	0.00	0	0
143	0.00	0.00	0	0
144	0.00	0.00	0	0
145	0.04	0.03	0	0
146	0.00	0.00	0	0
147	0.00	0.00	0	0
148	0.00	0.00	0	0

149	0.00	0.00	0	0
150	0.00	0.00	0	0
151	0.00	0.00	0	0
152	0.00	0.00	0	0
153	0.00	0.00	0	0
154	0.00	0.00	0	0
155	0.00	0.00	0	0
156	0.00	0.00	0	0
157	0.00	0.00	0	0
158	0.00	0.00	0	0
159	0.00	0.00	0	0
160	0.00	0.00	0	0
161	0.46	0.27	0	0
162	0.00	0.00	0	0
163	0.01	0.01	0	0
164	0.00	0.00	0	0
165	0.02	0.01	0	0
166	0.00	0.00	0	0
167	0.00	0.00	0	0
168	0.01	0.01	0	0
169	0.00	0.00	0	0
170	0.00	0.00	0	0
171	0.00	0.00	0	0
172	0.00	0.00	0	0
173	0.00	0.00	0	0
174	0.00	0.00	0	0
175	0.01	0.01	0	0
176	0.12	0.09	0	0
177	0.00	0.00	0	0
178	0.00	0.00	0	0
179	0.00	0.00	0	0
180	0.00	0.00	0	0
181	0.00	0.00	0	0
182	0.00	0.00	0	0
183	0.00	0.00	0	0
184	0.03	0.02	0	0
185	0.00	0.00	0	0
186	0.00	0.00	0	0
187	0.00	0.00	0	0
188	0.01	0.00	0	0
189	0.00	0.00	0	0
190	0.01	0.01	0	0
191	0.02	0.01	0	0
192	0.33	0.23	0	0

193	1.23	0.72	0	0
194	0.00	0.00	0	0
195	0.03	0.02	0	0
196	0.00	0.00	0	0
197	0.06	0.03	0	0
198	0.00	0.00	0	0
199	0.00	0.00	0	0
200	0.01	0.01	0	0
201	0.00	0.00	0	0
202	0.00	0.00	0	0
203	0.00	0.00	0	0
204	0.00	0.00	0	0
205	0.00	0.00	0	0
206	0.00	0.00	0	0
207	0.01	0.01	0	0
208	0.12	0.09	0	0
209	0.00	0.00	0	0
210	0.00	0.00	0	0
211	0.00	0.00	0	0
212	0.00	0.00	0	0
213	0.00	0.00	0	0
214	0.00	0.00	0	0
215	0.00	0.00	0	0
216	0.03	0.02	0	0
217	0.00	0.00	0	0
218	0.00	0.00	0	0
219	0.00	0.00	0	0
220	0.01	0.00	0	0
221	0.00	0.00	0	0
222	0.01	0.01	0	0
223	0.02	0.01	0	0
224	0.33	0.23	0	0
225	0.03	0.02	0	0
226	0.00	0.00	0	0
227	0.00	0.00	0	0
228	0.02	0.02	0	0
229	0.00	0.00	0	0
230	0.04	0.03	0	0
231	0.07	0.05	0	0
232	1.16	0.83	0	0
233	0.00	0.00	0	0
234	0.01	0.01	0	0
235	0.01	0.01	0	0
236	0.24	0.17	0	0

237	0.03	0.02	0	0
238	0.49	0.35	0	0
239	0.75	0.54	0	0
240	12.71	9.08	0	0
241	0.00	0.00	0	0
242	0.00	0.00	0	0
243	0.00	0.00	0	0
244	0.06	0.04	0	0
245	0.01	0.01	0	0
246	0.12	0.09	0	0
247	0.19	0.13	0	0
248	3.15	2.25	0	0
249	0.00	0.00	0	0
250	0.03	0.02	0	0
251	0.04	0.03	0	0
252	0.66	0.47	0	0
253	0.08	0.06	0	0
254	1.33	0.95	0	0
255	2.04	1.46	0	0
256	34.59	24.71	0	0

\$Test

Max LogLikelihood: Achievable Obtained Deviance d.f. p value
-541.8923 -857.8069 631.8292 492 1.938381e-05

\$Commentary

[1] "The model does not fit: Assumptions may be not justified"

Residual correlations between test

	Corr1-2	Corr1-3	Corr1-4	Corr1-5	Corr1-6	
pop 1 :	0.02966545	0.02748822	-0.04146374	0.02335436	-0.007204700	
pop 2 :	0.03676788	-0.01556004	0.04393568	0.01275859	0.008905438	
	Corr1-7	Corr1-8	Corr2-3	Corr2-4	Corr2-5	Corr2-6
pop 1 :	0.0003667263	0.06535817	0.04146528	-0.01234181	0.03409523	0.12179659
pop 2 :	-0.0023577244	0.05369714	-0.02207883	0.01071037	0.01003414	0.08589943
	Corr2-7	Corr2-8	Corr3-4	Corr3-5	Corr3-6	Corr3-7
pop 1 :	0.01363943	0.1267421	-0.01302279	0.03147218	0.007666826	0.1957482
pop 2 :	-0.03651559	0.1205577	0.01348754	-0.03281901	-0.024305582	0.1829973
	Corr3-8	Corr4-5	Corr4-6	Corr4-7	Corr4-8	Corr5-6
pop 1 :	0.2007084	0.11974713	-0.009463266	-0.001237415	-0.01300391	0.01924680
pop 2 :	0.1809753	0.01560615	0.039334803	0.025655508	0.02724110	-0.03674752
	Corr5-7	Corr5-8	Corr6-7	Corr6-8	Corr7-8	

```
pop 1 : 0.02362124 0.018492288 0.01576406 0.1127440 0.2519878  
pop 2 : -0.04385803 0.004320285 -0.01371877 0.0908689 0.2325526
```

\$Commentary

[1] "The residuals should be randomly distributed around 0"

BOOTSTRAP CONFIDENCE INTERVALS : 5000 samples

 pre1 pre2 Sp1 Sp2 Sp3 Sp4 Sp5 Sp6 Sp7 Sp8 Se1

 Clinf 0.1456 0.1612 1 0.9618 0.9329 1 0.993 0.9618 0.9169 0.8894 0.8969

 Clsup 0.2278 0.2781 1 0.9907 0.9726 1 1.000 0.9907 0.9614 0.9424 0.9825

 Se2 Se3 Se4 Se5 Se6 Se7 Se8

 Clinf 0.9244 0.9524 0.8600 0.6442 0.9699 0.9696 0.8859

 Clsup 0.9915 1.0000 0.9658 0.8130 1.0000 1.0000 0.9798

your data are stored in "new_8_double"

Note that you'll have to Save workspace image before leaving R if you want to use it in a new R session

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