# Sequential Implementation of Monte Carlo Tests with Uniformly Bounded Resampling Risk

#### Axel Gandy

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- ► Test statistic *T*, reject for large values.
- ▶ Observation: t.
- p-value:

$$p = P(T \ge t)$$

Often not available in closed form.

Monte Carlo Test:

$$\hat{p}_{\text{naive}} = \frac{1}{n} \sum_{i=1}^{n} I(T_i \geq t),$$

where  $T, T_1, \ldots T_n$  i.i.d.

- ► Examples:
  - ▶ Bootstrap,
  - Permutation tests.
- Goal: Estimate p using few  $X_i$

Mainly interested in deciding if  $p \leq \alpha$  for some  $\alpha$ .



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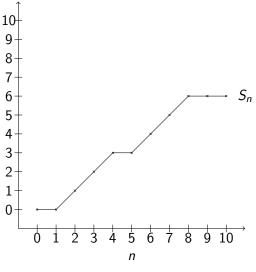
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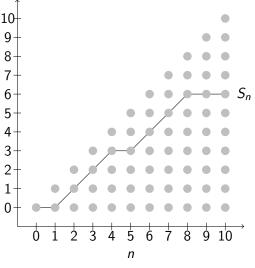


# Sequential approaches based on $S_n = \sum_{i=1}^n X_i$



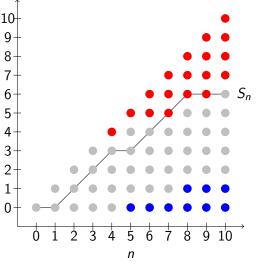
- Stop once  $S_n \ge U_n$  or  $S_n < L_n$
- ightharpoonup au: hitting time
- ► Compute  $\hat{p}$  based on  $S_{\tau}$  and  $\tau$ .
- ▶ Hit  $B_U$ : decide  $p > \alpha$ ,
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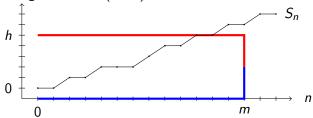
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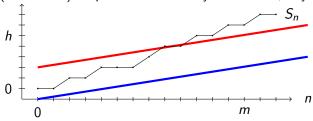
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#### Previous Approaches

► Besag & Clifford (1991):



► (Truncated) Sequential Probability Ratio Test, Fay et al. (2007)



► R-package MChtest.



Is 
$$p \leq \alpha$$
?

Two individuals using the same statistical method on the same data should arrive at the same conclusion.

First law of applied statistics, Gleser (1996)

Consider the resampling risk

$$RR_{p}(\hat{p}) \equiv \begin{cases} P_{p}(\hat{p} > \alpha) & \text{if } p \leq \alpha, \\ P_{p}(\hat{p} \leq \alpha) & \text{if } p > \alpha. \end{cases}$$

Want

$$\sup_{p \in [0,1]} \mathsf{RR}_p(\hat{p}) \leq \epsilon$$

for some (small)  $\epsilon>0$ . For Besag & Clifford (1991), SPRT:  $\mathsf{sup}_p\,\mathsf{RR}_P\geq0.5$ 



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Want:

$$\sup_{p} \mathsf{RR}_{p}(\hat{p}) \leq \epsilon$$

$$\mathsf{P}_{lpha}(\mathsf{hit}\;B_{oldsymbol{U}}) \leq \epsilon \ \mathsf{P}_{lpha}(\mathsf{hit}\;B_{oldsymbol{L}}) \leq \epsilon \ .$$

$$P_{\alpha}(\text{hit }B_{U} \text{ until } n) \leq \epsilon_{n}$$

$$P_{\alpha}(\text{hit }B_{L} \text{ until } n) \leq \epsilon_{r}$$



Want:

$$\sup_{p} \mathsf{RR}_{p}(\hat{p}) \leq \epsilon$$

Suffices to ensure

$$P_{\alpha}(\text{hit }B_{U}) \leq \epsilon$$
 $P_{\alpha}(\text{hit }B_{L}) \leq \epsilon$ 

#### Recursive definition:

Given  $U_1, \ldots, U_{n-1}$  and  $L_1, \ldots, L_{n-1}$ , define

▶ U<sub>n</sub> as the minimal value such that

$$\mathsf{P}_{\!lpha}(\mathsf{hit}\; extstyle{B_U}\; \mathsf{until}\; n) \leq \epsilon_n$$

▶ and L<sub>n</sub> as the maximal value such that

$$P_{\alpha}(\text{hit }B_{L} \text{ until }n) \leq \epsilon_{n}$$

where  $\epsilon_n \geq 0$  with  $\epsilon_n \nearrow \epsilon$  (spending sequence).



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- $\alpha = 0.2, \ \epsilon_n = 0.4 \frac{n}{5+n}.$
- $\triangleright$   $U_n$ =the minimal value such that

$$P_{\alpha}(\text{hit }B_{U}\text{ until }n) \leq \epsilon_{n}$$

$$P_{\alpha}(\text{hit }B_{L} \text{ until }n) \leq \epsilon_{n}$$

	n =
$P_{\alpha}(S_n = k, \tau \geq n)$	0
k= 3	
k= 2	
k=1	
k= 0	1
$\epsilon_n$	0
$\overline{U_n}$	1
Ln	-1

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			n =
$P_{\alpha}(S_n = k, \tau \geq n)$	0	1	
k= 3			
k=2			
k=1		.2	
k= 0	1	.8	
$\epsilon_n$	0	.07	
$\overline{U_n}$	1	2	
$L_n$	-1	-1	

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				n =
$P_{\alpha}(S_n = k, \tau \geq n)$	0	1	2	
k= 3				
k=2			.04	
k=1		.2	.32	
k= 0	1	.8	.64	
$\epsilon_n$	0	.07	.11	
$\overline{U_n}$	1	2	2	
$L_n$	-1	-1	-1	

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					n =
$P_{\alpha}(S_n = k, \tau \geq n)$	0	1	2	3	
k= 3					
k=2			.04	.06	
k=1		.2	.32	.38	
k= 0	1	.8	.64	.51	
$\epsilon_n$	0	.07	.11	.15	
$U_n$	1	2	2	2	
$L_n$	-1	-1	-1	-1	

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					n =	
$P_{\alpha}(S_n = k, \tau \geq n)$	0	1	2	3	4	
k= 3						
k= 2			.04	.06	.08	
k=1		.2	.32	.38	.41	
k= 0	1	.8	.64	.51	.41	
$\epsilon_n$	0	.07	.11	.15	.18	
$\overline{U_n}$	1	2	2	2	3	
$L_n$	-1	-1	-1	-1	-1	

- $\alpha = 0.2, \ \epsilon_n = 0.4 \frac{n}{5+n}.$
- $\triangleright$   $U_n$ =the minimal value such that

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$$P_{\alpha}(\text{hit }B_{L} \text{ until }n) \leq \epsilon_{n}$$

					n =		
$P_{\alpha}(S_n = k, \tau \geq n)$	0	1	2	3	4	5	
k= 3						.02	
k= 2			.04	.06	.08	.14	
k=1		.2	.32	.38	.41	.41	
k= 0	1	.8	.64	.51	.41	.33	
$\epsilon_n$	0	.07	.11	.15	.18	.20	
$\overline{U_n}$	1	2	2	2	3	3	
Ln	-1	-1	-1	-1	-1	-1	

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	n =								
$P_{\alpha}(S_n = k, \tau \geq n)$	0	1	2	3	4	5	6		
k= 3						.02	.03		
k= 2			.04	.06	.08	.14	.20		
k=1		.2	.32	.38	.41	.41	.39		
k= 0	1	.8	.64	.51	.41	.33	.26		
$\epsilon_n$	0	.07	.11	.15	.18	.20	.22		
$U_n$	1	2	2	2	3	3	3		
$L_n$	-1	-1	-1	-1	-1	-1	-1		

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	n =										
$P_{\alpha}(S_n = k, \tau \geq n)$	0	1	2	3	4	5	6	7			
k= 3						.02	.03	.04			
k= 2			.04	.06	.08	.14	.20	.24			
k=1		.2	.32	.38	.41	.41	.39	.37			
k= 0	1	.8	.64	.51	.41	.33	.26	.21			
$\epsilon_n$	0	.07	.11	.15	.18	.20	.22	.23			
$\overline{U_n}$	1	2	2	2	3	3	3	3			
$L_n$	-1	-1	-1	-1	-1	-1	-1	0			

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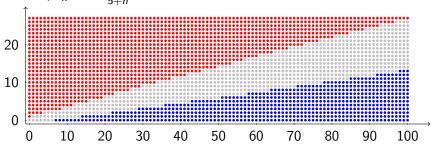
$$P_{\alpha}(\text{hit }B_{U}\text{ until }n) \leq \epsilon_{n}$$

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	n =										
$P_{\alpha}(S_n = k, \tau \geq n)$	0	1	2	3	4	5	6	7	8		
k= 3						.02	.03	.04	.05		
k= 2			.04	.06	.08	.14	.20	.24	.26		
k=1		.2	.32	.38	.41	.41	.39	.37	.29		
k= 0	1	.8	.64	.51	.41	.33	.26	.21			
$\epsilon_n$	0	.07	.11	.15	.18	.20	.22	.23	.25		
$\overline{U_n}$	1	2	2	2	3	3	3	3	3		
$L_n$	-1	-1	-1	-1	-1	-1	-1	0	0		

# Sequential Decision Procedure - Example

$$\alpha = 0.2$$
,  $\epsilon_n = 0.4 \frac{n}{5+n}$ .

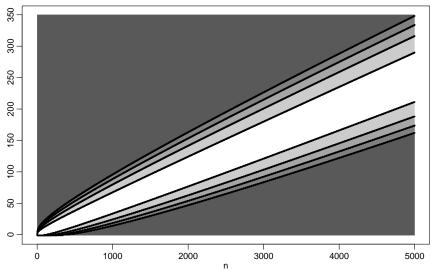


n



#### Influence of $\epsilon$ on the stopping rule

$$\epsilon = 0.1, \ 0.001, \ 10^{-5}, \ 10^{-7}; \ \epsilon_n = \epsilon \frac{n}{1000 + n}$$



#### Sequential Estimation based on the MLE

$$\hat{p} = \begin{cases} \frac{S_{\tau}}{\tau}, & \tau < \infty \\ \alpha, & \tau = \infty, \end{cases}$$

- One can show:
  - hitting the upper boundary implies  $\hat{p} > \alpha$ ,
  - hitting the lower boundary implies  $\hat{p} < \alpha$ .

Hence,

$$\sup_{p} \mathsf{RR}_{p}(\hat{p}) \leq \epsilon$$

- Furthermore,  $\exists$  random interval  $I_n$  s.t.
  - $ightharpoonup I_n$  only depends on  $X_1, \ldots, X_n$
  - $\hat{p} \in I_n$ .



# Example - Two-way sparse contingency table

- $\blacktriangleright$   $H_0$ : variables are independent.
- Reject for large values of the likelihood ratio test statistic T
- ▶  $T \xrightarrow{d} \chi^2_{(7-1)(5-1)}$  under  $H_0$ . Based on this: p = 0.031.
- Matrix sparse approximation poor?
- Use parametric bootstrap based on row and column sums.
- Naive test statistic  $\hat{p}_{naive}$  with n = 1,000 replicates p = 0.041 < 0.05.



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#### Example - Bootstrap and Sequential Algorithm

```
> dat <- matrix(c(1,2,2,1,1,0,1, 2,0,0,2,3,0,0, 0,1,1,1,2,7,3, 1,1,2,0,0,0,1,
                  0,1,1,1,1,0,0), nrow=5,ncol=7,byrow=TRUE)
> loglikrat <- function(data){
+ cs <- colSums(data); rs <- rowSums(data); mu <- outer(rs,cs)/sum(rs)
   2*sum(ifelse(data<=0.5, 0,data*log(data/mu)))
+ }
> resample <- function(data){</pre>
+ cs <- colSums(data);rs <- rowSums(data); n <- sum(rs)
   mu <- outer(rs,cs)/n/n
   matrix(rmultinom(1,n,c(mu)),nrow=dim(data)[1],ncol=dim(data)[2])
+ }
> t <- loglikrat(dat);</pre>
> library(simctest)
> res <- simctest(function(){loglikrat(resample(dat))>=t},maxsteps=1000)
> res
No decision reached.
Final estimate will be in [ 0.02859135 , 0.07965451 ]
Current estimate of the p.value: 0.041
Number of samples: 1000
> cont(res, steps=10000)
p.value: 0.04035456
Number of samples: 8574
```

**4** 🗇 ▶

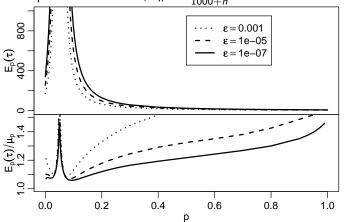
#### Further Uses of the Algorithm

- Simulation study to evaluate whether a test is liberal/conservative.
- ▶ Determining the sample size to achieve a certain power.
- ▶ Iterated Use:
  - Determining the power of a bootstrap test.
  - Simulation study to evaluate whether a bootstrap test is liberal/conservative.
  - Double bootstrap test.

#### **Expected Hitting Time**

Result:  $E_p(\tau) < \infty \ \forall p \neq \alpha$ 

Example with  $\alpha = 0.05$ ,  $\epsilon_n = \epsilon \frac{n}{1000+n}$ :



 $\mu_p$  = theoretical lower bound on  $E_p(\tau)$ .

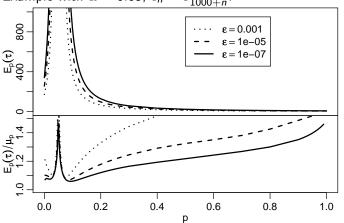
- Note:  $\int_0^1 \mu_p dp = \infty$ ;
- ▶ for iterated use: Need to limit the number of steps.



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#### Summary

- Sequential implementation of Monte Carlo Tests and computation of p-values.
- Useful when implementing tests in packages.
- After a finite number of steps:
  - $\triangleright$   $\hat{p}$  or
  - ▶ interval  $[\hat{p}_n^L, \hat{p}_n^U]$  in which  $\hat{p}$  will lie.
- Guarantee (up to a very small error probability):

 $\hat{p}$  is on the "correct side" of  $\alpha$ .

- R-package simctest available on CRAN. (efficient implementation with C-code)
- ► For details see Gandy (2009).

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