The Test Function

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```
autocor_checker<-function(y,k,i){</pre>
  # y is the vector for which we are keen on checking for randomness.
  # k is the lag.
  # i is the starting index.
  #The original test for checking if the numbers have a similar structure to a Uniform(0,1) distrubtion
  uniscaler <- function(y) {
  y \leftarrow ecdf(y)(y)
  }
#I am subsetting in following way to create the two sequences as described in the test.
subsetter_1<-seq(from=i,to=length(y),by=k)</pre>
subsetter_2<-seq(from=i+k, to=length(y),by=k)</pre>
#prath is short for Pratham which means First in Sanskrit
#dwi is short for "Dwitiya" which means Second in Sanskrit
prathseq <- uniscaler(y)[subsetter_1]</pre>
dwiseq <- uniscaler(y)[subsetter 2]</pre>
#I forcefully added 0 to dwiseq, because i wanted to see it in a Dataframe.
dwiseq[length(dwiseq)+1]<-0</pre>
dat <- data_frame(prathseq,dwiseq)</pre>
dat <- dat %>% mutate(prod=prathseq*dwiseq)
M \leftarrow floor(((length(y)-i)/k)-1)
rho_ik <- ((sum(dat$prod))/(M+1))-0.25
sig_ik \leftarrow (sqrt((13*M)+7))/(12*(M+1))
z <- rho_ik/sig_ik
test<-ifelse(2*(1-pnorm(abs(z)))<0.05,yes="Doh! No random data",no="I did not identify any dependency f
return(list(conclusion = test, p_value = 2*(1-pnorm(abs(z)))))
}
```

Using the Homework Example

```
cordat<-arima.sim(model = list(ar=0.98),n=100)</pre>
autocor_checker(cordat,k=1,i=3)
## $conclusion
## [1] "Doh! No random data"
##
## $p_value
## [1] 0.00948119
randbin < -rbinom(n=1001,80,.45)
autocor_checker(randbin,k=3,i=10)
## $conclusion
## [1] "I did not identify any dependency for this lag"
##
## $p_value
## [1] 0.2784444
randnorm < -rnorm(80, -1, 4)
autocor_checker(randnorm,k=4,i=6)
## $conclusion
## [1] "I did not identify any dependency for this lag"
## $p_value
## [1] 0.1905168
randexp \leftarrow rexp(1000, rate = 4.5)
autocor_checker(randexp,k=5,i=20)
## $conclusion
## [1] "I did not identify any dependency for this lag"
##
## $p_value
## [1] 0.067313
randt<-rt(1000,350)
autocor_checker(randt,12,1)
## $conclusion
## [1] "I did not identify any dependency for this lag"
##
## $p_value
## [1] 0.2599424
randbeta <- rbeta (1000,4,4)
autocor_checker(randbeta, k=30, i=1)
## $conclusion
## [1] "I did not identify any dependency for this lag"
##
## $p_value
## [1] 0.7065641
```