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naic.arima.3 = function (model, nruns, t, block) {

# function runs "nruns" simulations

# model needs to be of the Arima type

# starts at an arbitrary place on the actual data and then bootstraps from centered
residuals

# assumes data is log transformed

# extract and center residuals
resid_cent = model$residuals - mean(model$residuals)
min_start = max(model$arima)+2
max_start = length(model$x)

# initiate result matrix
sim_normal = vector(length=nruns)
sim_results = matrix(NA, nrow = nruns, ncol = t)

for (i in 1:nruns) {

#re-initializing vectors

    init_location = 0
    resid_vector = 0
    short_data = 0
    stack = ceiling(t/block)

#creating run-specific

#random selection of simulation starting location is at the core of "through
the cycle"

    init_location = sample((min_start:max_start), size=1)
    init_location2 = init_location-1

```

```

resid_vector = vector()

for (j in 1:stack) {
  # samples blocks of residuals to create a new vector for simulation

  resid = 0
  resid = sample((max_start-block), size= 1)
  resid_vector = append(resid_vector, resid_cent[resid:(resid+block)])
}

resid_vector = resid_vector[1:t] # truncates vector so that length = t
short_data = model$x[1:init_location2]

# create an Arima object and simulate

sim_arima = Arima(short_data, model = model) # version 1
sim_normal[i] = model$x[init_location]
sim_results [i,] = simulate (sim_arima, nsim = t, bootstrap = FALSE, innov =
resid_vector)

}

results = cbind(sim_normal,sim_results)
results = exp(results) # assuming log model
results = results/results[,1]
return (results)

}

```