



# An Open Source Software Reliability Tool: A Guide for Contributors

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## Abstract

This document describes a detailed procedure to integrate a new Software Reliability Growth Model into an Open Source Software Reliability Tool named SRT. SRT is designed in a modular approach to integrate any new or existing contributed model. Most part of the Model Architecture is loosely coupled with the Tool architecture, making it easier for the model contributors to focus on mathematical details of the model rather than on the design elements of tool architecture. The tool architecture validates the model and integrates it transparently. Contributors can visualize the application of their model on the failure data immediately after their model is validated. This flexible architecture allows reliability researchers and practitioner to integrate their own models and use the tool under Open-Source license agreement. The contributors can also compare and contrast their models with other models based on the robustness in terms of data handling and efficiency in terms time complexity. For the model to be valid the contributors should follow few procedural steps and conventions which are essential for model integration. The following sections serves as a guide to contributors of model with necessary details of overall tool architecture.

## Index Terms

GitHub, R statistical programming language, software reliability, software reliability growth model

## Acronyms

BM	Bisection method
CASRE	Computer-Aided Software Reliability Estimation tool
DSS	Delayed S-shaped
EM	Expectation-maximization algorithm
FC	Failure counts
FI	Failure intensity
FT	Failure times
GM	Geometric model
GO	Goel-Okumoto model
IF	Interfailure times
JM	Jelinski-Moranda
LL	Log-likelihood function
MTTF	Mean time to failure
MVF	Mean value function
NHPP	Non-homogeneous Poisson process
NM	Newton's method
SMERFS	Statistical Modeling and Estimation of Reliability Functions for Software
SRGM	Software reliability growth models
SRT	Software reliability tool
Wei	Weibull SRGM

## Notation

$m(t)$	MVF of NHPP
$\lambda(t)$	Instantaneous failure rate
$F(t)$	Cumulative distribution function of software fault detection process
$a$	Number of latent faults at start of testing
$b$	Scale parameter of Weibull SRGM testing
$n$	Observed number of faults
$t_i$	Time of the $i^{th}$ failure
$t_n$	$n$ th observed failure

## I. INTRODUCTION

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**T**HE modern society is highly dependent on software critical systems for safety and functionality. Therefore, these software intensive systems must be highly reliable to ensure safety. In other words, software reliability is a key to success of software systems, which is defined as the probability of failure free operation during a specific period of time in a specific environment. Typically software reliability growth models (SRM) are used to fit the observed data, which enables important metrics such as number of remaining faults, reliability, mean time to failure (MTTF), optimal release time and so on. Software reliability engineering [1] is a well established field, which is over four decades old. In this period, researchers have made significant contribution to the software reliability by developing large number of models and methods in response to the growing needs of the practitioner community. However, research has not always reached its target audience, namely software and reliability engineers. Often, these audiences are reluctant to apply software reliability models because they lack either the knowledge of the underlying mathematics or the time to develop expertise and implement models in their work.

Several automated tools developed for software reliability engineering include: SMERFS (Statistical Modeling and Estimation of Reliability Functions for Software) [2] which was incorporated into CASRE (Computer-Aided Software Reliability Estimation tool) [3] and SRATS (Software Reliability Assessment Tool on Spreadsheet) [?]. Although popular among the user community, the primary shortcoming of these previous tools is that they were implemented as desktop applications by a single researcher, which limited the models incorporated to those the individual researcher was familiar with. Another shortcoming was the need of more polished user interface to allow the user with insufficient software reliability knowledge to access the tool without much difficulty. Also, these tools are dependent on certain operating systems, which may throw some challenges in terms of stability and robustness of the tool.

To overcome the limitation of previous software reliability tools, an open source software reliability tool (SRT) is presented to promote collaboration among members of the international software reliability research community and users from industry and government organizations. The application is implemented in the R programming language, an open source environment for statistical computing and graphics. It is also accessible through a web-enabled framework web at [sasdlc.org](http://sasdlc.org) (System and software development life cycle). The code is accessible through the GitHub repository at [github.com/lfiordella/SRT](https://github.com/lfiordella/SRT). Its architecture will enable incorporation of existing software reliability models into a single tool, enabling more systematic comparison of models than ever before. Presently, interfailure time (IF), failure time (FT), and failure count (FC) data formats are supported. Implemented functionality includes: two trend tests for reliability growth, two failure rate, Jelinski-Moranda (JM) [1] and geometric (GM) [1], and three failure counting models, Goel-Okumoto (GO) [4], [1], delayed S-shaped (DSS) [5], and Weibull (Wei) [6] models as well as two measures of goodness of fit. Inferences such as the time to achieve a target reliability or detect a specified number of additional failures have also been implemented.

This paper explains the tool architecture and describes the steps involved in adding a new model to the tool. The purpose of this document is to serve as a handbook for reliability researchers and practitioner community to help them integrate their model without much difficulty. The paper is organized as follows: Section II presents the tool architecture, Section III presents the model architecture, Section sec:modelSub describes the model submission process, Section V includes details about the model testing, Section VI illustrates the model integration process through an example, and Section VII concludes the document.

## II. INPUT DATA FORMAT

The actual Implementation is simplified in the below diagram to let the contributors focus on the model implementation.

Data Format:

The input to the tool is in the form file. The file can be either CSV or an excel sheet. Both CSV and excel sheets will have header row. The tool scans for this header row with column names “FT”, “FN”, “IF”, “CFC”, “TI”. The failure data can be of either failure times type or failure counts type. For failure type data the header should have “FN” and either “IF” or “IF”. On the other hand for the failure count type data, the header should have columns with names “TI” and either “CFC” or “FC”. The user should make sure that the headers are formatted so for the tool to process the failure data.

A sample data set below is a standard dataset ‘SYS1’ from hand-book of Software Reliability which is a cumulative failure time data\_set.

---

SYS1						
[1]	3	33	146	227	342	...
[13]	836	860	968	1056	1726	...
[25]	3098	3278	3288	4434	5034	...
...	...	...	...	...	...	...
[133]	81542	82702	84566	88682		

---

This data-set is in the form of list data structure. When a data-sheet which is formatted according to tool requirements is uploaded then the data takes up the below format.

This data should be formatted with “FT” column for failure times and “FN” column for failure count. “FN” here is optional. The tool when provided with such input data file, will convert it to the following format for computations. The additional columns are computed and a dataframe is generated with columns with all “FN”, “IF” and “FT” columns as show below.

---

```

> data <- data.frame("FT"=SYS1)
> generate_dataframe(data)
      FT  IF  FN
1      3   3   1
2     33  30   2
3    146 113   3
...    ...  ...  ...

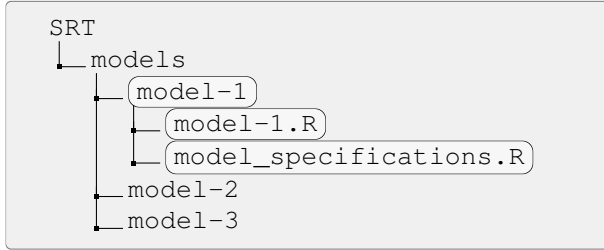
```

---

### III. MODEL ARCHITECTURE

This section describes how the software reliability growth models can be incorporated into the tool architecture. To ensure modularity, the model architecture has been decoupled from the tool using an interface called model-specifications, which is defined for each model in a separate file named 'model\_specifications.R'. This model specifications interface provides the details required for computations whenever a user request is made through the application's graphical user interface. The server then delegates the request to the specific model. Model functionality is defined in a file named MODEL METHOD TYPE.R. The bounded boxes STYLE1 here represents the placeholder. First bounding box here represents MODEL. The MODEL could be replaced with possible short name of a model i.e. GO. Likewise the second bounding box represents the METHOD, which could take values like BM, EM etc. The third bounding box here represents the TYPE of the input model can take, can be either FT or FC. Similar bounding box convention with different style STYLE2 is used in the model specification file and the model implementation file. The parameters set in the model specifications file are substituted in the implementation details of the model for computation along with the request object parameters by server on each request.

Two kinds of placeholders for parameters, one for input request object and other for Model specifications parameters. User request object is sent from the client to the shiny end as input object and the parameters such as MODEL, TODO: stands for Model request.



- 1) Model configuration.
- 2) Model design.

#### A. Model Configuration

This section describes the interface file model\_specifications.R. The model specifications file defines the model properties which will be used by server to autofill the implied additional details of the request for computation. The below code section is a templated version of the actual format and serves as the reference for the new model contributors. The code section contains the placeholders of STYLE1 and STYLE2. STYLE1 is the input request object placeholder. In other words its a client request for a specific model eg. GO and the STYLE1 placeholder will be replaced with the GO in the server code and further details of the model is obtained by sourcing the model\_specifications.R file. Sourcing is a means to place objects in the global environment for later reference. This way when there is request the server knows exactly what user wants and fills in the additional details in the model implementation function invocations.

---

```

# New model
MODEL_input <- c("FT"/"FC"/"IF")
MODEL_methods <- c("BM"/"NM"/...)
MODEL_params <- c("a1", "a2", ..., "an")
MODEL_type <- c("NHPP", "Exp", "S-shaped")
MODEL_numfailsparm <- c(Index of failure counts parameter)
MODEL_fullname <- c(string MODEL FULL NAME)
MODEL_plotcolor <- c(string COLOR)
MODEL_Finite <- bool TRUE/FALSE
# MODEL_prefer_method <- c("BM")

```

The following section explains the purpose of the each specification. `(MODEL)_input` defines the type of input model can take. For example GO Model requires the Failure Time(FT) data. So when user uploads the data and requests for the GO implementation for model fit, the server fills the detail that the GO requires FT data and the `data$FT` column is passed as a list instead of any other possibility. This reduces the overhead of users to specify what type of data should be passed for each model. This also enabled the tool framework to give simultaneous results for the list of models.

`(MODEL)_method` defines the list of models the given method can run. The efficiency of the model depends on the type of algorithm used to obtain a solution. The `GO_methods` parameter uses the bisection method to converge on to solution. This parameter is used by parameter estimation function. The list also serves as reference for the researchers to know what kind of algorithms are implemented for a given model.

`(MODEL)_params` defines the list of names of the parameters of the given model. This parameter is important for server to pass the parameter list to required function invocation. A dataframe of the solutions with this list as names of the solution is used throughout the server implementation for convenience. So the model implementation heavily relies on the naming convention as declared here. The `GO_params` list is defined as list of `amle`, `bmle`, `cmle`. So the model implementation should use this fact whenever they need parameter substitution. For example `GO_MVF` function defined in the `GO_BM_FT.R` file will use `params$amle` whenever it needs to substitute `amle` parameter in an equation. The server uses these facts and fills in the values for computation.

`(MODEL)_type` defines the type of model. Its not a mandatory parameter. Its for future purposes.

`(MODEL)_numfailsparm` is an internal implementaion detail helping the server to know ahat of a model. ahat is a representation for the estimate of number of failures.

`(MODEL)_fullname` defines the fullname of the model. Through the server implementation short name/ abbreviated form is preferred. But the fullname declared here will be used to replace the sections where the naive user should know the fullname of the model. The fullnames are used in almost all the user interface elements for convenience but short forms are used for model implementation.

`(MODEL)_plotcolor` defined the color of the model result in plots of the model results. This overwrites the default implementation, which is random color, for consistency and is not a mandatory parameter.

`(MODEL)_FINITE` defines if the given model is finite or infinite model. Finite models have limited capability in predicting the next  $n$  failures, whereas INFINITE models could predict any number of next failures. This parameter is a boolean value TRUE or FALSE. If TRUE value is assigned then the model is finite model otherwise its not.

`(MODEL)_prefer_method` defines the preferred method among the defined list of methods. This is to override the default method. The default method is always the first in the list of `(MODEL)_methods`.

An example model-specifications file is shown below.

```
# Goel_okumoto model
GO_input <- c("FT")
GO_methods <- c("BM")
GO_params <- c("aMLE", "bMLE")
GO_type <- c("NHPP", "Exp")
GO_numfailsparm <- c(1)
GO_fullname <- c("Goel-Okumoto")
GO_plotcolor <- c("green")
GO_Finite <- FALSE
# GO_prefer_method <- c("BM")
```

## B. Model Design

This section describes the functions that needs to be defined in the `(MODEL).R` file. The check list of functions:

Example model specifications:

1) *Estimating parameters*: This section describes how to define a function which will evaluate estimates of the parameters for a given model. The template for this function is:

```
"MLE function"
(MODEL)_METHOD_MLE <- function((MODEL)_input) {
  ...
  (MODEL)_params <- data.frame(
    a1=xxx,
    a2=xxx,
```

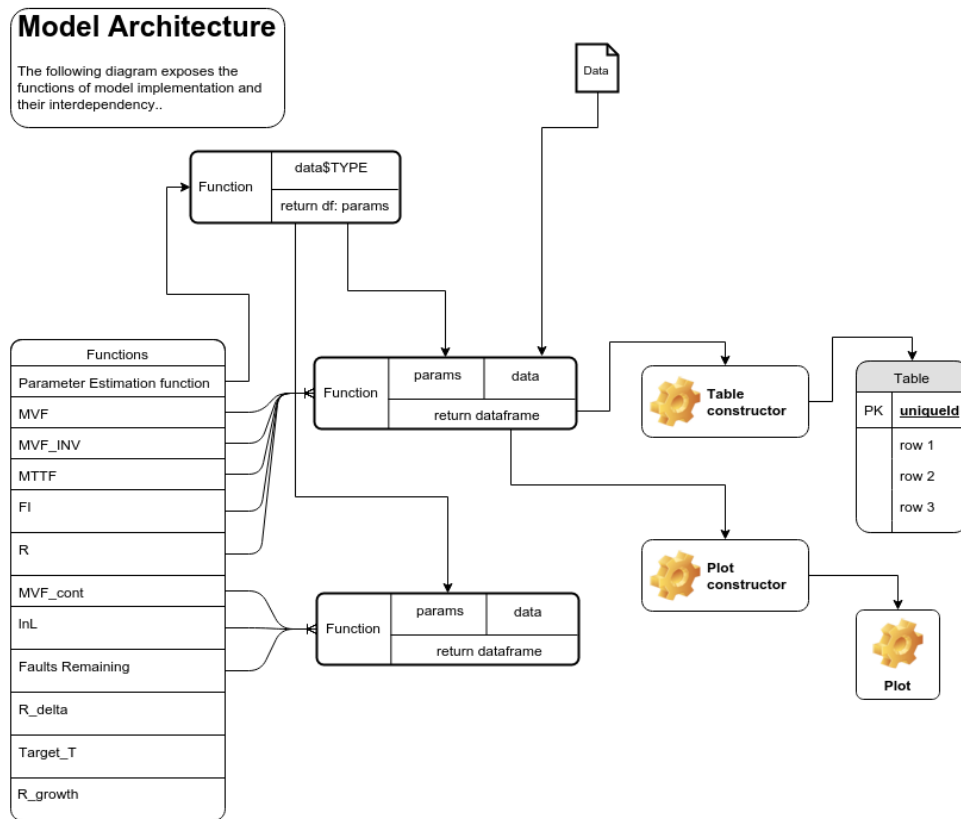


Figure 1: Model Architecture

```

    ... .. /
    an=xxx)
return (MODEL_params)
}

```

Example: GO - Goel-Okumoto

```

GO_BM_MLE <- function(x) {
# -----
# Maximum likelihood estimation method is used to estimate the parameters
# 'N0' and 'Phi'. The initial estimate leftinterval and right interval are expanded
# untill the zero of the equation is bracketed. The bisection method is used for
# bracketing the zero of equation hence the name BM in the name of function with
# '_' seperator. The zero of the expression is the maximum likelihood estimation of
# N0 and the corresponding Phi is derived.
# -----
#
# @params      : interFail (list)          Input interfailure vector
#
# @returns     : JM_params (data.frame) Dataframe of 'N0' and 'Phi'
#               JM_params$N0 refers to N0 and JM_params$Phi refers to Phi
# =====
...

GO_params <- data.frame(
  "GO_aMLE"=aMLE,
  "GO_bMLE"=bMLE)
return(GO_params)
}

```

}

a) *Naming convention:* The name of the function for parameter estimation function depends on the method of obtaining the parameters and the model itself. The method here is the algorithm to converge onto the solution for parameter estimation. “\_MLE” at the end of the function name is rather coined as Maximum Likelihood Estimation approach. Even this MLE could be variable in the next versions of the tool, but for version-1.0 of SRT the MLE is rather fixed since all models are based on Maximum Likelihood Estimation Approach.

b) *Input and return types:* The input for (MODEL)\_METHOD\_MLE is a list of inter failure data or cumulative failure time data or failure count data for inter failure count data. The type of input the model should take is defined in the model\_specifications.R file of the individual model folder. The server input the right type of data for the model on behalf of users. So the users duty to mention the right type of data in (MODEL)\_input is important part and rest is done by the tool. The return types for (MODEL)\_METHOD\_MLE function is dataframe with names of (MODEL)\_params and values of corresponding parameter. Order in which the parameters are named in a dataframe is not important. The values are retrieved by the names of the parameters rather than the by indexing.

2) *MVF:* Mean Value Function is function which estimates the number of failures at a given time. The path of the mean value function plot traces the original cumulative failures data.

```
(MODEL)_MVF <- function(params,(DATA))
{
  ...
  MVF_list <- c(...) # Numeric
  MVF <- data.frame(MVF_list,
                    (DATA)$TYPE,
                    rep((MODEL),n))
  names(MVF) <- c("Failure", "Time", "Model")
  return(MVF)
}
```

Example: GO - Goel-Okumoto

```
GO_MVF <- function(param,d) {
#-----
# This function computes the MVF data frame
# MVF - Mean Value Function
#-----
# @params      (data.frame)      Data.frame of parameters
# @d           (data.frame)      Data.frame of data FT,FC,FN,CFC,IF
#
# @returns     (data.frame)      data.frame of Failure, Time, Model columns
#-----

n <- length(d$FT)
r <- data.frame()
MVF <- param$GO_aMLE*(1-exp(-param$GO_bMLE*d$FT))
r <- data.frame(MVF,d$FT,rep("GO", n))
names(r) <- c("Failure", "Time", "Model")
return(r)
}
```

a) *Naming convention:* The name of the mean value function depends on the model name alone, but the name “Mean Value Function” is abbreviated as MVF instead for convenience. So in effect the naming convention for Mean Value Funtion (MVF) of a (MODEL) is: (MODEL)\_MVF

b) *Input and return types:* The input to the MVF function is a dataframe of parameters calculated by MLE function. The names of “params” is defined as in model\_specifications file. Along with the “params” parameter, Data is passed. The return of the MVF function should be dataframe with columns names as “Failure”, “Time”, “Model”

3) *MVF\_INV:* The MVF\_INV is the abbreviation for Inverse of Mean Value Function. The inverse of a function is a transpose of the domain and range of function. Equivalently a swap of X and Y axis of the plot.

```
(MODEL)_MVF_Inv <- function(params,(DATA))
{
  ...
  MVF_list <- c(...) # Numeric
  MVF_INV <- data.frame(MVF_list,
```

```

      (DATA)$ (TYPE),
      rep((MODEL), n) )
names(MVF_INV) <- c("Failure", "Time", "Model")
return(MVF_INV)
}

```

Example: GO - Goel-Okumoto

```

GO_MVF_inv <- function(param,d) {
#-----
# This does an "inverse" MVF function, solving for time given
# a specific value of MVF.
#-----

# @params      (data.frame)      Data.frame of parameters
# @d           (data.frame)      Data.frame of data FT,FC,FN,CFC,IF

# @returns     (data.frame)      data.frame of Failure, Time, Model columns
#-----

n <- length(d$FN)
r <- data.frame()
cumFailTimes <- -(log((param$GO_aMLE-d$FN)/param$GO_aMLE))/param$GO_bMLE
r <- data.frame(d$FN,cumFailTimes, rep("GO", n))
names(r) <- c("Failure", "Time", "Model")
return(r)
}

```

a) *Naming convention:* The naming convention is similar to MVF function with INV as abbreviation for inverse as :

(MODEL)\_MVF\_Inv

b) *Input and return types:* The input to the MVF\_Inv function is a dataframe of parameters calculated by MLE function. The names of “params” is defined as in model\_specifications file. Along with the “params” parameter, Data is passed. The return of the MVF\_Inv function should be dataframe with columns names as “Failure”, “Time”, “Model”

4) *MTTF:* The Mean Time To Failure function calculates the average number of failures that occurred in a given interval.

```

(MODEL)_MTTF <- function(params,(DATA))
{
  ...
  MTTF_list <- c(...) # Numeric
  MTTF <- data.frame(MVF_list,
    (DATA)$ (TYPE),
    rep((MODEL), n) )
  names(MTTF) <- c("Failure", "Time", "Model")
  return(MTTF)
}

```

Example: GO - Goel-Okumoto

```

GO_MTTF <- function(param,d) {
#-----
# This function MTTF of given d with parameters
# MTTF is Mean Time To Failure
#-----

# @params      (data.frame)      Data.frame of parameters
# @d           (data.frame)      Data.frame of data FT,FC,FN,CFC,IF

# @returns     (data.frame)      data.frame of Failure, MTTF, Model columns
#-----

n <- length(d$FT)
r <- data.frame()

```



```

currentTimes <- utils::tail(d, length(d$FT)-1)
prevTimes <- utils::head(d, length(d$FT)-1)
currentFailNums <- c(2:n)
prevFailNums <- c(1:(n-1))
IFTimes <- ((currentFailNums*currentTimes$FT)
            / (param$GO_aMLE*
              (1-exp(-param$GO_bMLE*currentTimes$FT))))
            - ((prevFailNums*prevTimes$FT)
              / (param$GO_aMLE*(1-exp(-param$GO_bMLE*prevTimes$FT))))
r <- data.frame(c(1, currentFailNums),
               c(((d$FT[1]) / (param$GO_aMLE*(1-exp(-param$GO_bMLE*d$FT[1])))),
                 IFTimes),
               rep("GO", n))
names(r) <- c("Failure_Number", "MTTF", "Model")
return(r)
}

```

a) *Naming convention:* The naming convention of this function is dependent on model name appended with abbreviated form of Mean Time To Failure as follows: (MODEL)\_MTTF

b) *Input and return types:* The input to the MTTF function is a dataframe of parameters calculated by MLE function. The names of “params” is defined as in model\_specifications file. Along with the “params” parameter, Data is passed. The return of the MTTF function should be dataframe with columns names as “Failure”, “MTTF”, “Model”

5) *FI:* The failure intensity function defines the number of failures in a given interval.

```

(MODEL)_FI <- function(params, (DATA))
{
  \ldots
  MTTF_list <- c(...) # Numeric
  MTTF <- data.frame(MVF_list,
                    (DATA)$ (TYPE),
                    rep((MODEL), n))
  names(MTTF) <- c("Failure", "Time", "Model")
  return(MTTF)
}

```

Example: GO - Goel-Okumoto

```

GO_FI <- function(param,d) {
#-----
# This function computes the Failure Intensity for a given data
# with parameters 'param' of a given data
#-----
# @params      (data.frame)      Data.frame of parameters
# @d           (data.frame)      Data.frame of data FT,FC,FN,CFC,IF
#
# @returns     (data.frame)      data.frame of Failure Count, Failure_Rate, Model columns
#-----

n <- length(d$FT)
r <- data.frame()
fail_number <- c(1:n)
failIntensity <- param$GO_aMLE*param$GO_bMLE*exp(-param$GO_bMLE*d$FT)
r <- data.frame(fail_number,failIntensity, rep("GO", n))
names(r) <- c("Failure_Count", "Failure_Rate", "Model")
return(r)
}

```

a) *Naming convention:* The naming convention of Failure Intensity function is model name appended with abbreviated form of Failure Intensity(FI) as below: (MODEL)\_FI

b) *Input and return types:* The input to the FI function is a dataframe of parameters calculated by MLE function. The names of “params” is defined as in model\_specifications file. Along with the “params” parameter, Data is passed. The return of the FI function should be dataframe with columns names as “Failure\_Count”, “Failure\_Rate”, “Model”

6) *lnL*: The *lnL* stands for loglikelihood function.

```
MODEL_lnL <- function((DATA),params)
{
  ...
  lnL <- numeric
  return(lnL)
}
```

Example: GO - Goel-Okumoto

```
GO_lnL <- function(x,params) {

#-----
# This computes Log-Likelihood for a given data x and parameters
#-----
# @params      (data.frame)      Data.frame of parameters
# @x           (data.frame)      Data.frame of data FT,FC,FN,CFC,IF

# @returns     (numeric)         Numeric value of lnL
#-----

n <- length(x)
tn <- x[n]
firstSumTerm <- 0
for(i in 1:n){
  firstSumTerm = firstSumTerm + (-params$GO_bMLE*x[i])
}
lnL <- -(params$GO_aMLE)*(1-exp(-params$GO_bMLE*tn))
      + n*(log(params$GO_aMLE)) +n*log(params$GO_bMLE)
      + firstSumTerm
return(lnL)
}
```

a) *Naming convention*: The naming convention for *lnL* function is model name appended with “\_lnL” shown as below:

MODEL\_lnL

b) *Input and return types*: The input to the *lnL* function is a dataframe of parameters calculated by MLE function. The names of “params” is defined as in model\_specifications file. Along with the “params” parameter, Data is passed. The return of the *lnL* function should be a numeric data type.

7) *MVF\_cont*: The *mvf\_cont* is a continuous form of Mean value function. Unlike MVF function which return a dataframe of mvf values “mvf\_cont” returns mvf value at a particular time.

```
MODEL_MVF_cont <- function(params,t) {
  return(MVF(t))
}
```

Example: GO - Goel-Okumoto

```
GO_MVF_cont <- function(params,t) {

#-----
# This function computes MVF at a particular time
# This is a continuous function of time hence the name 'cont'
#-----
# @params      (data.frame)      Data.frame of parameters
# @t           (numeric)         time

# @returns     (numeric)         MVF value at time t
#-----

return(params$GO_aMLE*(1-exp(-params$GO_bMLE*t)))
}
```

a) *Naming convention:* The mvf\_cont is similar MVF function except this is appended with \_cont defining that this is a continuous form a MVF function. `(MODEL)_MVF_cont`

b) *Input and return types:* The input to the mvf\_cont function is a dataframe of parameters calculated by MLE function. The names of “params” is defined as in model\_specifications file. Along with the “params” parameter, time at which mvf should be calculated is passed. The return of the mvf\_conf function should be a numeric data type.

8) *R\_delta:* This function computes the change in reliability with change in time.

```
(MODEL)_R_delta <- function(params,cur_time,delta){
  return(exp(-(MODEL)_MVF_cont(params,(cur_time+delta))
             -(MODEL)_MVF_cont(params,cur_time))))
}
```

Example: GO - Goel-Okumoto

```
GO_R_delta <- function(params,cur_time,delta){
  #-----
  # This function computes the Change in Reliability with delta change in time
  #-----
  # @params      (data.frame)      Data.frame of parameters
  # @cur_time    (numeric)         current time -> time at which reliability is calculated
  # @delta       (numeric)         delta time -> (t' - t), t' is delta away from current time

  # @returns     (numeric)         Change in reliability with delta change in time
  #-----

  return(exp(-(GO_MVF_cont(params,(cur_time+delta))
                -GO_MVF_cont(params,cur_time))))
}
```

a) *Naming convention:* The naming convention is as follows: `(MODEL)_R_delta.R`

b) *Input and return types:* The input to the R\_delta function is a dataframe of parameters calculated by MLE function. The names of “params” is defined as in model\_specifications file. Along with the “params” parameter, time at which reliability should be calculated is passed and delta change in time is passed. The return of the R\_delta function should be a numeric data type.

9) *R\_Root:* Root finding utility to determine the target reliability.

```
(MODEL)_R_MLE_root <- function(params,cur_time,delta, reliability){
  root_equation <- reliability - R
  return(root_equation)
}
```

Example: GO - Goel-Okumoto

```
GO_R_BM_root <- function(params,cur_time,delta, reliability){
  #-----
  # This defines the function required for root finding target reliability
  #-----
  # @params      (data.frame)      Data.frame of parameters
  # @cur_time    (numeric)         current time -> time at which reliability is calculated
  # @delta       (numeric)         delta time -> (t' - t), t' is delta away from current time
  # reliability   (numeric)         reliability

  # @returns     (function)        Return a function for uniroot evaluation used by JM_Target_T
  #-----

  root_equation <- reliability -
    exp(params$GO_aMLE*(1-exp(-params$GO_bMLE*cur_time))
        -params$GO_aMLE*(1-exp(-params$GO_bMLE*(cur_time+delta))))
  return(root_equation)
}
```

a) *Naming convention:* The naming convention is not strictly defined since this is an internal function to the model, but the model\_name should be used at the beginning (MODEL\_R\_Root

b) *Input and return types:* The input to the R\_Root function is a dataframe of parameters calculated by MLE function. The names of “params” is defined as in model\_specifications file. Along with the “params” parameter, time at which reliability should be calculated is passed and delta change in time is passed. The return of the R\_Root function should be a numeric data type.

10) *Target\_T:* This

```
(MODEL)_Target_T <- function(params,cur_time,delta, reliability){
  return(sol)
}
```

Example: GO - Goel-Okumoto

```
GO_Target_T <- function(params,cur_time,delta, reliability){
  #-----
  # This computes the time it takes to achieve the target reliability
  #-----
  # @params      (data.frame)      Data.frame of parameters
  # @cur_time    (numeric)         current time -> time at which reliability is calculated
  # @delta       (numeric)         delta time -> (t' - t), t' is delta away from current time
  # @reliability (numeric)         reliability

  # @returns     (numeric)/        time it takes to achieve the target reliability
  #              (character)       string message if target reliability is already achieved
  #-----

  f <- function(t){
    return(GO_R_MLE_root(params,t,delta, reliability))
  }

  current_rel <- GO_R_delta(params,cur_time,delta)
  if(current_rel < reliability){
    # Bound the estimation interval

    sol <- 0
    interval_left <- cur_time
    interval_right <- 2*interval_left
    local_rel <- GO_R_delta(params,interval_right,delta)
    while (local_rel <= reliability) {
      interval_right <- 2*interval_right
      if(local_rel == reliability) {
        interval_right <- 2.25*interval_right
      }
      if (is.infinite(interval_right)) {
        break
      }
      local_rel <- GO_R_delta(params,interval_right,delta)
    }
    if(is.finite(interval_right) && is.finite(local_rel) && (local_rel < 1)) {
      while (GO_R_delta(params,
        (interval_left +
          (interval_right-interval_left)/2),
        delta) < reliability) {
        interval_left <- interval_left + (interval_right-interval_left)/2
      }
    } else {
      sol <- Inf
    }
  }
}
```

```

}

if (is.finite(interval_right) && is.finite(sol)) {
  sol <- tryCatch(
    stats::uniroot(f,
      c(interval_left, interval_right),
      extendInt="yes",
      maxiter=maxiter,
      tol=1e-10)$root,
    warning = function(w) {
      #print(f.lower)
      if(length(grep("_NOT_ converged",w[1]))>0) {
        maxiter <- floor(maxiter*1.5)
        print(paste("recursive", maxiter, sep='_'))
        GO_Target_T(a,b,cur_time,delta, reliability)
      }
    },
    error = function(e) {
      print(e)
      #return(e)
    })
  } else {
    sol <- Inf
  }
} else {
  sol <- "Target reliability already achieved"
}

return(sol)
}

```

11) *R\_growth*: This section describes the inverse of mean value function definition.

```

MODEL_R_growth <- function(params,data,delta) {
  g <- data.frame(r[1],r[2],r[3])
  names(g) <- c("Time", "Reliability_Growth", "Model")
  return(g)
}

```

Example: GO - Goel-Okumoto

```

GO_R_growth <- function(params,d,delta) {
  #-----
  # This function computes the reliability growth
  #-----

  # @params      (data.frame)      Data.frame of parameters
  # @d           (data.frame)      Data.frame of data FT,FC,CFC,IF
  # @delta       (numeric)         delta time -> (t' - t), t' is delta away from current time

  # @returns     (data.frame)      Data frame of Time,Reliability Growth, Model
  #-----
  r <- data.frame()
  for(i in 1:length(d$FT)) {
    r[i,1] <- d$FT[i]
    temp <- GO_R_delta(params,d$FT[i],delta)
    if(typeof(temp) != typeof("character")) {
      r[i,2] <- temp
      r[i,3] <- "GO"
    }
  }
  else{

```

```

        r[i,2] <- "NA"
        r[i,3] <- "GO"
    }
}
g <- data.frame(r[1],r[2],r[3])
names(g) <- c("Time", "Reliability_Growth", "Model")
#print(g)
return(g)
}

```

#### IV. MODEL SUBMISSION

A model can be submitted to SRT through GitHub, which implements a series of model validation phases to provide the contributor with feedback so that they can understand if all of their modules have been accepted. To begin submission the contributor must be a GitHub user. From GitHub perspective the contributor should fork srt-repository. After a successful fork process, the forked repository will be visible within the contributor's profile. This forked repository is your own personal repository and changes made to this repository will not affect the original srt-repository, so a user does not need to worry that experimentation will destabilize the tool. The next step is to clone this forked repository to your local machine. Once the forked repository is cloned to your local machine, you should be able to see the models directory which is similar in structure mentioned in (ref). As a model contributor, any additions or modifications required should be performed within this models directory. To contribute a new model, you will need to create a new folder with name same as your model's short name. Make sure that your model-design file is named with the model shortname and possesses an 'R' extension at the end.

In addition to the model-design file, the tool requires the model-specifications file. These model specifications and model-design files should be within the model (MODEL) folder. After performing these steps, you can submit the model for integration. To submit the model, the changes made must first be pushed to your own repository. This can be accomplished in two ways, namely the Terminal friendly or GUI approach.

1) *Terminal approach*: Execute the following sequence of commands to push the code. Through the command line, change to the srt clone-directory. Then, check for git status with:

```
git status
```

This command lists the files changed so far. Next, add the changes you made to your local machine repository with:

```
git add MODEL_METHOD_APPROACH.R
git add model_specifications.R
```

To make sure these files are added to your local repository, use the git status command as above.

The next step is to commit the changes made to push them to remote forked-repository with the following command:

```
git commit -am 'Message for easy trace back'
```

Finally, push the code to remote repository.

```
git push origin master
TODO: remotes add
```

This pushes your local machine code to your online repository. Be sure to check this operation is reflected in your online forked repository. Refer to the pull-request section to submit your code to the original repository.

2) *Web-GUI approach*: The new model should be tested against the standard data-sets and the results of the data-sets should be saved in the same folder. This helps other contributors ensure that the tool is working as defined. The GitHub procedure is as follows: TODO.

##### A. Pull-Request

In the GitHub GUI browse to the forked repository as shown in figure. TODO: Needs different image and correct annotation.

Checklist : TODO

#### V. MODEL TESTING

After successfully submitting a model, the third party continuous integration service Travis-CI will test the models with basic unit testing and then enters the integration testing phase. Unit testing makes ensures proper naming conventions are followed and that required functions are and proper specifications are defined. Integration testing ensures that the functions

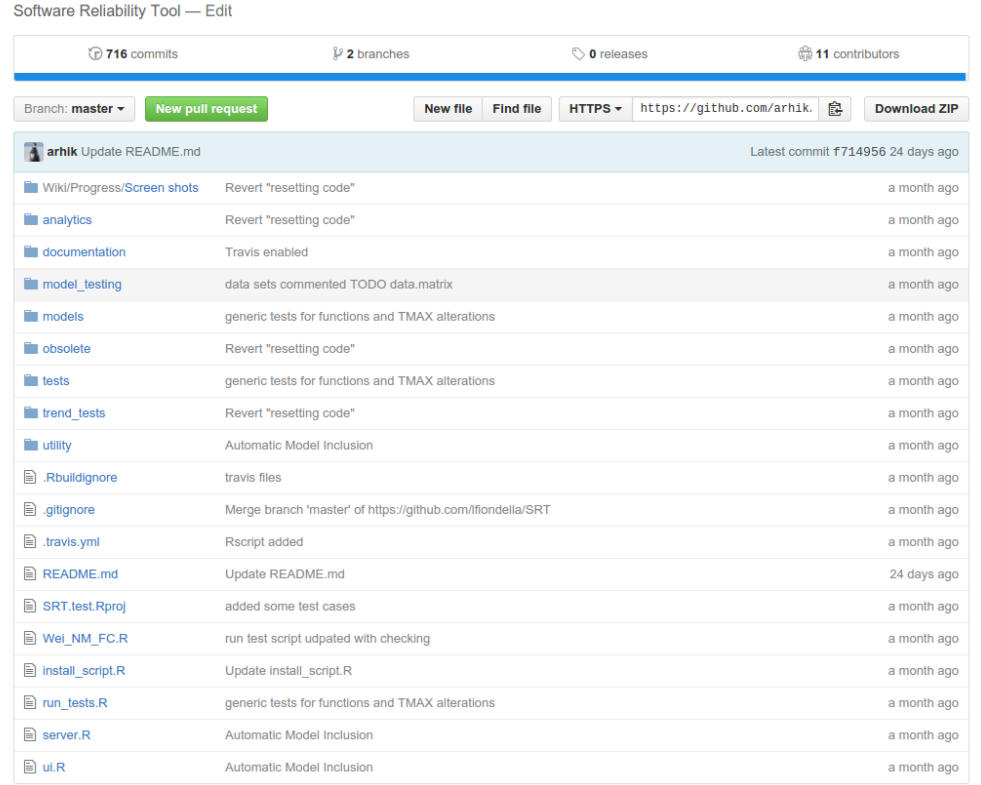


Figure 2: GUI interface

defined behave as expected. The return types of each function are tested to verify that they produce the expected data-structures as well as values which are valid for computations to carry on.

Click on the travis reference for additional details.

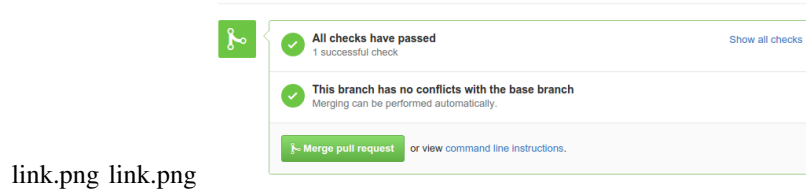


Figure 3: Travis link

This redirects you to the build phase of the model. The interface contains the build log of the tool.

The build log helps you to debug your model in case testing fails. Figure 5 shows an example build log when the test is passed.

If any of the test fails then the build log will contain FAIL statements which require further inspection in the model design or model specifications to resolve.

## VI. ILLUSTRATION

This section illustrates the model integration procedure described in Section III in the context of Goel-Okumoto model considering failure times data, which is one of the model implemented in the software reliability tool. GO SRGM is a finite exponential type nonhomogeneous Poisson process (NHPP) software reliability model. We have chosen bisection method (BM) [7] to estimate the model parameters for the sake of illustration.

### A. Goel-Okumoto NHPP SRGM

The nonhomogeneous Poisson process is a stochastic process [8] that counts the number of events that occur by time  $t$ . The expected value of an NHPP is characterized by the mean value function (MVF), denoted  $m(t)$ . The MVF can take many

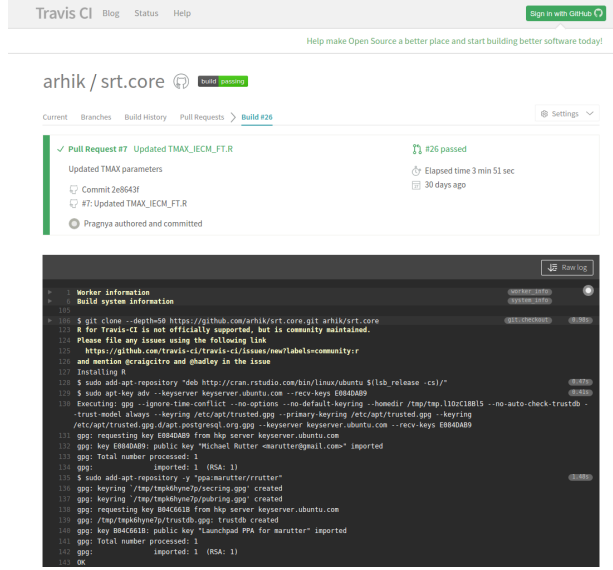


Figure 4: Travis Interface

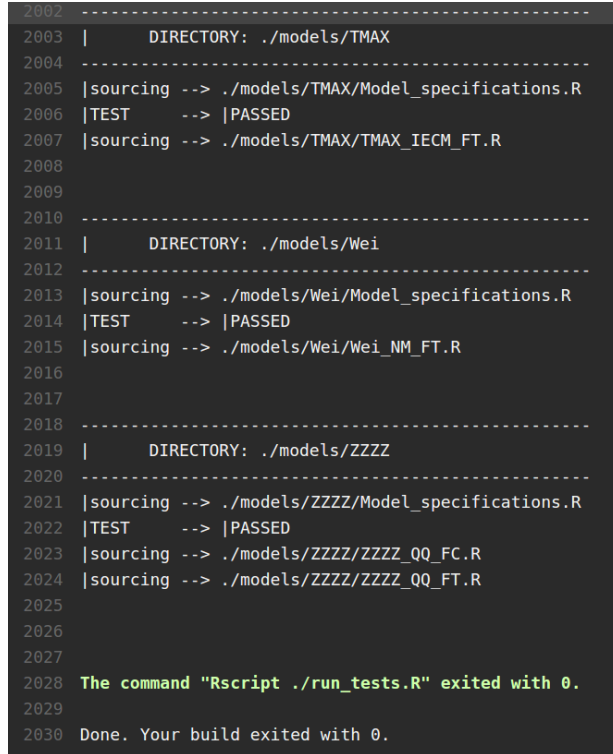


Figure 5: Travis Build Log

functional forms. In the context of software reliability, the NHPP counts the number of faults detected after the software has been tested for a given period of time. Okamura *et al.* [9] noted that the MVF of several SRGM can be written as

$$m(t) = a \times F(t), \quad (1)$$

where  $a$  denotes the number of faults to be detected with infinite testing and  $F(t)$  is the CDF of a continuous probability distribution, characterizing the software fault detection process.

The rate of occurrence of failures is time varying with instantaneous failure rate, which is known as failure intensity (FI)

$$\lambda(t) = \frac{dm(t)}{dt}. \quad (2)$$



1	FN	IF	FT
2		1	3
3		2	30
4		3	113
5		4	81
6		5	115
7	...	...	...

Figure 6: Excel input file format

Using Equation (1), the MVF of the Goel-Okumoto model, which was originally proposed by Goel and Okumoto [10] is

$$m(t) = a(1 - e^{-bt}), \quad (3)$$

where  $a$  is the number of faults present in software at the beginning of testing and  $b$  is the fault detection rate.

### B. Input data

Consider a failure times data such as SYS1 data set [1] from the Handbook of software reliability, which is of the form

---

```

SYS1
[1]      3      33      146      227      342      ...
[13]    836     860     968    1056    1726      ...
[25]   3098    3278    3288    4434    5034      ...
...      ...      ...      ...      ...      ...
[133] 81542  82702  84566  88682

```

---

The raw data set is in the form of a list and this should be formatted to make it compatible with the tool, which is of the format

---

```

> data <- data.frame("FT"=SYS1)
> generate_dataframe(data)
      FT  IF  FN
1      3   3   1
2     33  30   2
3    146 113   3
...    ...  ...

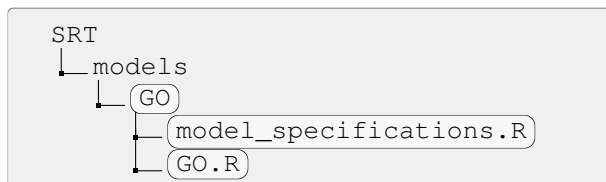
```

---

However, it is sufficient to specify the "FT" column as it is a failure times data and this will be automatically converted to the failure count and interfailure times format by the tool. The "IF" and "FN" column shows an example of the two other types of input data sets. The data format will be identified by the tool by reading the column headings such as "FT". Figure 6 shows a screenshot of the data format in excel spreadsheet. However, data can also be formatted as a csv (comma separated) file.

### C. Model Design

In the SRT architecture, every model is programmed using two functional files as shown below.



**Configure and Apply Models**

Specify the number of failures for which the models will make predictions

Specify for how many failures into the future the models will predict

1

Choose one or more models to run, or exclude one or more models.

Goel-Okumoto

Run Selected Models

Figure 7: Goel-Okumoto model in user interface

1) *model\_specifications.R*: The "model\_specifications.R" file for GO SRGM is as shown below

---

```
GO_input <- c("FT")
GO_methods <- c("BM")
GO_params <- c("aMLE", "bMLE")
GO_type <- c("NHPP", "Exp")
GO_numfailsparm <- c(1)
GO_fullname <- c("Goel-Okumoto")
GO_plotcolor <- c("green")
GO_Finite <- TRUE
# GO_prefer_method <- c("BM")
```

---

The detailed description of this file is described here;

- First line specifies the input file format, which is failure times for GO, indicated as "FT" formatted as specified in Section VI-B.
- Second line specifies the parameter estimation method, which is in this case is bisection method, indicated as "BM".
- Third line specifies the model parameters defined as *aMLE* and *bMLE*, as GO model is characterized by two model parameters *a* and *b* as in Equation (1).
- Fourth line specifies that GO is "NHPP" type exponential ("EXP") model.
- Fifth line indicates the parameter representing the initial number of faults, which is *a*, indicated as *aMLE* in the specifications.
- Sixth line specifies the full name of the model, which will be displayed in the user interface in the second tab as shown in Figure 7 below .
- Seventh line specifies green color for GO model, which is optional.
- Eight line indicates that GO is a "finite" model.
- Last line indicates the preferred model evaluation method if there were more than method specifies in second line.

2) *GO.R*: This section describes all the functions that needs to be written for any model to fulfill the basic functionalities incorporated in the tool. These routines will be syntactically compatible with the SRT naming conventions to get through the travis-CI test. These functions are:

- **GO\_BM\_MLE.R**: This routine estimates the model parameters *a* and *b* for GO model using bisection method utilizing the MLE equations given by

$$aMLE = \frac{n}{1 - e^{-bt}}$$

and,

$$bMLE = \frac{n}{aMLE t_n e^{-bt_n} + \sum_{i=1}^n t_i}.$$

These equations are implemented in R language as

```
GO_BM_MLE <- function(x) {
  ...
  GO_params <- data.frame(
    "GO_aMLE"=aMLE,
    "GO_bMLE"=bMLE)
```

```

    return(GO_params)
}

```

However, the BM implementation is shown by ... here and once the parameters are estimated, it will be assigned to variables prefixed by model name such as  $GO_aMLE$  and  $GO_bMLE$ , which will be returned to the called function.

- **Mean value function:** The MVF of the model is implemented through three different routines:

- 1) **GO\_MVF.R:** routine implements the mean value function of the GO model given in Equation (3) as

```

GO_MVF <- function(param,d) {
  n <- length(d$FT)
  r <- data.frame()
  MVF <- param$GO_aMLE*(1-exp(-param$GO_bMLE*d$FT))
  r <- data.frame(MVF,d$FT,rep("GO", n))
  names(r) <- c("Failure", "Time", "Model")
  return(r)
}

```

- 2) **GO\_MVF\_inv.R:** This routine transposes the observed failure data and returns a data frame, which will be compared with the estimated values graphically. This is implemented in R as

```

GO_MVF_inv <- function(param,d) {
  n <- length(d$FN)
  r <- data.frame()
  cumFailTimes <- -(log((param$GO_aMLE-d$FN)/param$GO_aMLE))/param$GO_bMLE
  r <- data.frame(d$FN,cumFailTimes, rep("GO", n))
  names(r) <- c("Failure", "Time", "Model")
  return(r)
}

```

- 3) **GO\_MVF\_cont.R:** This routine returns the MVF given in Equation (3) with the estimated parameter values from the GO\_BM\_MLE.R routine. This will be compared against the original data computed in GO\_MVF\_inv.R routine. This routine is implemented in R as

```

GO_MVF_cont <- function(params,t){
  return(params$GO_aMLE*(1-exp(-params$GO_bMLE*t)))
}

```

These three functions will enable comparison of the observed data with the fitted MVF in Tab 2 of the tool graphically as shown in Figure 8 for SYS1 data set.

- **GO\_lnL.R:** This routine implements the log-likelihood function of the GO SRGM given by

$$LL(a,b|\mathbf{T}) = -aMLE(1 - e^{-bMLEt_n}) + \sum_{i=1}^n \log(aMLEbMLEe^{-bMLEt_i}). \quad (4)$$

The log-likelihood function in R is written as

```

GO_lnL <- function(x,params){
  n <- length(x)
  tn <- x[n]
  firstSumTerm <- 0
  for(i in 1:n){
    firstSumTerm = firstSumTerm + (-params$GO_bMLE*x[i])
  }
  lnL <- -(params$GO_aMLE)*(1-exp(-params$GO_bMLE*tn))
  + n*(log(params$GO_aMLE)) + n*log(params$GO_bMLE)
  + firstSumTerm
  return(lnL)
}

```

- **GO\_FL.R:** implements the failure intensity of the GO model given by

$$\lambda(t) = abe^{-bt} \quad (5)$$

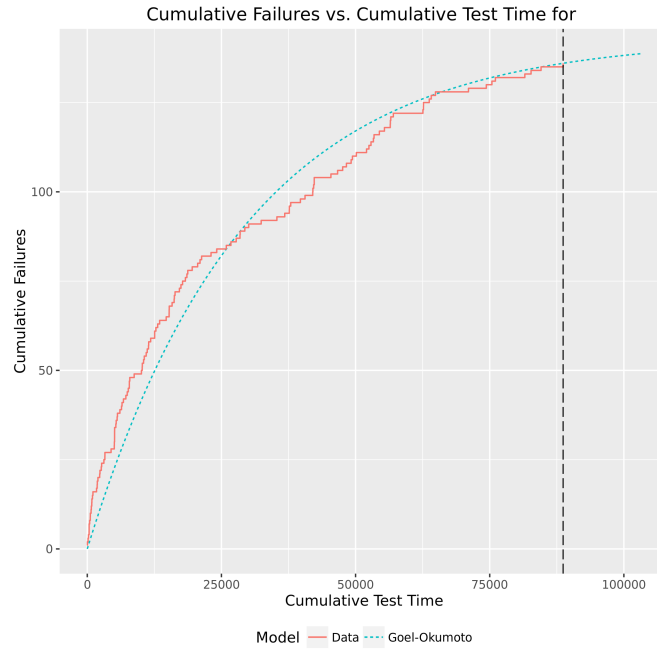


Figure 8: SYS1 cumulative failure and model fit

The failure intensity function is written in R as

```
GO_FI <- function(param,d) {
  n <- length(d$FT)
  r <- data.frame()
  fail_number <- c(1:n)
  failIntensity <- param$GO_aMLE*param$GO_bMLE*exp(-param$GO_bMLE*d$FT)
  r <- data.frame(fail_number,failIntensity, rep("GO", n))
  names(r) <- c("Failure_Count", "Failure_Rate", "Model")
  return(r)
}
```

This is displayed in the second tab of the tool as shown in Figure 9

- **GO\_MTTF**: This routine implements the mean time to failure of the GO model given by

$$\begin{aligned} MTTF &= \int_0^{\infty} R(t)dt \\ &= \frac{1}{b} \end{aligned} \quad (6)$$

This is implemented in R as follows

```
GO_MTTF <- function(param,d) {
  n <- length(d$FT)
  r <- data.frame()
  currentTimes <- utils::tail(d, length(d$FT)-1)
  prevTimes <- utils::head(d, length(d$FT)-1)
  currentFailNums <- c(2:n)
  prevFailNums <- c(1:(n-1))
  IFTimes <- ((currentFailNums*currentTimes$FT)
    / (param$GO_aMLE*(1-exp(-param$GO_bMLE*currentTimes$FT)))
    - ((prevFailNums*prevTimes$FT)
    / (param$GO_aMLE*(1-exp(-param$GO_bMLE*prevTimes$FT))))
  r <- data.frame(
    c(1, currentFailNums),
```

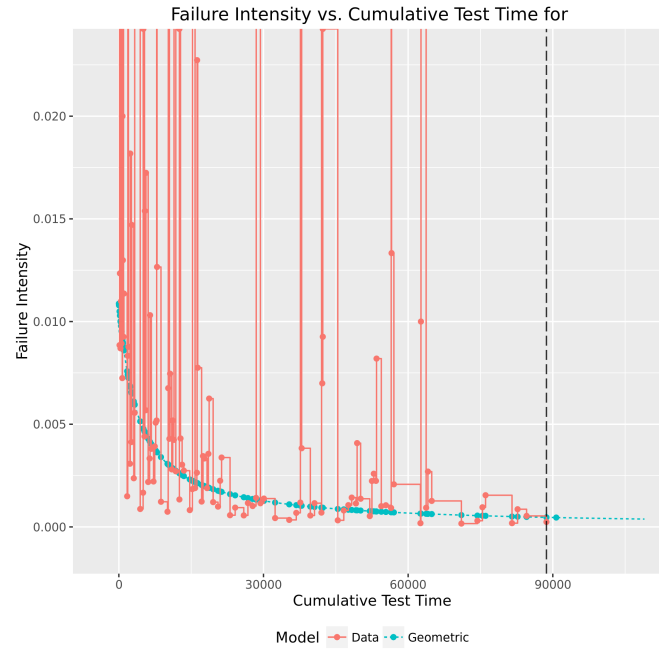


Figure 9: Failure intensity for SYS1 data set

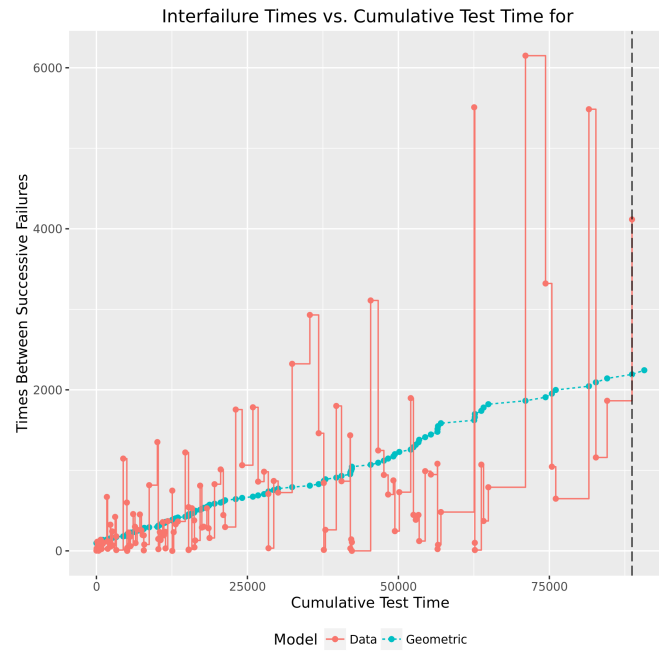


Figure 10: Time between failures for SYS1 data set

```

    c(((d$FT[1]) / (param$GO_aMLE * (1 - exp(-param$GO_bMLE * d$FT[1])))),
      IFTimes),
    rep("GO", n))
names(r) <- c("Failure_Number", "MTTF", "Model")
return(r)
}

```

This is displayed in the second tab of the tool as shown in Figure 10

- 1) **GO\_R\_delta**: This implements the reliability of the GO SRGM given by

$$R(\delta) = e^{-m(t_n + \delta) - m(t_n)} \quad (7)$$

where  $t_n$  indicates the  $n^{th}$  failure data and  $m(t)$  indicates the mean value function as in Equation (3). This is implemented in R as

```
GO_R_delta <- function(params,cur_time,delta){
  return(exp(-(GO_MVF_cont(params,(cur_time+delta))-GO_MVF_cont(params,cur_time))))
}
```

- 2) **GO\_R\_MLE\_root**: This is an intermediate routine written to return the reliability equation to the called function to compute the reliability growth value. In R, it is implemented as

```
GO_R_MLE_root <- function(params,cur_time,delta, reliability){
  root_equation <- reliability -
    exp(params$GO_aMLE*(1-exp(-params$GO_bMLE*cur_time))
    -params$GO_aMLE*(1-exp(-params$GO_bMLE*(cur_time+delta))))
  return(root_equation)
}
```

- 3) **Go\_R\_growth**: calls the GO\_R\_delta routine to compute the reliability growth values, which is implemented as

```
GO_R_growth <- function(params,d,delta){
  r <-data.frame()
  for(i in 1:length(d$FT)){
    r[i,1] <- d$FT[i]
    temp <- GO_R_delta(params,d$FT[i],delta)
    if(typeof(temp) != typeof("character")){
      r[i,2] <- temp
      r[i,3] <- "GO"
    }
    else{
      r[i,2] <- "NA"
      r[i,3] <- "GO"
    }
  }
  g <- data.frame(r[1],r[2],r[3])
  names(g) <- c("Time","Reliability_Growth","Model")
  #print(g)
  g
}
```

This is displayed in the second tab of the tool as shown in Figure 11

- **GO\_Target\_T**: computes the time required to reach the target reliability, which is implemented as below

```
GO_Target_T <- function(params,cur_time,delta, reliability){
  f <- function(t){
    return(GO_R_MLE_root(params,t,delta, reliability))
  }

  current_rel <- GO_R_delta(params,cur_time,delta)
  if(current_rel < reliability){
    # Bound the estimation interval

    sol <- 0
    interval_left <- cur_time
    interval_right <- 2*interval_left
    local_rel <- GO_R_delta(params,interval_right,delta)
    while (local_rel <= reliability) {
      interval_right <- 2*interval_right
    }
  }
}
```

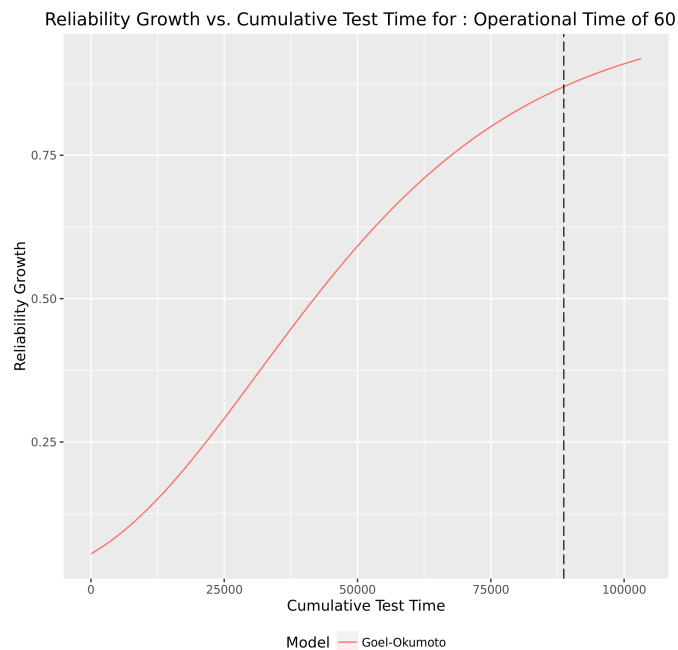


Figure 11: Time between failures for SYS1 data set

```

if(local_rel == reliability) {
  interval_right <- 2.25*interval_right
}
if (is.infinite(interval_right)) {
  break
}
local_rel <- GO_R_delta(params,interval_right,delta)
}
if(is.finite(interval_right) && is.finite(local_rel) && (local_rel < 1)) {
  while (
    GO_R_delta(
      params,
      (interval_left + (interval_right-interval_left)/2),
      delta) < reliability) {
    interval_left <- interval_left + (interval_right-interval_left)/2
  }
} else {
  sol <- Inf
}

if (is.finite(interval_right) && is.finite(sol)) {
  sol <- tryCatch(
    stats::uniroot(f, c(interval_left, interval_right),
      extendInt="yes", maxiter=maxiter, tol=1e-10)$root,
    warning = function(w) {
      #print(f.lower)
      if(length(grep("_NOT_ converged",w[1]))>0) {
        maxiter <- floor(maxiter*1.5)
        print(paste("recursive", maxiter,sep='_'))
        GO_Target_T(a,b,cur_time,delta, reliability)
      }
    },
    error = function(e) {

```

```

        print(e)
        #return(e)
    })
} else {
    sol <- Inf
}
} else {
    sol <- "Target reliability already achieved"
}
return(sol)
}

```

## VII. CONCLUSION

The tool is designed to encapsulate a wide variety of software reliability models. It is anticipated that the tool architecture and corresponding work flow will evolve to incorporate more method to assess software reliability throughout the life cycle. The open-source nature of the tool provides a platform for the international software reliability researcher community to incorporate results from past decades as well as to define a research agenda for the future.

## ACKNOWLEDGMENT

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