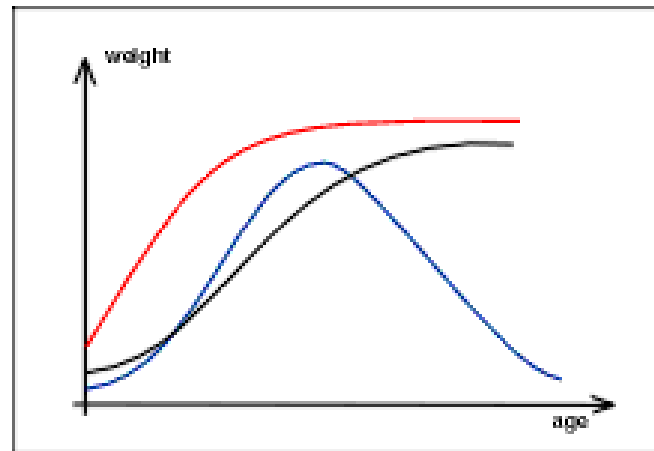


Nonlinear Statistical models for Pest Populations- A case study of Aphid



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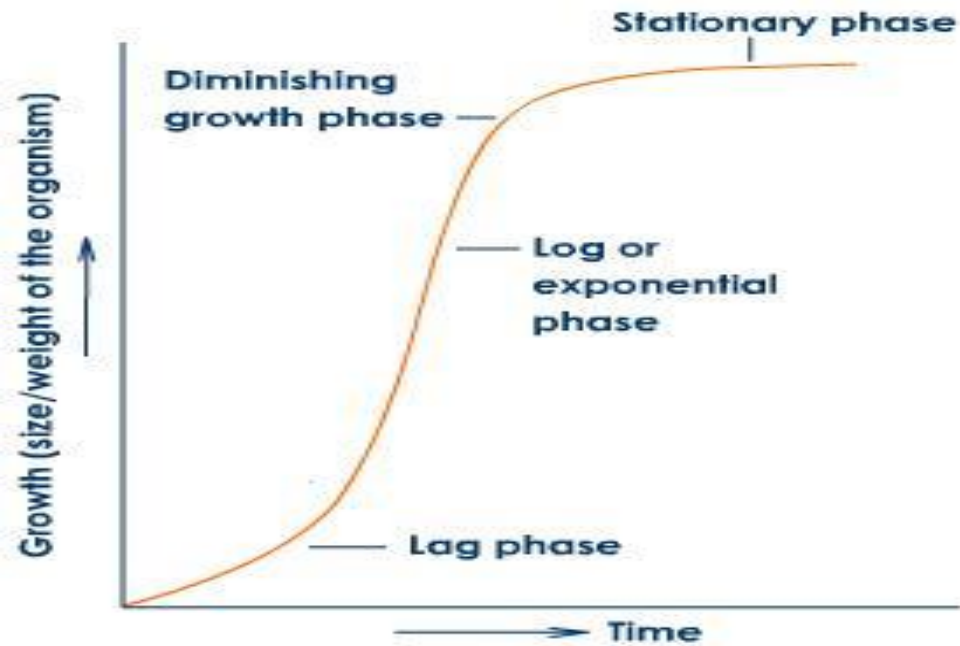
Directorate of Oilseeds Research

Hyderabad

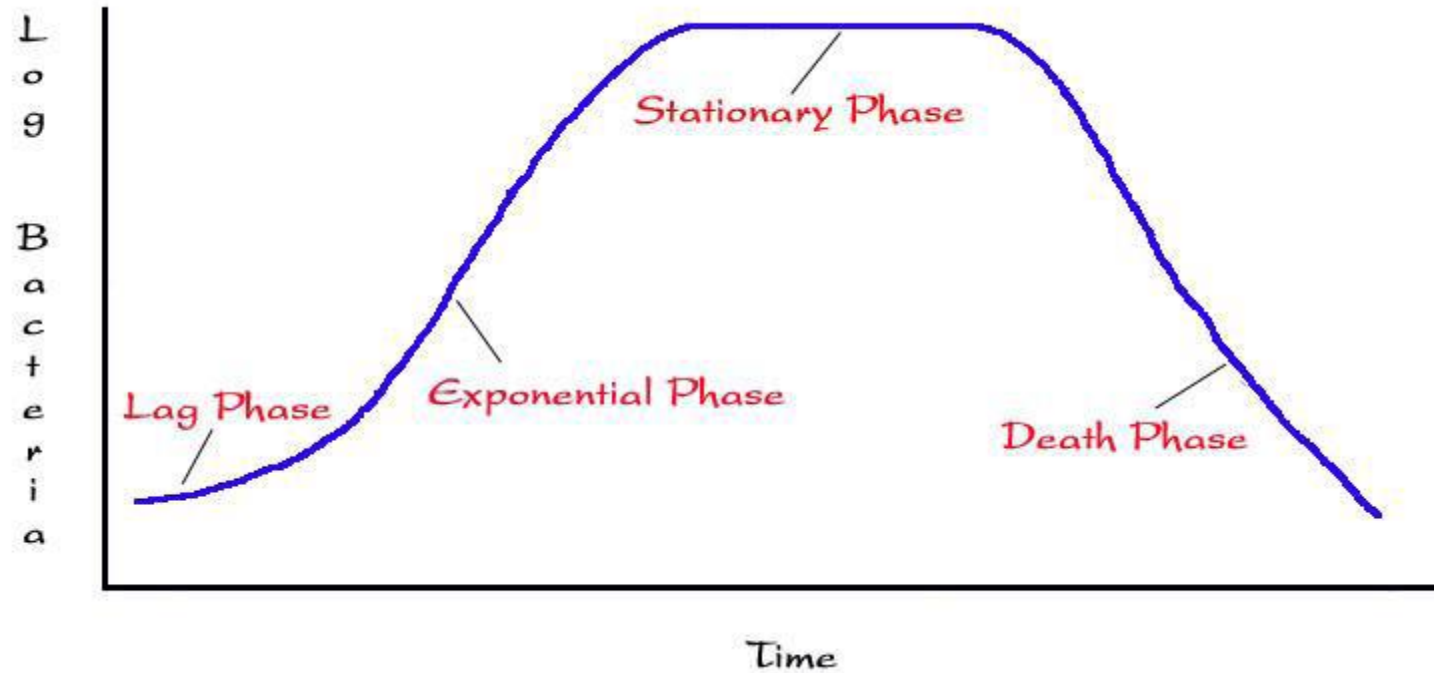
saradac@yhoo.com



Plant / Animal Growth Curve

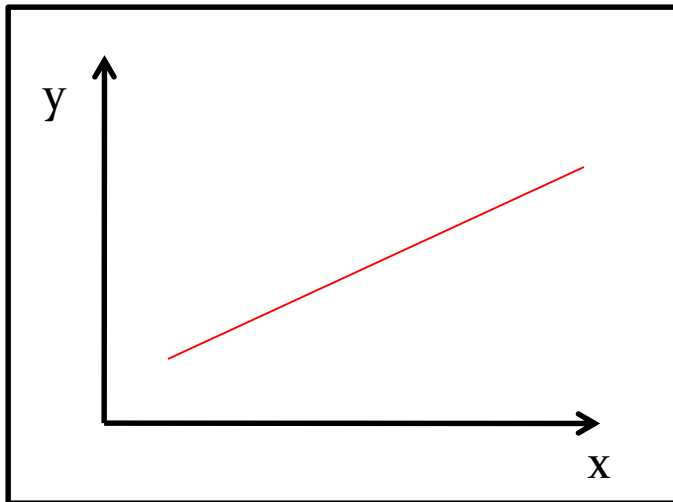


Bacterial Growth Curves



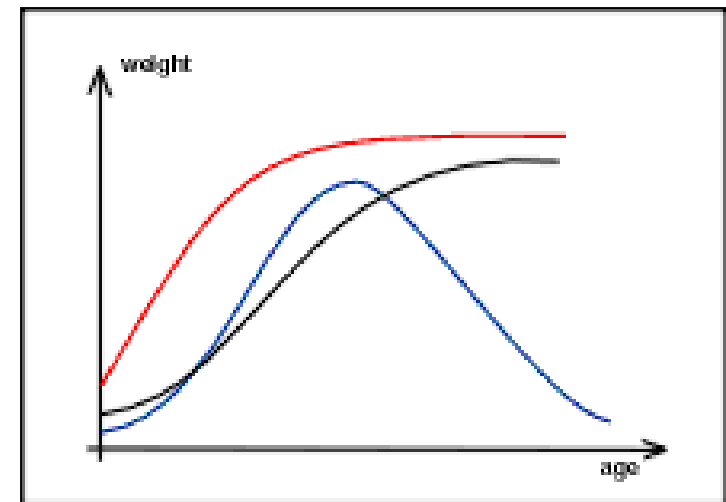
Regression Models

Linear model



$$Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \dots + \beta_p X_p + \varepsilon$$

Nonlinear model



$$Y = f(X_1, X_2, \dots, X_p | \theta_1, \theta_2, \dots, \theta_q) + \varepsilon$$

Nonlinear Model

- A non-linear regression model is one in which at least one of the parameters appear non-linearly.
- The derivatives of the model with respect to the model parameters depends on one or more parameters.
- **Linear Model** $y = b_0 + b_1x + b_2x^2 + e$
- $dy/db_0 = 1$ $dy/db_1 = x$ $dy/db_2 = x^2$
- **Nonlinear Model** $Y = a X^b + e$
- $dy/da = X^b$
 $dy/db = a X^b (\log (X))$

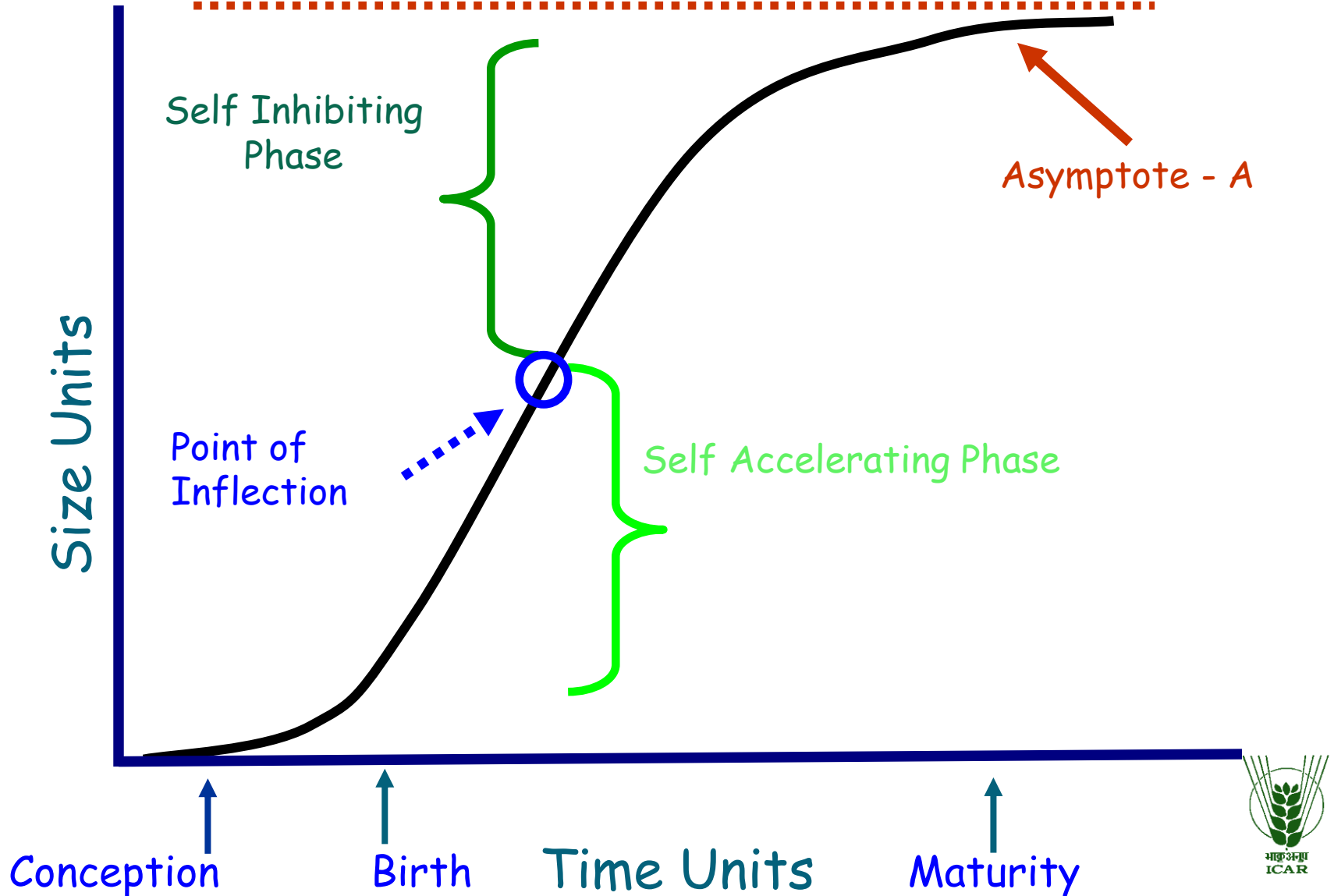
Properties

- Nonlinear models are often derived on the basis of physical and/or biological considerations,.
- The parameters of a nonlinear model usually have direct interpretation in terms of the process under study.
- Constraints can be built into a nonlinear model easily and are harder to enforce for linear models.

Model Development

- Development
 - Statistical model
 - Assumptions
 - Type of Variables considered
- Validation
 - Goodness of fit
 - Model diagnostics
 - Modification of the Model and revalidation
- Implementation
 - Experimental trials
 - Advisory systems / DSS

Phases of the Growth Curve



Some Important Nonlinear Growth Models

Logistic	$Y_t = A/(1 + Be^{-kt}) + \varepsilon$
Gompertz	$Y_t = A \exp(-Be^{-kt}) + \varepsilon$
Von Bertalanffy	$Y_t = A(1 - Be^{-kt}) + \varepsilon$
Mercer –Foldin	$Y_t = (AB + kt^D)/(B + t^D) + \varepsilon$
Wiebull	$Y_t = A - B \exp(-kt^D) + \varepsilon$
Richards	$Y_t = A/(1 + \exp(b - kt))^{\frac{1}{D}} + \varepsilon$

Y_t = Yield /growth /length/weight /biomass /Population etc at time period t

A = Asymptote or mature weight at age t approaches infinity/carrying capacity

B = Initial yield/growth/length/weight/population

k = Growth rate

D = Shape parameter

Fitting of Nonlinear Models

- Parameter Estimation
- Choice of Initial Values
- Goodness of Fit of a Model
- Examination of Residuals

Parameter Estimation

- Nonlinear Least Squares
- Normal equations are nonlinear in parameters
- Iterative procedures
(requires supplying of initial values)
 - Gauss –Newton Method
 - Newton Method
 - Levenberg –Marquardt’s Method
 - Steepest Descent (Gradient Method)

Choice of Initial Values

- Making Initial Guess
- Estimates calculated from previous experiments
- Known values for similar systems
- Values computed from theoretical considerations

- Linearization
- Solving system of equations
- Using properties of the model
- Graphical method

Goodness of Fit statistics

Coefficient of Determination

$$R^2 = 1 - \frac{\sum (Y_i - \hat{Y}_i)^2}{\sum (Y_i - \bar{Y})^2}$$

$$\text{Mean Square Error (MSE)} = \frac{\sum (Y_i - \hat{Y}_i)^2}{(n - p)}$$

$$\text{Mean Absolute Error (MAE)} = \frac{\sum |Y_i - \hat{Y}_i|}{(n)}$$

Examination of Residuals

- Assumptions
 - Errors are independent
 - Plotting of residuals
 - Tests for independence of errors (Runs Test)
 - Errors are normally distributed
 - Plotting Normal probability plot
 - Tests for Normality (Shapiro-Wilk test)

Wald – Wolfowitz Runs Test

- H_0 : Errors are independent
- H_1 : Errors are not independent

$$\text{Mean } (\mu) = 2mn/(m+n)+1$$

$$\text{Variance} = 2mn(2mn-m-n)/(m+n)^2(m+n-1)^{-1}$$

$$Z = (r-h-\mu)/\sigma$$

Where,

$$h = \begin{cases} 0.5, & \text{if } r < \mu \\ -0.5, & \text{if } r > \mu \end{cases}$$

Test for Normality (Shapiro-Wilk test)

- H_0 : Errors are normally distributed
- H_1 : Errors are not normally distributed.

$$S^2 = \sum a(k) \{X_{(n+1-k)} - X_{(k)}\}, b = \sum (X_i - \bar{X})^2$$

Where,

$$K = \begin{cases} 1, 2, \dots, n/2 \text{ when } n \text{ is even} \\ 1, 2, \dots, (n-1)/2 \text{ when } n \text{ is odd} \end{cases}$$

SOFTWARE

- SAS – PROC NLIN
- SPSS – NLR
- S-PLUS / R - NLS
- SAS – Enterprise guide
- SPSS – Nonlinear Regression
- JMP

Model Development

- Development
 - Statistical model
 - Assumptions
 - Type of Variables considered
- Validation
 - Goodness of fit
 - Model diagnostics
 - Modification of the Model and revalidation
- Implementation
 - Experimental trials
 - *Advisory systems / DSS*

Model Diagnostics

$$N(t) = e^{a+bt} (1 + e^{d+bt})^{-2} + \epsilon$$

- Measure of Nonlinearity of parameters
 - Intrinsic Nonlinearity (IN) - Low value indicates unbiased prediction
 - Parameter –effects nonlinearity (PE)
 - Low value indicates good Confidence region /Symmetric confidence intervals
 - Marginal Curvatures -
 - Properties of each individual parameter
 - Profile t – plots - Close – to – linear
 - Skewness, Kurtosis , Bias
 - Reparameterization of the parameters
 - *Simulation* - 1000 data sets - Parameter estimates
 - Histograms – Long right hand tail – exponential of parameter
Long left hand tail - logarithm of the parameter

Development of Aphid Population density Model

$$N(t) = ae^{bt} (1 + de^{bt})^{-2} + \epsilon$$

Prajneshu, (1998)

Model diagnostics

Simulation studies (1000 data sets)

$$N(t) = e^{a+bt} (1 + e^{d+bt})^{-2} + \epsilon$$

Sarada.C and Prajneshu
(2005)

Model Properties

Ross.G, Prajneshu and Sarada.C (2010)

Reparametrized Aphid growth model

C.Sarada *et al.* (2012)

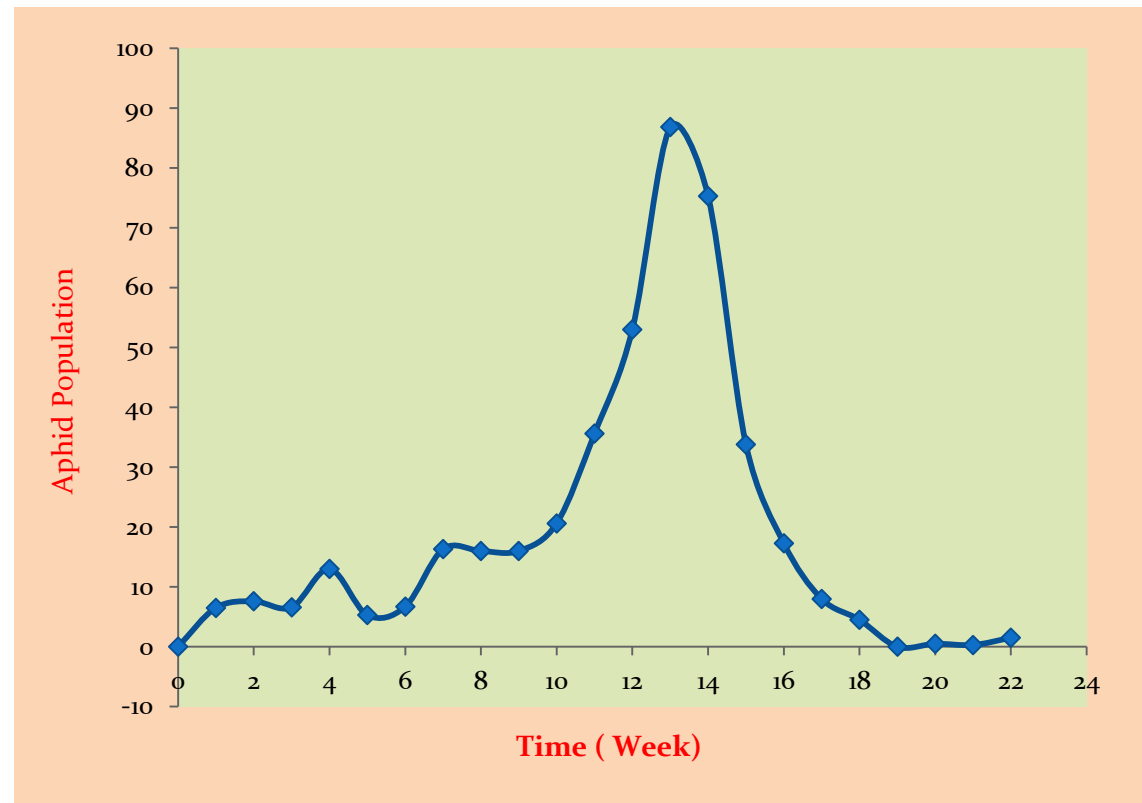
Safflower Aphid population

Development of Aphid Population density Model

Time (weeks)	Aphid Population per 100 leaves
0	0.0
1	6.5
2	7.6
3	6.6
4	13.0
5	5.3
6	6.7
7	16.3
8	16.0
9	16.0
10	20.6
11	35.6
12	53.0
13	86.8
14	75.3
15	33.8
16	17.3
17	8.0
18	4.5
19	0.0
20	0.5
21	0.3
22	1.5

$$N(t) = ae^{bt} (1 + de^{bt})^{-2} + \epsilon$$

Prajneshu, (1998)



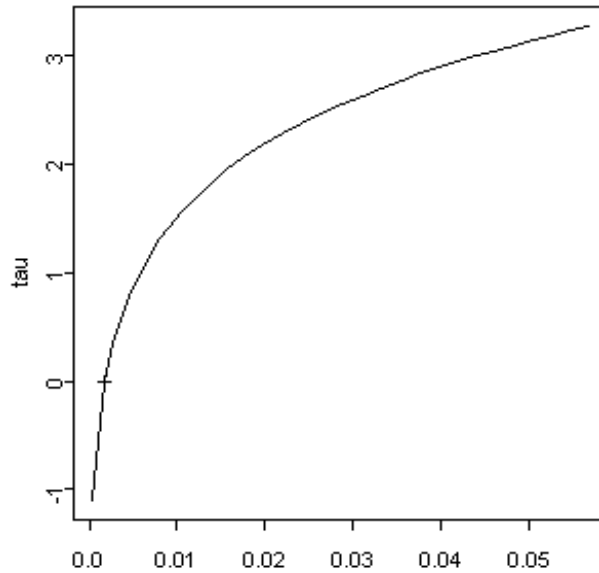
Verma, K.D. and Parihar, S.B.S. (1991). Build up of the vector *Aphis gossypii* glover on potato. *J. Aphidol.*, **5**, 16 - 18.

Model Diagnostics

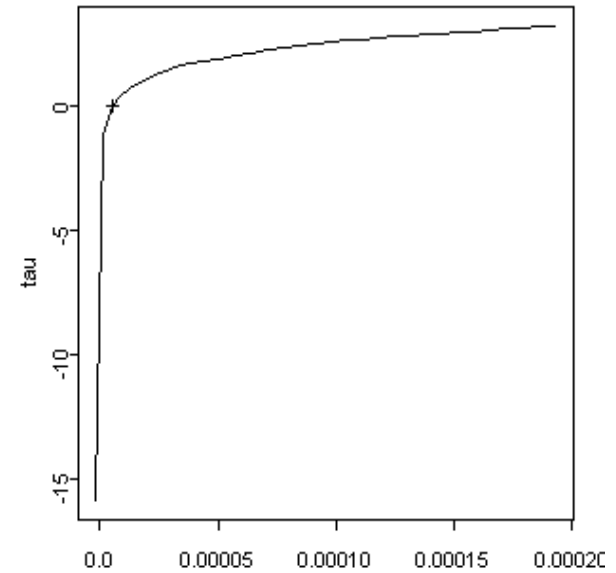
Statistic	Coefficeint	
i. Parameters		
<i>a</i>	0.0018 (0.0017)	
<i>b</i>	0.92 (0.08)	
<i>d</i>	0.0 ⁵ 55 (0.0 ⁵ 59)	
ii Confidence Intervals	Lower	Upper
<i>a</i>	0.001	0.007
<i>b</i>	0.8	1.07
<i>d</i>	0.000	0.0002
iii Measures of Nonlinearity		
IN	0.21	
PE	<u>12.23</u>	
iii. Marginal Curvatures		
<i>a</i>	0.78	
<i>b</i>	0.07	
<i>d</i>	0.82	

Profile t-plots

Parameter - a

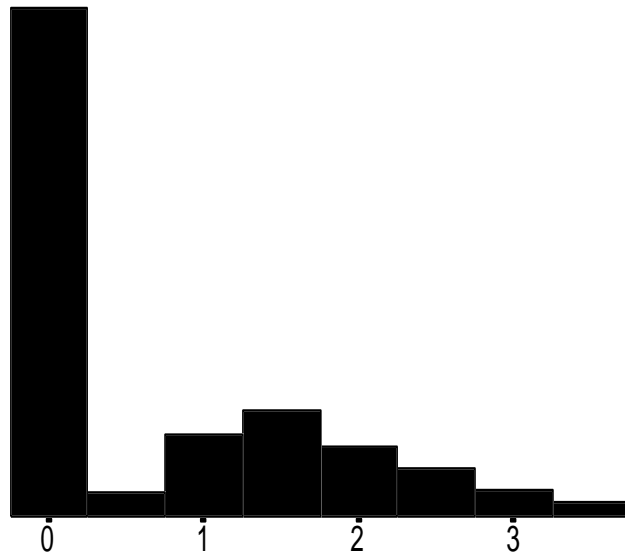


Parameter - d

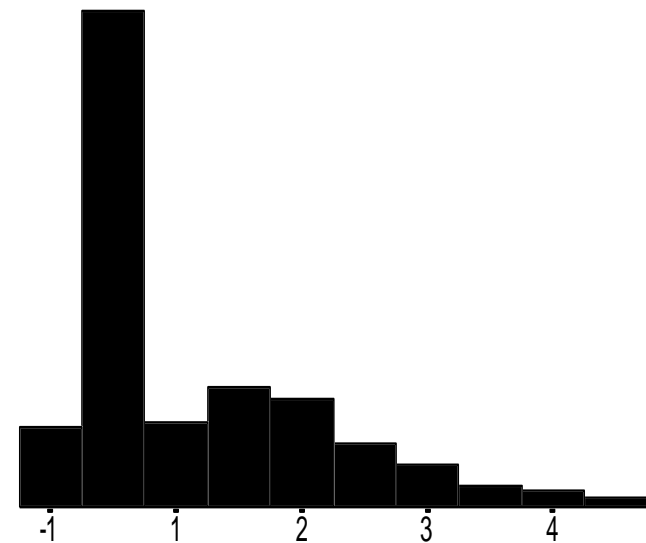


Histograms of 1000 simulated standardized parameter estimates

Parameter - a



Parameter - d



Reparameterization of Aphid Model

Model - I

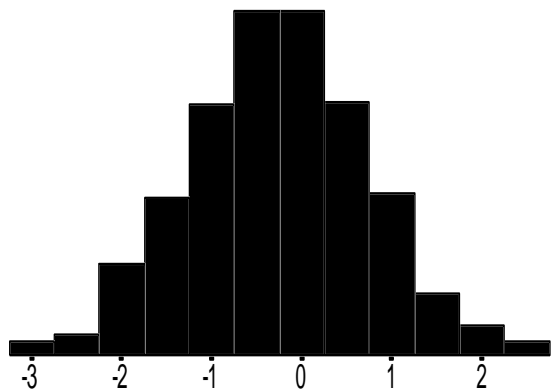
$$N(t) = ae^{bt} (1 + de^{bt})^{-2} + \epsilon$$

Model - II

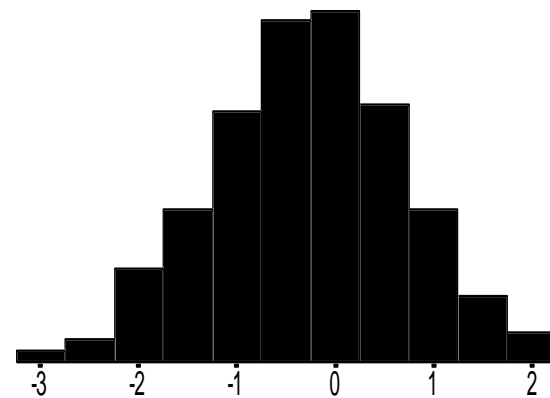
$$N(t) = e^{a+bt} (1 + e^{d+bt})^{-2} + \epsilon$$

Histograms of 1000 simulated standardized parameter estimates for Model -II

Parameter - a

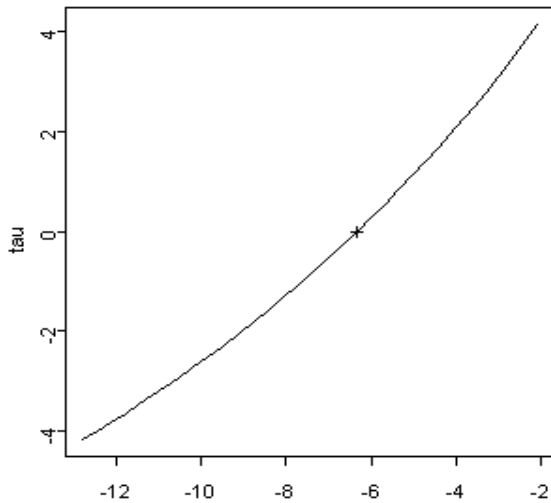


Parameter - d

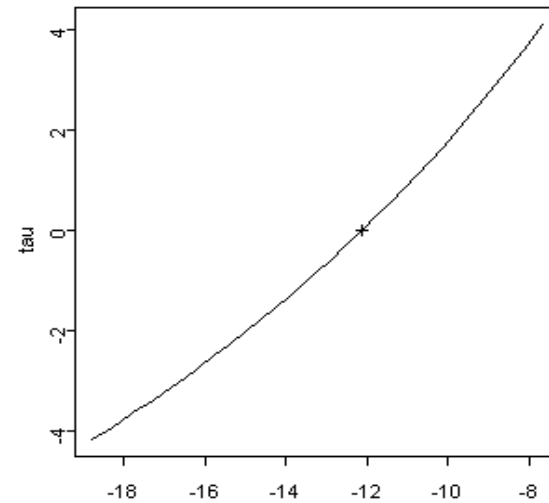


Profile t-plots Model -II

Parameter - a



Parameter - d



Parameter estimation, rms curvatures measures, marginal curvature, measures of nonlinearity and simulation studies

	Model I	Model II		
(i) Parameter estimation:				
a	0.0018 (0.0017)	-6.34 (0.99)		
b	0.92 (0.08)	0.92 (0.08)		
d	0.0 ⁵ 55 (0.0 ⁵ 59)	-12.13 (1.67)		
(ii) 95% Confidence-intervals:				
	Lower	Upper	Lower	Upper
a	0.001	0.0070	-8.17	-4.76
b	0.800	1.0700	0.80	1.07
d	0.000	0.0002	-14.03	-10.04
(iii) rms Curvature effects:				
(IN) $\sqrt{F_{3,20}(0.05)}$	0.21	0.21		
(PE) $\sqrt{F_{3,20}(0.05)}$	12.23	0.24		
(iv) (Marginal curvatures) $t_{20}(0.05)$:				
a	0.78	0.08		
b	0.07	0.07		
d	0.82	0.07		
(v) Simulation Studies:				
Skewness:				
a	0.61**	-0.18 ^{NS}		
d	0.61**	-0.10 ^{NS}		
Kurtosis:				
a	-1.06**	0.29 ^{NS}		
d	1.05**	0.27 ^{NS}		
% Bias:				
a	-23.30**	0.02 ^{NS}		
d	-24.02**	0.01 ^{NS}		

Development of Safflower Aphid Population Model

$$N(t) = e^{a+bt} (1 + e^{d+bt})^{-2} + \epsilon$$

$$C = a(2b^2 d)^{-1}$$

$$N_0 = a(1 + d)^{-2}$$

$$r = (b^2 - 2N_0 C^{-1})^{1/2}$$

N_0 = Initial population

C = carrying capacity

$N(t)$ = population at time t

r = growth rate

Data

- **Safflower variety : CO-1**
- **Years : 2005-10**
- **Region : Solapur**
- **Frequency : Weekly Aphid Population**
- **Source : Safflower Annual Reports, DOR, Hyderabad**

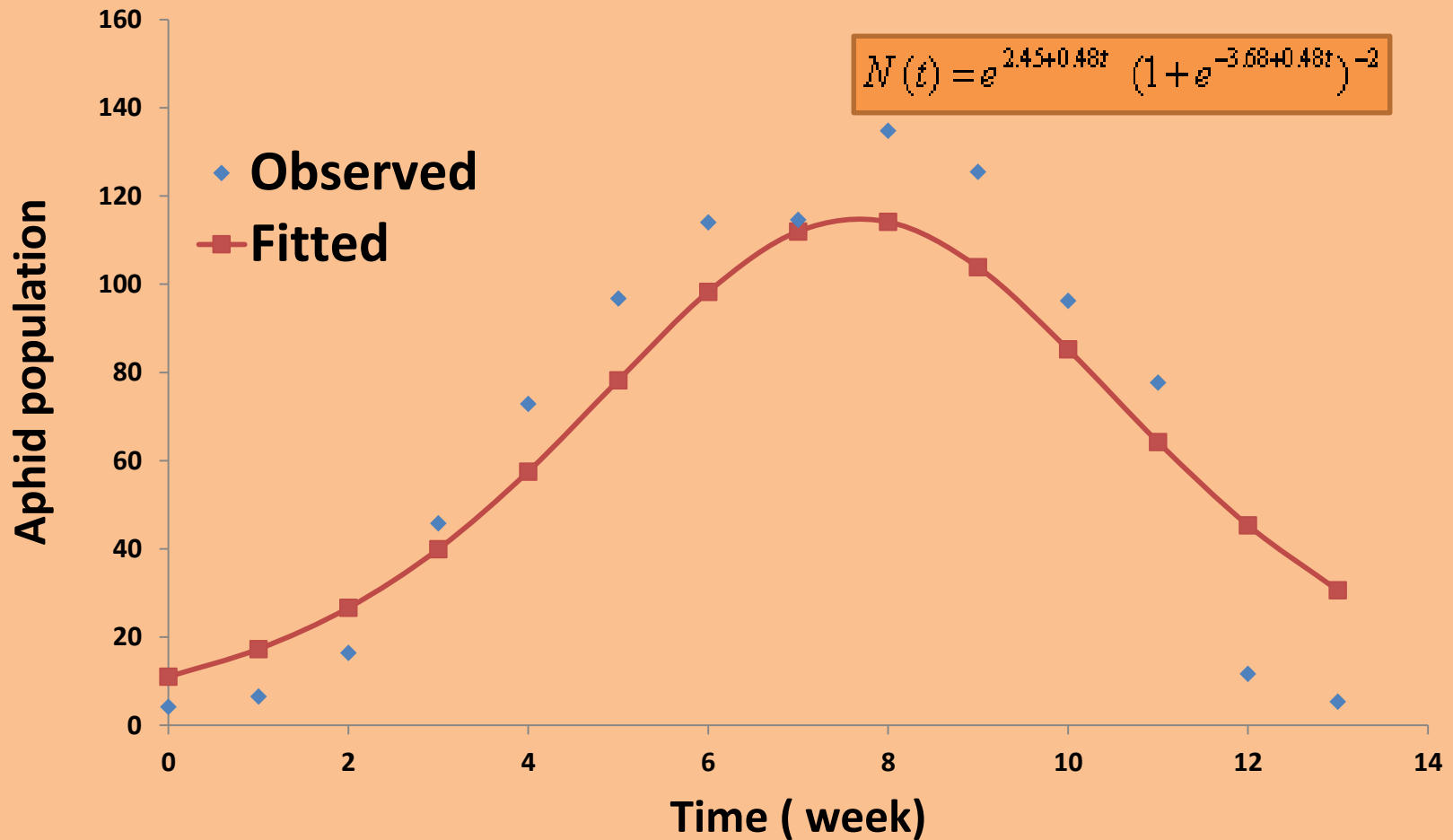
Fitting of Nonlinear Model

$$N(t) = e^{a+bt} (1 + e^{d+bt})^{-2} + \epsilon$$

Estimate the parameters	:	a, b, d
Iterative procedure	:	Levenberg-Marquardt
Goodness of fit statistic	:	Root mean square error
Examination of Residuals	:	Run test Shapiro-Wilk test

Software : SAS v 9.2 / Proc : NLIN

Fitted Nonlinear Aphid population model



Results

Statistic	Coefficeint
i. Parameters	
<i>a</i>	2.45(0.35)
<i>b</i>	0.48(0.05)
<i>d</i>	-3.68(0.40)
ii. Examination of Residuals	
Run test Z	1.2
Shapiro –Wilk test (W)	0.96
iii. Goodness of fit statistic	
Root mean square error	54.52

Summary

- **Polynomial functions are not nonlinear functions**
- Nonlinear function is one in which the parameters appear nonlinearly
- Biological Interpretation of the data can be understood by Nonlinear models
- Choosing the suitable model and initial guess values are important for obtaining the better/ suitable model for the data under consideration.
- Software : SAS , JMP , R , S -Plus



Thank you