SDS | Page No. Bayerian Competing Namer Breksha A. Patel sapid = 60004210126 Branch - Computer Engineering Experiment no. 4 Aim : To implement a model for a bioassay experiment: consider data from Gelman at al. (2003), where are observers a dose level (in 10 g g/ml) The number of asimals and the number of deaths for each of four groups. Multipaxameter Models :-These are statistical models that involve multiple parameters or unknown quantities. These models are commonly used in reaxions fields, including statistics, machine learning and Scientific research, to represent complex relationships between reasiables and to make predictions or inferences about the parameters. In Bayesian computing, multiparameter models are analyzed using Bayesian methods which incorporate prior beliefs about the parameters into the analysis and update their beliefs based on observed data. Bayesian inference provides comprehensive framework for handling uncertainty multiparameter models Logistic Regression: Logistic Regression is a statistical model used for binary elassification, where the goal is to predict a binary outcome leg, yes or no, successor failure etc. based on a set of independent variables

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In Bayesian logistic regression, prior beliefs about the model parameters, such as the xigression coeffecients, or incorporated into the analysis using probability distribution. These priors are updated ming Bayes rule band on observed data resulting in posterior distributions that reflect the updated beliefs about the parameters A Rioassay Experiment & Bioassay experiments are used to determine the dose-response relationship for a biological agents such as drug or toxin, Thue experiments typically involve exposing a group of organism to different doses of the agent of measuring the biological responses, such as survival or Leath Logistic Regression can be used in a Rayerian framework to analyze bioassay data by modeling probability of a positive response is a function of the dose The model paxameters can be intexpreted as the dose of at which circtain proportion of population is expected to experience positive response Conclusion 5 In this experiment, we implemented a Bayesian model for a

In this experiment, we implemented a Bayesian model for a binary experiment using R. The model fitted the date ruell, capturing the uncertainty in the parameter estimate. The sesues demonstrated the utility of Bayesian methods in interpreting bioassay data and the influence of prior information on the posterior distribution.

In conclusion, the experiment provided valuable insights into Bayesian computing and its application in bioassay analysis by modeling of analyzing a bioassay experiment suring Bayesian logistic regression



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Experiment No. 4

Aim:

Implement a model for A Bioassay Experiment: consider data from Gelman et al. (2003), where one observes a dose level (in log g/ml), the number of animals, and the number of deaths for each of four groups.

Code:

Importing Libraries

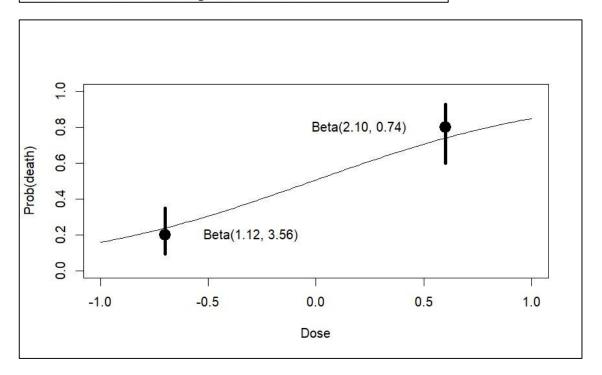
library('LearnBayes')

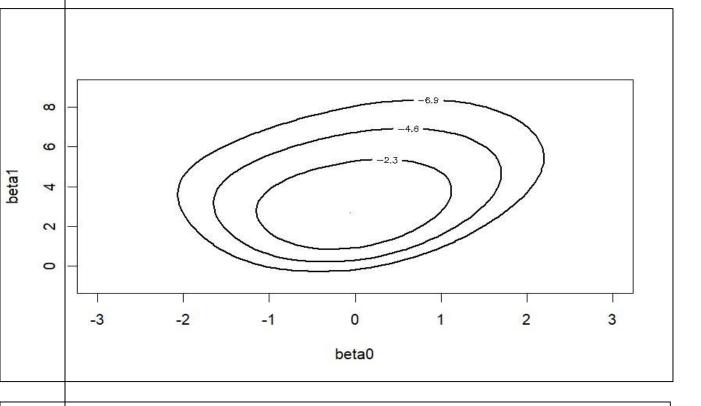
Bayesian fitting of a logistic model using data from a dose-response experiment.

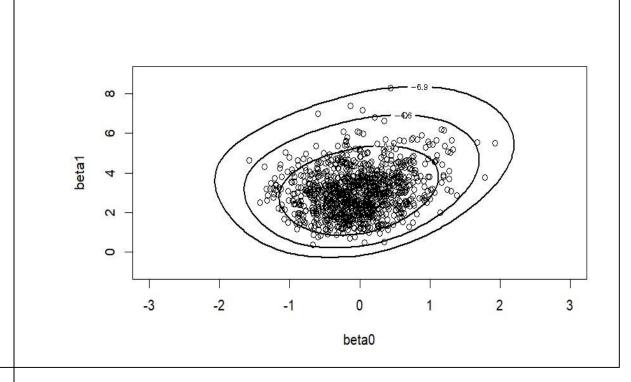
```
list(p=.9, x=.98))
prior <- rbind(c(-0.7, 4.68, 1.12),
c(0.6, 2.10, 0.74)) data.new <-
rbind(data, prior)
plot(c(-1,1), c(0, 1), type="n",
xlab="Dose", ylab="Prob(death)") lines(-
0.7 * c(1, 1), qbeta(c(.25, .75),
a1.b1[1], a1.b1[2]), lwd=4) lines(0.6 *
c(1, 1), qbeta(c(.25, .75),
                               a2.b2[1],
a2.b2[2]), lwd=4) points(c(-0.7, 0.6),
qbeta(.5, c(a1.b1[1],
                          a2.b2[1]),
c(a1.b1[2], a2.b2[2])),
                              pch=19,
cex=2) text(-0.3, .2, "Beta(1.12, 3.56)")
text(.2, .8, "Beta(2.10, 0.74)") response <-
rbind(a1.b1, a2.b2) x <- c(-0.7, 0.6) fit <-
glm(response \sim x, family = binomial)
curve(exp(fit\$coef[1] + fit\$coef[2] * x) /
(1 + \exp(\text{fit}\cdot\cos[1] + \text{fit}\cdot\cos[2] * x)),
add=T)
```

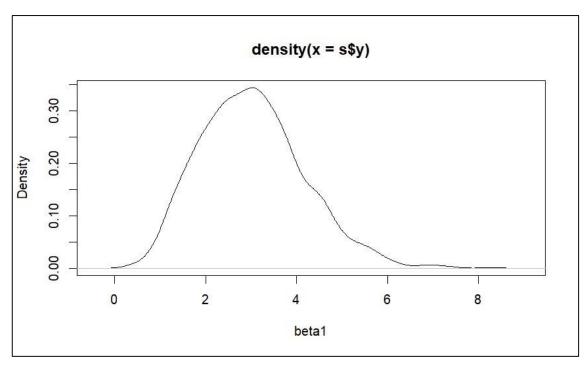
Output:

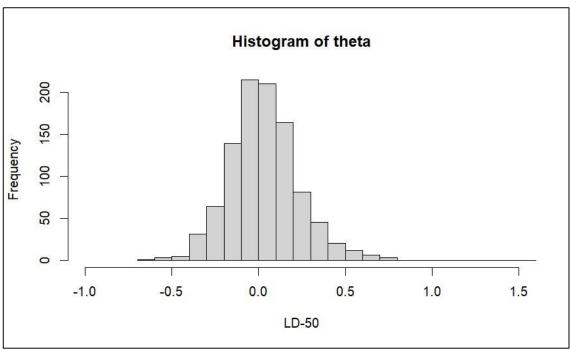
```
glm(formula = glmdata \sim x, family = binomial)
Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept)
              0.8466
                         1.0191
                                  0.831
                                           0.406
              7.7488
                         4.8728
                                  1.590
                                           0.112
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 15.791412 on 3 degrees of freedom
Residual deviance: 0.054742 on 2 degrees of freedom
AIC: 7.9648
Number of Fisher Scoring iterations: 7
```











2.5% 97.5% -0.3251418 0.4707840