

The Analysis of Biological Data - Whitlock and Schluter

Solutions to practice problems

Chapter 1

1. **(a)** Ordinal **(b)** Nominal **(c)** Nominal **(d)** Ordinal
2. **(a)** Observational (cats were observed to fall a given number of stories—the researchers did not assign the number of stories fallen to the cats). **(b)** The number of floors fallen. **(c)** The number of injuries per cat.
3. **(a)** Discrete **(b)** Continuous **(c)** Discrete **(d)** Continuous
4. **(a)** Collectors do not usually sample randomly, but prefer rare and unusual specimens. Therefore, the moths are probably not a random sample. **(b)** A sample of convenience. **(c)** Bias (rare color types are likely to be over-represented in the sample compared with the population).
5. Accuracy.
6. **(a)** Categorical **(b)** No random sampling procedure was followed so the survey is not random; the respondents were volunteers. **(c)** Volunteer bias (those most interested in the issue were more likely to respond).
7. **(a)** US army personnel stationed in Iraq. **(b)** Yes. The number of subjects interviewed was a random sample of only 100 from the population, therefore the responses of the particular individuals who happened to be sampled will differ from the population of interest by chance. **(c)** The advantage of random sampling is that it minimizes bias and allows the precision of estimates of stress levels to be measured.
8. **(a)** The population being estimated is all the small mammals of Kruger National Park. **(b)** No, the sample is not likely to be random. In a random sample, every individual has the same chance of being selected. But some small mammals might be easier to trap than others (for example, trapping only at night might miss all the mammals active only in the day time). In a random sample individuals are selected independently. Multiple animals caught in the same trap might not be independent if they are related or live near one another (this is harder to judge). **(c)** The number of species in the sample might underestimate the number in the Park if sampling was not random (e.g., if daytime mammals were missed), or if rare species happened to avoid capture. In this case m would be a biased estimate of M .
9. Chicks were not sampled randomly. The multiple chicks from the same nest are not independent. Chicks from the same nest are likely to be more similar to one another in their measurements than chicks chosen randomly from the population. Ignoring this problem will lead to erroneous measurements of the precision of survival estimates, sampled randomly from the population.

Chapter 2

1. **(a)** 25% **(b)** 33% **(c)** 33%

Analysis Of Biological Data Assignment Problem Answers

Didier Musso



Analysis Of Biological Data Assignment Problem Answers:

The Analysis of Biological Data Michael Whitlock, Dolph Schluter, 2009 This text emphasizes intuitive understanding rather than an over reliance on formulas The focus is on data and graphical displays rather than the mathematical foundations of statistics and students do not need knowledge of mathematics beyond simple algebra *Encyclopedia of Bioinformatics and Computational Biology*, 2018-08-21 Encyclopedia of Bioinformatics and Computational Biology ABC of Bioinformatics Three Volume Set combines elements of computer science information technology mathematics statistics and biotechnology providing the methodology and in silico solutions to mine biological data and processes The book covers Theory Topics and Applications with a special focus on Integrative omics and Systems Biology The theoretical methodological underpinnings of BCB including phylogeny are covered as are more current areas of focus such as translational bioinformatics cheminformatics and environmental informatics Finally Applications provide guidance for commonly asked questions This major reference work spans basic and cutting edge methodologies authored by leaders in the field providing an invaluable resource for students scientists professionals in research institutes and a broad swath of researchers in biotechnology and the biomedical and pharmaceutical industries Brings together information from computer science information technology mathematics statistics and biotechnology Written and reviewed by leading experts in the field providing a unique and authoritative resource Focuses on the main theoretical and methodological concepts before expanding on specific topics and applications Includes interactive images multimedia tools and crosslinking to further resources and databases **Building Bioinformatics Solutions 2nd Edition** Conrad Bessant, Darren Oakley, Ian Shadforth, 2014 This book introduces the reader to all the key concepts and technologies needed to begin developing their own bioinformatics tools The new edition includes more bioinformatics specific content and a new chapter on good software engineering practices to help people working in teams *Brain-Inspired Computing: From Neuroscience to Neuromorphic Electronics driving new forms of Artificial Intelligence* Jonathan Mapelli, Giacomo Indiveri, Angelo Arleo, 2022-03-08 *Bulletin of the Electrical Engineering and Electronic Engineering Departments*, 1977 Pacific Symposium on Biocomputing 2008 Russ Altman, 2008 The Pacific Symposium on Biocomputing PSB 2008 is an international multidisciplinary conference for the presentation and discussion of current research in the theory and application of computational methods in problems of biological significance Presentations are rigorously peer reviewed and are published in an archival proceedings volume PSB 2008 will be held on January 4th 2008 at the Fairmont Orchid Big Island of Hawaii Tutorials will be offered prior to the start of the conference PSB 2008 will bring together top researchers from the US the Asian Pacific nations and around the world to exchange research results and address open issues in all aspects of computational biology It is a forum for the presentation of work in databases algorithms interfaces visualization modeling and other computational methods as applied to biological problems with emphasis on applications in data rich areas of

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conceptions of race **Macroinvertebrate Data Interpretation Guidance Manual** Richard Allan Lillie, Stanley W. Szczytko, Michael A. Miller (Ecologist), 2003 **Journal of Mammalogy**, 1929 **Report** United States. Bureau of Fisheries, 1914

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