

and are forging ahead!

SSA Biostatistics & Bioinformatics Newsletter Q2 2020

The Biostatistics and Bioinformatics section committee hope you, your family, and your friends are keeping well, in particular those who are in Melbourne Lockdown 2.0, selfisolating or self-quarantining. Many of you have probably adapted to the new "normal"

This newsletter contains updates on our mentoring program pilot and statistical consultancy workshop. Also, we have included some resources that may be of interest for those whose research may have been impacted as a result of this

Welcome to the our second newsletter of 2020!

pandemic. Finally, to help you "stay at home", check-out our selection of favourite statistical podcasts. If you have any ideas to support you better during these challenging times, please let us know, we are there for you!

The SSA Biostatistics and Bioinformatics section committee

SSA Biostatistics & Bioinformatics section mentoring update



currently collating responses from the surveys and learning from the experiences of our mentors and mentees to build on this program for 2021 so watch this space! In the meantime, if you want any information about the mentoring program or have any suggestions on how you would like the SSA mentoring program to look or wish to help with future mentoring programs, please feel free to contact our mentoring coordinator, Karen Lamb, at klamb@unimelb.edu.au. In addition to coordinating the mentoring program, one of our committee members, Karen Lamb, took part in the SSA Victorian Branch mentoring evening on 23rd June. This twohour event, conducted online via Zoom, consisted of short career presentations from a diverse group of senior mentors, as well as one-to-one discussion sessions between an early-career statistician and more experienced statistician to allow junior members to learn from more senior members. These conversations were conducted in a "speed

dating" format to enable mentees to meet multiple mentors over the course of the evening. Karen really enjoyed the opportunity to meet with early career statisticians to discuss their career challenges, aspirations and to share her own experiences. **Resources related to COVID-19** You may be engaged and involved in the consequences of the pandemic on clinical trials and observational studies either via ongoing or new projects. Our Section is actively collating resources on the methodological aspects that you may have to tackle, for

• When discussing the impact of COVID-19 on ongoing clinical trials, the European

example:

adaptive designs,

Medicines Agency has finalised a point to consider document and the Food and Drug Administration (FDA) has finalised a guidance document. Furthermore, a publication on statistical issues and recommendations for and challenges in assessing the impact on integrity and interpretability of clinical trials. • When designing trials of interventions for COVID-19, a publication on efficient

- Finally, the SSA has compiled some general resources on COVID-19. Meet our mailing list

This space is rapidly evolving. We welcome input from our member to help us collate a list

methodology to deal with the impact of the pandemic on both ongoing and new trials and

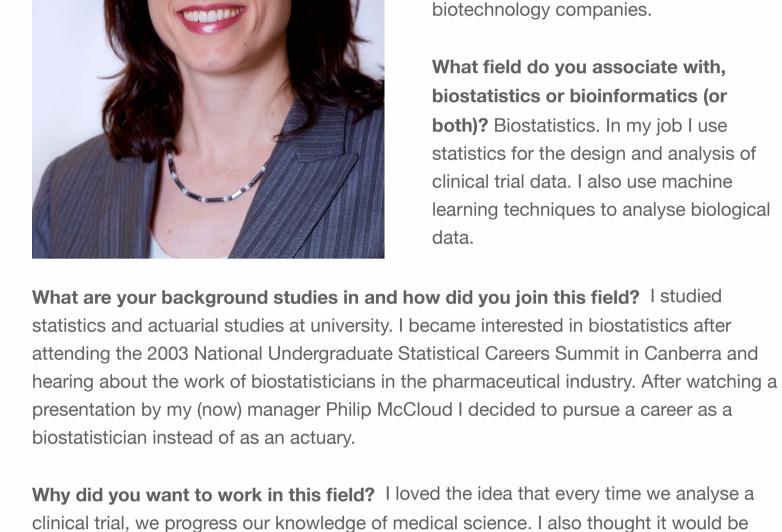
studies. Please contact Sabine Braat or Karen Lamb if there is any resource you would

of documents and publications on the topic of biostatistics and bioinformatics

like to share with other members of the Biostatistics and Bioinformatics Section.

Where do you currently work? I work as a statistician at McCloud Consulting Group, a contract research organisation

biotechnology companies.



(or bad) to the stakeholders.

profiles. Its a great way to get some

interested, please contact Nicole De La

exposure to our >200 readers. If

biostatistics or bioinformatics (or both)? Biostatistics. In my job I use statistics for the design and analysis of clinical trial data. I also use machine learning techniques to analyse biological data.

What field do you associate with,

providing biometric services to medical

researchers, pharmaceutical and

Name: Sandra Louw

exciting to be the first person to see the results of a clinical trial and break the good news What is the most exciting concept/idea upcoming in your work or field? It's exciting to see the combination of statistics and machine learning techniques being used to solve medical problems, for example for prediction in wide datasets with lots of patient features.

Want to be featured in the next member profile? We are always in need of more member

If you didn't pursue biostatistics and/or bioinformatics, what would have been your

alternate career? If I wasn't a statistician I would like to be a fiction writer. I can imagine

spending my days reading and writing, discussing books at writers' conferences, and

attending the movie sets of my books as they get turned into movies!

Mata. **SSA** webinars and events

How to make a learning healthcare system work: from linking data, machine



come from and how it will be used by clinicians in managing care. She provided insight into how routine care data can be used to improve care, how data sources can be linked to understand health and how challenging it can be working with data not designed for research purposes. The seminar ended with breakout sessions to enable attendees to chat to one another and Katrien to answer questions in small groups. Any SSA members who missed the webinar can access it online here.

through a biostatistician's lens by Dr Elasma Milanzi

The R Podcast This podcast provides advice, information and tips to tap into the power of R. There are 33 episodes you can listen to here that cover a range of topics including R markdown, Shiny R, reproducible analyses, ggplot and more. Eric Nantz is the

creator of this podcast as well as a youtube channel and

"Statistics need Stories to give them meaning. Stories need

Statistics to give them credibility." What a beautiful blend of

Do you have a favourite statistical podcast or blog?

so, please contact Karen Lamb with the details.

website resources to shape you into an R expert.

microbiome analysis explaining the processing of raw reads, alpha and beta diversity, exploration and visualisation of microbiome data, and statistical methods for analysis of microbiome data in R. The webinar was attended by both members and non-members of the SSA. The recording will become available soon, please check out the SSA website.

Two webinars on bioinformatics topics will be hosted later this year, including Professor

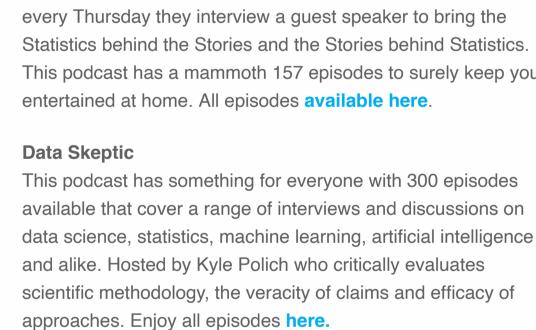
Jean Yang (University of Sydney) and Professor Melanie Bahlo (Walter and Eliza Institute

Podcasts

of Medical Research). All upcoming SSA webinars and events can be found here.

On 19th June, Dr Elasma Milanzi shared her journey as a biostatistician into the field of

Getting to know our "cell mates": a practical approach to microbiome analysis



Stats + Stories

concepts to bring life to statistics and its purpose. Hosted by John Bailer, Richard Campbell and Rosemary Pennington, every Thursday they interview a guest speaker to bring the Statistics behind the Stories and the Stories behind Statistics. This podcast has a mammoth 157 episodes to surely keep you

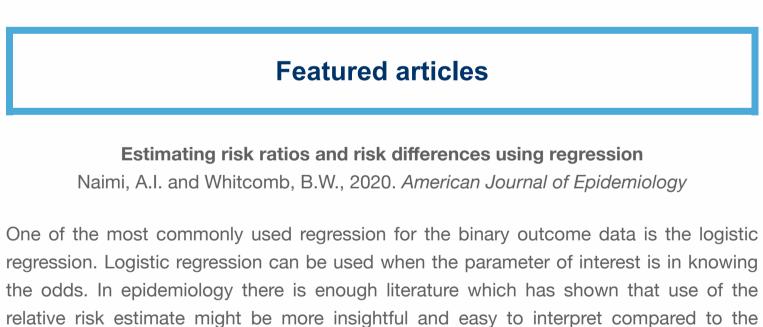
TIC-

Our previous newsletters have covered a range of

biostatistics and bioinformatics blogs and podcasts. In

this newsletter, we are highlighting a few podcasts that

might interest our mailing list readers.



estimate from the logistic regression. However, when it comes to the estimation of the

relative risk one may compute this using the log-binomial regression in any standard

statistical software. However, a common issue with the use of the log-binomial regression

is the convergence issue. Many researchers have shown how to use the log-linear

regression to estimate the relative risk as such, but, not much focus is given to the

estimation of the standard errors. In this paper by Naimi and Whitcomb (2020) they

describe how to estimate the relative risk using the standard glm method and also how to

estimate the standard errors. For example, the glm method in Stata can be specified as

"glm y x c, family(poisson) link(log) vce(robust)". The glm is the generalised linear model

command with the outcome variable (y), the exposure (x) and the set of confounders (c).

The command, family, indicates that one is using, Poisson, with the link function being,

We'd love to have your input to grow our lists. Perhaps you would be happy to provide a

brief overview of your favourite statistical podcast or blog to feature on our mailing list. If

log, and the variance covariance matrix being specified using the vce option (robust) where the standard errors are estimated using the Sandwich estimator. Recent publication that you'd like to share? We love learning about all things related to biostatistics and bioinformatics. If you have a recent publication (your own or other) that you'd like to have featured in our newsletter, please contact Nicole De La Mata with the details. **Statistical Consultancy Workshop**

We are going online with our workshop on statistical consultancy!

of our experts! This workshop is aimed at statisticians who provide statistical advice, analysis, and training to non-statisticians. This workshop will be of greatest interest to recent graduates and early career researchers who are keen to develop their skills in this area.

website as soon as possible.

Joint Statistical Meeting 2020

Virtual conference, 2nd-6th Aug 2020

The workshop is designed to provide participants with an understanding of some of the key

skills involved in biostatistical consultancy. This will cover a range of topics including how

consultancy, as well as an interactive panel session to allow participants to pick the brains

Preparations are ongoing and will be announced via the SSA weekly newsletter and

Other upcoming events

While there is generally a range of other events of interest for our biostatisticians and

bioinformaticians, the schedule of events have been substantially impacted by the COVID

pandemic and many events have been cancelled. Please always check out the

conference website to ensure you have the latest information on the conference. On the

other hand, this has lead to the creation of many virtual events that are often more

to run a consultancy service and how to reach and interact with different clients. The

workshop will feature presentations from statisticians experienced in biostatistical

affordable and accessible. **STATA** conference 2020 Virtual conference, 30th-31st Jul 2020

Virtual conference (previously Krakow, Poland) 23th-27th Aug 2020 **Royal Statistical Society International Conference** Virtual conference, Week beginning 7th Sep 2020

41st Annual Conference of International Society for Clinical Biostatistics

If you would like to advertise any upcoming biostatistics or bioinformatics events to our mailing list, please contact one of our co-Chairs, Sabine Braat or Jaimi Greenslade **Closing comments** Our Section Committee is interested in biostatistics and bioinformatics and we

bioinformatics. We are interested in hearing from our members, especially regarding ideas for future events, presentations, webinars or workshops held by the SSA Biostatistics

are committed to assisting you with your

career in biostatistics or/and

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Please contact us via our section co-Chairs, either Sabine Braat or Jaimi Greenslade regarding these or any other related queries. If your colleagues would like to receive this newsletter, they can sign up via mailchimp

below. We actively promote upcoming events, job opportunities and relevant material. Our online community is growing with over 697 followers on Facebook and 446 on twitter. If you have any material you would like us to promote on our social media outlets or share with the fellow members, please let Sabine Braat know.

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and you wish to be removed, please contact us at j.greenslade@uq.edu.au or s.braat@unimelb.edu.au.

& Bioinformatics section. We are



