DJORDJE DJORDJEVIC

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Born: 5th April 1988

Objective: Pursue a career in the biological sciences where I can apply the power of computers to solving complex biological problems. Science is my passion and the power of computers to change the world is undeniable. I hope to contribute specifically to the fields of genotype / phenotype correlation and epigenetic control modelling to facilitate the future of synthetic and regenerative biology.

EDUCATION

Doctor of Philosophy (PhD) - 2017

Computational methods for analysing the regulation of genetic systems across species Faculty of Medicine - University of New South Wales

Bachelor of Engineering (Bioinformatics) (Honours) - 2013

Phenotype prediction using data mining and machine learning techniques

Faculty of Computer Science and Engineering - University of New South Wales

PUBLICATIONS

Citations: 46 h-index: 4

Light-focusing human micro-lenses generated from pluripotent stem cells model lens development and drug-induced cataract in vitro

Patricia Murphy, Md Humayun Kabir, Tarini Srivastava, Michele E Mason, Chitra U Dewi, Seakcheng Lim, Andrian Yang, Djordje Djordjevic, Murray C Killingsworth, Joshua WK Ho, David G Harman, Michael D O'Connor

Development 145 (1), dev155838

Identification of satellite cells from anole lizard muscle and demonstration of increased musculoskeletal potential

Joanna Palade*, Djordje Djordjevic*, Elizabeth D. Hutchins, Rajani M. George, John A. Cornelius, Alan Rawls, Joshua W.K. Ho, Kenro Kusumi, and Jeanne Wilson-Rawls (* indicates co first authors) **Developmental biology 433 (2), 344-356**

iSyTE 2.0: a database for expression-based gene discovery in the eye

Atul Kakrana, Andrian Yang, Deepti Anand, Djordje Djordjevic, Deepti Ramachandruni, Abhyudai Singh, Hongzhan Huang, Joshua W K Ho, Salil A Lachke

Nucleic acids research 46 (D1), D875-D885

XGSA: a statistical method for cross-species gene set analysis

Djordje Djordjevic, Kenro Kusumi, Joshua W. K. Ho

Bioinformatics (2016), 32 (17): i620-i628

hiHMM: Bayesian non-parametric joint inference of chromatin state maps

Kyung-Ah Sohn, Joshua W. K. Ho, Djordje Djordjevic, Hyun-hwan Jeong, Peter J. Park and Ju Han Kim

Bioinformatics (2015), 27/02/15, pii: btv117

Decoding the complex genetic causes of heart diseases using systems biology

Djordje Djordjevic, Vinita Deshpande, Tomasz Szczesnik, Andrian Yang, David T. Humphreys, Eleni Giannoulatou, Joshua W. K. Ho

Biophysical Reviews (2015), 10/12/14, 7:141159

How difficult is inference of mammalian causal gene regulatory networks?

Djordje Djordjevic, Andrian Yang, Armella Zadoorian, Kevin Rungrugeecharoen, Joshua W. K. Ho PLOS ONE (2014), 4/11/14, 9(11):e111661

Djordje Djordjevic Page 2

BOOK CHAPTERS

Epigenomic analysis of chromatin organization and DNA methylation

Xin Wang, Helen McCormick, Djordje Djordjevic, Eleni Giannoulatou, Catherine M Suter, Joshua WK Ho, in Computational Biology Bioinformatics Gene Regulation (2016) (Ed. Wong KC). CRC Press. ISBN 9781498724975 - CAT K25752

SUBMITTED MANUSCRIPTS

GEOracle: Mining perturbation experiments using free text metadata in Gene Expression **Omnibus**

Djordje Djordjevic, Yun Xin Chen, Shu Lun Shannon Kwan, Raymond WK Ling, Gordon Qian, Chelsea YY Woo, Samuel J Ellis, Joshua WK Ho

bioRxiv, 150896

CardiacProfileR: An R package for extraction and visualisation of heart rate profiles from wearable fitness trackers

Djordje Djordjevic, Beni K Cawood, Sabrina K Rispin, Leo HH Yim, Christopher S Hayward, Joshua WK Ho

bioRxiv, 324004

C3: An R package for cross-species compendium-based cell-type identification

Md Humayun Kabir, Djordje Djordjevic, Michael D O'Connor, Joshua Ho bioRxiv, 267880

CONFERENCE PRESENTATIONS

XGSA: a statistical method for cross-species gene set analysis

Djordje Djordjevic, Kenro Kusumi, Joshua W. K. Ho

European Conference on Computational Biology 2016 - Selected oral presentation and poster International Conference on Systems Biology 2016 - Poster

Australian Bioinformatics and Computational Biology Society Conference 2015 - Selected oral presentation and poster

GEOracle: Harnessing GEO to discover mammalian causal gene regulatory networks Djordje Djordjevic, Yun Xin Chen, Raymond Ling, Gordon Qian, Chelsea Woo, Joshua W. K. Ho Sydney Bioinformatics Research Symposium 2016 - Poster and short oral presentation

Harnessing a large collection of gene perturbation data to discover mammalian causal gene regulatory

Djordje Djordjevic, Andrian Yang, Shu Lun Shannon Kwan, Joshua W. K. Ho

Intelligent Systems for Molecular Biology / European Conference on Computational Biology 2015 -Poster

COMBINE Student Symposium 2015 - Poster

Comprehensive analysis of chromatin landscape in filamentous fungus Djordje Djordjevic, Vinita Deshpande, Kaeling Tan, Koon Ho Wong, Joshua W. K. Ho Australian Pathogen Bioinformatics Symposium 2014 - Selected oral presentation

Harnessing sparse gene perturbation data to discover causal gene regulatory networks Djordje Djordjevic, Andrian Yang, AmirHossein Kamali, Joshua W. K. Ho International Conference on Systems Biology 2014 - Poster

Constructing causal gene regulatory networks to investigate organ development and disease Djordje Djordjevic, Andrian Yang, Vinita Deshpande, Tomasz Szczesnik, Joshua W. K. Ho St Vincent's Research Symposium 2014 - Poster Australian Bioinformatics Conference 2014 - Poster

A systems biology framework for prioritisation of pathogenic genes Djordje Djordjevic, Joshua W. K. Ho

Sydney Bioinformatics Research Symposium 2014 - Poster and short oral presentation

Djordje Djordjevic Page 3

iSyte: a web-resource of curated lens gene regulatory networks reveals the interplay of diverse pathways in ocular development and disease

Deepti Anand, Djordje Djordjevic, Sylvie Smith, Joshua W K. Ho, Salil A. Lachke

Annual Meeting of the Association for Research in Vision and Ophthalmology (ARVO) 2014 - Poster

Transcriptional profiling and integrative analysis of cardiomyocyte stretch response

Djordje Djordjevic, Vesna Nikolova-Krstevski, Mirana Ramialison, Ashley Waardenberg, Joshua W. K. Ho, Diane Fatkin

St Vincent's Research Symposium 2013 - Poster

Chromatin state analysis in model organisms

Djordje Djordjevic, Chris Wong, Joshua W. K. Ho

Sydney Bioinformatics Research Symposium 2013 - Poster and short oral presentation

AWARDS

International Society for Computational Biology - Travel fellowship to attend ECCB 2016 - \leq 900 Victor Chang Cardiac Research Institute - Best student talk Paul Korner seminar series 2015 - \$500 Australian Pathogen Bioinformatics Conference - Travel fellowship 2014 - \$300

Australian Bioinformatics Conference - Travel bursary 2014 - \$400

Australian Bioinformatics Winter School - Travel bursary 2013 - \$250

SOFTWARE AND WEBSITES

GEOracle: Mining the GEO database for perturbation data sets - http://georacle.victorchang.edu.au/XGSA: a statistical method for cross-species gene set analysis - https://github.com/VCCRI/XGSA CardiacCode - http://cardiaccode.victorchang.edu.au/

iSyTE 2.0 - http://research.bioinformatics.udel.edu/iSyTE

 $\label{lem:cardiacProfiler:} Cardiac Profiler: \ Extraction \ and \ visualisation \ of \ heart \ rate \ profiles \ from \ wearable \ fitness \ trackers - https://github.com/VCCRI/CardiacProfileR$

C3: Cross-species compendium-based cell-type identification - https://github.com/VCCRI/C3

ONGOING COLLABORATIONS

Characterisation and prioritisation of pathogenic non-coding variants in the atrial fibrillation GWAS locus 4q25 - with Prof. Diane Fatkin and Prof. Richard Harvey, VCCRI / UNSW

Transcriptional and epigenomic characterisation of cardiac myocyte maturation in mice - with Prof Bob Graham, $VCCRI\ /\ UNSW$

SUPERVISORY ROLES

I have supervised three honours students and 9 summer scholarship students over the course of my research. Two of these students became co-authors on my fiirst publication and 8 others are co-authors on manuscripts currently under preparation.

SKILLS

I am a highly motivated learner and am always eager to learn new skills.

Bioinformatics / Data Analysis

I have a growing skillset in the field of Bioinformatics, developed over the course of my research at the Victor Chang Cardiac Research Institute. I have become proficient in comprehensive mRNA microarray and RNA-Seq analysis, cross-species chromatin landscape and gene set comparisons, image analysis, disease gene prioritisation and gene regulatory network inference and analysis, amongst many other projects. I also have limited experience working with financial data.

I am adept at combining concepts from statistics with machine learning approaches and novel ideas to gain insights from large and complex data sets and solve real world data problems.

Computer Proficiency

I am a native R programmer, comfortable in Perl, Python, Shell, Latex, HTML, CSS, Javascript, and have limited experience with Java, OpenGL, C, PHP, Prolog and Basic. I have a working

Djordje Djordjevic Page 4

knowledge of relational-databases and biological knowledge bases. I am comfortable with Microsoft Office, Open Office and the Adobe software suites, and working in Windows, Mac OSX and Linux. I have well developed internet research and navigation skills.

Communication

I am experienced in applying effective communication techniques to maximise understanding, solve problems and resolve conflicts. I've gained these skills throughout my life as a Parkour instructor and community leader, a corporate workshop facilitator, a retail professional, a soccer referee and most recently a business owner and manager. Additionally my language and communication skills have developed through a passion for reading, growing up in a multi-lingual household and travelling through Asia and Europe extensively.

Throughout my academic career numerous presentations and projects have enabled me to further refine these skills. Specifically, I have authored scientific manuscripts, presented my research at national and international conferences as both oral and poster presentations, supervised 11 undergraduate research students and trained team members in new skills and concepts. I have delivered several award winning presentations during my PhD.

Laboratory Experience

I have over 200 hours of chemistry and biology laboratory practical work experience during which I learned many standard laboratory techniques. As a result I am comfortable handling potential biological hazards and working with common laboratory equipment.

Photography and video production

Since I was young I have enjoyed photography and this has developed into a passion during my lifetime. I have been involved in many photography and videography projects, paid and unpaid, including being the de-facto photographer for events at the Victor Chang Cardiac Research Institute, training the media staff to take photos, creating videos for the use of the institute, and many non-science related videography projects which have amassed millions of views on YouTube.

EMPLOYMENT

Research Assistant / PhD Student / Postdoctoral Scientist, Victor Chang Cardiac Research Institute, 2013 - 2018

My research so far has been with the Victor Chang Cardiac Research Institute, a partner of the University of New South Wales.

Facilitator, Developer, Decoded, 2015 - Present

I am a casual workshop facilitator and content developer for Decoded, specialising in delivering data analytics and coding foundational courses to commercial clients.

Managing Director, Jump Squad Pty Ltd, 2011 - Present

My company Jump Squad is Australia's leading provider of Parkour and Freerunning services, including performances, workshops, facility designs and training programmes.

Various, 2003 - 2011

PROFESSIONAL AFFILIATIONS

International Society for Computational Biology (ISCB) (Member) Australian Bioinformatics and Computational Biology Society (Member) COMBINE / ISCB Students Australia (Former committee Member)

Non-science related Australian Parkour Association (Founding committee member) Parkour New South Wales (Founding member) Djordje Djordjevic Page 5

OTHER QUALIFICATIONS

Construction Induction (White Card), 2010 Working with Children Check, 2010 Responsible Service of Alcohol, 2010 Responsible Conduct of Gambling, 2010 Senior First Aid (Level 2), 2009

REFERENCES

A/Prof. Joshua Ho, Head Bioinformatics and Systems Medicine Laboratory, VCCRI +61 2 9295 8645, j.ho@victorchang.edu.au

Prof. Diane Fatkin, Head Inherited Heart Diseases Laboratory, VCCRI +61 2 9295 8618, d.fatkin@victorchang.edu.au

Prof. Kenro Kusumi, Associate Dean of Research, College of Liberal Arts and Sciences, ASU +1480~727~0530, Kenro.Kusumi@asu.edu