

Curriculum Vitae - Bioinformatics

Dr. Jeremy D. Parsons

Personal Details

Name: Jeremy David Parsons **Home Address:** 5 Roughdown Villas Road,
Phone: +44 7913 024857 Hemel Hempstead,
+63 945 1940 708 Herts.
Personal email: jp @ jeremyparsons .com HP3 0AX
Consulting website: http://www.littlest.co.uk/ England
Marital Status: Single
Nationality: British

Academic Qualifications

- University of Cambridge, Department of Medicine Ph.D.
- University of Kent at Canterbury, Computer Science MSc.
- Sheffield University Special Honours Microbiology BSc. class 2:1

Technical Skills

- **Languages:** Java, 'C', Perl, Python, PostScript, Bash UNIX shell script, occam2, Motorola 68000 assembler, C++, NewtonScript, R
- **Libraries and Patterns:** Parallel processing and multi-threading, Swing, regular expressions, XML, JMF, image processing, Standard design patterns J2SE, RMI, CORBA
- **Web Programming:** Servlets, JSP, Spring, JSTL, PHP, Apache and Tomcat maintenance
- **Databases:** JDBC Programming, PostgreSQL, MySQL, and Oracle.
- **System Administration:** Unix (15+ years with Linux and Solaris), Macs (infrequently over 10 years), Firewalls, LDAP
- **Bioinformatics:** Genome & RNA assemblers & editors, Next Gen. mappers, functional, variant & network analyses

Career

2017 - 2019 Solar Power Developer and investor, Australia and Asia. www.filsolar.com

- Promotion and financing of cheap distributed renewable energy.

2015 - 2016 Garvan Institute of Medical Research, Immunology Dept., Australia

- Differential Expression Analyses in mice. UNIX scripting and RNAseq pipelines.

- Big data processing, R presentation graphics and signal compression for heatmaps.
- Endogenous Retrovirus (ERV) expression
- Transcriptional control of Immune Response (IR) genes by ERV TFBS motifs.

2011 - University of Queensland Bioinformatics with QFAB and Phil Hugenholtz, Brisbane,
2014 Australia

- SQL and Java Web databases, Bioinformatics support and analysis, pipeline scripting and Java Programming
- Metagenomic analyses of mouse and sheep digestion
- Bacterial genome assembly including difficult 16S repeat regions
- Initial Macadamia and Koala genome assemblies and analyses.
- Transcriptomics and Differential Expression analyses in Sugar Cane and bacteria
- BRAEMBL project and many others

2010 - Bioinformatics research & genome comparisons: Byrappa Venkatesh, Singapore
2011

- DNA Assembly and analysis, pipeline scripting and Java Programming
- Studying Vertebrate evolution using Elephant Shark as a reference genome
- NGS Roche 454 Linker and adapter removal via sensitive parallel alignments

2008 - Cheminformatics at European Patent Office (EPO Den Haag) with Stephane Nauche
2009

- Java Programming and Natural Language Parsing
- Discovering new chemicals in EPO patent images and text
- Parsing scientific literature using OSCAR 3
- XML handling, databases and Java servlets

2007 ACI Worldwide - Java Programming

- Introduction to electronic payments
- User interface design in Java

2006 - Reuters - Java JRisk Programmer
2007

- Financial software training
- Java J2EE bugfixing

2006 - Building Research Establishment - Analyst and Programmer
Contract

- Image processing applications for visual analysis of timber
- Multi-threaded Java image analysis pipeline and GUI

2003 - Paradigm Therapeutics -Senior Bioinformatician (analyst/programmer)
2005

- Phenotype database development.
- Flexible data collection, querying and MIS report generation
- sostgreSQL DB with Java Beans, JSP, JSTL, and servlets

- Systems support including networking and centralized Linux servers for Macs and Linux desktops

2002 - Bioinformatics consulting at littlest.co.uk
2003

- General bioinformatics support
- Development of parallel text indexing and publishing tools - Java, AWT, Swing, Regular Expressions, PVM, C

2001 - Bioinformatics at Lion Bioscience Ltd, Cambridge, UK. with Thure Etzhold
2002

- Analyst and programmer for client/server systems including a graphical SRS PRISMA viewer - Java, Swing (large canvas component), XML specifications, Icarus
- Bioinformatics support and documentation

1999 - Bioinformatics at Cereon Genomics, Boston, MA, USA. with Stan Letovsky & David
2000 Bush

- Positional cloning and analysis of plant disease resistance genes - Databases, Perl, UNIX scripting, regular expressions
- Analysis database support, user training - Java, servlets

1997- Post-doctoral research at European Bioinformatics Institute with Tom Flores, Geoff
1999 Barton and Patricia Rodriguez-Tomé

- Sequence chromatogram (DNA trace file) browsing client/server systems
- Analyst and programmer: Java parsing, communication and graphics on scrolling canvas
- Linking Biological Databases using CORBA leading a five nation European consortium.
- Architect and programmer for a parallel distributed incremental DNA sequence alignment system that worked across 40 processors - 'C' and PVM
- Java client applications combining multiple data sources into graphical data summaries
- C++ server and CORBA wrapper for EST alignments and CORBA supercluster application

1996- Bioinformatics research and support for Zeneca Pharmaceuticals with Rakesh Anand
1997 and David Pioli

- Positional cloning support for arthritis research
- Bioinformatics programming including an email and web-based sequence alert system - Perl
- System support - Solaris and Linux
- Graphical client interfaces and general bioinformatics - Java

1993 - Post-doctoral research at Bob Waterston's Genome Sequencing Center, Washington
1995 University St. Louis. with LaDeana Hillier

- Development of sequence analysis and visualisation tools including ESTCluster and Miropeats(PrintRepeats)
- C, Perl and PostScript programming
- Managing and training new programmers for a joint project with David States' group at IBC
- System administration - Solaris and MacOS
- Studies of variables in DNA sequence production efficiency - Postgres DBMS and SQL querying
- Image analysis of sequence gels and lane tracking algorithms, EST clustering, DNA sequence assembly - 'C', and X11 graphics

- 1989-1993 Ph.D. in University of Cambridge Dept. of Medicine, MGU lead by Sydney Brenner (Nobel Laureate)
- Ph.D. Thesis title is "Computer Analysis of Molecular Sequences" supervised by Martin Bishop
 - Clustering partially sequenced cDNA sequences (ESTs) - 'C'
 - Visualisation of dipeptide frequencies. - UNIX shell and PostScript
- 1989 University of Cambridge Computing Service Research Associate: Colleges Project Officer.
- 1987-88 University of Kent at Canterbury - Two years of computer science including a parallel processing research project in communications entitled "Implementing the X-25 Data Link Protocol in Occam 2"
- 1984-1987 University of Sheffield studying microbiology, chemistry and basic computing
- 1984 Self-employed: writing graphical computer games in 6809 assembly language: one sold over 4000 copies.

Publications: Scopus: 7402708504 , ORCID: 0000-0003-1716-731X

- 2016 Paungfoo-Lonhienne, Chanyarat; Lonhienne, Thierry G. A.; Yeoh, Yun Kit; Donose, Bogdan C.; Webb, Richard I.; Parsons, Jeremy; Liao, Webber; Sagulenko, Evgeny; Lakshmanan, Prakash; Hugenholtz, Philip; Schmidt, Susanne; Ragan, Mark A., Crosstalk between sugarcane and a plant-growth promoting Burkholderia species, *Nature Scientific Reports* **6:37389** DOI: 10.1038/srep37389
- 2016 Kate L. Ormerod, Jeremy D. Parsons, Philip Hugenholtz et al., Genomic characterization of the uncultured Bacteroidales family S24-7 inhabiting the guts of homeothermic animals *Microbiome* **4:36** DOI: 10.1186/s40168-016-0181-2
- 2013 Mohammad Tawhidul Islam, Jeremy Parsons et al., Unlocking the Puzzling Biology of the Black Périgord Truffle Tuber melanosporum, *J. Proteome Res.*, **12**, pp 5349-5356., DOI: 10.1021/pr400650c
- 2001 Perret X., Parsons J., Viprey V., Reichwald K., Broughton W.J.; IRepeat sequences of genomes of Rhizobium and Sinorhizobium meliloti: a comparative analysis.; *Canadian Journal of Microbiology* (2001), **47(6)**, 548-558, DOI: 10.1139/cjm-47-6-548

- 2000 Parsons, J.D. and Rodriguez-Tomé P., JESAM: CORBA software components to create and publish EST alignments and clusters, *Bioinformatics*, **16**, 313-325.
- 1999 Parsons, Jeremy D., Buehler, Eugen, and Hillier, LaDeana, DNA Sequence Chromatogram Browsing Using JAVA and CORBA., *Genome Res.*, **9**, 277-281.
- 1996 Hillier, L, Parsons, J.D., et al., Generation and Analysis of 280,000 Human Expressed Sequence Tags, *Genome Research*, **6**, 807-828.
- 1996 Cooper, Matthew L., Maffitt, David R., Parsons, Jeremy.D., Hillier, LaDeana, and States, David J., Lane Tracking Software for Four-color Fluorescence-based Electrophoretic Gel Images, *Genome Res*, **6**, 1110-1117.
- 1995 Berks, M., and the *C. elegans* Genome Mapping and Sequencing Consortium, The *C. elegans* Genome Sequencing Project, *Genome Res.*, **5**,99-104.
- 1995 Parsons, J.D., Miropeats: Graphical DNA Sequence Comparisons, *Comput. Applic. Biosci.*, **11**, 615-619.
- 1995 Parsons, J.D., Improved Tools for DNA Comparison and Clustering, *Comput. Applic. Biosci.*, **11**, 603-613
- 1994 Johnston,M., Parsons,J., et al., Complete Nucleotide Sequence of *Saccharomyces Cerevisiae* Chromosome VIII. *Science*, **265**, 2077-2082
- 1992 Duff, K., Parsons J., and Hodgeman, T.C., Secondary structure analysis identifies a putative mouse protein demonstrating similarity to the repeat units found in CDC4, the G protein B subunits and related proteins., *J. DNA Seq. and Map.*, **3**, 213-220.
- 1992 Parsons, J.D., Brenner, S. and Bishop, M.J., (1992), Clustering cDNA Sequences, *Comput. Applic. Biosci.*, **8**, 461-466
- 1991 Poster at Cold Spring Harbor - Clustering DNA Sequences

Other Qualifications

- A Levels: Biology, Chemistry, Physics
- AO Level: Statistics.
- O Levels: English Language, Mathematics, Physics, Chemistry, Biology, French, Geography
- British and Australian driving licences

Interests and Activities

- Tourist-level German, and Spanish
- International travel and travel writing, hang gliding, cycletouring, hillwalking, skiing, sailing
- Renewable energy development, aircraft and rocket engineering.
- Photography and open source photographic software development