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EDUCATION

- 1998 - 2003 University of Cape Town, Cape Town, South Africa**
Degree course: **PhD in Molecular and Cell Biology**
Thesis title: **Development of candidate *Human papillomavirus* vaccines**
I investigated deleted and mutated gene products of the HPV-16 major capsid protein gene towards expressing candidate HPV vaccines in plants, with the final aim being the design and analysis of novel chimaeric HPV vaccines.
- 1993 - 1997 Loughborough University, Loughborough, England**
Degree Course: **Medicinal and Pharmaceutical Chemistry (BSc, DIS)**

WORK EXPERIENCE

07/16 – current Arizona State University, Tempe, USA

Associate professor in Center of Fundamental and Applied Microbiomics, The Biodesign Institute and the School of Life Sciences.

02/09 – 07/16 University of Canterbury, Christchurch, New Zealand

Senior lecturer/ researcher in molecular biology / virology

I taught virology, general molecular and microbiology to 1st, 2nd, 3rd and 4th year students and ran an active research program in DNA virus discovery and evolution.

07/03 – 09/08 University of Cape Town, Cape Town, South Africa

Lecturer for the Masters program in Structural Biology at the University of Cape Town. Funded (5 years) by the Carnegie Corporation of New York, this was the first structural biology initiative in Africa. I lectured on basic molecular biology, protein expression and purification, virus structure, aspects of structural bioinformatics, molecular visualisation and cryo-electron microscopy. In addition to this I provided molecular biology support for most of the projects undertaken by the students as part of their MSc projects.

1995 - 1996 Astra Charnwood (Astra Zeneca Pharmaceuticals), Loughborough, England

I worked as a pre-formulation research scientist in the pharmaceutical sciences department. My work mainly involved taste masking of a research compound using various ion-exchange resins and alternative salt forms. All analysis was done using HPLC, UV spectrophotometry and IR spectrophotometry in compliance with company GLP and SOPs. In addition to this, I did a fair amount of thermal degradation work using TGA, DSC, and hot stage microscopy.

ASSOCIATE RESEARCH POSITIONS

- **Honorary research associate** (2009 – current)
Structural Biology Research Unit, University of Cape Town, Cape Town, South Africa
- **Courtesy research professor** (2014 – current)
Department of Plant Pathology and Emerging Pathogens Institute, University of Florida, Gainesville, USA
- **Adjunct Associate professor** (2017 – current)
School of Biological Sciences, University of Canterbury, Christchurch, New Zealand

AWARDS AND PRIZES (ACADEMIC)

- Patent incentive award from the South African Department of Science and Technology (DST) of US\$ 2000 (2008)
- Patent incentive award from the South African Department of Science and Technology (DST) of US\$ 2000 (2007)

RESEARCH

I work across ecosystems, from plants to animals, and from the tropics to the Antarctic with a strong focus on viral evolution and dynamics, and viral metagenomics. My research uses a combination of traditional virology, microscopy (including transmission electron microscopy), molecular and cellular biology techniques in conjunction with modern techniques including next-generation sequencing, synthetic biology and bioinformatics. A brief overview of my current research projects is provided below.

- Darren Martin (author of the most widely used recombination detection software; South Africa) and I have set up a wide collaborative network of scientists world-wide to study viral evolution through recombination, viral phylogeography and global viral movement patterns of the plant-infecting geminiviruses and nanoviruses. While initially our research was based in Africa and South America, recently we have also focused on the Pacific region. As a team, Philippe Roumagnac, Pierre Lefeuvre, Darren Martin and I are attempting to identify drivers of plant viral emergence in ecosystems.
- In collaboration with Mya Breitbart, Karyna Rosario and Milen Marinov, I have embarked on a unique concept of using top end insect predators to sample viromes in various ecosystems. Additionally, my group has recently started looking at human and animal viruses vectored by black flies and mosquitoes in New Zealand and the Pacific islands.
- I have an active avian circovirus disease research program with the Department of Conservation of New Zealand, Parc Zoologique et Forestier (New Caledonia), Bethany Jackson (Murdoch University, Australia), Tomasz Stenzel (University of Warmia and Mazury, Poland) and Tomasz Piasecki (Wrocław University of Environmental and Life Sciences, Poland).
- I am running a variety of viral discovery projects associated with bark beetles in the US, fruit bats in the Pacific island of Tonga, various fungi, farm animals, insects and molluscs in New Zealand.
- In a collaborative effort, David Ainley (penguin ecologist, USA), Grant Ballard (penguin ecologist, USA), Stacey Kim (marine ecologist, USA), Jennifer Burns (seal ecologist, USA), Melanie Massaro (avian ecologist, Australia), Joseph Levy (permafrost geologist, USA), William Davison (fish physiologist, New Zealand) and I are exploring the diversity of viral communities in the Antarctic.

RESEARCH COLLABORATORS

- **Darren Martin**, University of Cape Town, Cape Town, South Africa
- **Gordon Harkins**, University of the Western Cape, Bellville, South Africa
- **Dionne Shepherd**, University of Cape Town, Cape Town, South Africa
- **Trevor Sewell**, University of Cape Town, Cape Town, South Africa
- **Edward Rybicki**, University of Cape Town, Cape Town, South Africa
- **Salem Samtally**, Mauritian Sugar Industry Research Institute, Reduit, Mauritius
- **Sunday Oluwafemi**, Bowen University, Iwo, Nigeria
- **Innocent Zinga**, University of Bangui, Bangui, Central African Republic
- **Appolinaire Tange**, IRAD Yaounde, Cameroon
- **Fidèle Tiendrebeogo**, Université de Ouagadougou, Ouagadougou, Burkina Faso
- **Jahangir Heydarnejad**, Shahid Bahonar University of Kerman, Kerman, Iran
- **Safaa Kumari**, ICARDA, Tunis, Tunisia
- **Asma Najar**, National Institute of Agricultural Research, Tunisia
- **Rob Briddon**, National Institute for Biotechnology and Genetic Engineering, Faisalabad, Pakistan
- **Michel Peterschmitt**, CIRAD, Montpellier, France
- **Stephan Blanc**, INRA, Montpellier, France
- **Jean-Michel Lett**, CIRAD, Saint Pierre, La Réunion, France
- **Pierre Lefeuvre**, CIRAD, Saint Pierre, La Réunion, France
- **Phillipe Roumagnac**, CIRAD, Montpellier, France
- **Phillipe Lemey**, Katholieke Universiteit Leuven, Leuven, Belgium
- **Tomasz Piasecki**, Wrocław University of Environmental and Life Sciences, Wrocław, Poland
- **Tomasz Stenzel**, University of Warmia, Olsztyn, Poland
- **Keizo Nagasaki**, Fisheries Research Agency, Hiroshima, Japan
- **Murillo Zerbini**, Universidade Federal de Viçosa, Viçosa, Brazil
- **Alice Inoue-Nagata**, Embrapa Hortaliç, Brasília, Brazil
- **Simone da Graça Ribeiro**, Embrapa Hortaliç, Brasília, Brazil
- **John Thomas**, University of Queensland, Brisbane, Australia
- **Andrew Geering**, University of Queensland, Brisbane, Australia
- **Almudena Lorenzo**, Direction de l'Environnement de la Province Sud, Nouméa, New Caledonia
- **Liping Pang**, Environmental Science and Research, Christchurch, New Zealand
- **Kate McInnes**, Department of Conservation, New Zealand
- **Jon Harding**, University of Canterbury, New Zealand
- **Sharyn Goldstien**, University of Canterbury, New Zealand

- **Peyman Zawar Reza**, University of Canterbury, New Zealand
- **Christopher Gomez**, University of Canterbury, New Zealand
- **William Davison**, University of Canterbury, New Zealand
- **Christopher Buck**, National Institute of Health, USA
- **Koenraad van Doorslaer**, University of Arizona, USA
- **Jane Polston**, University of Florida, Gainesville, USA
- **Terry Ng**, Centre for Disease Control and Prevention, Atlanta, USA
- **Eric Delwart**, Blood Systems Research Institute, San Francisco, USA
- **Joseph Levy**, Institute for Geophysics, University of Texas, USA
- **David G Ainley**, HT Harvey and Associates, Los Gatos, USA
- **Grant Ballard**, Point Blue Conservation Science, Petaluma, USA
- **Stacy Kim**, Moss Landing Laboratories, Moss Landing, USA
- **Jennifer Burns**, University of Alaska, Anchorage, USA
- **Mya Breitbart**, University of South Florida, USA
- **Brian Smith**, Arizona State University, USA
- **Alexander Lucas**, Arizona State University, USA
- **Anne Stone**, Arizona State University, USA
- **Melissa Wilson-Sayres**, Arizona State University, USA

STUDENT SUPERVISION

University of Canterbury, New Zealand

PhD

Ms Daisy Stanton, **PhD in Microbiology** (2010-2015) – Primary supervisor

Towards understanding the evolution of Banana bunchy top virus and the detection of associated badnaviruses

Ms Anisha Dayaram, **PhD in Microbiology** (2011-2015) – Primary supervisor

Diversity of replication associated protein encoding viruses amongst top end insect predators and concentrators in ecosystems

Ms Simona Kraberger, **PhD in Microbiology** (2010-2015) - Primary supervisor

Towards understanding mastrevirus dynamics and the use of viral metagenomic approaches to identify novel gemini-like circular DNA viruses

Ms Kata Farkas, **PhD in Microbiology** (2011-2014) - Primary supervisor

Mimicking virus retention and transport in groundwater using surface charge-modified, DNA-labelled silica nanobeads

MSc

Ms Zoe Smeele, **MSc in Microbiology** (2017) – Primary supervisor

Identification of papillomaviruses in Weddell seals

Ms Judith Nonis, **MSc in Microbiology** (2015) – Co-supervisor

Bacteriophage as a biocontrol tool for foodborne pathogens

Ms Olivia Steel, **MSc in Microbiology** (2015) - Primary supervisor

Exploring the diversity of CRESS DNA viruses associated with the faecal matter of wild and domestic animals in New Zealand

Ms Maketalena Male, **MSc in Microbiology** (2015) - Primary supervisor

Identification of CRESS DNA viruses in faeces of Pacific flying foxes in the Tongan archipelago

Ms Alyssa Sikorski, **MSc in Microbiology** (2012 -2013) – Primary supervisor

Molecular characterisation of novel single-stranded DNA viruses recovered from animal faeces

Ms Carina Davis, **MSc in Microbiology** (2012-2013) – Co-supervisor

Metagenomic approaches to microbial source tracking

Ms Laurel Julian, **MSc in Microbiology** (2011-2012) - Primary supervisor

Analysis of genetic diversity and evolution through recombination of beak and feather disease virus

Mr Robert Lawry, **MSc in Microbiology** (2009-2010) - Primary supervisor

Diversity and evolution of geminiviruses

University of Cape Town, South Africa

PhD

Mr Adérito Monjane, **PhD in Molecular and Cell Biology** (2007- 2012) – Co-supervisor
Analysis of recombination and evolution of maize streak virus

MSc

Mr Kyle Dent, **MSc in Structural Biology** (2007-2009) – Co-supervisor
3D reconstruction of maize streak virus structure

Ms Jennifer Miller, **MSc in Structural Biology** (2006-2008) – Co-supervisor
Structural studies on *Heterocapsa circularisquama* RNA virus

Honours

Mr Alistair Gray, **Honours in Molecular and Cell Biology** (2008) – Co-supervisor
Characterisation of the D surface of nitrile hydratase of *Bacillus pumilus*

Mr Oliver Windram, **Honours in Molecular and Cell Biology** (2006) – Primary Supervisor
Preliminary investigation into the use of Human papillomavirus type 16 virus like particles as a delivery vector system for foreign proteins

Murdoch University, Australia

PhD

Ms Bethany Jackson, **PhD in Wildlife Diseases** (2011-2014) – Co-supervisor
Health and disease in kakariki (*Cyanoramphus novaezelandiae*) on Tiritiri Matangi Island: Implications of Beak and feather disease virus infection for conservation management

PATENTS

International patents:

1. Rybicki E.P., **Varsani A.D.** (2010) *Chimaeric Human Papillomavirus 16 L1 Virus Like Particles and a Method for Preparing the Particles*. Patent No. 24614765, Japan.
2. Rybicki E.P., **Varsani A.D.** (2009) *Chimaeric Human Papillomavirus 16 L1 Virus Like Particles and a Method for Preparing the Particles*. Patent No. 2003232951, Australia.
3. Rybicki E.P., **Varsani A.D.** (2009) *Chimaeric Human Papillomavirus 16 L1 Virus Like Particles and a Method for Preparing the Particles*. Patent No. 1506222, Europe (EU).
4. **Varsani, A.D.** and Rybicki, E.P. (2008) *Chimaeric Human Papillomavirus 16 L1 Virus Like Particles and a Method for Preparing the Particles*. Patent No. 7407807, US.
5. Rybicki E.P., **Varsani A.D.** and Williamson A-L. (2008) *Pharmaceutical Compositions, a Method of Preparing and Isolating Said Pharmaceutical Compositions for Prophylactic Treatment of Lesions and Carcinomas*. Patent No. 221817, India.
6. Rybicki E.P., **Varsani A.D.** and Williamson A-L. (2008) *Pharmaceutical Compositions, a Method of Preparing and Isolating Said Pharmaceutical Compositions for Prophylactic Treatment of Lesions and Carcinomas*. Patent No. 02821542.7, China.
7. Rybicki E.P., **Varsani A.D.** (2006) *Chimaeric Human Papillomavirus 16 L1 Virus Like Particles and a Method for Preparing the Particles*. Patent No. 2004/10137, South Africa.
8. Rybicki E.P., **Varsani A.D.** and Williamson A-L. (2006) *Vectors and/or Constructs, and Transgenic Organisms*. Patent No. 02821507.9, China.
9. Rybicki E.P., **Varsani A.D.** and Williamson A-L. (2005) *Pharmaceutical Compositions, a Method of Preparing and Isolating Said Pharmaceutical Compositions for Prophylactic Treatment of Lesions and Carcinomas*. Patent No. 2004/2505, South Africa.
10. Rybicki E.P., **Varsani A.D.** and Williamson A-L. (2005) *Vectors and/or Constructs, and Transgenic Organisms*. Patent No. 2004/2504, South Africa.

PUBLICATIONS

H-Index: 38; i10-index: 106

1. Fahsbender, E., Burns, J.M., Kim, S., Kraberger, S., Frankfurter, G., Eilers, A., Shero, M., Beltran, R., Kirkham, A., McCorkell, R., Bergart, R., Male, M.F., Ballard, G., Ainley, D.G., Breitbart, M., **Varsani, A.** (2017) *Diverse and highly recombinant anelloviruses associated with Weddell seals in Antarctica*. *Virus Evolution*. 3:vex017
2. Nonis, J., Premaratne, A., Billington, C., **Varsani, A.** (2017) *Genome of a podovirus (AAPEc6) isolated from wastewater in New Zealand that infects *Escherichia coli* O45:H10*. *Genome Announcements*. 5: e00462-17
3. Boukari, W., Alcalá-Briseño, R.I., Kraberger, S., Fernandez, E., Filloux, D., Daugrois, J-H., Comstock, J.C., Lett, J-M., Martin, D.P., **Varsani, A.**, Roumagnac, P., Polston, J.E., Rott, P.C. (2017) *Occurrence of a novel*

mastrevirus in sugarcane germplasm collections in Florida, Guadeloupe and Réunion. *Virology Journal*. 14:e146

4. Kraberger, S., Saumtally, S., Pande, D., Khoodoo, M.H.R., Dhayan, S., Dookun-Saumtally, A., Shepherd, D.N., Hartnady, P., Atkinson, A., Lakay, F., Hanson, B., Redhi, D., Monjane, A.L., Windram, O., Walter, M., Oluwafemi, O., Michel-Lett, J., Lefeurve, P., Martin, D.P., **Varsani, A.** (2017) *Molecular diversity, geographic distribution and host range of monocot-infecting mastreviruses in Africa and surrounding islands*. *Virus Research*. 238: 171-178
5. Kraberger, S., Geering, A.D.W., Walters, M., Martin, D.P., **Varsani, A.** (2017) *Novel mastreviruses identified in Australian wild rice*. *Virus Research*. 238: 193-197
6. Adams, M.J., Lefkowitz, E.J. King, A.M.Q., Harrach, B., Harrison, R., Knowles, N.J., Kropinski, A.M., Krupovic, M., Kuhn, J.H., Mushegian, A.R., Nibert, M., Sabanadzovic, S., Sanfaçon, H., Siddell, S.G., Simmonds, P., **Varsani, A.**, Zerbini, F.M., Gorbalenya, A.E., Davison, A.J. (2017) *Changes to taxonomy and the International Code of Virus Classification and Nomenclature ratified by the International Committee on Taxonomy of Viruses*. *Archives of Virology*. 162:2505–2538
7. Susi, H., Laine, A-L., Filloux, D., Kraberger, S., Farkas, K., Bernardo, P., Frilander, M.J., Martin, D.P., **Varsani, A.**, Roumagnac, P. (2017) *Genome sequences of a capulavirus infecting Plantago lanceolata in the Åland archipelago of Finland*. *Archives of Virology*. 162: 2041-2045
8. **Varsani, A.**, Roumagnac, P., Fuchs, M., Navas-Castillo, J., Moriones, E., Idris, A., Briddon, R.W., Rivera-Bustamante, R., Zerbini, F.M., Martin, D.P. (2017) *Capulavirus and Grablovirus: Two new genera in the family Geminiviridae*. *Archives of Virology*. 162: 1819-1831
9. Kraberger, S., Polston, J.E., Capobianco, H.M., Alcalá-Briseño, R.I., Fontenele, R.S., **Varsani, A.** (2017) *Genomovirus genomes recovered from Echinothrips americanus sampled in Florida, USA*. *Genome Announcements*. 5:e00445-17
10. Ouattara, A., Tiendrébéogo, F., Lefeuve, P., Hoareau, M., Traoré, E.V., Barro, N., Traoré, O., **Varsani, A.**, Lett, J-M. (2017) *New strains of chickpea chlorotic dwarf virus discovered on papaya and tomato in Burkina Faso*. *Archives of Virology*. 162: 1791-1794
11. Najar, A., Hamdi, I., **Varsani, A.** (2017) *Barley yellow dwarf virus in barley crops in Tunisia: prevalence and molecular characterization*. *Phytopathologia Mediterranea*. 56, 111–118
12. Postler, T.S., Clawson, A.N., Amarasinghe, G.K., Basler, C.F., Bavari, S., Benkő, M., Blasdel, K.R., Briese, T., Buchmeier, M.J., Bukreyev, A., Calisher, C.H., Chandran, K., Charrel, R., Clegg, C.S., Collins, P.L., de la Torre, J. C., DeRisi, J.L., Dietzgen, R.G., Dolnik, O., Dürrwald, R., Dye, J.M., Easton, A J., Emonet, S., Formenty, P., Fouchier, R. A. M., Ghedin, E., Gonzalez, J.-P., Harrach, B., Hewson, R., Horie, M., Jiāng, D., Kobinger, G., Kondo, H., Kropinski, A.M., Krupovic, M., Kurath, G., Lamb, R.A., Leroy, E.M., Lukashevich, I.S., Maisner, A., Mushegian, A.R., Netesov, S.V., Nowotny, N., Patterson, J.L., Payne, S.L., Paweska, J.T., Peters, C.J., Radoshitzky, S.R., Rima, B.K., Romanowski, V., Rubbenstroth, D., Sabanadzovic, S., Sanfaçon, H., Salvato, M. S., Schwemmle, M., Smither, S.J., Stenglein, M.D., Stone, D.M., Takada, A., Tesh, R.B., Tomonaga, K., Tordo, N., Towner, J.S., Vasilakis, N., Volchkov, V.E., Wahl-Jensen, V., Walker, P.J., Wang, L.F., **Varsani, A.**, Whitfield, A.E., Zerbini, F.M., Kuhn, J.H. (2017). *Possibility and challenges of conversion of current virus species names to Linnaean binomials*. *Systematic Biology*. 66 (3): 463-473
13. Heydarnejad, J., Kamali, M., Massumi, H., Kvarnheden, A., Male, M.F., Kraberger. S., Stainton, D., Martin, D.P., **Varsani, A.** (2017) *Identification of a nanovirus-aphasatellite complex in Sophora alopecuroides*. *Virus Research*. 235: 24-32
14. Adams, M.J., Lefkowitz, E.J. King, A.M.Q., Harrach, B., Harrison, R., Knowles, N.J., Kropinski, A.M., Krupovic, M., Kuhn, J.H., Mushegian, A.R., Nibert, M., Sabanadzovic, S., Sanfaçon, H., Siddell, S.G., Simmonds, P., **Varsani, A.**, Zerbini, F.M., Orton, R.J., Smith, D.B., Gorbalenya, A.E., Davison, A.J. (2017) *50 years of formal virus taxonomy: overview and prospects*. *Archives of Virology*. 162: 1441-1446
15. **Varsani, A.**, Frankfurter, G., Stainton, D., Male, M.F., Kraberger, S., Burns, J.M. (2017) *Identification of a polyomavirus in a Weddell seal (Leptonychotes weddellii) from the Ross Sea (Antarctica)*. *Archives of Virology* 162: 1403-1407
16. Rosario, K., Breitbart, M., Harrach, B., Segalés, J., Delwart, E., Biagini, P., **Varsani, A.** (2017) *Revisiting the taxonomy of the family Circoviridae: Establishment of the genus Cyclovirus and removal of the genus Gyrovirus*. *Archives of Virology* 162:1447-1463
17. Godinhoa, M.T., Paula, D.P., **Varsani, A.**, Ribeiro, S.G. (2017) *Genome sequence of Cauliflower mosaic virus identified in earwigs (Doru luteipes) through a metagenomic approach*. *Genome Announcements*. 11: e00043-17
18. Simmonds, P., Adams, M. J., Benkő, M., Breitbart, M., Brister, J. R., Carstens, E. B., Davison, A. J., Delwart, E., Gorbalenya, A. E., Harrach, B., Hull, R., King, A. M. Q., Koonin, E. V., Krupovic, M., Kuhn, J. H., Lefkowitz, E. J., Nibert, M. L., Orton, R., Roossinck, M. J., Sabanadzovic, S., Sullivan, M. B., Suttle, C. A., Tesh, R. B., van der Vlugt, R. A., **Varsani, A.**, Zerbini, F. M. (2017). *Virus taxonomy in the age of metagenomics*. *Nature Reviews Microbiology*. 15: 161-168
19. Pang, L., Robson, B., Farkas, K., McGill, E., **Varsani, A.**, Gillot, L., Li, J., Abraham, P. (2017) *Tracking effluent discharges in undisturbed stony soil and alluvial gravel aquifer using synthetic DNA tracers*. *Science of the Total Environment*. 492: 144-152

20. Zerbini, F.M., Briddon, R.W., Idris, A., Martin, D.P., Moriones, E., Navas-Castillo, J., Rivera-Bustamante, R., Rougmanac, P., **Varsani, A.**, ICTV Report Consortium (2017) *ICTV Virus Taxonomy Profile: Geminiviridae*. Journal of General virology. 98:131–133
21. Stainton, D., Martin, D.P., Collings, D.A., **Varsani, A.** (2017) *Comparative analyses of common regions found in babuviruses and alphasatellite molecules*. Archives of Virology. 162:849-855
22. Walters, M., Bawuro, M., Christopher, A., Knight, A., Kraberger, S., Stainton, D., Chapman, H., **Varsani, A.** (2017) *Novel single-stranded DNA virus genomes recovered from chimpanzee faeces sampled from Mambilla Plateau in Nigeria*. Genome Announcements. 5: e01715-16
23. Kamali, M., Heydarnejad, J., Pouramini, N., Masumi, H., Farkas, K., Kraberger, S., **Varsani, A.** (2017) *Genome sequences of Beet curly top Iran virus, Oat dwarf virus, Turnip curly top virus, and Wheat dwarf virus identified in leafhoppers*. Genome Announcements. 5: e01674-16.
24. Kazlauskas, D., Dayaram, A., Kraberger, S., Goldstien, S., **Varsani, A.**, Krupovic, M. (2017) *Evolutionary history of ssDNA bacilladnaviruses features horizontal acquisition of the capsid protein gene from ssRNA nodaviruses*. Virology. 204: 114-121
25. Pande, D., Madzokere, E., Hartnady, P., Kraberger, S., Hadfield, J., Rosario, K., Jäschke, A., Monjane, A.L., Dida, M.M., Shepherd, D.N., Martin, D.P., **Varsani, A.**, Harkins, G.W. (2017) *The role of Kenya in the trans-African spread of maize streak virus strain A*. Virus Research. 232: 69-76
26. **Varsani, A.**, Krupovic, M. (2017) *Sequence-based taxonomic framework for the classification of uncultured single-stranded DNA viruses of the family Genomoviridae*. Virus Evolution. 3: vew037
27. Mabvakure, B., Martin, D.P., Kraberger, S., Cloete, L., van Brunschot, S., Geering, A.D., Thomas, J.E., Bananej, K., Lett, J-M., Lefevre, P., **Varsani, A.**, Harkins, G.W. (2016) *Ongoing geographical spread of Tomato yellow leaf curl virus*. Virology. 498: 257–264
28. Lamas, N.S., Fontenele, R.S., Melo, F.L., Costa, A.F., **Varsani, A.**, Ribeiro, S.G. (2016) *Complete sequence of a genomovirus associated with common bean plant leaves in Brazil*. Genome Announcements. 4:e01247-16
29. Bananej, K., Kraberger, K., **Varsani, A.** (2016) *Okra enation leaf curl virus in papaya from Iran displaying severe leaf curl symptoms*. Journal of Plant Pathology 98, 637-639
30. Frost, C.M., Peralta, G., Rand, T.A., Didham, R.K., **Varsani, A.**, Tylianakis, J. (2016) *Apparent competition drives community-wide attack rates across ecosystem boundaries*. Nature Communications. 7: e12644
31. De Bruyn, A., Harimalala, M., Zinga, I., Mabvakure, B.M., Hoareau, M., Ravigné, V., Walters, M., Reynaud, B., **Varsani, A.**, Harkins, G.W., Martin, D.P., Lett, J-M., Lefevre, P. (2016) *Divergent evolutionary and epidemiological dynamics of cassava mosaic geminiviruses in Madagascar*. BMC Evolutionary Biology. 16: e182
32. Adams, M.J., Lefkowitz, E.J., King, A.M.Q., Harrach, B., Harrison, R.L., Knowles, N.J., Kropinski, A.M., Krupovic, M., Kuhn, J.H., Mushegian, A.R., Nibert, M., Sabanadzovic, S., Sanfaçon, H., Siddell, S.G., Simmonds, P., **Varsani, A.**, Zerbini, F.M., Gorbalenya, A.E., Davison, A.J. (2016) *Ratification vote on taxonomic proposals to the International Committee on Taxonomy of Viruses (2016)*. Archives of Virology. 161:2921–2949
33. Krupovic, M., Ghabrial, S.A., Jiang, D., **Varsani, A.** (2016) *Genomoviridae: a new family of widespread single-stranded DNA viruses*. Archives of virology. 161:2633–2643
34. Steel, O., Kraberger, S., Sikorski, A., Young, L.M., Catchpole, R.J., Stevens, A.J., Ladley, J.J., Coray, D.S., Stainton, D., Dayaram, A., Julian, L., van Bysterveldt, K., **Varsani, A.** (2016) *Circular replication-associated protein encoding DNA viruses identified in the faecal matter of various animals in New Zealand*. Infection, Genetics and Evolution. 43, 151-164
35. Khalifa, M., **Varsani, A.**, Ganley, A.R.D., Pearson, M.N. (2016) *Comparison of Illumina de novo assembled and Sanger sequenced viral genomes: A case study for RNA viruses recovered from the plant pathogenic fungus Sclerotinia sclerotiorum*. Virus Research. 219, 51-57
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- 2) Committee member of the “International Committee on Taxonomy of Viruses” geminivirus working group (2010 – current)
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- 4) Advisory board for the journal Archives of Virology (2009 – 2016)
- 5) Advisory board of the journal Virology (2015 – current)
- 6) Referee for the following journals
 - Nature Microbiology Review (NPG)
 - ISME (NPG)
 - Scientific Reports (NPG)
 - Emerging Infectious Diseases (Centers for Disease Control and Prevention)
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