Mohammad Ali Nilforooshan

CONTACT LIC

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Hamilton 3240, New Zealand

E-mail: mohammad.nilforooshan@lic.co.nz Website: https://nilforooshan.github.io GitHub: https://github.com/nilforooshan

Google Scholar: https://goo.gl/iKbCoU

Interests Matrix Algebra

Statistical methods applied to livestock genetic and genomic evaluation

Putting biology into equations and equations into program codes

Single-step genomic evaluation

Computational biology

Traveling

EDUCATION Swedish University of Agricultural Sciences, Dept. Animal Breeding and Genetics

PhD in Animal Breeding and Genetics – Quantitative Genetics, June 2007 – May 2011

AWARDS • Grant from Interbull Centre for participation in the 9th World Congress on Genetics Applied to Livestock Production (2010), Leipzig, Germany

• Grant from FUR foundation (SLU funds for internationalization of postgraduate studies) for participation in the 60th Annual meeting of the European Association for Animal Production (2009), Barcelona, Spain

• Scholarship from the Guest Scholarship Program (2008) of the Swedish Institute

• Grant from FUR foundation (SLU funds for internationalization of postgraduate studies) for participation in the 59th Annual meeting of the European Association for Animal Production (2008), Vilnius, Lithuania

 ${\rm Skills}$

Scientific: Matrix Algebra, Quantitative Genetics and General Animal Breeding, Statistics, "Big Data", Occasional reviewer for some scientific journals

Programming: R, SAS, Fortran, Linux Shell scripting (Sed and Awk), Python, Markdown and RMarkdown, Git and GitHub, Docker, Julia (beginner), and C (beginner)

Analytics: Designing, streamlining and documenting genetic/genomic evaluation pipelines; experienced in several computer packages such as ASReml, DFReml, MiX99 and BLUPF90

Professional Expenses

LIC

Experience Scientist - Quantitative Genetics

May 2020 – Present

June 2012 - October 2015

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Postdoctoral Research Fellow in Quantitative Genetics December 2018 – May 2020

University of Otago, Dept. Mathematics and Statistics

Postdoctoral Fellow in Quantitative Genetics November 2015 – October 2018

Swedish University of Agricultural Sciences,

Dept. Animal Breeding and Genetics, Interbull Centre

Geneticist

University of Nebraska-Lincoln, Dept. Statistics

Postdoctoral Research Fellow in Computational Biology September 2011 – May 2012

Isfahan Science and Technology Town, Animal Science Core

Research Assistant September 2004 – June 2006

TEACHING

- Supervising a visiting PhD student to the Department of Mathematics and Statistics, University of Otago, August – October, 2018.
- Teaching some lectures of the postgraduate course in Quantitative Genetics, held at the University of Otago, Dunedin, New Zealand, July – October, 2018.
- Teaching some lectures of the postgraduate course in Quantitative Genetics, held at the University of Otago, Dunedin, New Zealand, July – October, 2017.
- Teaching R programming, and giving several other talks in the University of Chapingo, Texcoco, Mexico, April 2015.
- Teaching "Dairy Cattle Husbandry" in Kabootar-Abad school of Agriculture, Isfahan, Iran, September 2004 – June 2005.

Software

Nilforooshan, M.A. 2020. R package pedSimulate: Pedigree, Genetic Merit and Phenotype Simulation.

Nilforooshan, M.A. 2019. R package mbend: Matrix Bending.

Nilforooshan, M.A. 2019. R package ggroups: Pedigree and Genetic Groups.

PUBLICATIONS

- Nilforooshan, M.A. 2020. mbend: An R package for bending non-positive-definite symmetric matrices to positive-definite. *BMC Genet.* 21: 97. doi:10.1186/s12863-020-00881-z
- Nilforooshan, M.A., and L.A. Saavedra-Jiménez. 2020. ggroups: An R package for pedigree and genetic groups data. *Hereditas* 157: 17. 10.1186/s41065-020-00124-2
- Nilforooshan, M.A. 2020. Memory-efficient self-cross-product for large matrices using R and Python. J. Math. Comput. Sci. 10(3): 497–506. doi:10.28919/jmcs/4457
- Nilforooshan, M.A. 2020. Updating genetic relationship matrices and their inverses: A methodology note. Can. J. Anim. Sci. 100(2): 292–298. doi:10.1139/CJAS-2019-0106
- Nilforooshan, M.A. 2020. Application of single-step GBLUP in New Zealand Romney sheep. *Anim. Prod. Sci.* 60(9): 1136–1144. doi:10.1071/AN19315
- Nilforooshan, M.A., and M. Lee. 2019. The quality of the algorithm for proven and young with various sets of core animals in a multi-breed sheep population. *J. Anim. Sci.* 97(3): 1090–1100. doi:10.1093/jas/skz010
- Nilforooshan, M.A., J.H. Jakobsen, W.F. Fikse, B. Berglund, and H. Jorjani. 2014. Multiple-trait multiple country genetic evaluation of Holstein bulls for female fertility and milk production traits. *animal* 8(6) 887–894. doi:10.1017/S1751731114000895
- Behdad, S., M.A. Edriss, M.A. Nilforooshan, and A.H. Movassagh. 2013. Wool characteristics
 of Bakhtiari sheep and the correlated responses to selection for pre-weaning daily gain. *Indian*J. Anim. Sci. 83(1): 82–85.
- Nilforooshan, M.A. 2011. Multiple-trait multiple country genetic evaluation of fertility traits in dairy cattle. Uppsala: Sveriges lantbruksuniv., *Acta Universitatis Agriculturae Sueciae*, 1652-6880; 2011:31. ISBN 978-91-576-7566-8 [Doctoral thesis].
- Nilforooshan, M.A., W.F. Fikse, B. Berglund, J.H. Jakobsen, and H. Jorjani. 2011. Short communication: Quantifying bias in a single-trait international model ignoring covariances from multiple-trait national models. *J. Dairy Sci.* 94: 2631–2636. doi:10.3168/jds.2010-3863
- Nilforooshan, M.A., J.H. Jakobsen, W.F. Fikse, B. Berglund, and H. Jorjani. 2010. Application of a multiple-trait multiple-country genetic evaluation model for female fertility traits. *J. Dairy Sci.* 93: 5977–5986. doi:10.3168/jds.2010-3437
- Nilforooshan, M.A. 2010. Contemporary grouping in mixed size dairy herds experiencing four seasons. *Turk. J. Vet. Anim. Sci.* 34: 129–135. doi:10.3906/vet-0710-18
- Nilforooshan, M.A., A. Khazaeli, and M.A. Edriss. 2008. Effects of missing pedigree information on dairy cattle genetic evaluations. *Arch. Tierz.*, *Dummerstorf* 51: 99–110. doi:10.5194/aab-51-99-2008
- Edriss, M.A., P. Hosseinia, M. Edrisi, H.R. Rahmani, and M.A. Nilforooshan. 2008. Prediction of second parity milk performance of dairy cows from first parity information using artificial neural network and multiple linear regression methods. *Asian J. Anim. Vet. Adv.* 3(4): 222–229. doi:10.3923/ajava.2008.222.229
- Hosseinia, P., M. Edrisi, M.A. Edriss, and M.A. Nilforooshan. 2007. Prediction of second

- parity milk yield and fat percentage of dairy cows based on first parity information using neural network system. J. Applied Sci. 7(21): 3274–3279. doi:10.3923/jas.2007.3274.3279
- Edriss, M.A., G. Dashab, A.A. Ghare Aghaji, M.A. Nilforooshan, and H. Movasagh. 2007. A study on some physical attributes of Naeini sheep wool for textile industry. *Pak. J. Biol. Sci.* 10(3): 415–420. doi:10.3923/pjbs.2007.415.420
- Nilforooshan, M.A., and M.A. Edriss. 2007. Comparison of Holstein bull semen sources on milk traits in Isfahan province in Iran. *Arch. Tierz.*, *Dummerstorf* 50: 71–83. doi:10.5194/aab-50-71-2007
- Edriss, M.A., M.A. Nilforooshan, and J.M. Sadeghi. 2006. Estimation of direct genetic and maternal effects for production traits of Iranian Holstein cows using different animal models. *Pak. J. Biol. Sci.* 9(4): 636–640. doi:10.3923/pjbs.2006.636.640
- Dashab, G., M.A. Edriss, A.A. Ghare Aghaji, H. Movasagh, and M.A. Nilforooshan. 2006.
 Wool fiber quality of Naeini sheep. Pak. J. Biol. Sci. 9(2): 270–276. doi:10.3923/pjbs.2006.270.276
- Nilforooshan, M.A., and M.A. Edriss. 2004. Effect of age at first calving on some productive and longevity traits in Iranian Holsteins of the Isfahan province. *J. Dairy Sci.* 87: 2130–2135. doi:10.3168/jds.S0022-0302(04)70032-6
- Nilforooshan, M.A. 2003. Genetic Evaluation and Comparison between Domestic and Imported (Canada, USA, Europe) Sperms for Milk Yield, Fat Yield, and Fat Percentage in Some Dairy Farms of Isfahan Province. [MSc thesis].

Proceedings

- Nilforooshan, M.A., B. Auvray, M. Lee. 2018. Animal-specific combining of the GRM and the NRM in single-step GBLUP. 11th World Congress on Genetics Applied to Livestock Production, Auckland, New Zealand. February, 11–16, 2018. [poster file, paper]
- Pabiou, T., M.A. Nilforooshan, E. Venot and K. Moore. 2015. Interbeef presents new opportunities for beef farmers. *British Cattle Conference*, Telford, England. January 19–21, 2015.
- Pabiou, T., M.A. Nilforooshan, D. Laloe, E. Hjerpe, and E. Venot. 2014. Across-Country Genetic Parameters in Beef Cattle for Interbeef Weaning Weight Genetic Evaluation. 10th World Congress on Genetics Applied to Livestock Production, Vancouver, Canada. August, 17–22, 2014. [poster file]
- Venot, E., T. Pabiou, E. Hjerpe, M.A. Nilforooshan, A. Launay, and B. Wickham. 2014. Benefits of Interbeef International Genetic Evaluations for Weaning Weight. 10th World Congress on Genetics Applied to Livestock Production, Vancouver, Canada. August, 17–22, 2014.
- Mäntysaari, E. and M.A. Nilforooshan. 2014. Effect of the size of the reference population on the validation reliability of national genomic evaluations. *Interbull Meeting*, Berlin, Germany. May 20–21, 2014.
- Nilforooshan, M.A. 2014. A review of the validation of national genomic evaluations. *Interbull Bulletin*. 48: 111–117. *Interbull Meeting*, Berlin, Germany. May 20–21, 2014. [slides file]
- Nilforooshan, M.A. and E. Hjerpe. 2014. Interbeef report on weaning genetic evaluation, performed in March 2014 for Charolais and Limousin. *Interbeef Technical Meeting*, Uppsala, Sweden. March 10, 2014.
- Braillon, L. and M.A. Nilforooshan. 2013. Review of the Interbeef proof publication proposal. *Interbeef Technical Meeting*, Nantes, France. August 27, 2013.
- Nilforooshan, M.A., E. Hjerpe, V. Palucci and J. Dürr. 2013. Interbeef report on weaning genetic evaluation for Charolais and Limousin. *Interbeef Technical Meeting*, Aarhus, Denmark. March 27–28, 2013.
- Nilforooshan, M.A. 2012. Report on the weaning weight phenotypic data editing for Charolais and Limousin. *Interbeef Technical Meeting*, Uppsala, Sweden. December 6–7, 2012.
- Jorjani, H., J. Jakobsen, M.A. Nilforooshan, E. Hjerpe, B. Zumbach, V. Palucci and J. Dürr. 2011. Genomic Evaluation of BSW Populations. InterGenomics: Results and Deliverables. Interbull Bulletin. 43: 5–8. Interbull Technical Workshop, Guelph, Ontario, Canada. February 27–28, 2011. [slides file]
- Nilforooshan, M.A., L. Rönnegård and H. Jorjani. 2010. Various statistical models for prediction of SNP effects from a multi-national/breed reference population. 9th World Congress on Genetics Applied to Livestock Production, Leipzig, Germany. August 1–6, 2010. [poster file]

- Nilforooshan, M.A., B. Zumbach, J. Jakobsen, A. Loberg, H. Jorjani, and J. Dürr. 2010. Validation of national genomic evaluations. *Interbull Bulletin*. 42: 56–61. *Interbull Meeting*, Riga, Latvia. May 31 to June 4, 2010. [slides file]
- Nilforooshan, M.A. 2009. Developing the method of estimating genetic similarity between populations. 60th Annual meeting of the European Association for Animal Production, Barcelona, Spain, Book of Abstracts No. 15: 610. [slides file]
- Nilforooshan, M.A., J. Jakobsen, F. Fikse, B. Berglund and H. Jorjani. 2009. MT–MACE for female fertility and milk yield. *Interbull Bulletin*. 40: 68–71. *Interbull Meeting*, Barcelona, Spain. August 21–24, 2009. [slides file]
- Nilforooshan, M.A. 2009. International genetic evaluations of fertility traits using MT–MACE. Faculty research day, Faculty of Veterinary Medicine and Animal Science, Swedish University of Agricultural Sciences, Uppsala, Sweden. December 7, 2009.
- Nilforooshan, M.A. 2009. International genetic evaluations of fertility traits using MT-MACE. 80th Anniversary of the Department of Animal Breeding and Genetics, Swedish University of Agricultural Sciences, Uppsala, Sweden, June 4, 2009.
- Nilforooshan, M.A., J. Jakobsen, F.W. Fikse, B. Berglund, and H. Jorjani. 2009. International genetic evaluations of fertility traits using multi-trait MACE. *Interbull Bulletin*. 39:99–102. Interbull Workshop on Genomic Information in Genetic Evaluations, Uppsala, Sweden. January 26–29, 2009. [slides file]
- Nilforooshan, M.A., F.W. Fikse and H. Jorjani. 2008. International genetic evaluations of fertility traits considering more than one trait per country. 59th Annual meeting of the European Association for Animal Production, Vilnius, Lithuania, Book of Abstracts No. 14: 268. [slides file]

Extension Papers in Farsi