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Main Areas of Research

My focus has been to understand **hybrid zones**: narrow regions in which distinct populations meet, mate and produce hybrids. I have worked on a variety of field systems, including grasshoppers (*Podisma pedestris*), toads (*Bombina bombina/variegata*), butterflies (*Heliconius erato* and *H.melpomene*), and snapdragons (*Antirrhinum majus*). This led naturally to development of **mathematical models** for how populations evolve when they extend through space, and when large numbers of genes interact. These models have been applied to a wide range of questions, and in particular, to **speciation**, and to the **quantitative genetics** of complex traits. Key components of my present work are: i) a long-term study of the Antirrhinum hybrid zone, and ii) development of methods for inferring population structure and selection from DNA sequence variation, from both natural and experimental populations.

Academic Career

2015 – present	Dean of Graduate School Institute of Science and Technology Austria (IST), Austria
2008 – present	Professor , Institute of Science and Technology (IST), Austria
2000 - 2010	Personal Chair , Institute of Evolutionary Biology University of Edinburgh, UK
1990 – 2000	Darwin Trust Fellow , Institute of Evolutionary Biology University of Edinburgh, UK
1982 – 1990	Lecturer/Reader, Department of Genetics, University College London, UK
1979 – 1982	Research Fellowship, Girton College, UK
1976 – 1979	Ph.D. in Biology (Supervised by Dr. G.M. Hewitt) <i>University of East Anglia</i> , UK Thesis: A narrow hybrid zone in the alpine grasshopper <i>Podisma pedestris</i>
1973 - 1976	B.A. (First Class) in Natural Sciences (Genetics) University of Cambridge, UK

Additional research achievements

- Scientific awards: Erwin Schrodinger Prize (2013), Austrian Academy of Sciences; Mendel Medal (2013), Leopoldina, Darwin-Wallace Medal (2009), Linnean Society of London; Darwin Medal (2006), The Royal Society; Zoological Society Scientific Medal (1992); American Society of Naturalists President's Award (joint with Mark Kirkpatrick) (1998); David Starr Jordan Prize (joint with S. Pacala), (1994); Linnean Society Bicentennial Medal (1985).
- 2. Academic Memberships and community service: Elected President, Society for the Study of Evolution (2001, on Council 2000-2002); Elected Vice-President, Society for the Study of Evolution (1989); Chair, Human Frontiers Fellowship panel(2013-2016); Chair, ERC Starting Grants Panel LS8 (2013 present); Elected President, Royal Society, Biological Sciences Awards Committee (2009-2011), Co-Chair, Portuguese Foundation for Science and Technology (FCT), Chair, Royal Society Research Grants Committee (Board H) (2004 2006).
- 3. **Editorships**: Senior Editor, *Genetics* (2016 present), Associate Editor, *Molecular Ecology* 013 present], Associate Editor, *Genetics* (2013 present), Editorial Board, *Philosophical Transactions of the Royal Society B* (2010 2015), Handling Editor, *Evolution* (2008-2011), Editorial Board, PLoS Biology (2004 2007), Editorial Board, *American Naturalist* (2003), Editorial Board, *Journal of Evolutionary Biology* (1997-2000).

4. **Recent Grants**: 2010-2015 – ERC Advanced Grant "Information and Evolution", €1.9M, 2010-2014 – Natural Selection in Spatially Structured Populations, (EPSRC 1013091 €215K), with Alison Etheridge, 2014-2018 – FP7 FET grant SAGE "Speed of Adaptation in Population Genetics and Evolutionary Algorithms", grant with T. Paixao (IST), Per Kristian Lehre (Nottingham), Tobias Friedrich (Jena), Dirk Sudholt (Sheffield). (€366K)

Publications

10 Most Important Scientific Publications

- 1. Barton, N.H. 2017. How does epistasis influence the response to selection? Heredity 118.1 (2017): 96-109. doi:10.1038/hdy.2016.109
- 2. Charlesworth, D., Barton, N.H., Charlesworth, B. 2017. The sources of adaptive evolution. Proc. Roy. Soc. (Lond.) B 284: 2016.2864. doi:doi: 10.1098/rspb.2016.2864
- 3. Ringbauer, H., Coop, G., Barton, N. H.. (2017). Inferring recent demography from isolation by distance of long shared sequence blocks. Genetics 205(3):1335-1351 doi:10.1534/genetics.116.196220
- 4. Paixao, T., Barton, N.H. 2016. The effect of gene interactions on the long-term response to selection. PNAS 113: 4422-4427 doi:10.1073/pnas.1518830113
- Paixao, T., Badkobehe, G., Barton, N.H., Dolgan, C., Dang, D.C., Friedrich, T., Lehre, P.K., Sudholt, D., Trubenova, B. 2015. Towards a unifying framework for evolutionary processes. J. Theor. Biol. 383: 28-43. doi:10.1016/j.jtbi.2015.07.011
- Barton, N.H. (2009) Why sex and recombination? Cold Spring Harbor Symposia Quant. Biol. 74 doi: 10.1101/sqb.2009.74.030
- 7. Barton, N.H., Briggs, D.E.G., Eisen, J.A., Goldstein, D.B., Patel, N.H. (2007) Evolution. Cold Spring Harbor Laboratory Press. ISBN-13: 9780879696849
- 8. Kirkpatrick, M., Johnson, T. Barton, N.H. (2002) General models of multilocus evolution. Genetics 161: 1727-1750 PMID: 12196414
- 9. Coyne, J.A., N.H. Barton, and M. Turelli. (1997) A critique of Wright's shifting balance theory of evolution. Evolution 51: 643-671. doi:10.1111/j.1558-5646.1997.tb03650.x
- 10. Barton, N. H. (1995) Linkage and the limits to natural selection. Genetics 140: 821-841. PMID: 7498757