### **Human Evolution**

Genes, Genealogies and Phylogenies

Controversy over human evolution remains widespread. However, the Human Genome Project and genetic sequencing of many other species have provided myriad precise and unambiguous genetic markers that establish our evolutionary relationships with other mammals. *Human Evolution* identifies and explains these identifiable rare and complex markers, including endogenous retroviruses, genome-modifying transposable elements, genedisabling mutations, segmental duplications, and gene-enabling mutations. The new genetic tools also provide fascinating insights into when, and how, many features of human biology arose: from aspects of placental structure; vitamin C-dependence and trichromatic vision; to tendencies to gout, cardiovascular disease and cancer.

Bringing together a decade's worth of research and tying it together to provide an overwhelming argument for the mammalian ancestry of the human species, this book will be of interest to professional scientists and students in both the biological and biomedical sciences.

GRAEME FINLAY is Senior Lecturer in Scientific Pathology at the Department of Molecular Medicine and Pathology, and Honorary Senior Research Fellow at the Auckland Cancer Society Research Centre, University of Auckland, New Zealand. Cambridge University Press 978-1-107-04012-0 - Human Evolution: Genes, Genealogies and Phylogenies Graeme Finlay Frontmatter More information

# **Human Evolution**

## Genes, Genealogies and Phylogenies

GRAEME FINLAY

Department of Molecular Medicine and Pathology, Auckland Cancer Society Research Centre, University of Auckland, New Zealand



#### CAMBRIDGE UNIVERSITY PRESS

University Printing House, Cambridge CB2 8BS, United Kingdom

Cambridge University Press is part of the University of Cambridge.

It furthers the University's mission by disseminating knowledge in the pursuit of education, learning and research at the highest international levels of excellence.

www.cambridge.org Information on this title: www.cambridge.org/9781107040120

© G. Finlay 2013

This publication is in copyright. Subject to statutory exception and to the provisions of relevant collective licensing agreements, no reproduction of any part may take place without the written permission of Cambridge University Press.

First published 2013 3rd printing 2014

Printed in the United Kingdom by Clays, St Ives plc

A catalogue record for this publication is available from the British Library

Library of Congress Cataloguing in Publication data
Finlay, Graeme, 1953–
Human evolution : genes, genealogies and phylogenies / Graeme Finlay,
Department of Molecular Medicine and Pathology, Auckland Cancer Society
Research Centre, University of Auckland, New Zealand.
pages cm
Includes bibliographical references and index.
ISBN 978-1-107-04012-0 (hardback)
1. Human evolution. 2. Human population genetics. 3. Evolutionary
genetics. 4. Genetic genealogy. I. Title.
GN281.F54 2013
599.93'8-dc23 2013015863

ISBN 978-1-107-04012-0 Hardback

Cambridge University Press has no responsibility for the persistence or accuracy of URLs for external or third-party internet websites referred to in this publication, and does not guarantee that any content on such websites is, or will remain, accurate or appropriate.

### Contents

	Pref	page ix			
	Prologue				
	1	Darwin's science			
	2	Genet	ics arrives on the scene	4	
	3	Theol	6		
	4	Interpretations of evolution today			
	5	Evolut	tion and the genome revolution	12	
	6	The so	cope of this book	18	
1	Retroviral genealogy			21	
	1.1	The retroviral life cycle			
	1.2	Retroviruses and the monoclonality of tumours			
	1.3	Endog			
		of spee	cies	32	
	1.4	.4 Natural selection at work: genes from junk		47	
		1.4.1	ERVs and the placenta	48	
		1.4.2	ERVs that contribute to gene content	55	
	1.5	Natural selection at work: regulatory networks			
	1.6	Are there alternative interpretations of the data?			
	1.7	Conclusion: a definitive retroviral genealogy for similar			
		prima	tes	68	
2	Jumping genealogy			70	
	2.1	The activities of retroelements		73	
		2.1.1	LINE-1 elements	74	
		2.1.2	Alu elements	77	
		2.1.3	SVA elements	78	

v

### vi contents

	2.2	Retroelements and human disease		78	
	2.3	Retroelements and primate evolution		84	
		2.3.1	LINE-1 elements	84	
		2.3.2	Alu elements	88	
		2.3.3	Retroelements and phylogeny: validation	97	
	2.4	More ancient elements and mammalian evolution		101	
		2.4.1	Euarchontoglires: the primate-rodent group	103	
		2.4.2	Boreoeutheria: incorporating the		
			primate-rodent group and the Laurasian beasts	105	
		2.4.3	Eutheria	107	
		2.4.4	Mammals	111	
		2.4.5	TE stories on other branches of the tree of life	114	
	2.5	Exapta	ation of TEs	116	
		2.5.1	Raw material for new genes	117	
		2.5.2	Raw material for new exons	118	
		2.5.3	Raw material for new regulatory modules	120	
	2.6	The e	The evolutionary significance of TEs		
		2.6.1	TEs, genomic reorganisation and speciation	124	
		2.6.2	TEs and evolvability	128	
3	Pse	eudogenealogy			
	3.1	Mutations and the monoclonal origins of cancers			
	3.2	Old so	cars on DNA	138	
		3.2.1	Classical marks of NHEJ	139	
		3.2.2	LINEs and Alus	141	
		3.2.3	NUMTs	142	
		3.2.4	Interstitial telomeric sequences	145	
	3.3	_		148	
		3.3.1	Human-specific pseudogenes	152	
		3.3.2	Ape-specific pseudogenes	157	
		3.3.3	Simian-specific pseudogenes	163	
		3.3.4	Pseudogenes and sensory perception	172	
		3.3.5	Pseudogenes from further afield	180	
	3.4	Processed pseudogenes		183	

CAMBRIDGE

### contents vii

	3.5	5 Rare mutations that conserve protein-coding function		
	3.6	Conclusions	189	
4	The origins of new genes			
	4.1	New genes in cancer		
	4.2	Copy number variants		
	4.3	.3 Segmental duplications		
		4.3.1 Some early pointers	201	
		4.3.2 Systematic studies of SDs	203	
	4.4	New genes		
		4.4.1 Reproduction	207	
		4.4.2 Hydrolytic enzymes	218	
		4.4.3 Neural systems	220	
		4.4.4 Blood	224	
		4.4.5 Immunity	228	
		4.4.6 Master regulators of the genome	236	
	4.5	Retrogenealogy	238	
		4.5.1 Reverse-transcribed genes in primates	239	
		4.5.2 Reverse-transcribed genes in mammals	246	
	4.6	DNA transposons		
	4.7	De novo origins of genes		
	4.8	Generating genes and genealogies		
	Epilogue: what really makes us human			
	1	Immune systems		
	2	2 Nervous systems		
		2.1 Critical periods	272	
		2.2 Learning from neglect	273	
	3	Features of personhood		
	4	Stories and narrative identity		
	Refe	erences	284	
	Inde	ex	351	

Cambridge University Press 978-1-107-04012-0 - Human Evolution: Genes, Genealogies and Phylogenies Graeme Finlay Frontmatter More information

### Preface

Histories are subject to different interpretations. We would expect biological history to conform to this variety of understandings. But the strange thing is that the very existence of biological history is denied in some quarters. This field of science has acquired a 'more than scientific' aura to it. People argue about it as if it were an ideology. Vast resources, including a lot of goodwill, have been expended in the debate. To have achieved this notoriety, we must conclude that biological history (or evolutionary biology) is widely misunderstood. But the evidence for it is there; and a vast volume of fresh genetic data has been added recently. Such data are compelling.

This is a history book, and for two reasons. It attempts to describe, in a very limited and situated sense, a spectacular period in the history of science. Its timeframe covers, with somewhat fuzzy edges, the first decade of the twenty-first century. This is the period during which the human genome sequencing project has been elaborated to ever increasing degrees of detail, and during which myriad fascinating insights into the biological basis of our humanness have been revealed.

Secondly, it describes the evolutionary history of our species, as inscribed in great detail in our genomes. The DNA that we carry around as part of our bodies is an extraordinary library of genetic information. But it is more than simply a blueprint for the human body plan; it also carries, inscribed in its base sequence, a record of its own formative history. Multiple other mammal and vertebrate genomes have also been sequenced over the last decade or so, and this means that we have access to their histories too. When our genomic history is laid out, side by side with those of other species, particular discrete changes in the historical records can be identified

ix

#### X PREFACE

in our genome and in the genomes of cohorts of other species. We can thus infer, unambiguously and with a great deal of confidence, that most of our genetic history has been shared with the genetic histories of other primates and, more inclusively, other mammals. Our evolutionary history is well documented.

Molecular evolution is at least as old as the work of Alan Wilson, who used molecular data to infer evolutionary relationships between organisms as long ago as the 1960s. Phylogenetic analyses of DNA and protein sequences have also been used to generate evolutionary trees. Such approaches require expertise in statistics and computation, and require specialist treatments. However, the novel and intuitively appealing approaches surveyed in this book are based, in general, on the identification of particular complex mutations. These arise in unique events. When any such mutation is found in multiple species, it is only because it has been inherited from the one ancestor in which the mutation arose. These are thus very powerful signatures of phylogenetic relatedness.

Along the way, we find out many fascinating things about our biology. We discover that our genome is an entire ecosystem in which semi-autonomous units of genetic material play out their own life cycles. We discover why some people have violent allergic reactions to eating certain animal products. We find out why we must have vitamin C in our diets, whereas other organisms lack this requirement. We learn of the basis of our tendency to suffer from gout. We find clues as to why humans may be particularly cancer-prone. We discover how three-colour vision arose. Indeed many processes through which new genetic functionality has been generated have been laid bare.

Everything that is presented herein is in the public domain. Anything that I have not reported accurately, or that calls for further elaboration, can be fully checked against the source literature. To me, as a cell biologist, the wonder of our DNA-inscribed history is that it requires no logic other than that which is fundamental to all genetics. (Perhaps if I were a palaeontologist, the study of fossils Cambridge University Press 978-1-107-04012-0 - Human Evolution: Genes, Genealogies and Phylogenies Graeme Finlay Frontmatter <u>More information</u>

PREFACE Xi

would be just as intuitively compelling! But I am not a palaeontologist and I suspect that far fewer people are knowledgeable about fossils than are knowledgeable about the basic mechanisms of heredity.) I believe that the logic of this book will be widely available, although it will require a modicum of biological literacy.

I am very grateful to my superiors in the University of Auckland and the Auckland Cancer Society Research Laboratory, Professors Peter Browett and Bruce Baguley, for allowing me the space and time to work on this book. I thank many senior colleagues who have provided kind and helpful advice: Professor Bill Wilson and Associate Professor Philip Pattemore, Associate Professor Andrew Shelling, Professors Wilf Malcolm, Richard Faull, Malcolm Jeeves and John McClure. Theological input has come from the late Dr Harold Turner, as well as Dr Bruce Nicholls and Dr Nicola Hoggard-Creegan. I am hugely indebted to personnel at the Faraday Institute for Science and Religion, St Edmunds College, University of Cambridge, including Dr Denis Alexander, for sharing their erudition and for their encouragement.

I am deeply grateful to the editorial staff at Cambridge University Press and Out of House Publishing for their unvarying courtesy, patience and helpfulness. It has been a pleasure to work with and learn from them.

I am also grateful to those who have given me scope to work out ideas and evolve ways of expressing them. In particular, I thank the editors of the Paternoster Press periodical *Science and Christian Belief*, and the multi-author book *Debating Darwin: Is Darwinism True & Does it Matter*? (2009). They have allowed me to explore, and reflect upon, earlier phases of an explosively expanding scientific field.