

Clonally transmissible cancers in dogs and Tasmanian devils



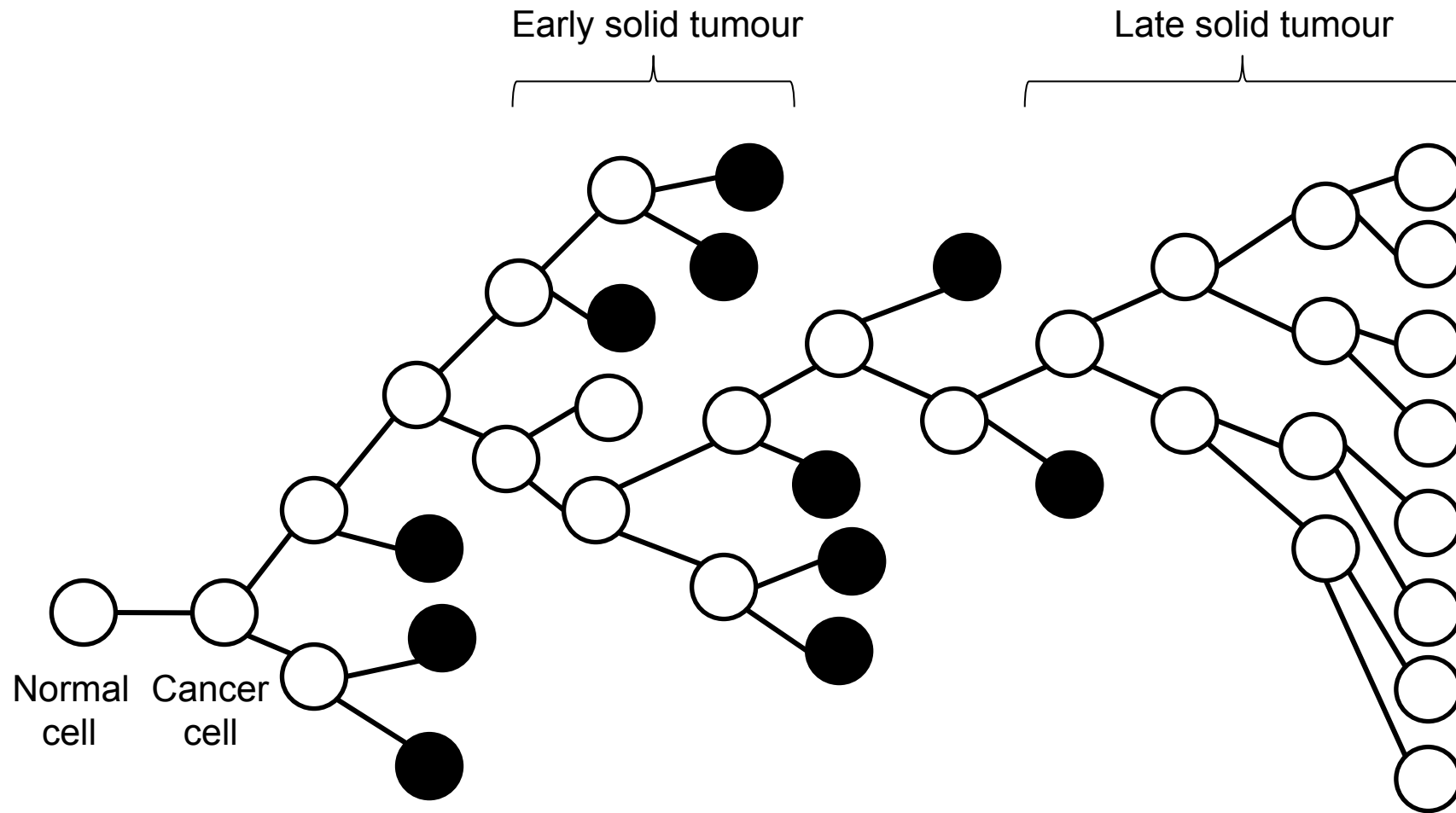
Photo: The Broad Institute



Photo: Christo Baars

Elizabeth Murchison, Wellcome Trust Sanger Institute, Cambridge, UK

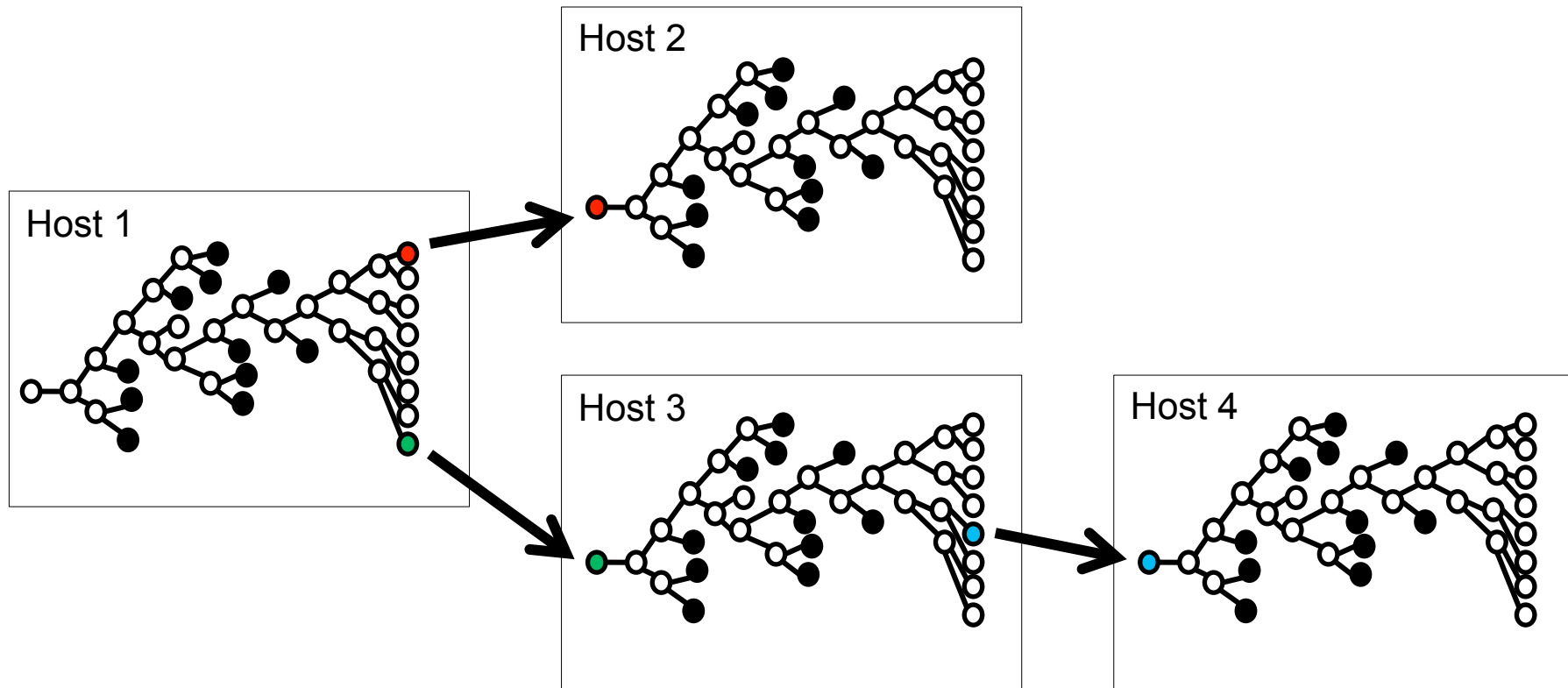
Clonal evolution of cancer



Adapted from Nowell, *Science*, 194 (4260):23-26, 1976

Clonally transmissible cancer

- Tasmanian devil facial tumour disease (DFTD)
- Canine transmissible venereal tumour (CTVT)



Transmissible cancers in devils and dogs

To use genomics to understand

- Origin
- Diversity
- History
- Evolution



Photo: Tasmanian Devil Conservation Park



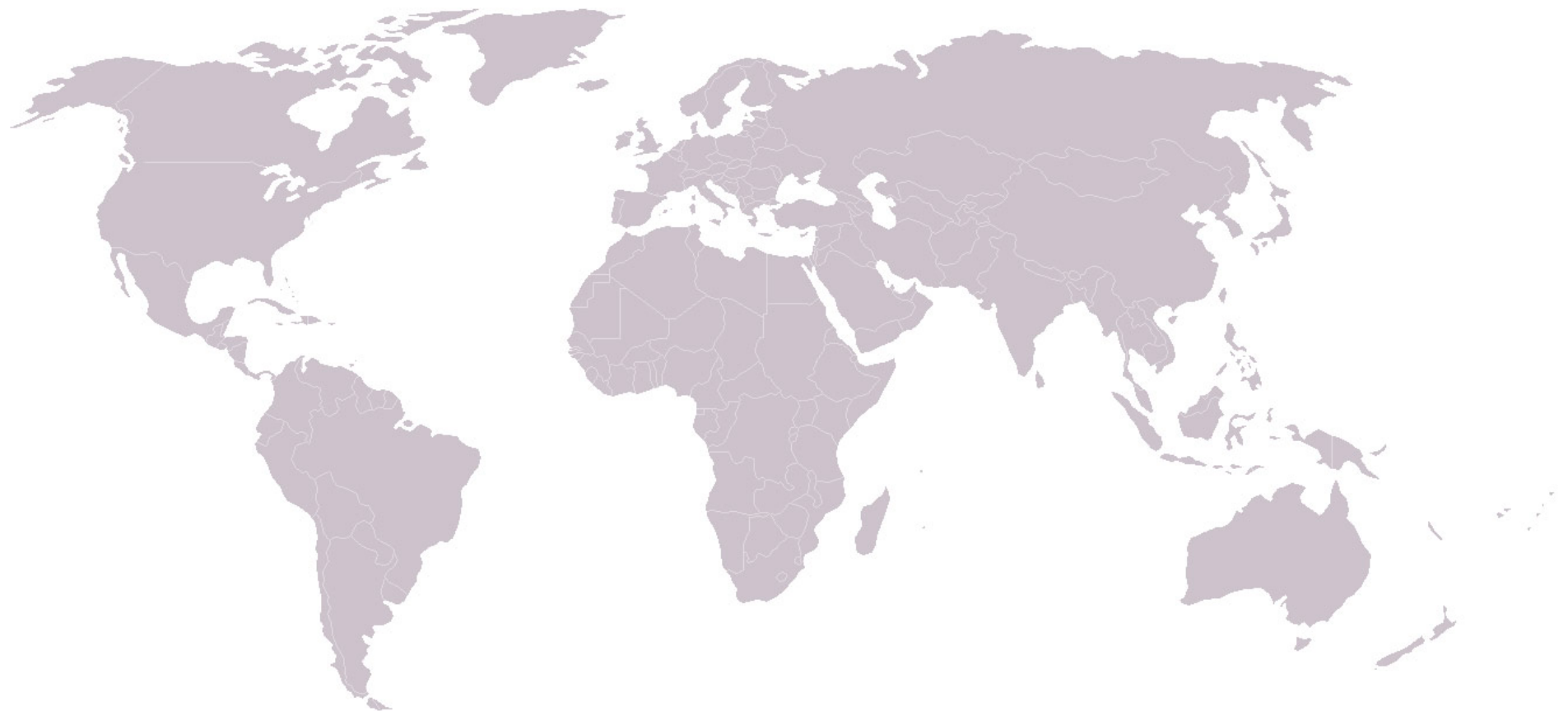
Photo: www.animalswallpaper.info

Tasmanian devil

Sarcophilus harrisii



Photo: Wayne McLean





Nick Mooney





Photo: Rodrigo Hamede



Photo: Tasmanian Department of Primary Industries, Parks, Water and the Environment



Photo: Christo Baars

Photo: Rodrigo Hamede

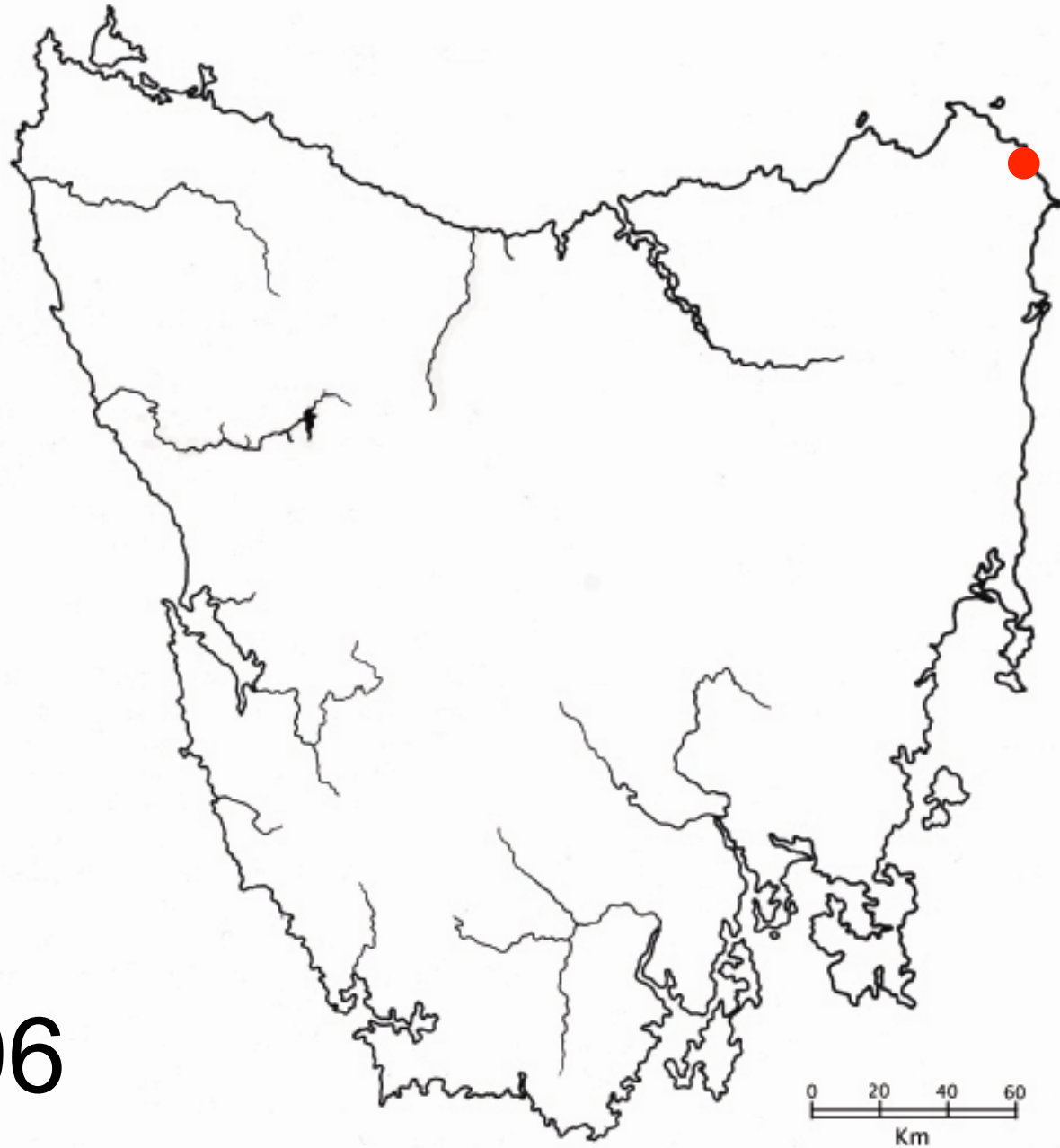






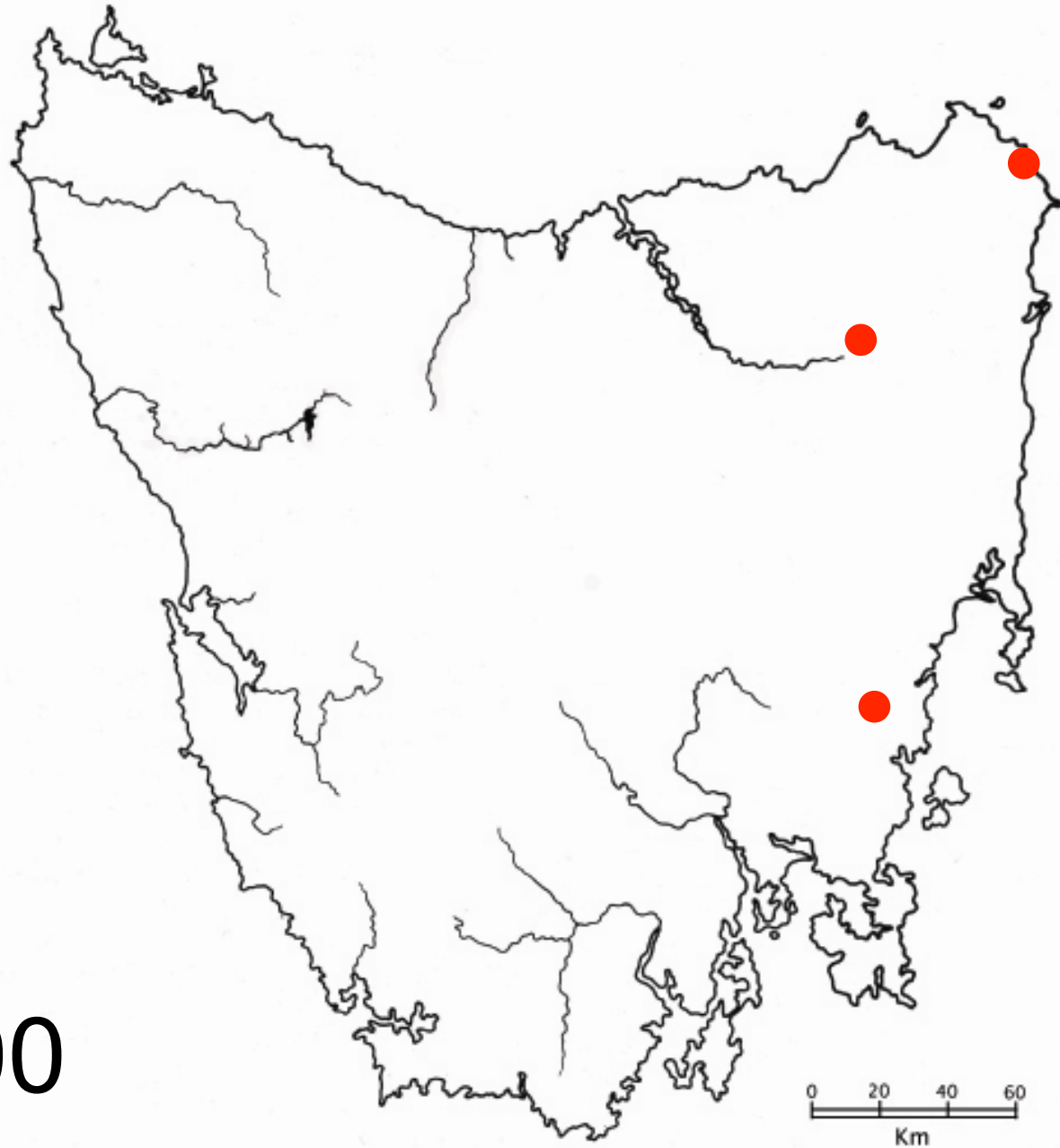
Photo: Christo Baars, 1996

1996



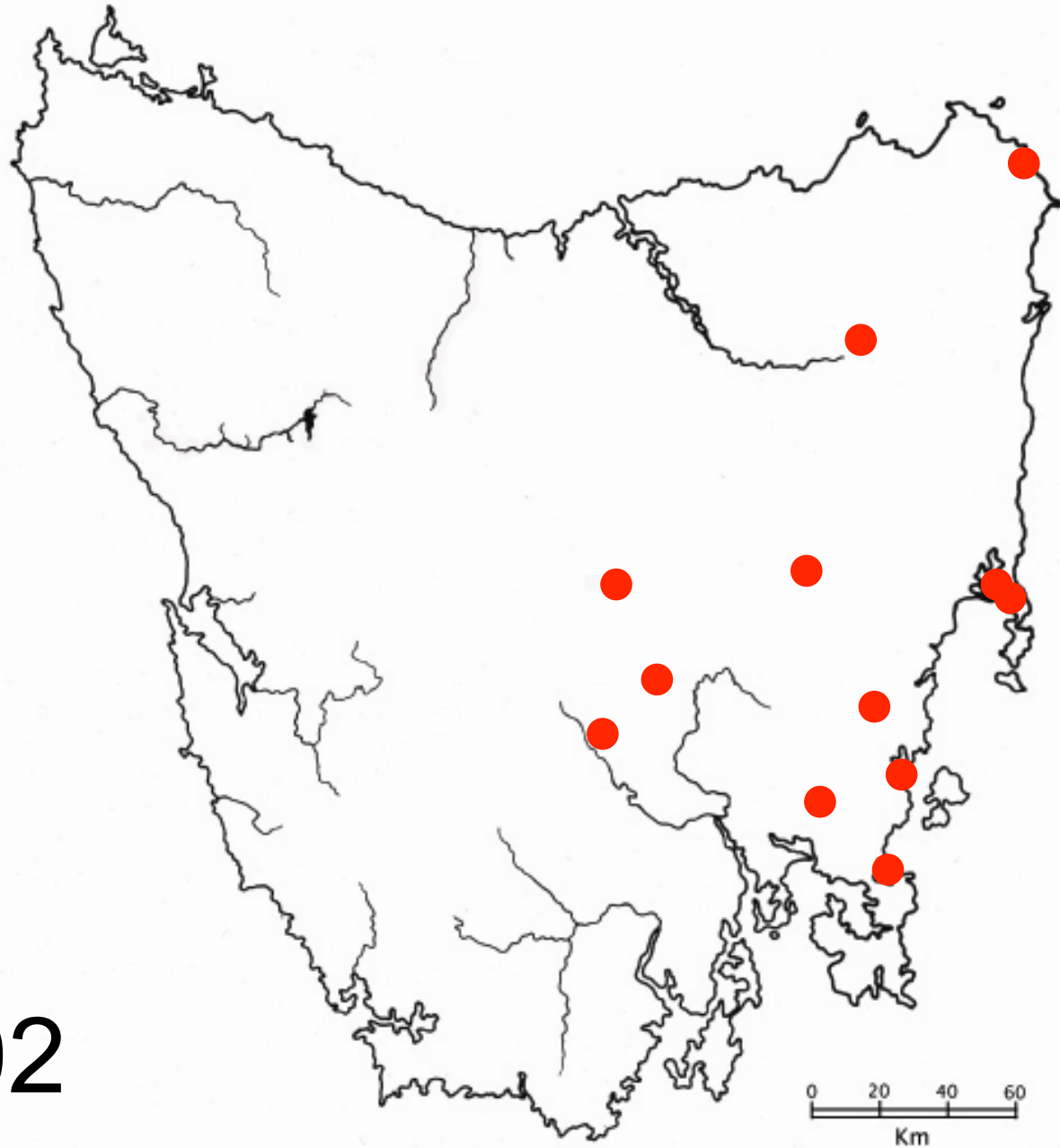
Adapted from Hawkins et al, *Biol Conserv* 131:307-24, 2006

2000



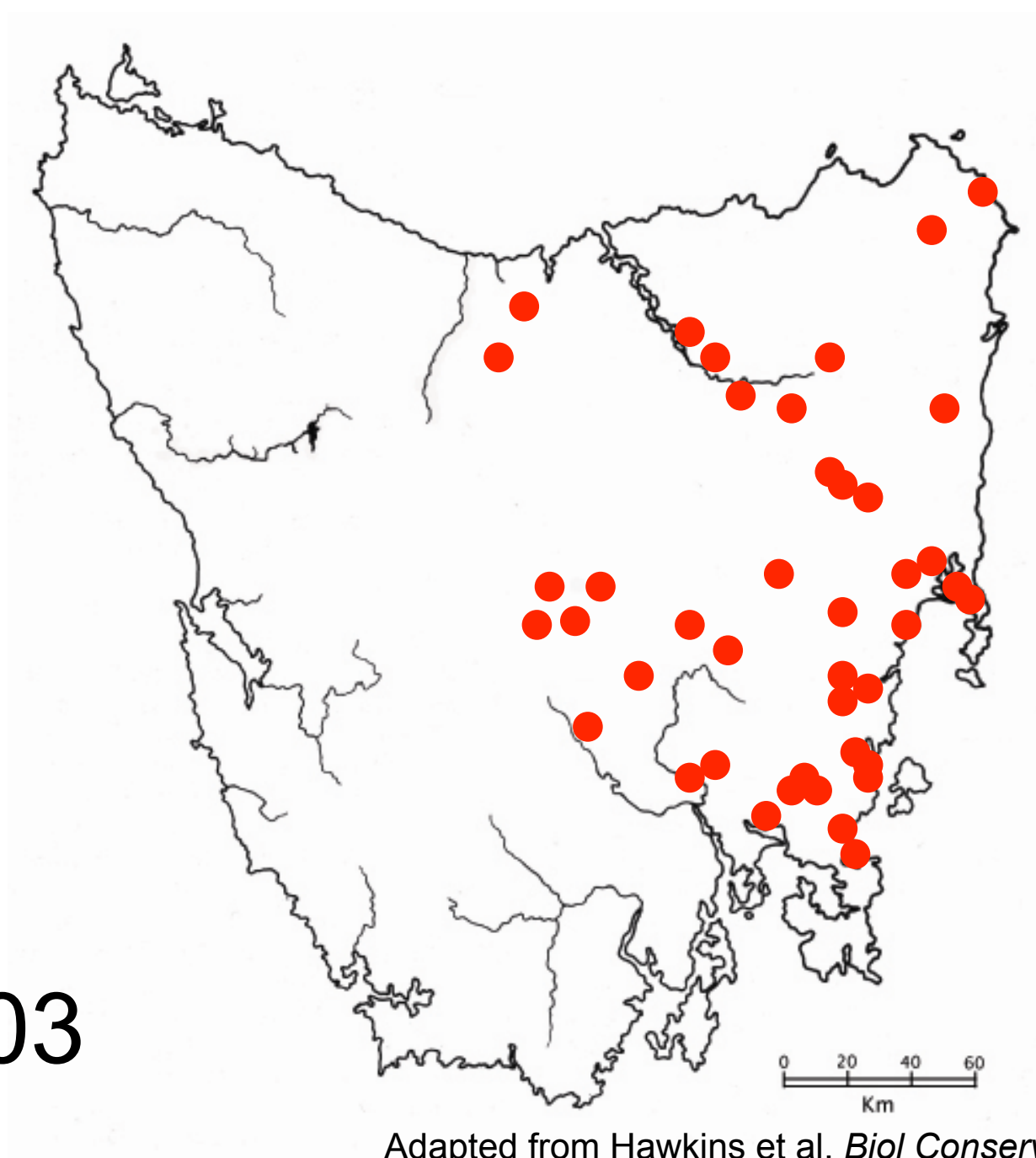
Adapted from Hawkins et al, *Biol Conserv* 131:307-24, 2006

2002



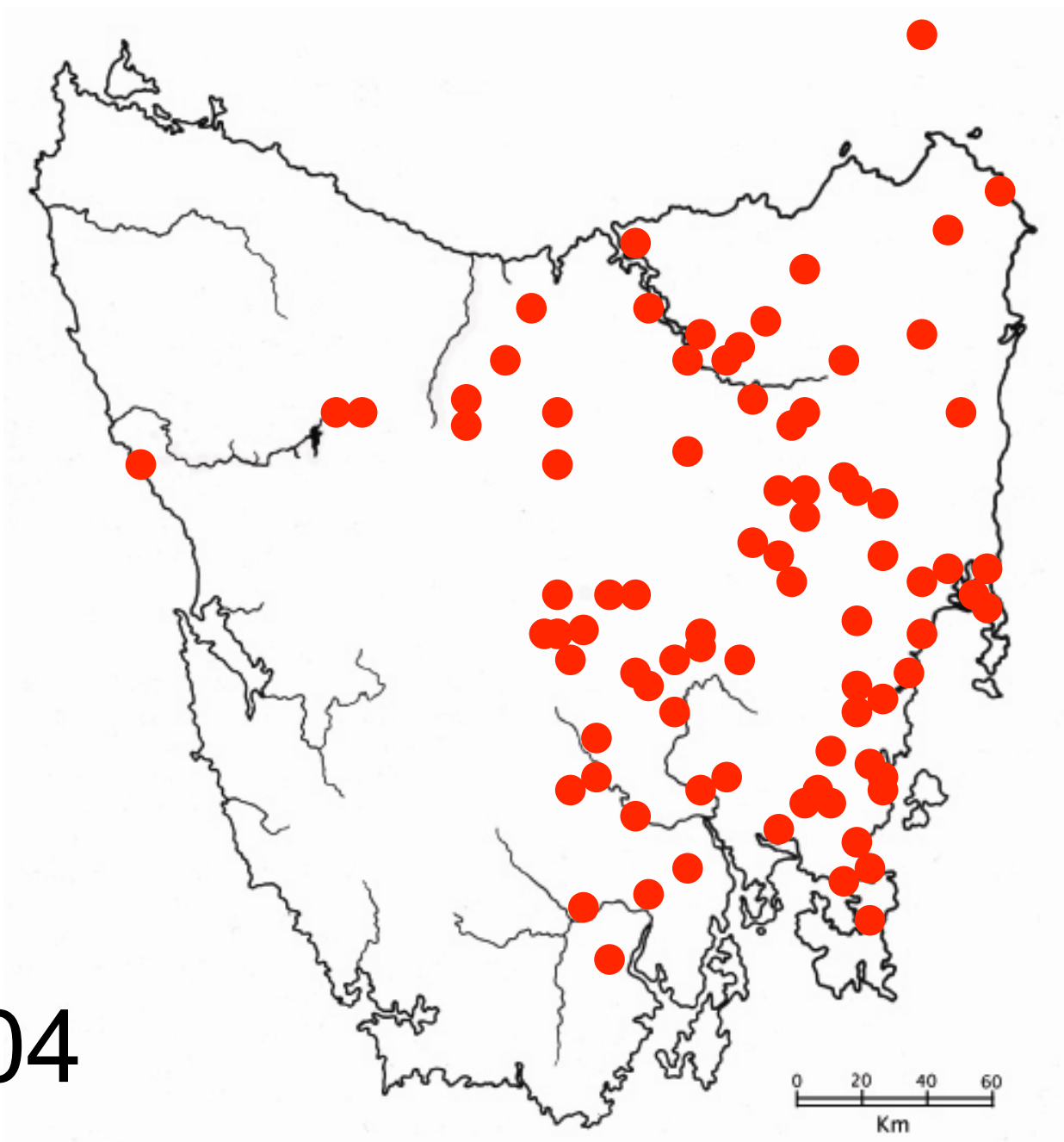
Adapted from Hawkins et al, *Biol Conserv* 131:307-24, 2006

2003



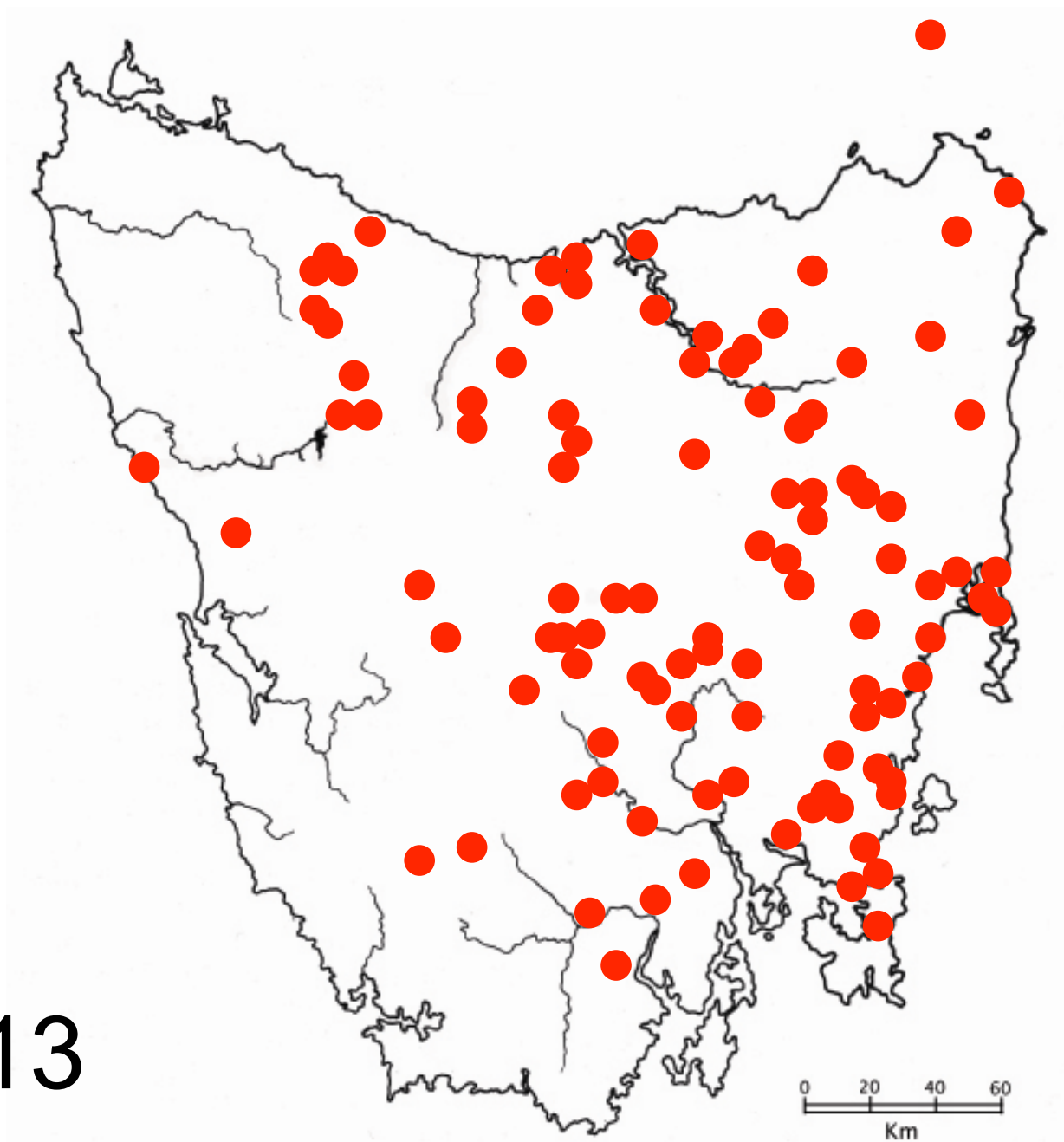
Adapted from Hawkins et al, *Biol Conserv* 131:307-24, 2006

2004



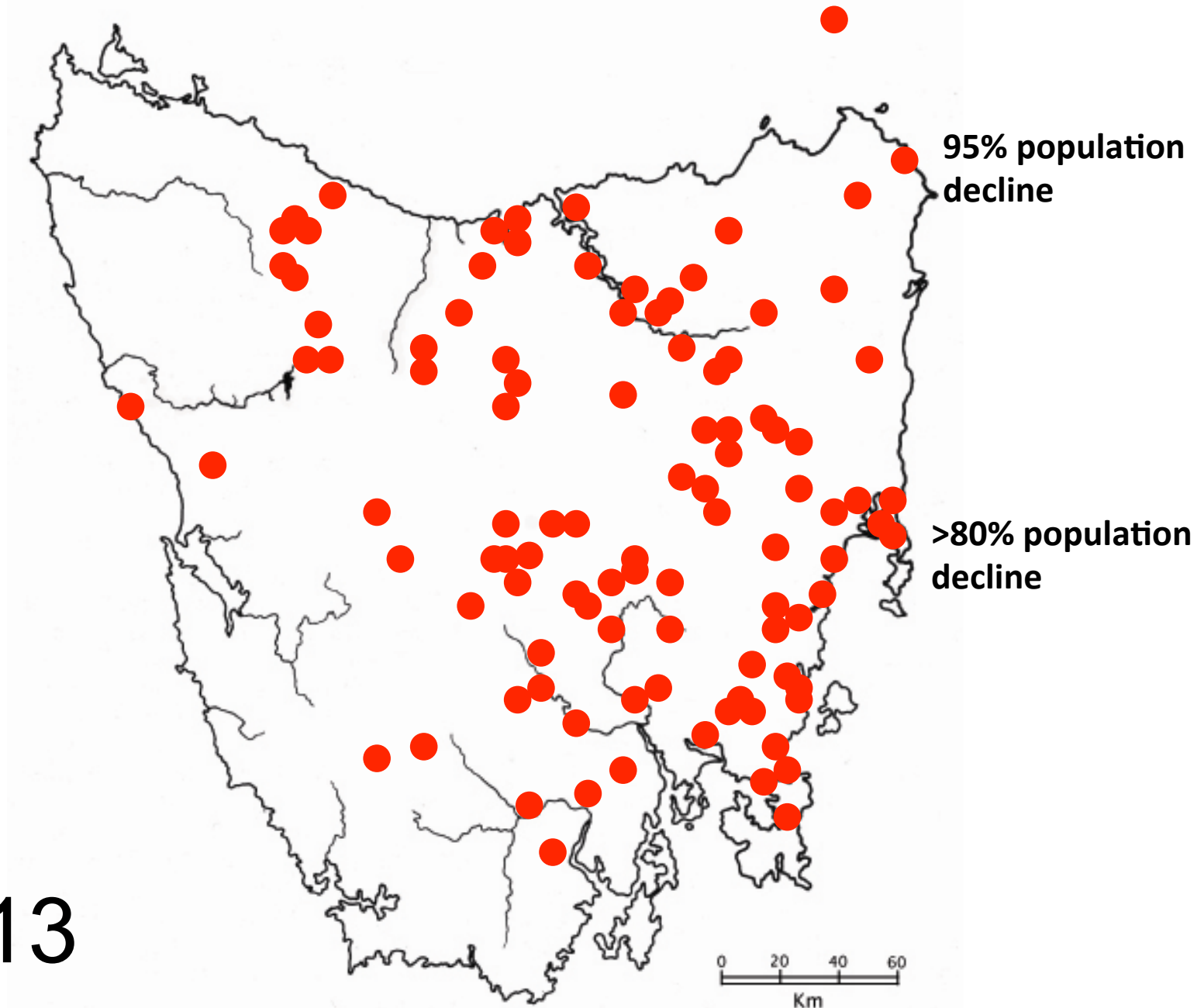
Adapted from Hawkins et al, *Biol Conserv* 131:307-24, 2006

2013



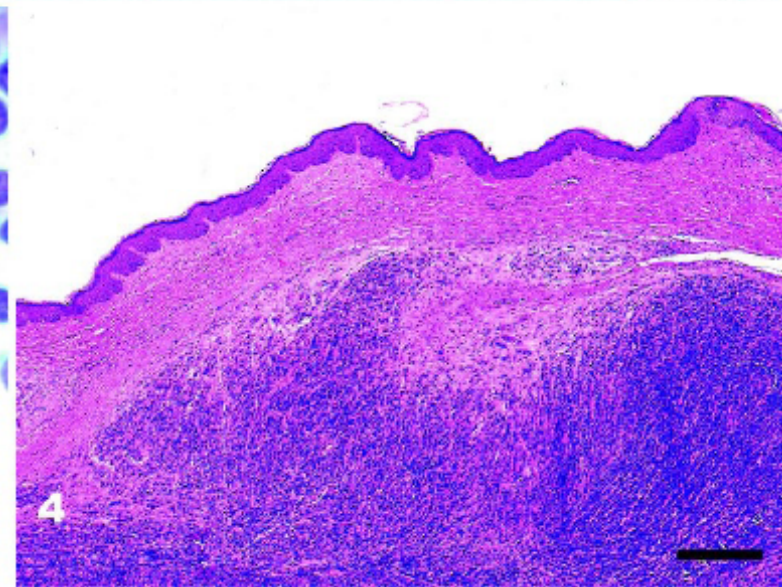
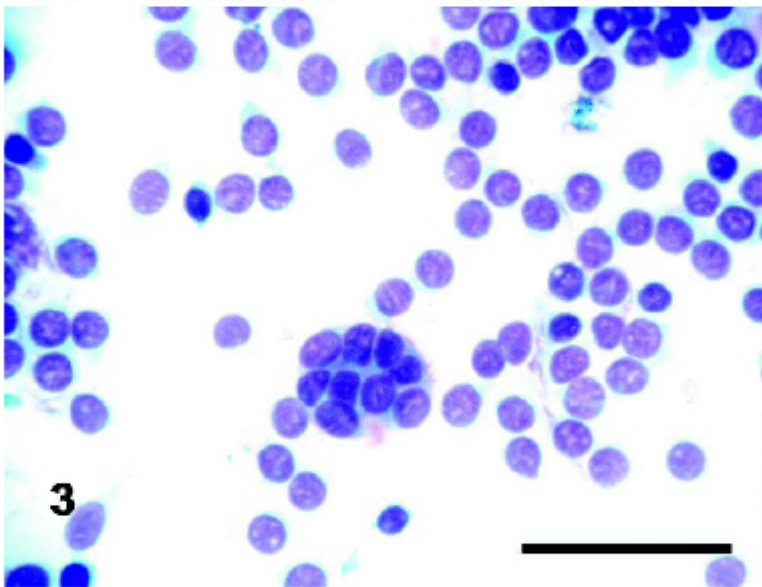
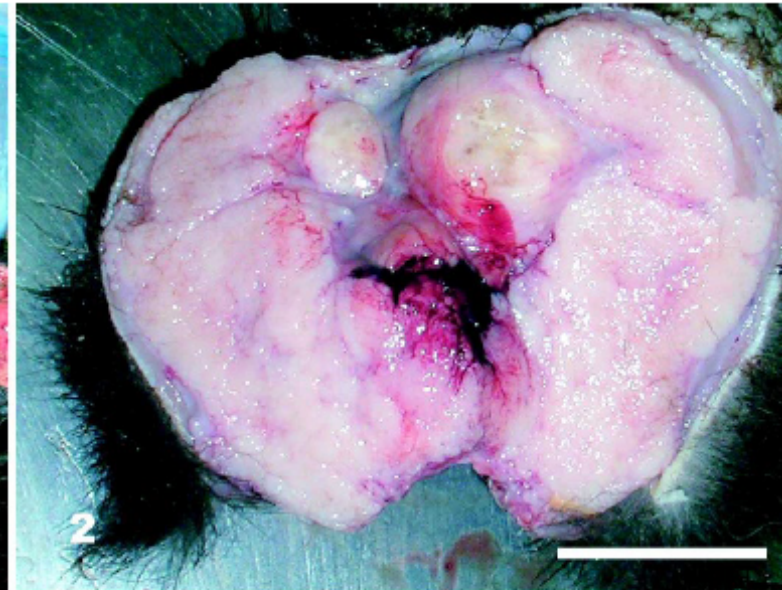
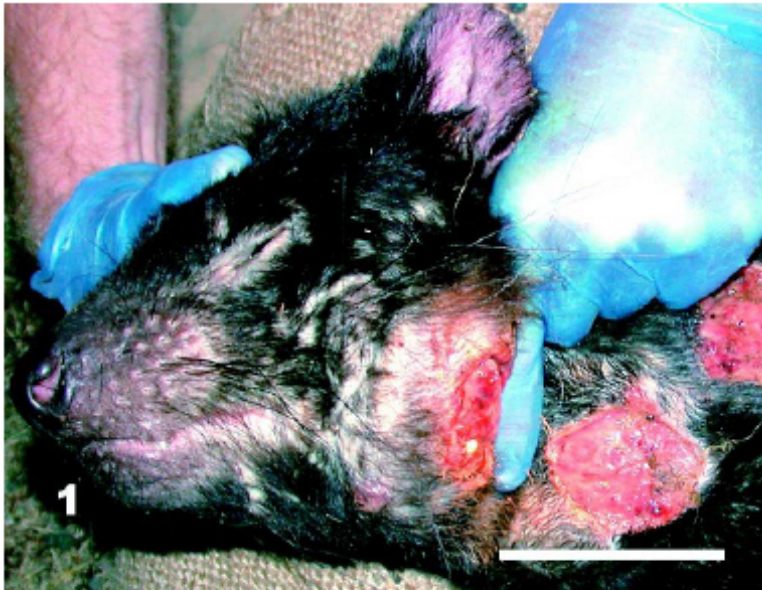
Adapted from Hawkins et al, *Biol Conserv* 131:307-24, 2006

2013

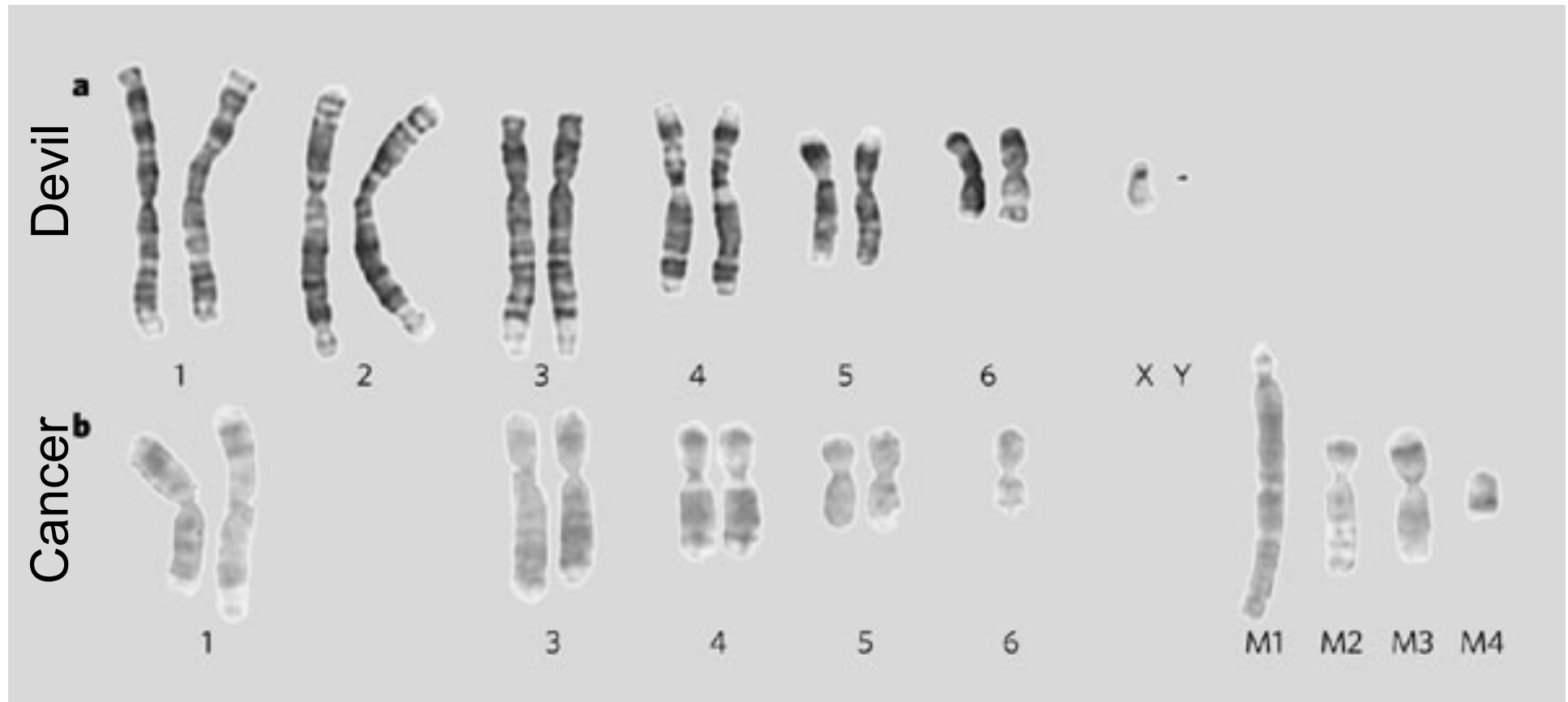


Adapted from Hawkins et al, *Biol Conserv* 131:307-24, 2006

“Tasmanian Devil Facial Tumour Disease” (DFTD)

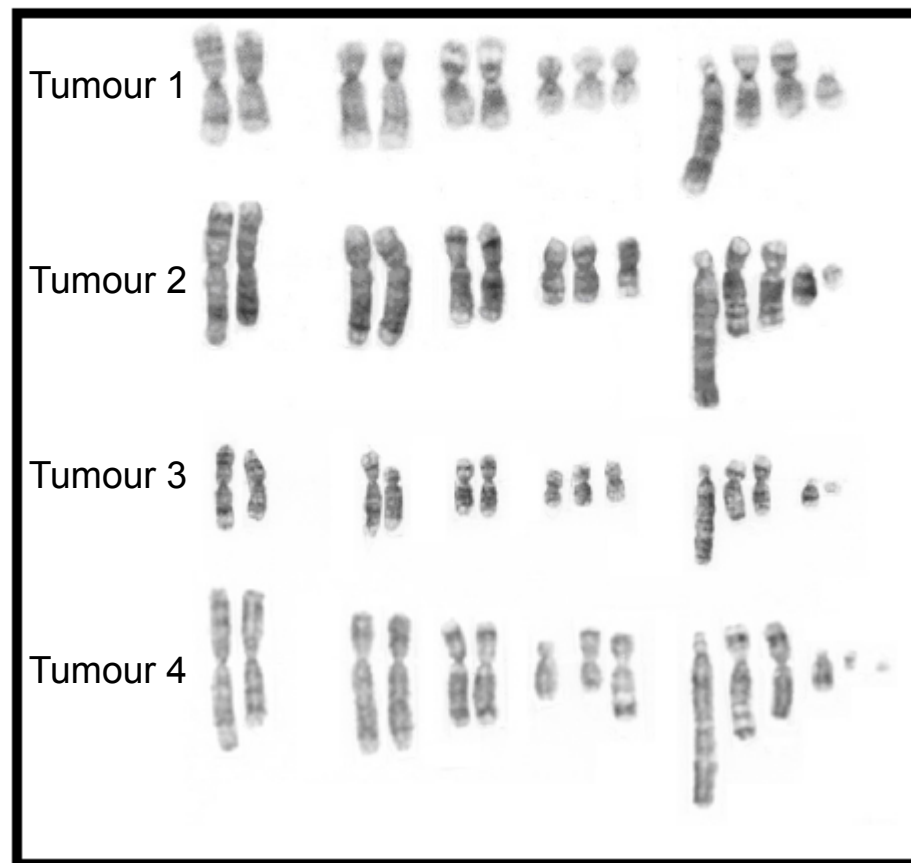
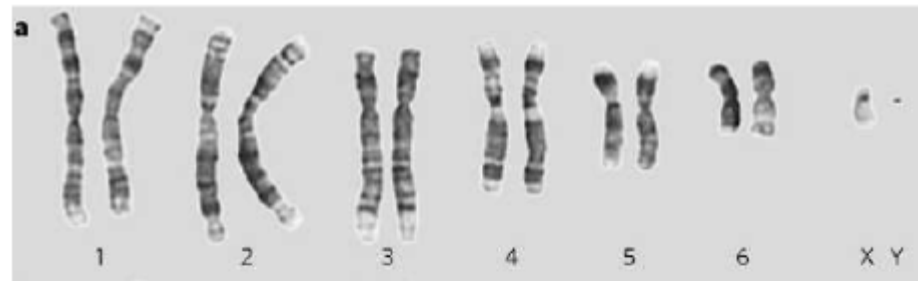


Devil cancer chromosomes



Pearse and Swift, *Nature*, 439 (7076):549, 2006

Normal devil



Anne-Maree Pearse, Unpublished



Photo: Christo Baars

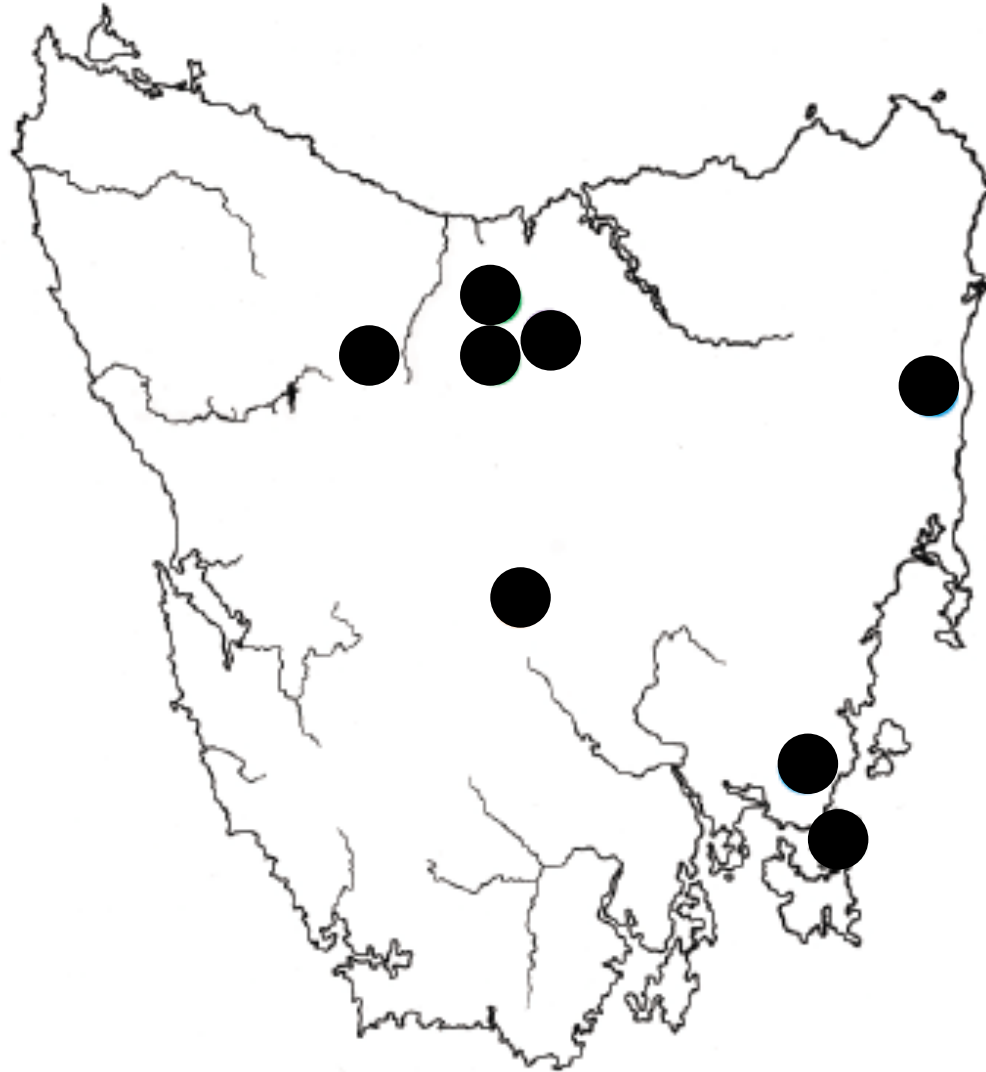
Predictions....

- If the devil's cancer is spread by living cancer cells then
 - All the cancers should be genetically identical
 - The cancers should be genetically different to their hosts

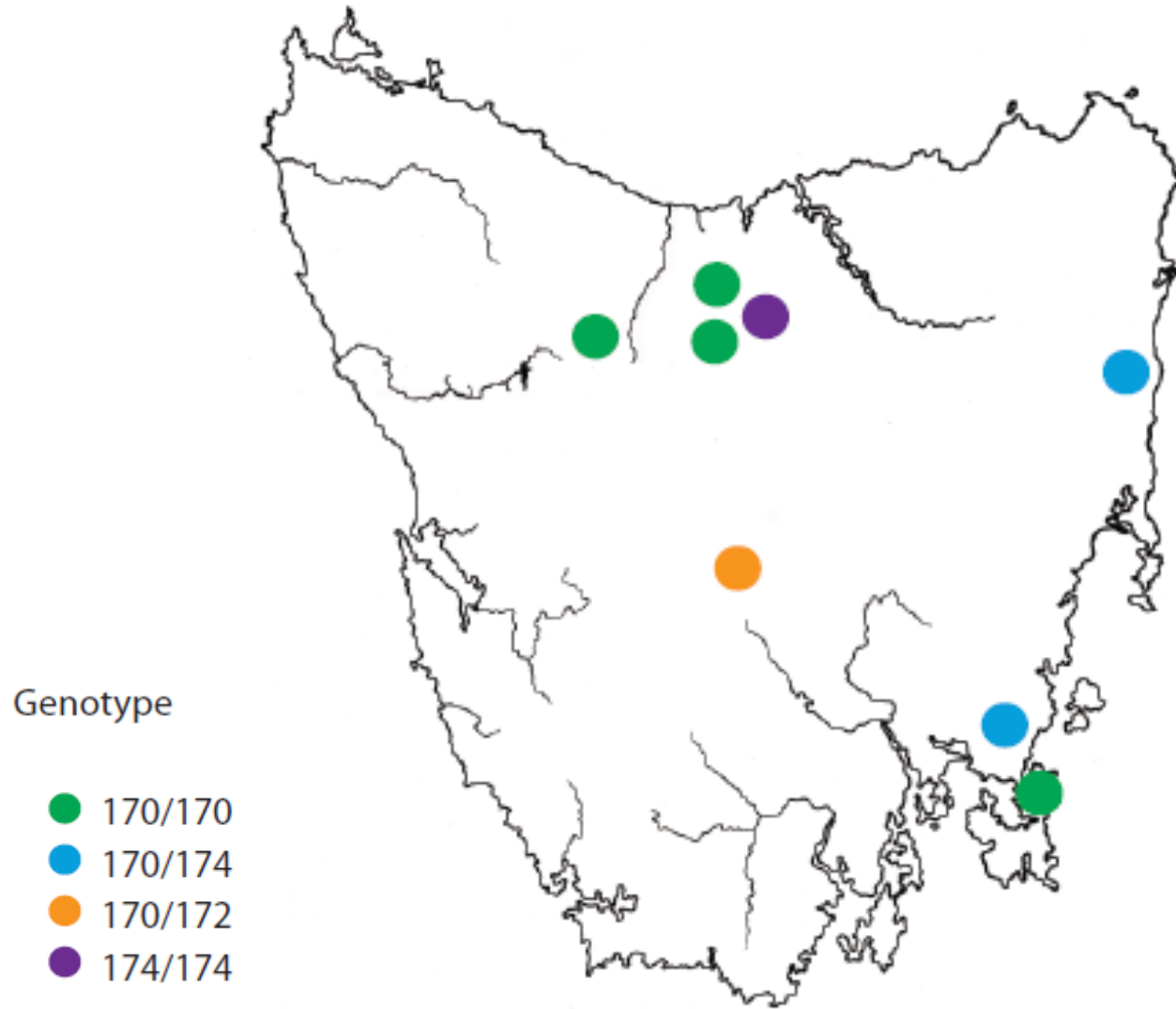


Photo: Hannah Bender

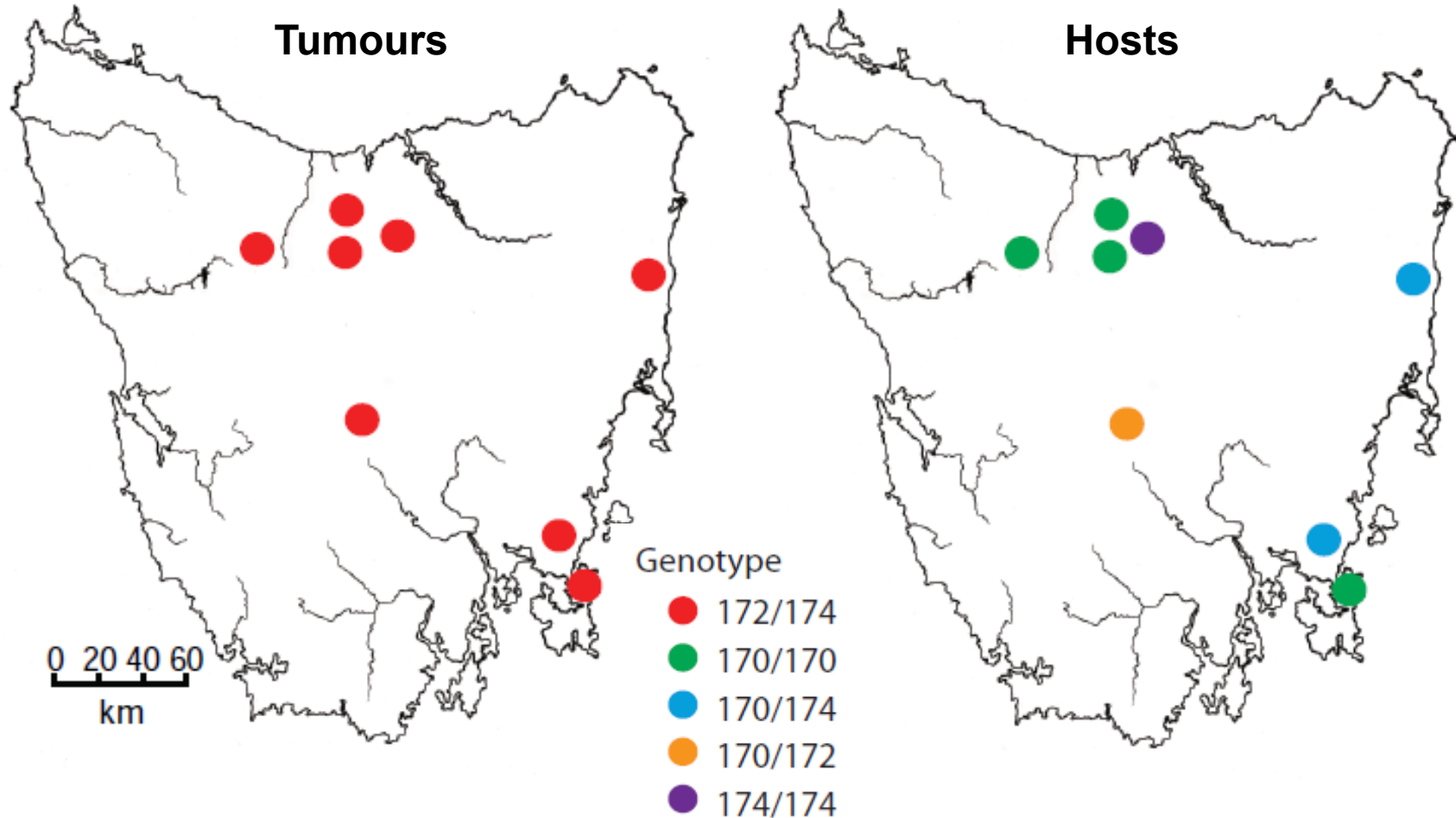
Microsatellite genotyping



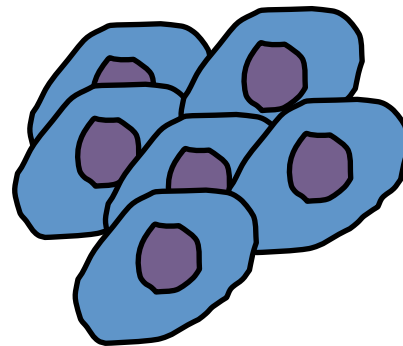
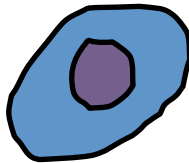
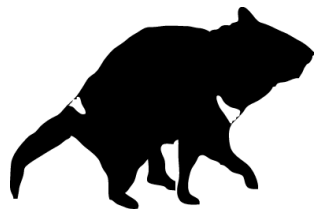
Microsatellite genotyping



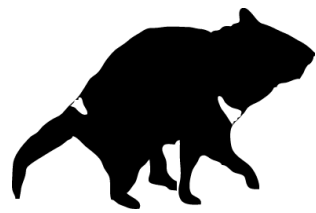
Devil tumours are genetically identical



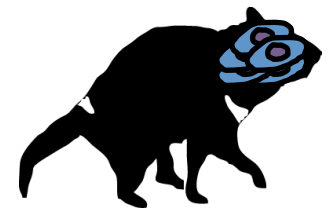
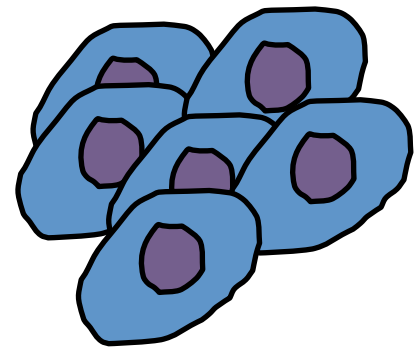
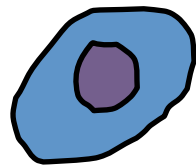
Murchison et al, *Science*, 327:84-7, 2010
Siddle et al, *PNAS*, 104 (41):16221-16226, 2007



DFTD founder devil

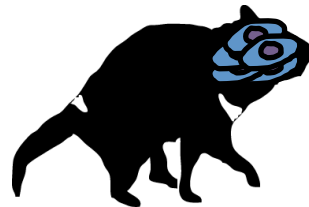
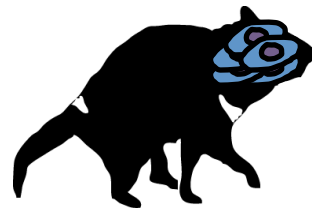
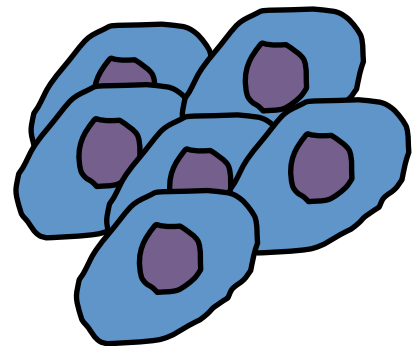
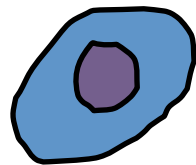


DFTD founder devil



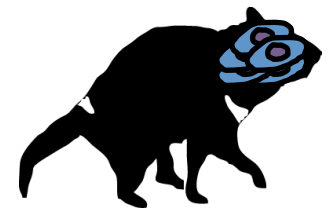
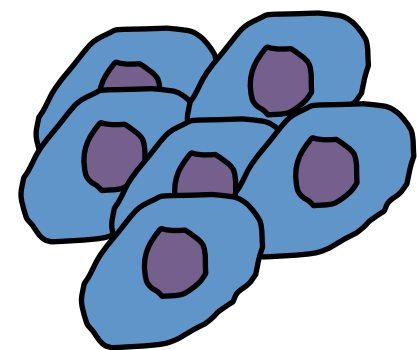
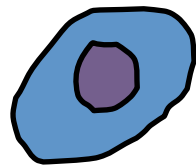


DFTD founder devil





DFTD founder devil



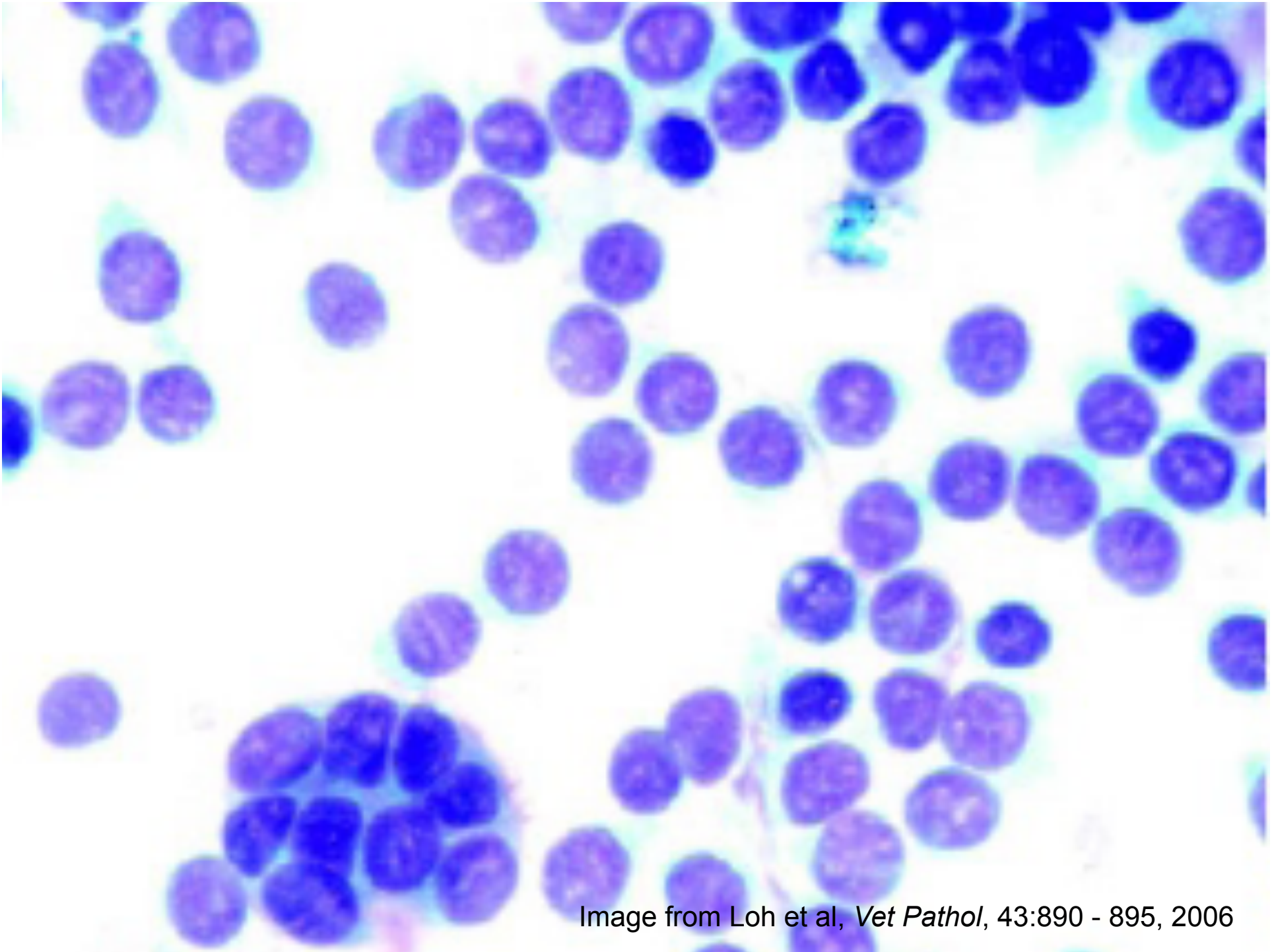


Image from Loh et al, *Vet Pathol*, 43:890 - 895, 2006

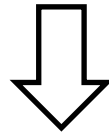


Tasmanian devil reference genome

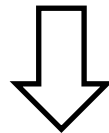
Photo: Taronga Zoo



“Salem”: Female Tasmanian devil



Illumina paired end sequencing



De novo assembly

Tasmanian devil (DEVIL7.0)

About this species

- Description
- Genome Statistics
 - Assembly and Genebuild
 - Top 40 InterPro hits
 - Top 500 InterPro hits
- What's New
- Sample entry points
 - Karyotype (not available)
 - Location (GL193649.1)
 - Gene (HSP1_SARHA)
 - Transcript (HSP1_SARHA)

- Configure this page
- Manage your data
- Export data
- Bookmark this page

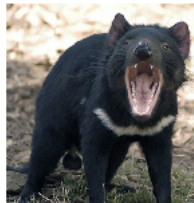
Search Ensembl Tasmanian devil

Search for:
e.g. HSP1_SARHA or GL193649.1 or keratin

Description

Tasmanian devil (*Sarcophilus harrisi*)

Assembly



This site displays the DEVIL7.0 assembly (GCA_000219685.1) of the Tasmanian devil (*Sarcophilus harrisi*) genome. The genome sequence and assembly are provided by the [Wellcome Trust Sanger Institute](#) and [Illumina](#). The N50 size is the length such that 50% of the assembled genome lies in blocks of the N50 size or longer. The N50 length for supercontigs is 1847.19 kb and is 20.13 kb for contigs. The total number of bases in supercontigs is 3.17 Gb and in contigs is 2.93 Gb.



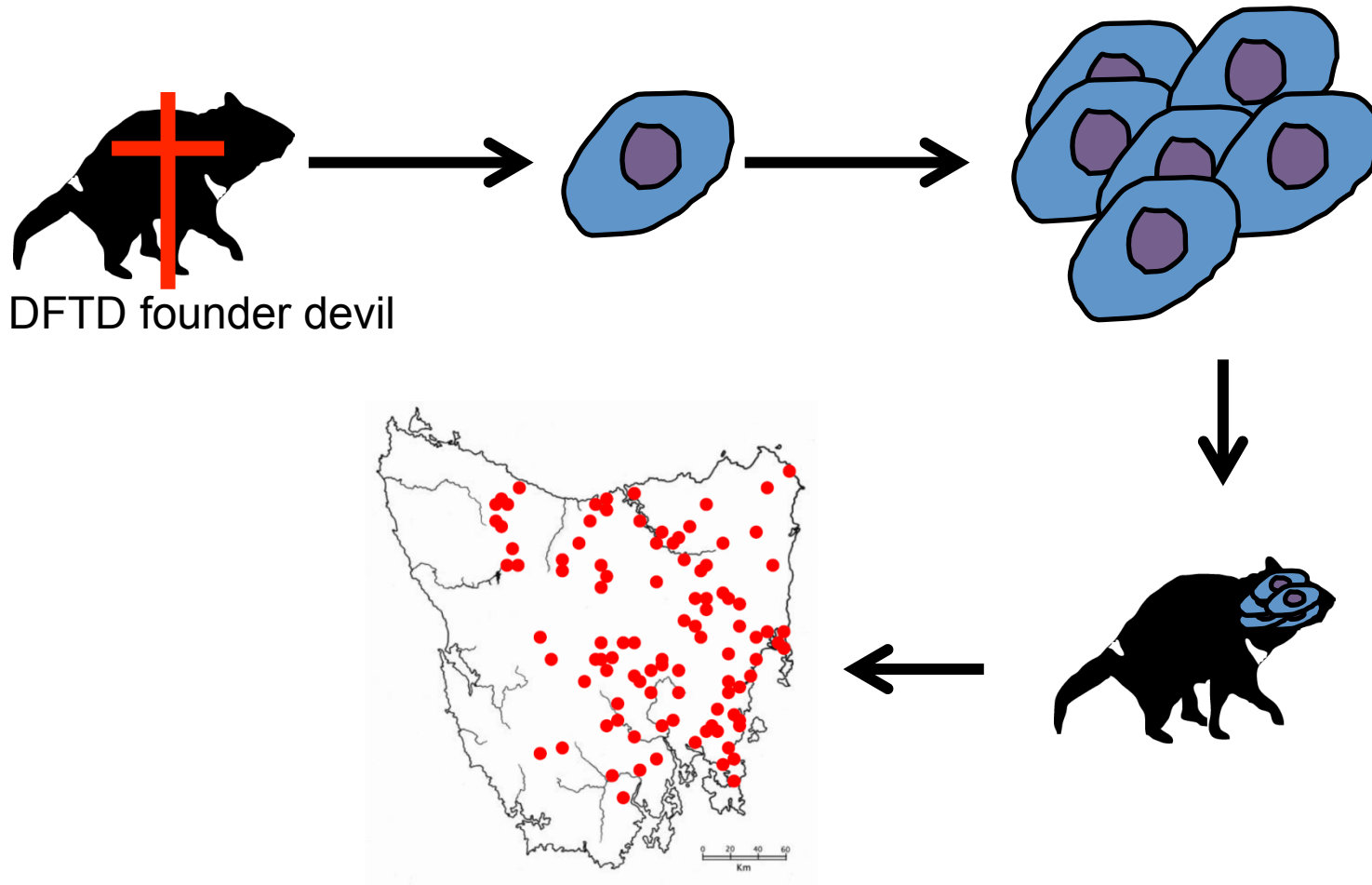
[Download Tasmanian devil genome sequence](#) (FASTA)

Annotation

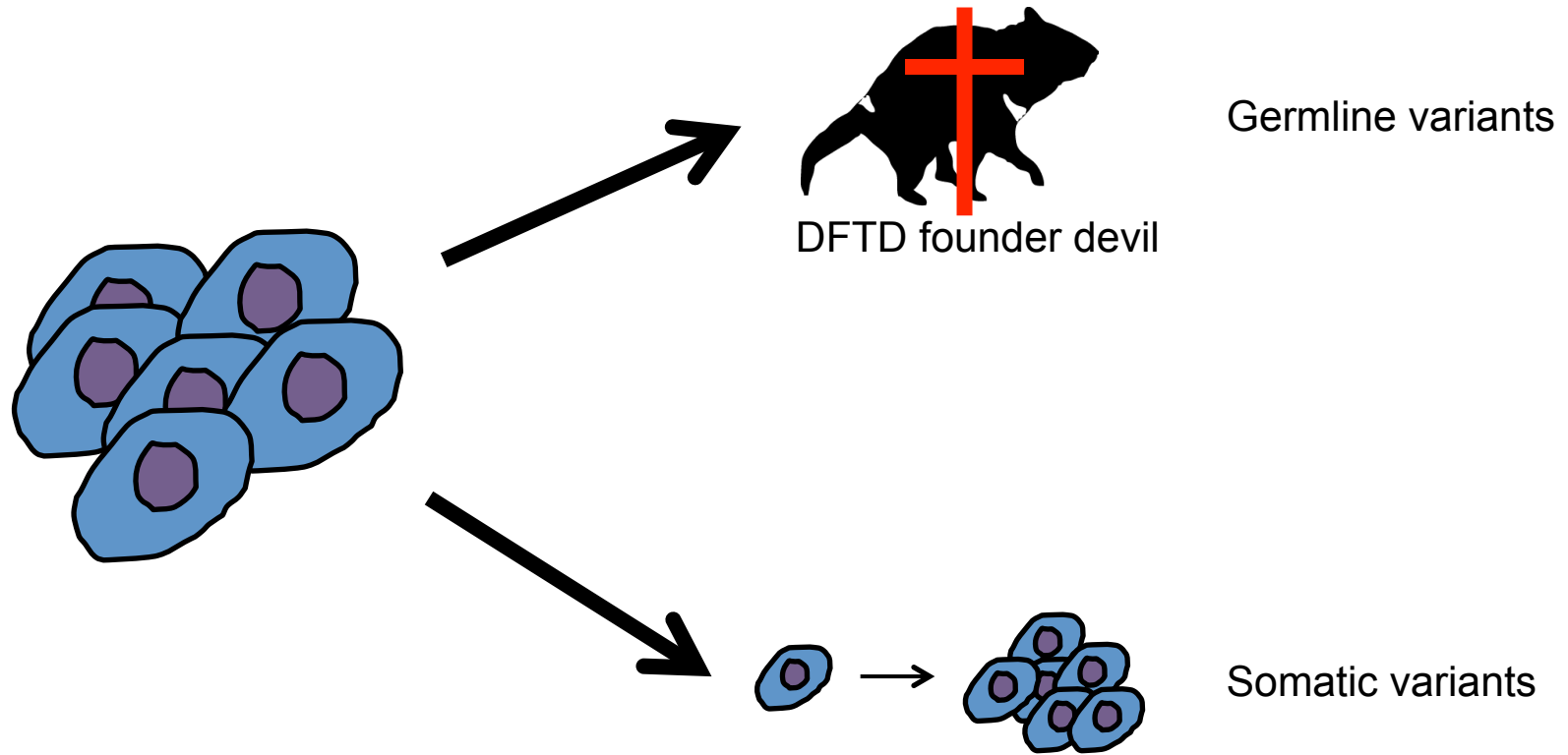
The Ensembl genome annotation pipeline was used to identify genes. Models built from Tasmanian devil proteins and cDNAs have been given priority over predictions from other vertebrate species. 5,663 transcript models made from paired end Illumina RNA-Seq were added into the gene set where they added a novel model or splice variant. RNA-Seq data was also used to add UTR to non species specific models. The total gene set contains 20,419 genes including pseudogenes and ncRNAs.

- [Detailed information on genebuild](#) (PDF)

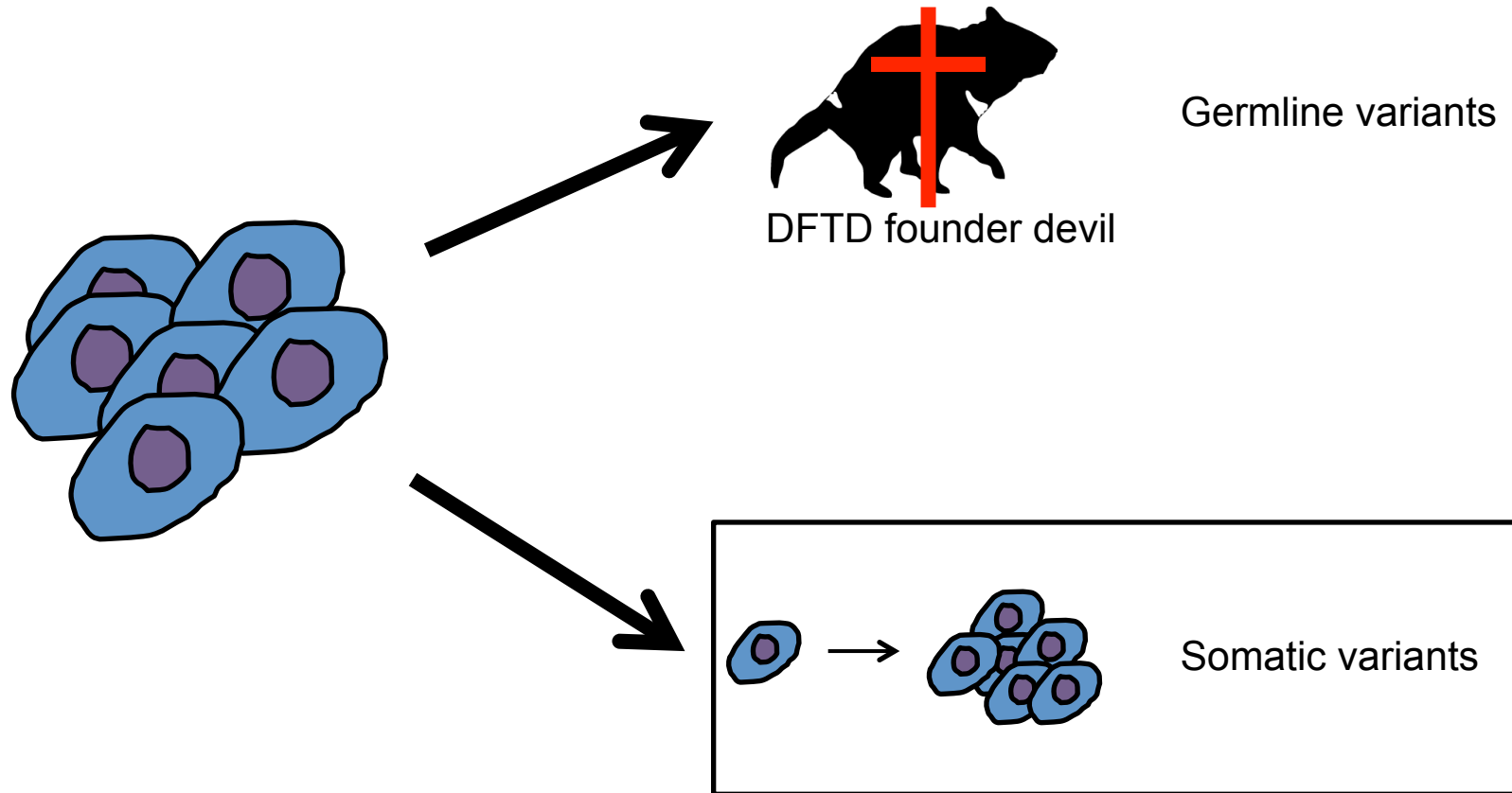
Sequencing the DFTD genome



The DFTD genome



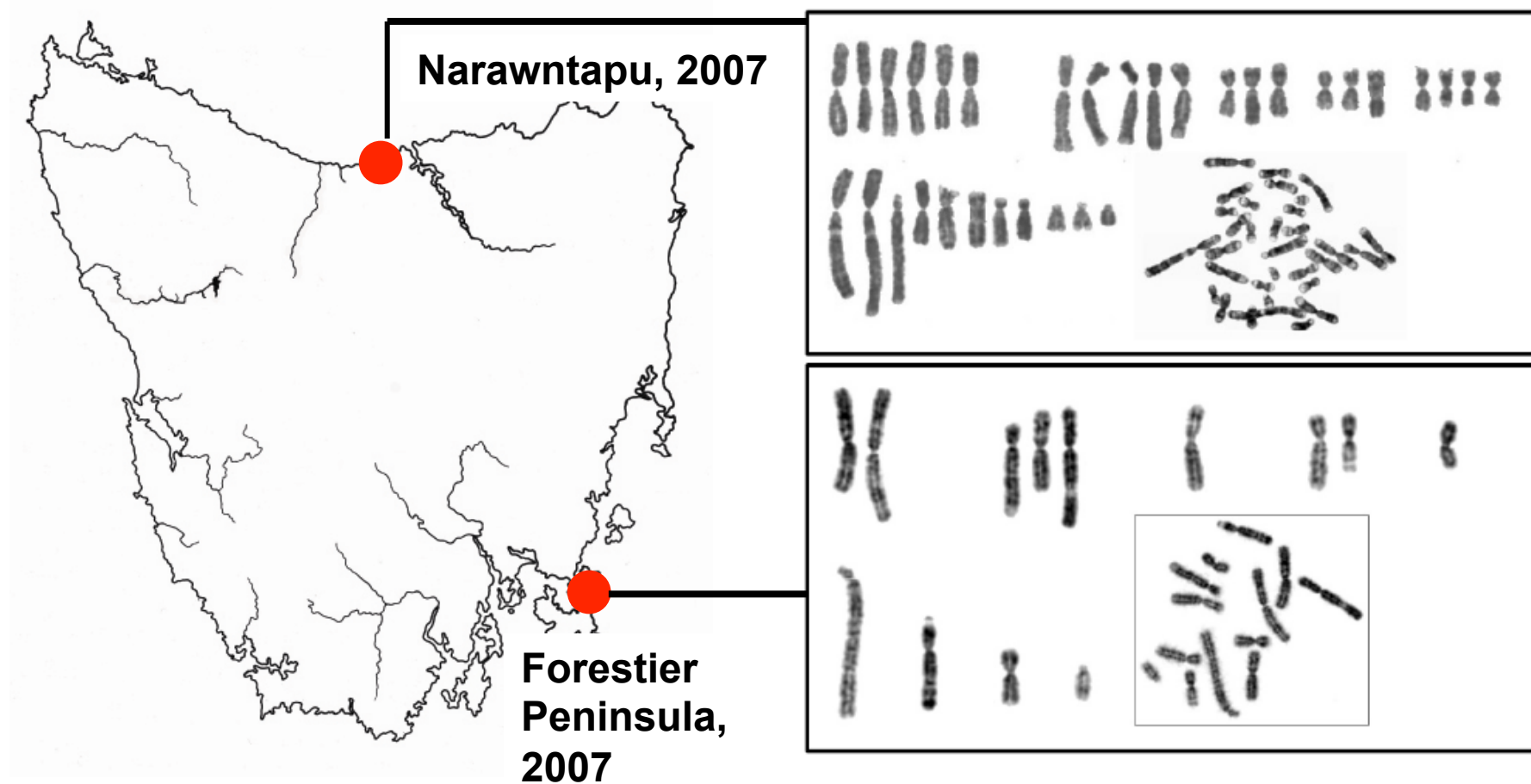
The DFTD genome



Phylogeography

How many somatic mutations are
present in DFTD?

Tasmanian devil cancer genome sequencing

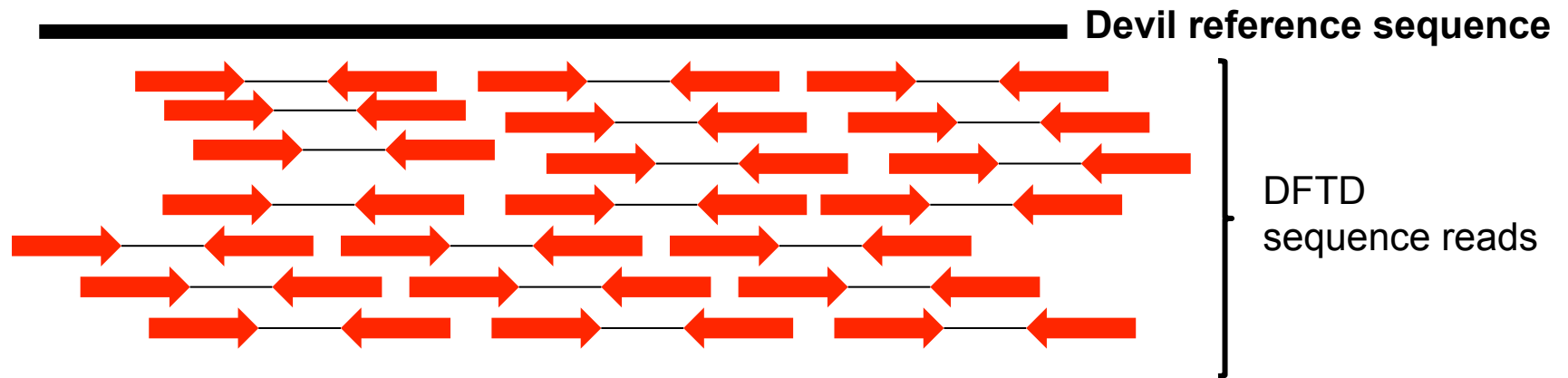


DFTD genome sequencing

$\sim 2 \times 10^9$ paired end sequence reads



DFTD genome analysis



DFTD variant calling

ATCTTCGGAGTACTTCTAGTACTAAGCGTAAGATAACAACAG

Devil reference sequence

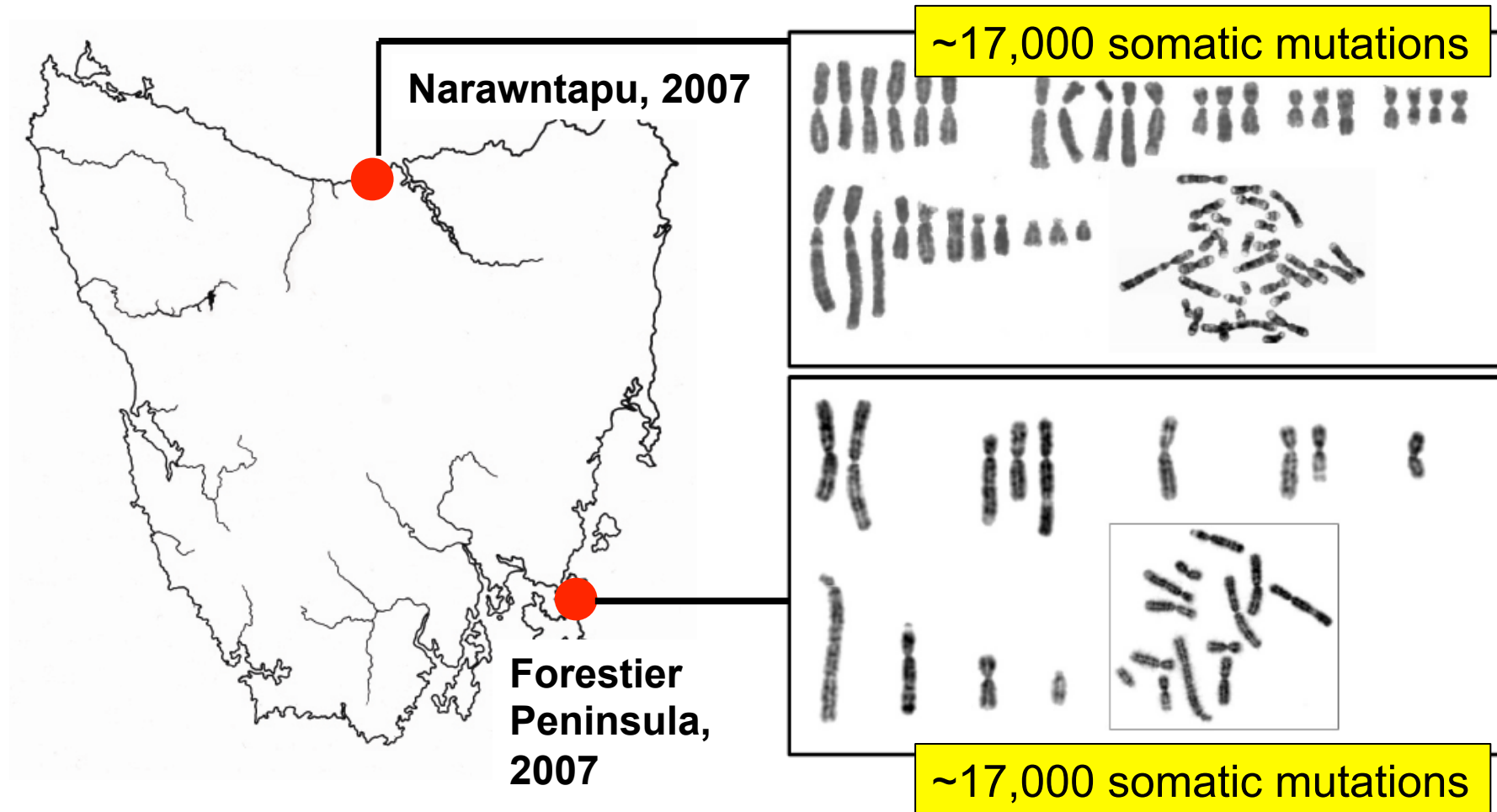
|||
ATCTTCGGAGTACTT**A**TAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTT**A**TAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTT**A**TAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTT**A**TAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTT**A**TAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTT**A**TAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTT**A**TAGTACTAAGCGTAAGATAACAACAG

Narawntapu DFTD
sequence reads

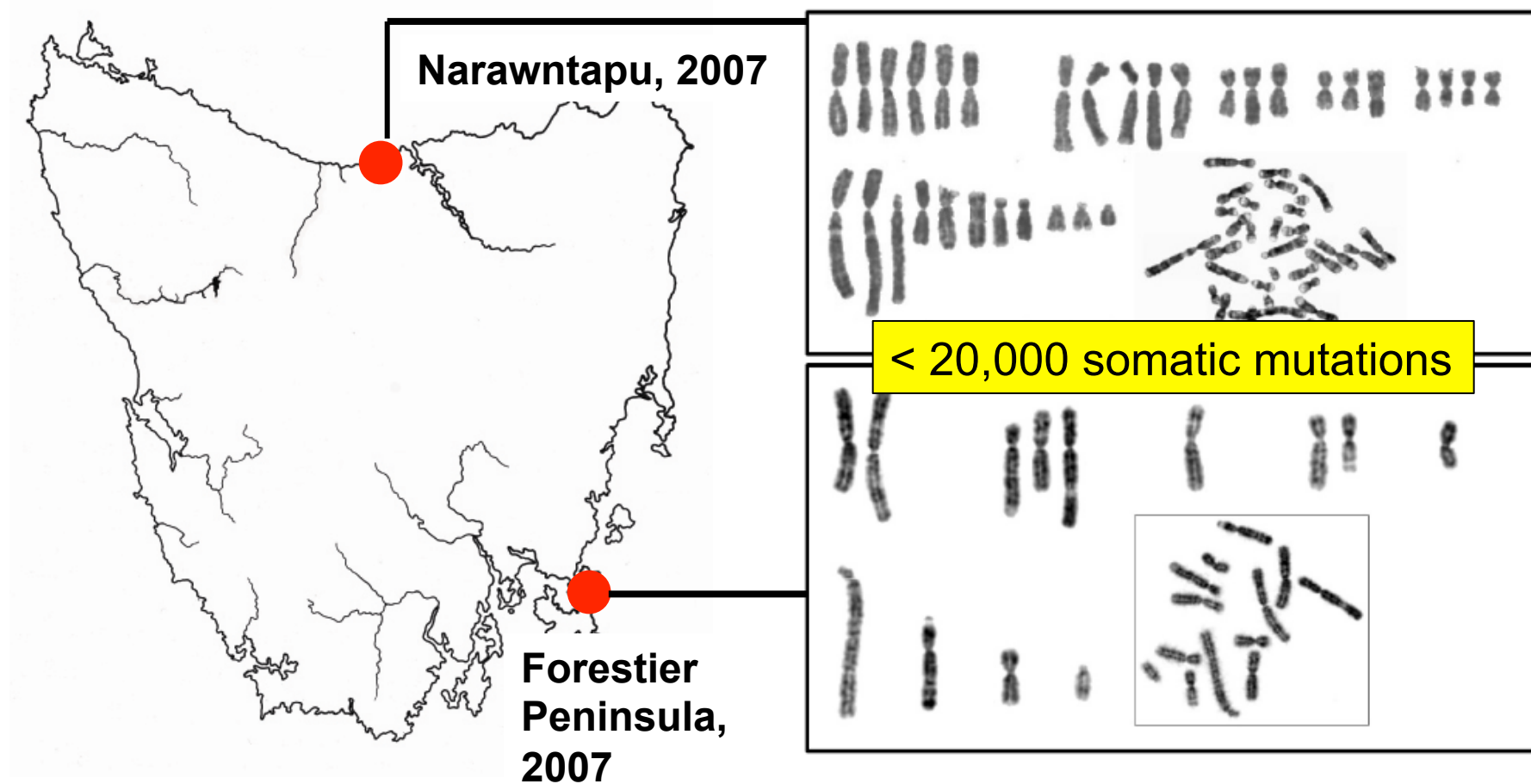
ATCTTCGGAGTACTTCTAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTTCTAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTTCTAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTTCTAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTTCTAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTTCTAGTACTAAGCGTAAGATAACAACAG
ATCTTCGGAGTACTTCTAGTACTAAGCGTAAGATAACAACAG

Forestier DFTD
sequence reads

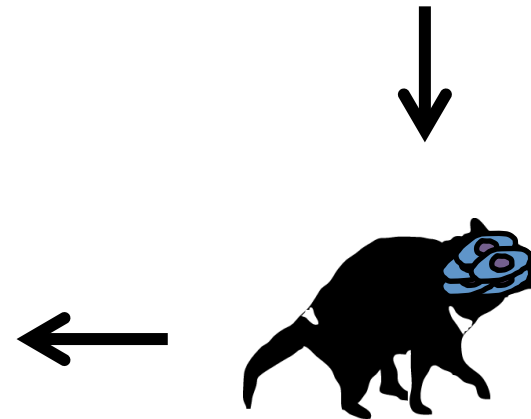
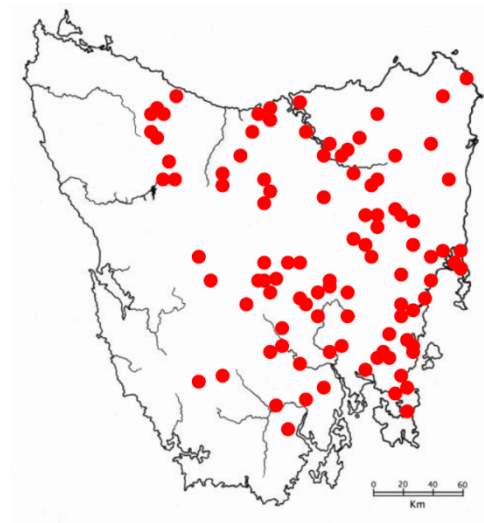
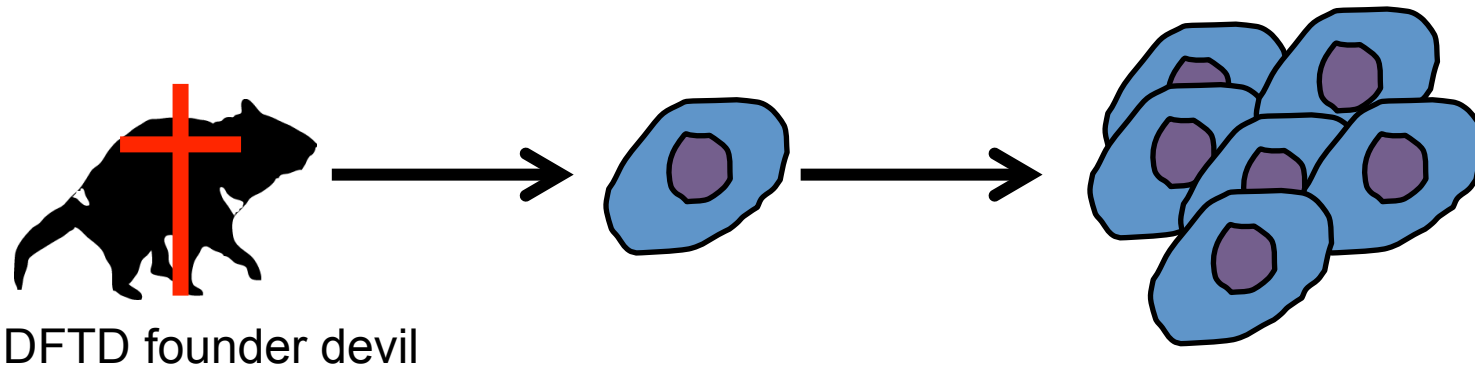
Tasmanian devil cancer genomes



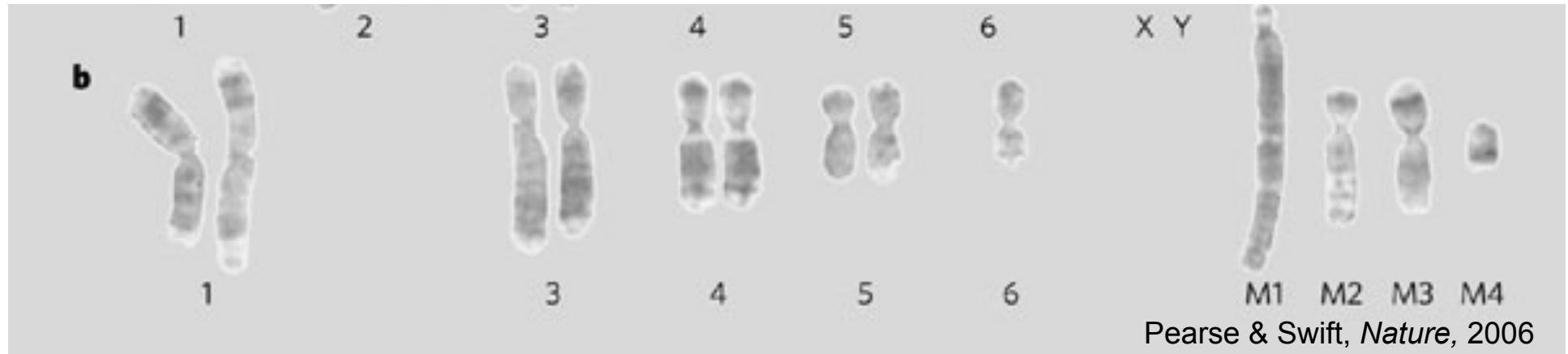
Tasmanian devil cancer genomes



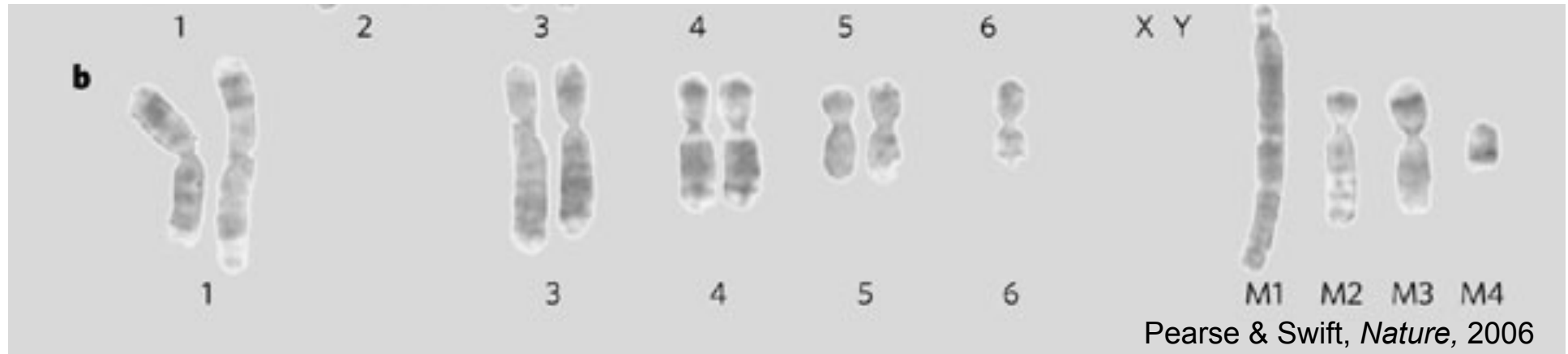
DFTD founder devil



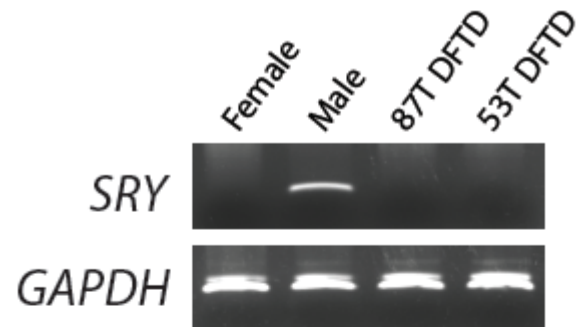
Gender of the founder devil



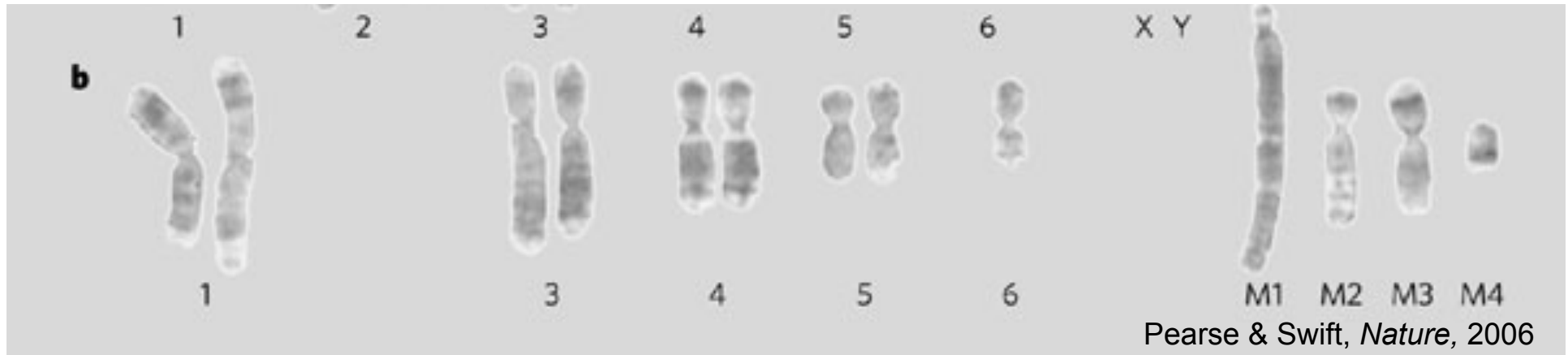
Gender of the founder devil



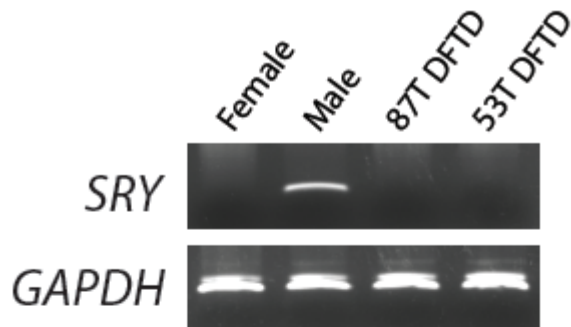
Y chromosome



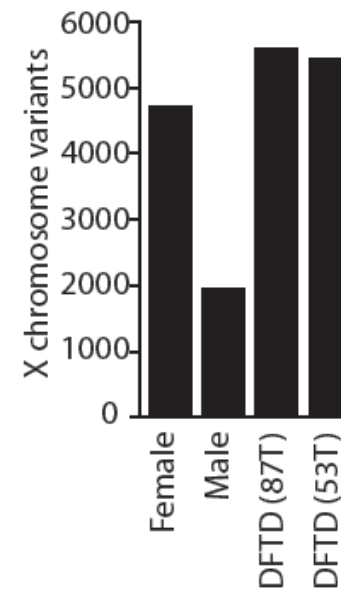
Gender of the founder devil



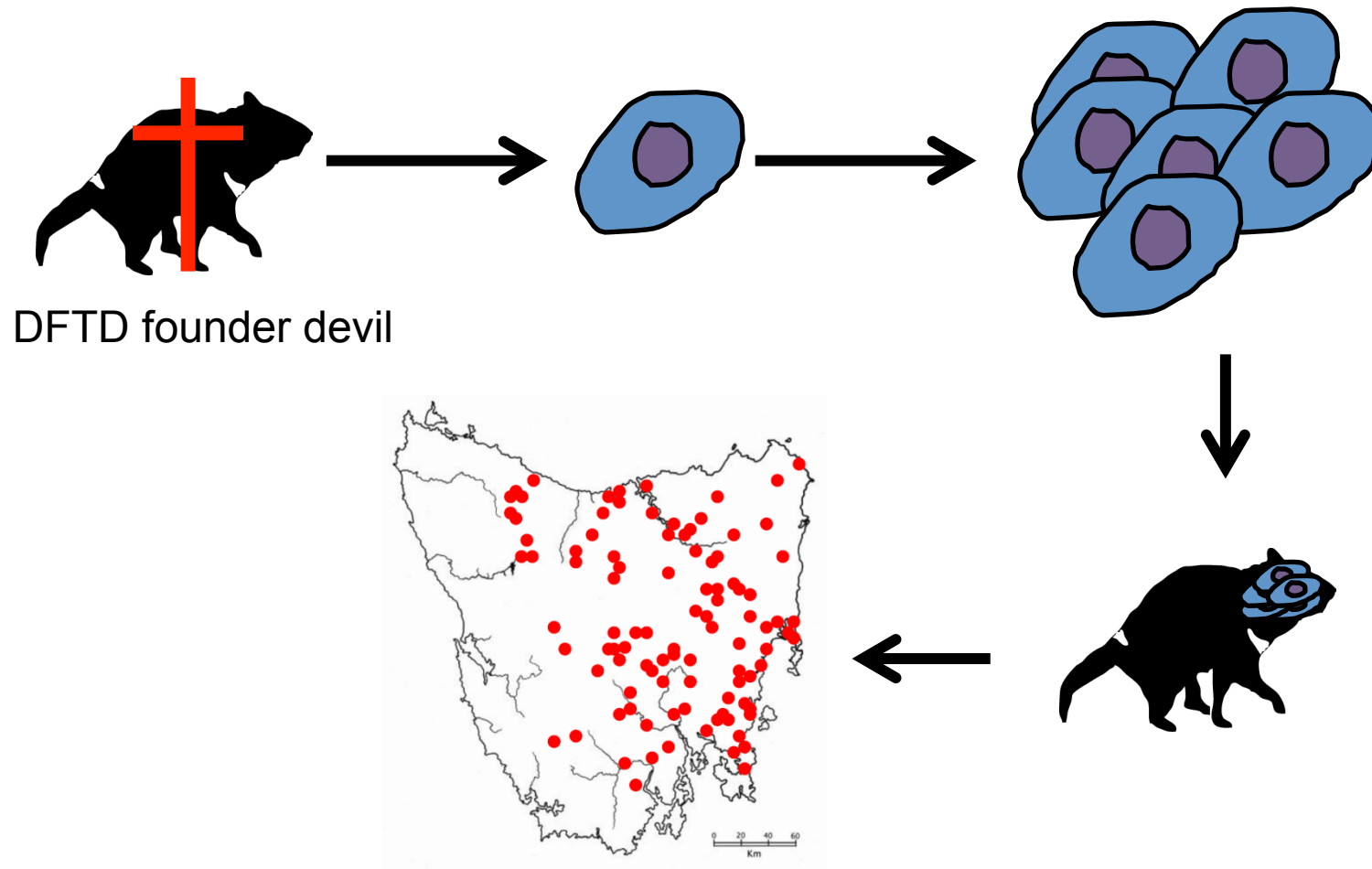
Y chromosome



X chromosome

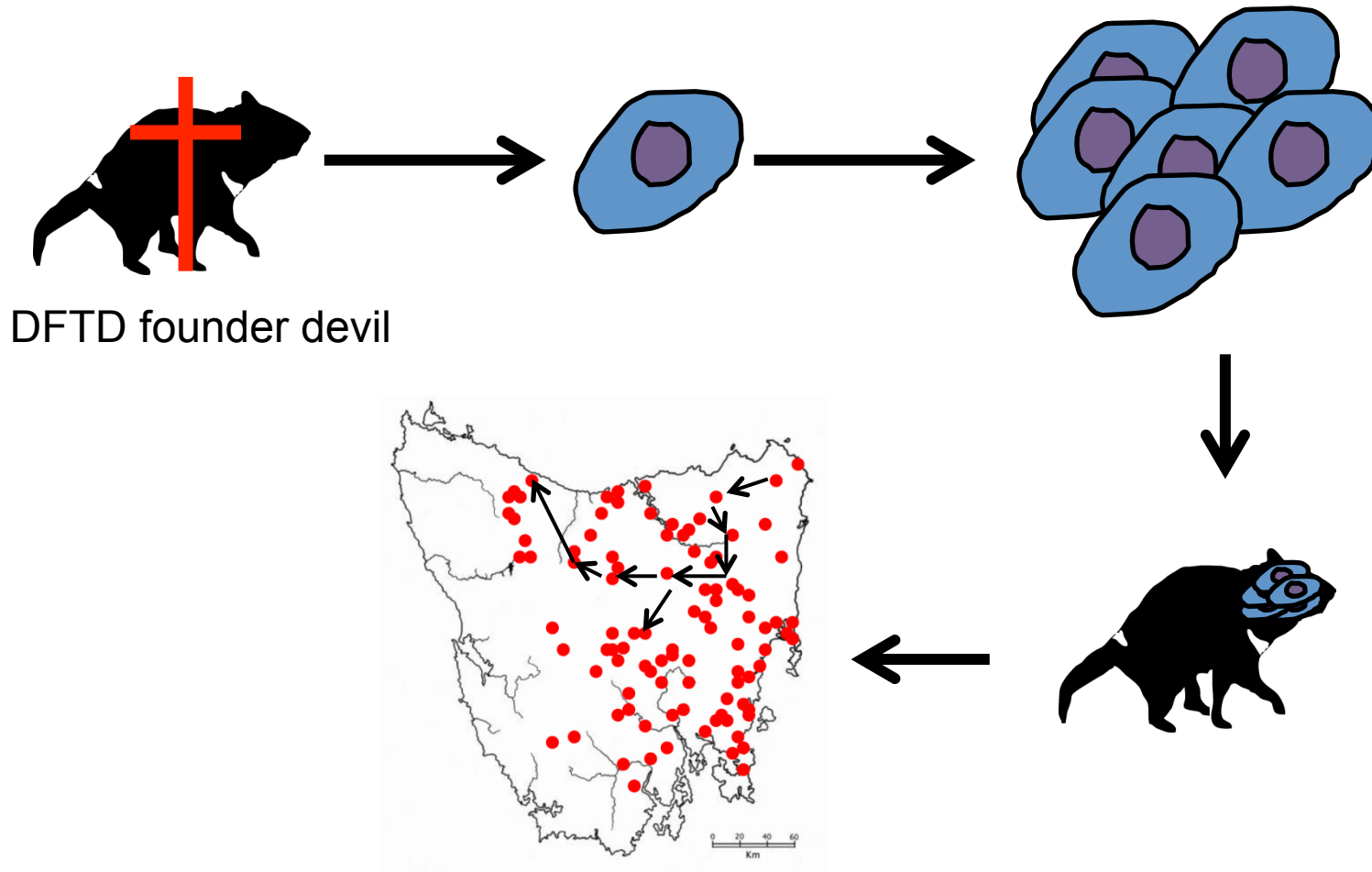


Summary



DFTD founder devil

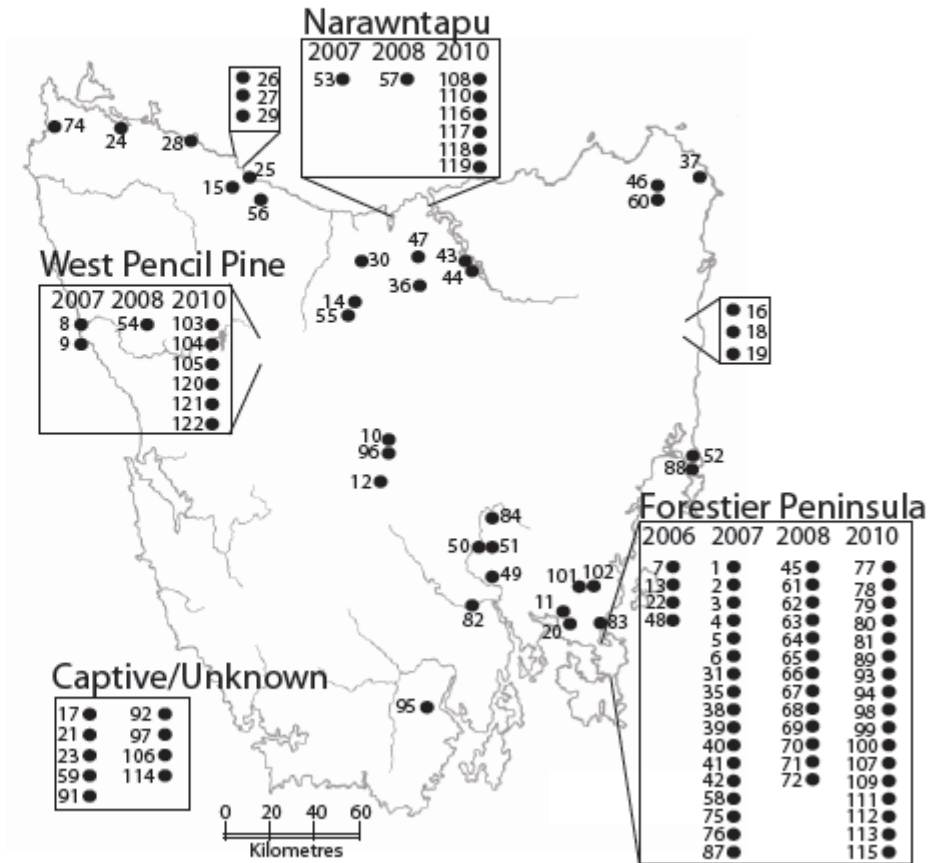
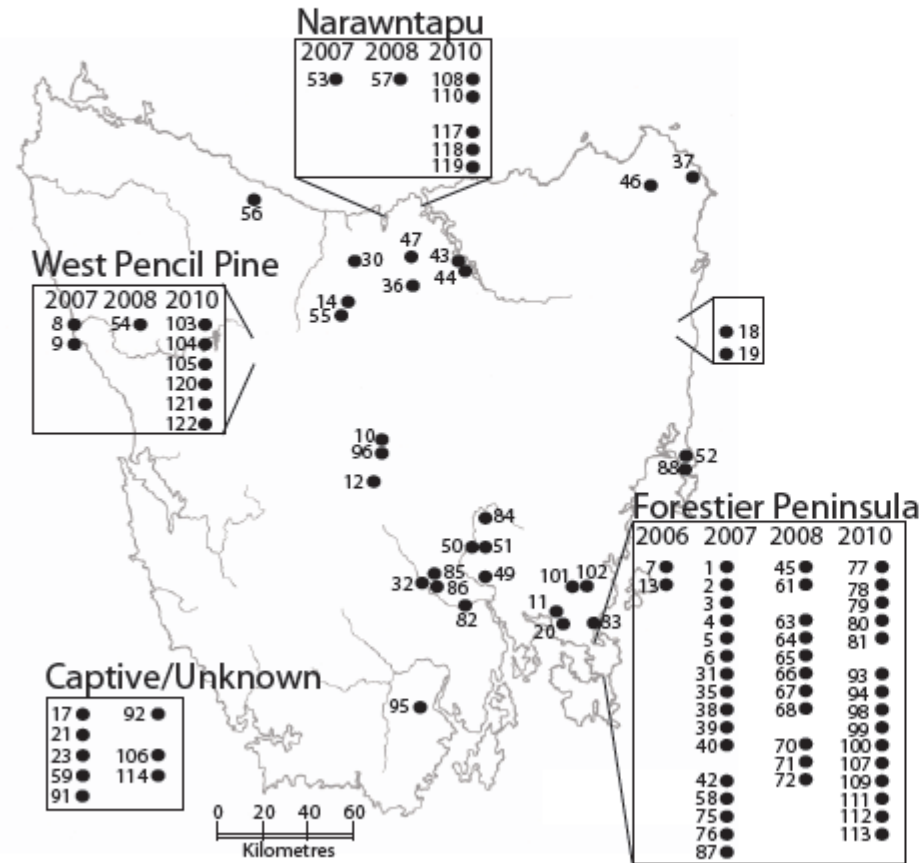
Cancer phylogeography



Devil samples

Tumour

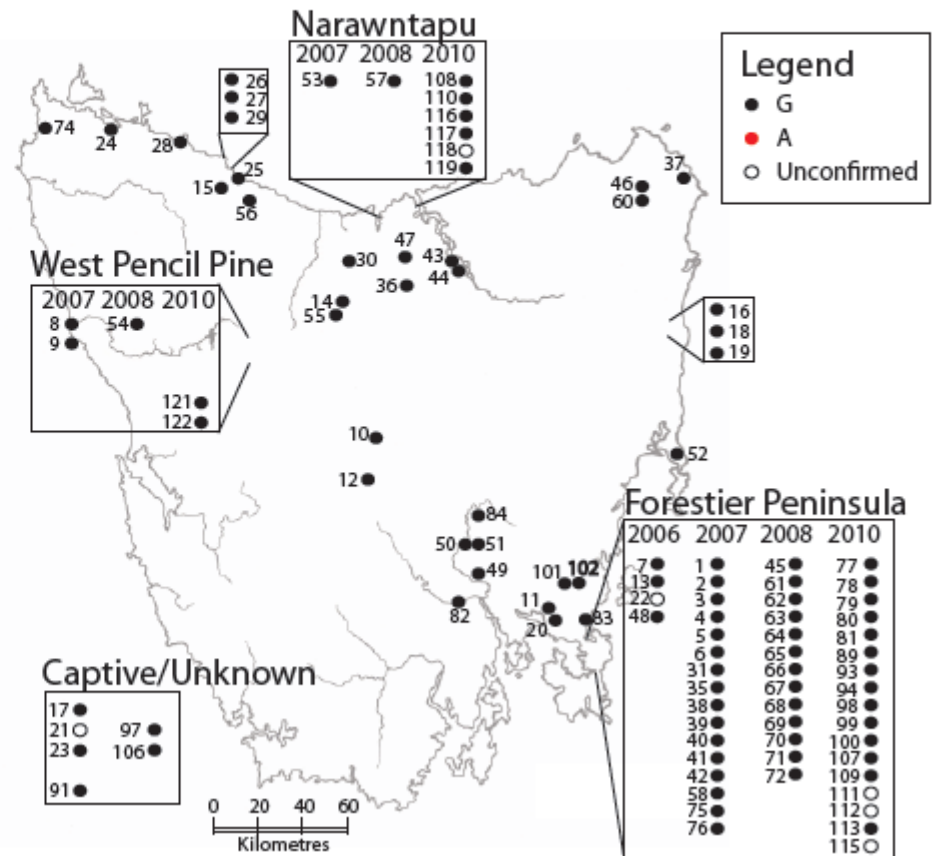
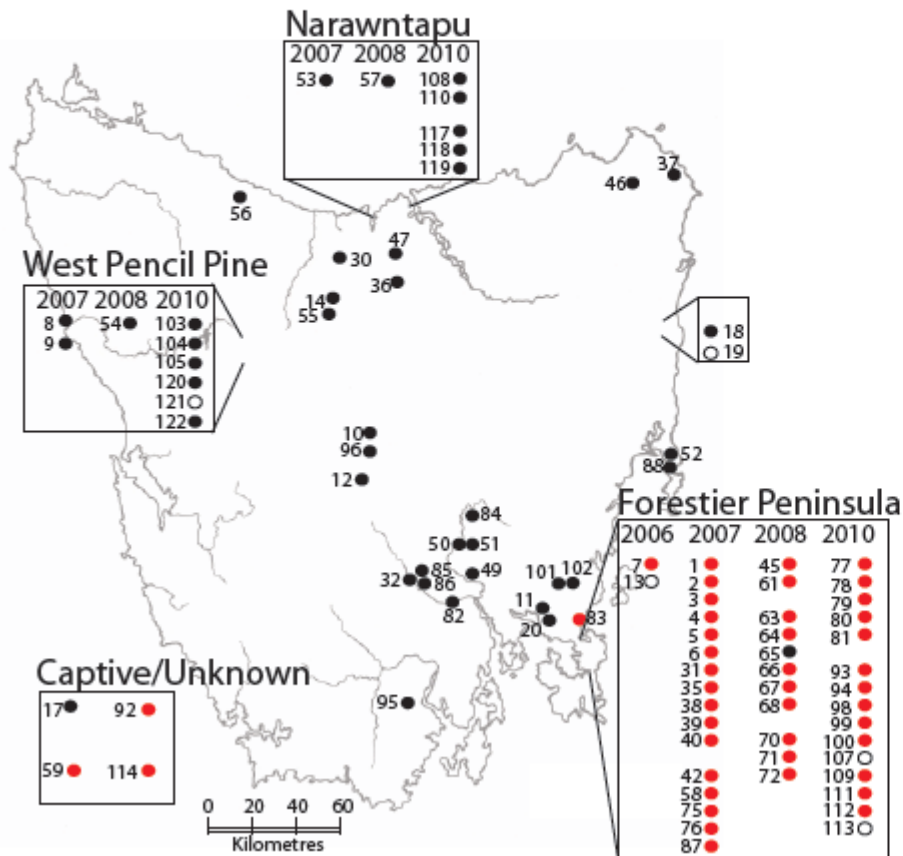
Host



MT:3297 G>A

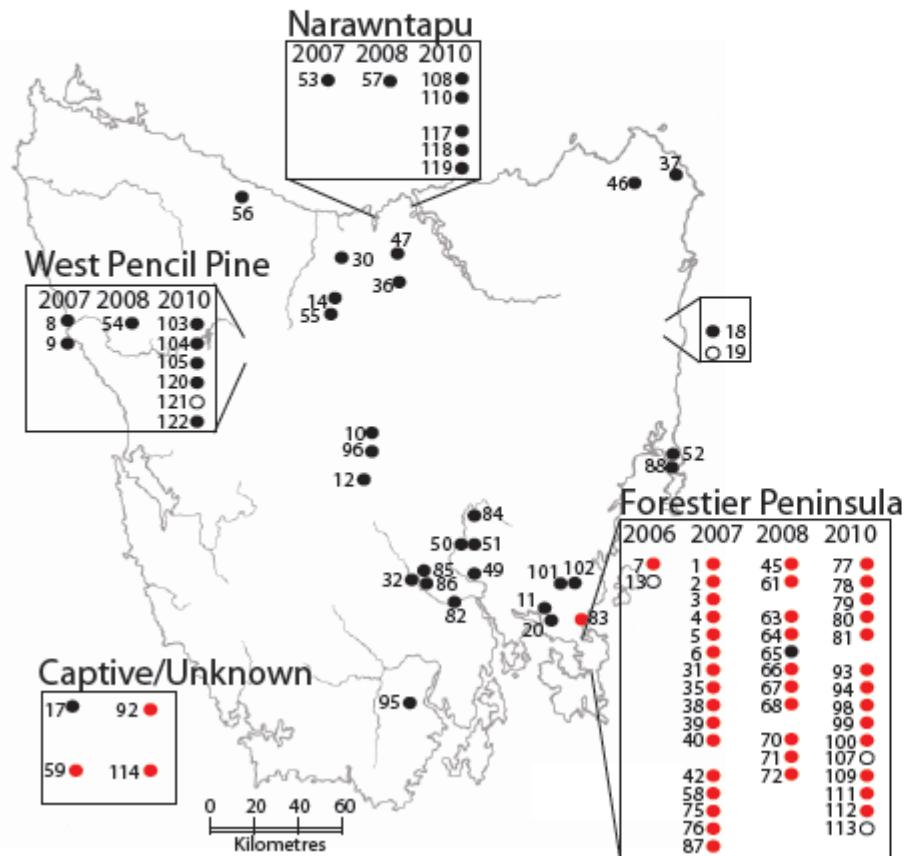
Tumour

Host



MT:3297 G>A

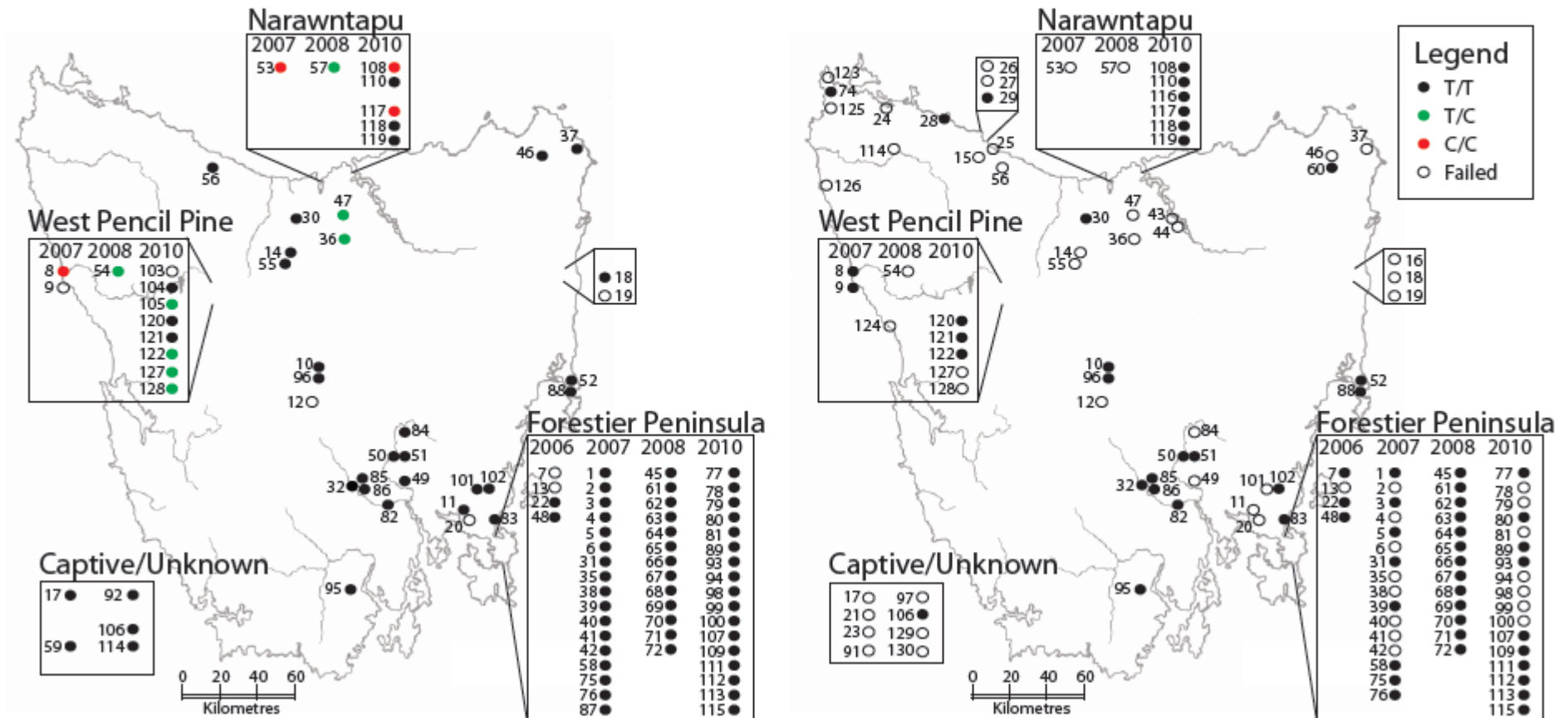
Tumour



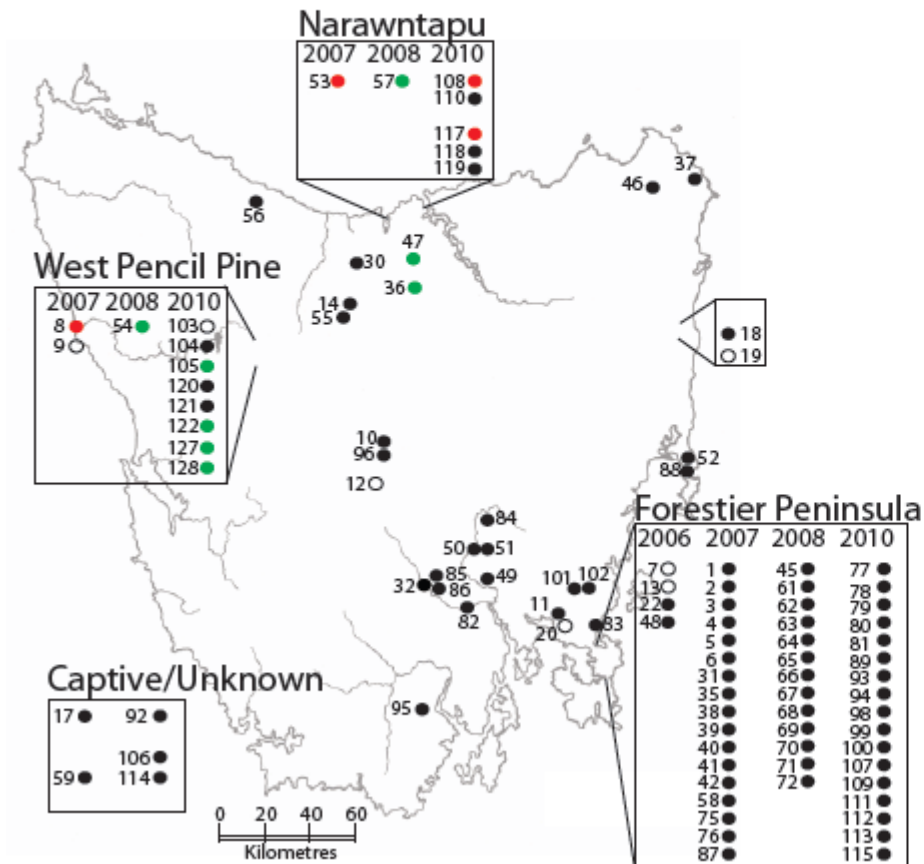
Chromosome 1_supercontig_000000246:95466 T>C

Tumour

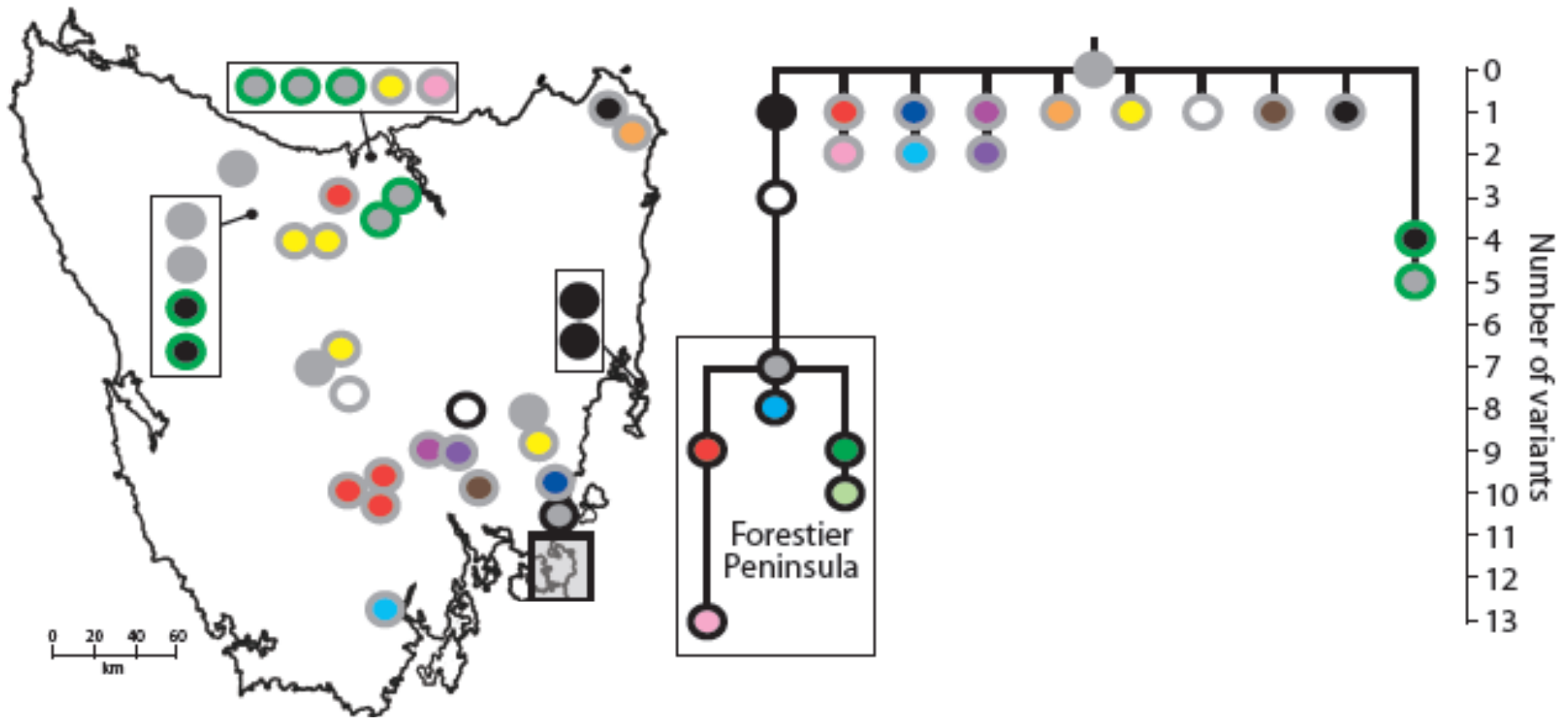
Host

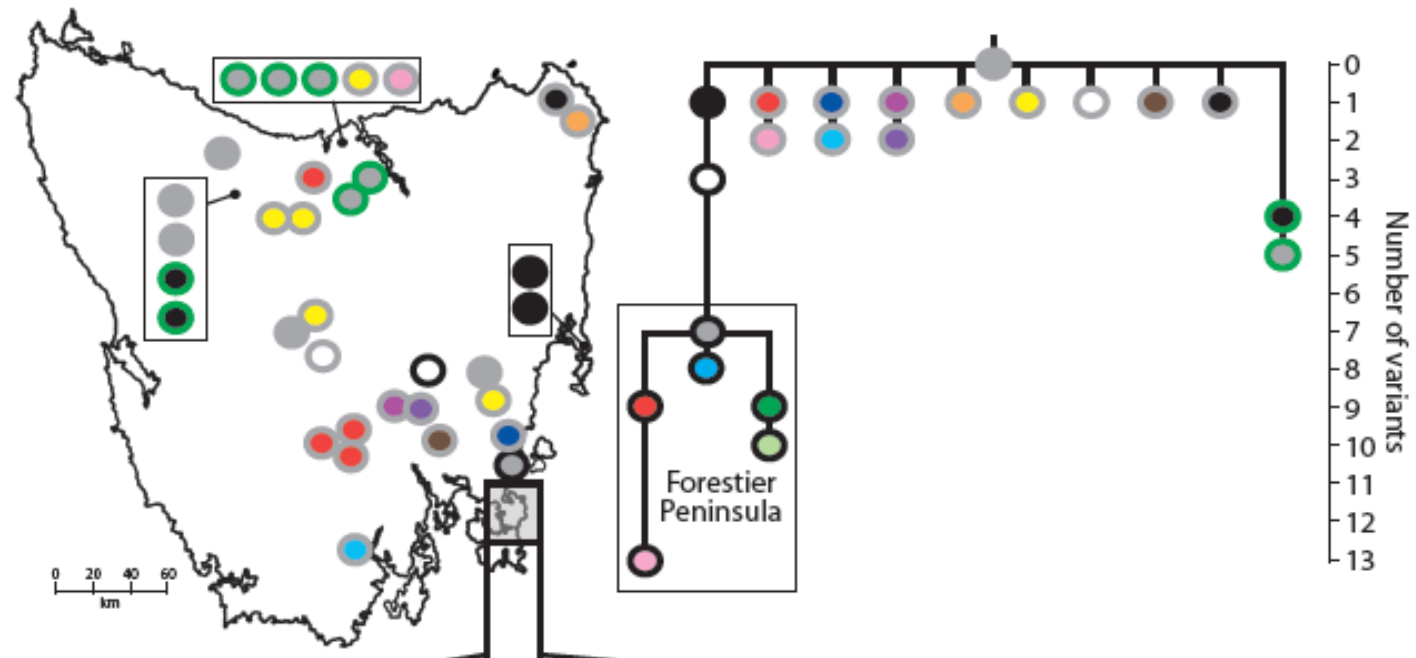


Chromosome 1_supercontig_000000246:95466 T>C Tumour

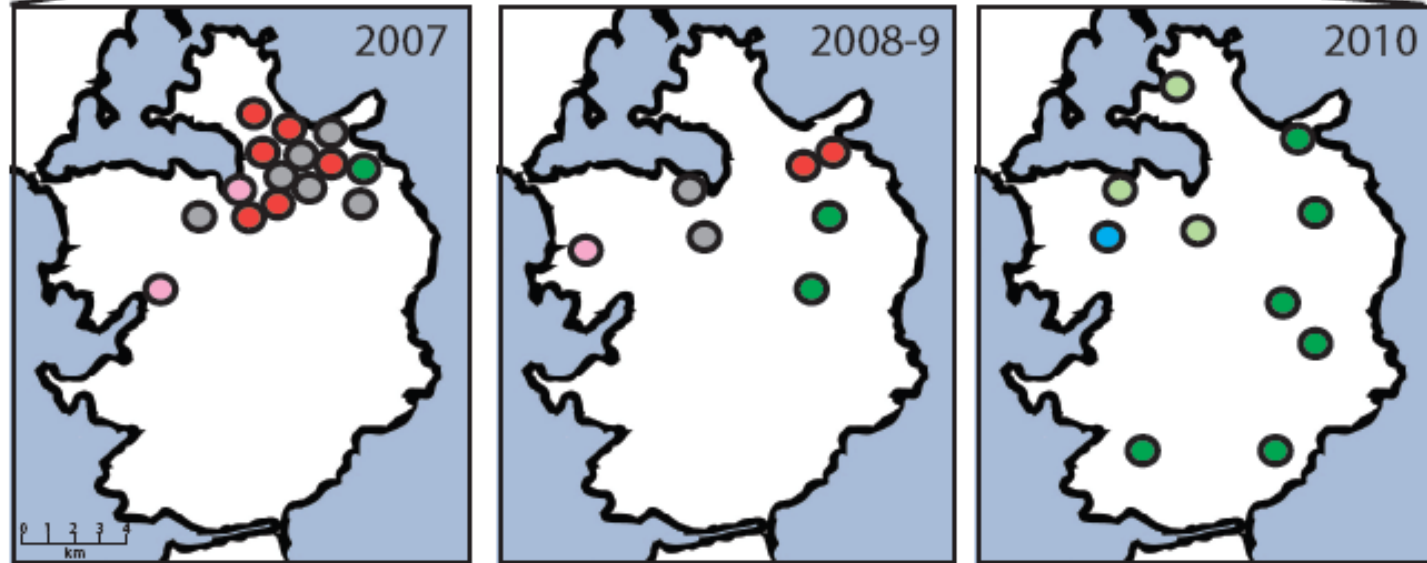


Summary





Forestier Peninsula



What's next for the Tasmanian devil?

- Captive breeding programs
- Restrict disease spread in the wild
- Future research
 - Monitor DFTD genetic diversity and evolution
 - DFTD host interaction
 - DFTD vaccines and therapies



Photo: Save the Tasmanian Devil Program

Canine transmissible venereal tumour



Photo: Ted Donelan

Canine transmissible venereal tumour (CTVT)



Gabriele Marino

Male



Ted Donelan

Female

CTVT worldwide distribution



Andrea Strakova

CTVT is a single somatic cell lineage with a global distribution

- Transplantation experiments

(Nowinsky, 1876; Sticker, 1904)

- Karyotype

(Murray et al, 1969; Oshimura et al 1973, Weber et al 1965, Wright et al, 1970)

- Genetics

- LINE-1 insertion upstream of *MYC*

(Katzir et al, 1987)

- Microsatellites, MHC

(Murgia et al, 2006; Rebbeck et al, 2009)

ON THE
DISEASES
OF
HORSES AND DOGS;

so conducted as to
ENABLE PERSONS TO PRACTISE WITH EASE AND SUCCESS

ON
THEIR OWN ANIMALS,
WITHOUT THE ASSISTANCE OF A FARRIER:

Including likewise the Natural Management, as Stabling,
Feeding, Exercise, &c.: together with the Outlines of
a Plan for the Establishment of Genuine Medicines for
these Animals throughout the Kingdom.



BY DELABERE BLAINE,

PROFESSOR OF ANIMAL MEDICINE;

Author of "The Anatomy of the Horse;" "Outlines of the Veterinary
Art;" "A Treatise on the Distemper in Dogs," &c. &c.

FOURTH EDITION,

WITH VERY LARGE ADDITIONS.

LONDON:

PRINTED FOR T. BOOSEY, 4, OLD BROAD STREET,
ROYAL EXCHANGE.

1810.

habitants of mountainous countries, and has been supposed to be dependant on some particular quality of the water in those vicinities. But in dogs no such peculiarity takes place: it does not appear in them indigenous to any particular soil, but almost peculiar to some particular species of dogs, though other dogs sometimes have it, as terriers; but it is much less frequent, and in the larger tribes is hardly ever seen. It comes on generally while very young, and continues to increase to a certain size, when it becomes stationary, seldom increasing to such a degree as to prove fatal. It is however troublesome, and in some measure hurtful, from the pressure it occasions on the surrounding parts. If an ointment is made with equal parts of mercurial and blistering ointment, and the swelling rubbed with it every day, avoiding salivation, it commonly lessens and frequently wholly removes it. But it is necessary also at the same time to give internal alteratives: four, five, six, or seven grains of burnt sponge, with half the quantity of nitre given every morning, will be found useful.

CANCER.

Two parts only in dogs are subject to a cancerous affection, and both these are organs concerned in generation. The teats of bitches become at times indurated, and swelled with a schirrous indolent tumour, which more commonly remains indolent and without ulcer: but now and then, when such a tumour has increased to a very considerable size, a small ulcer bursts out, which slowly increases to a very considerable surface. In the worst of these cases there is not present the virulent and horrible spreading of the

human cancer, nor does it appear to give much pain, or to injure the health; and dogs who are suffered to remain with it, live for a great length of time without much inconvenience to themselves. This state admits of only one cure, which is the complete removal of the whole tumour. The vagina or rather the womb of bitches also frequently takes on an ulcerous state, accompanied with a fungous excrescence, which is brought on oftentimes from the horrible brutality of boys who force dogs from bitches in the act of copulation. This complaint admits of no cure, that I have witnessed. In the penis of dogs also a similar fungous excrescence sometimes forms, but it does not appear to erode the neighbouring parts much: it increases rather than diminishes the size, till its offensiveness obliges the animal to be made away with.

habitants of mountainous countries, and has been supposed to be dependant on some particular quality of the water in those vicinities. But in dogs no such peculiarity takes place: it does not appear in them indigenous to any particular soil, but almost peculiar to some particular species of dogs, though other dogs sometimes have it, as terriers; but it is much less frequent, and in the larger tribes is hardly ever seen. It comes on generally while very young, and continues to increase to a certain size, when it becomes stationary, seldom increasing to such a degree as to prove fatal. It is however troublesome, and in some measure hurtful, from the pressure it occasions on the surrounding parts. If an ointment is made with equal parts of mercurial and blistering ointment, and the swelling rubbed with it every day, avoiding salivation, it commonly lessens and frequently wholly removes it. But it is necessary also at the same time to give internal alteratives: four, five, six, or seven grains of burnt sponge, with half the quantity of nitre given every morning, will be found useful.

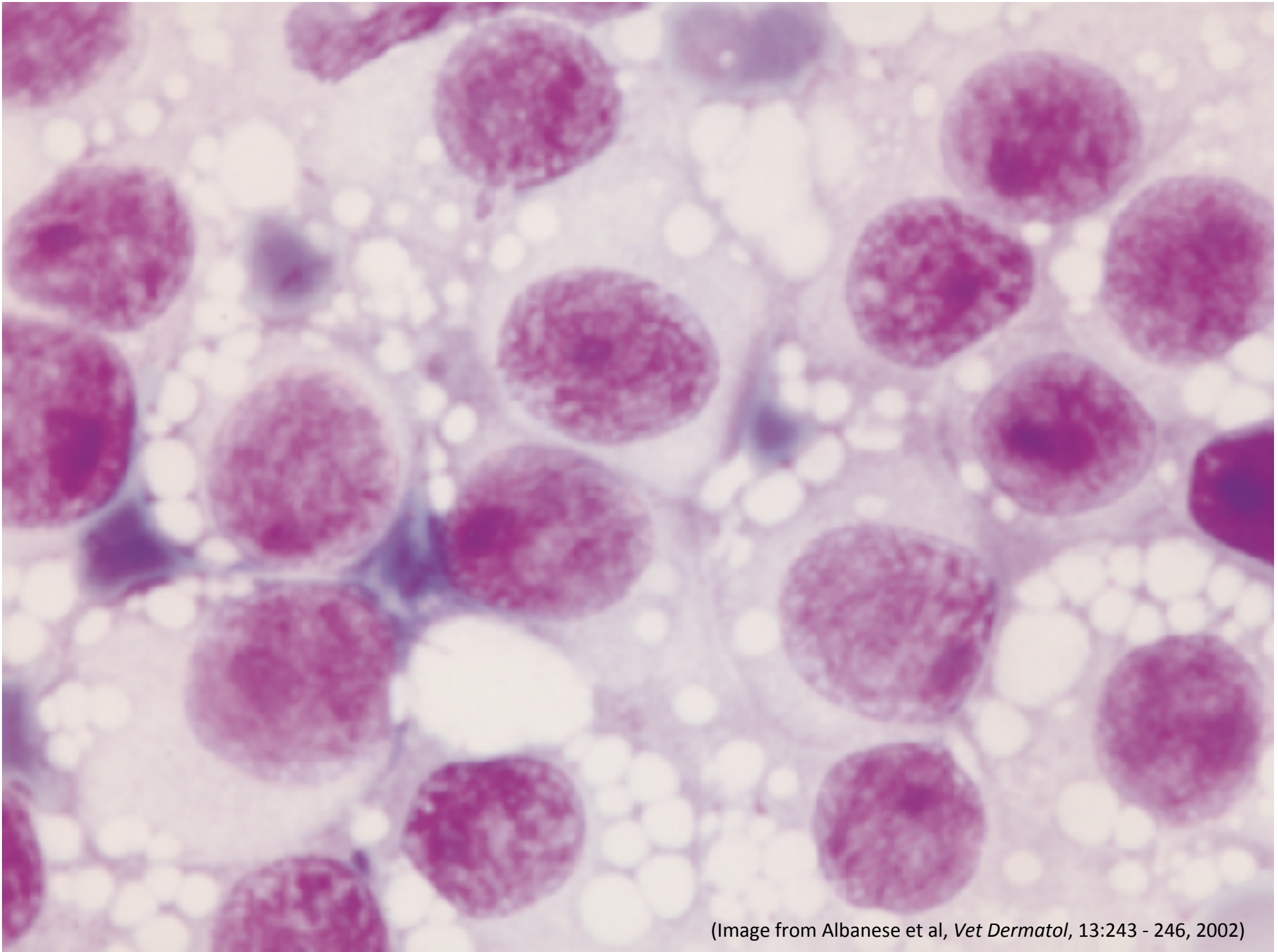
CANCER.

Two parts only in dogs are subject to a cancerous affection, and both these are organs concerned in generation. The teats of bitches become at times indurated, and swelled with a schirrous indolent tumour, which more commonly remains indolent and without ulcer: but now and then, when such a tumour has increased to a very considerable size, a small ulcer bursts out, which slowly increases to a very considerable surface. In the worst of these cases there is not present the virulent and horrible spreading of the

human cancer, nor does it appear to give much pain, or to injure the health; and dogs who are suffered to remain with it, live for a great length of time without much inconvenience to themselves. This state admits of only one cure, which is the complete removal of the whole tumour. The vagina or rather the womb of bitches also frequently takes on an ulcerous state, accompanied with a fungous excrescence, which is brought on oftentimes from the horrible brutality of boys who force dogs from bitches in the act of copulation. This complaint admits of no cure, that I have witnessed. In the penis of dogs also a similar fungous excrescence sometimes forms, but it does not appear to erode the neighbouring parts much: it increases rather than diminishes the size, till its offensiveness obliges the animal to be made away with.

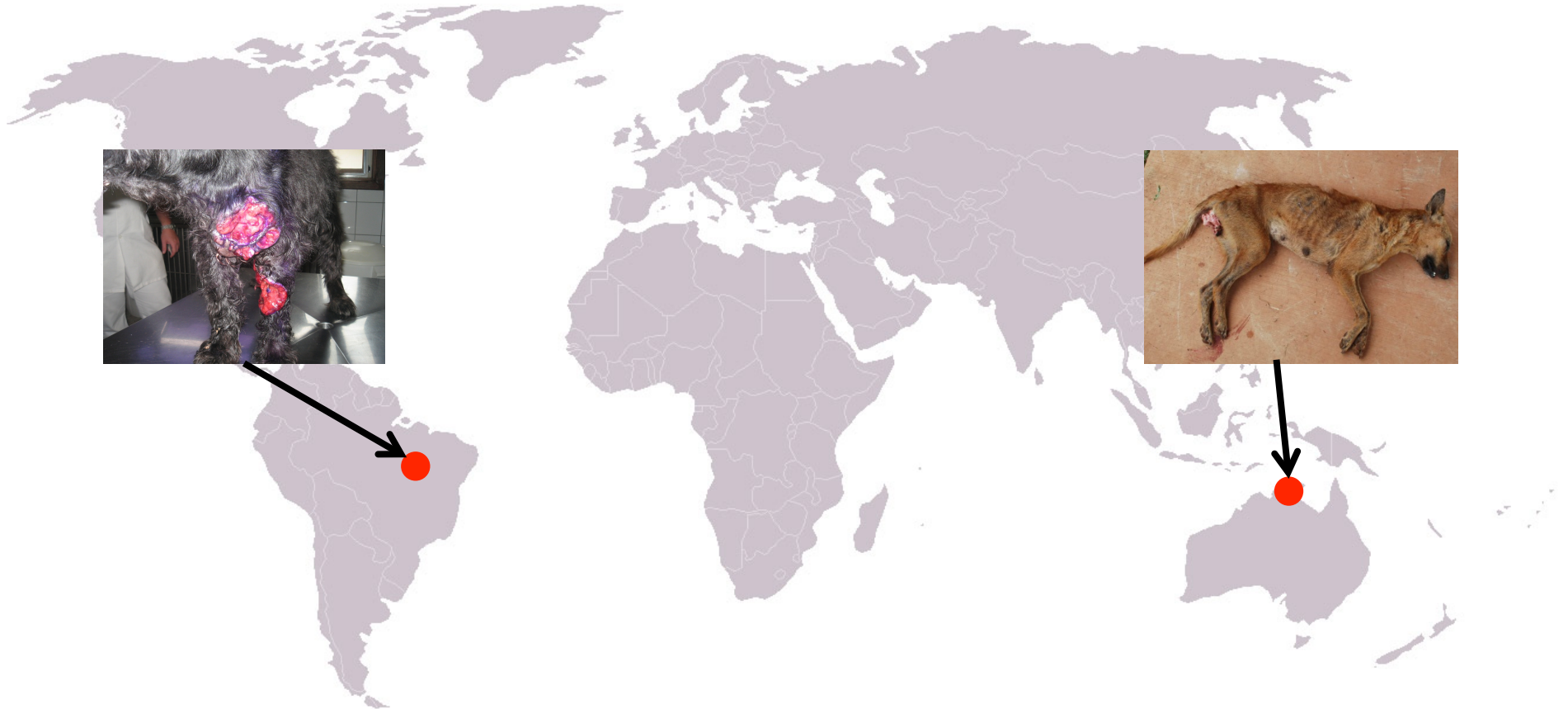
CTVT is the oldest known mammalian somatic cell lineage

- CTVT lineage is at least 200 years old
- CTVT may be much older than this

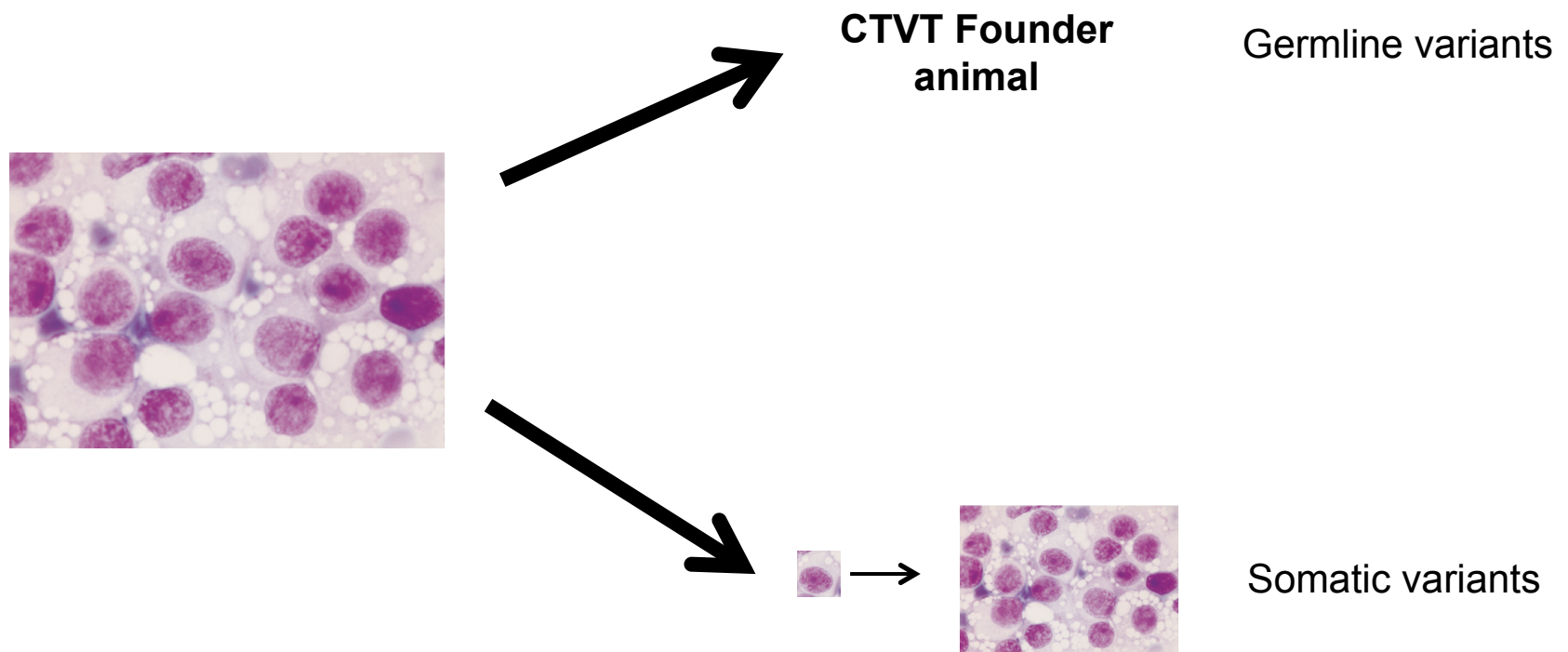


(Image from Albanese et al, *Vet Dermatol*, 13:243 - 246, 2002)

CTVT genome sequencing



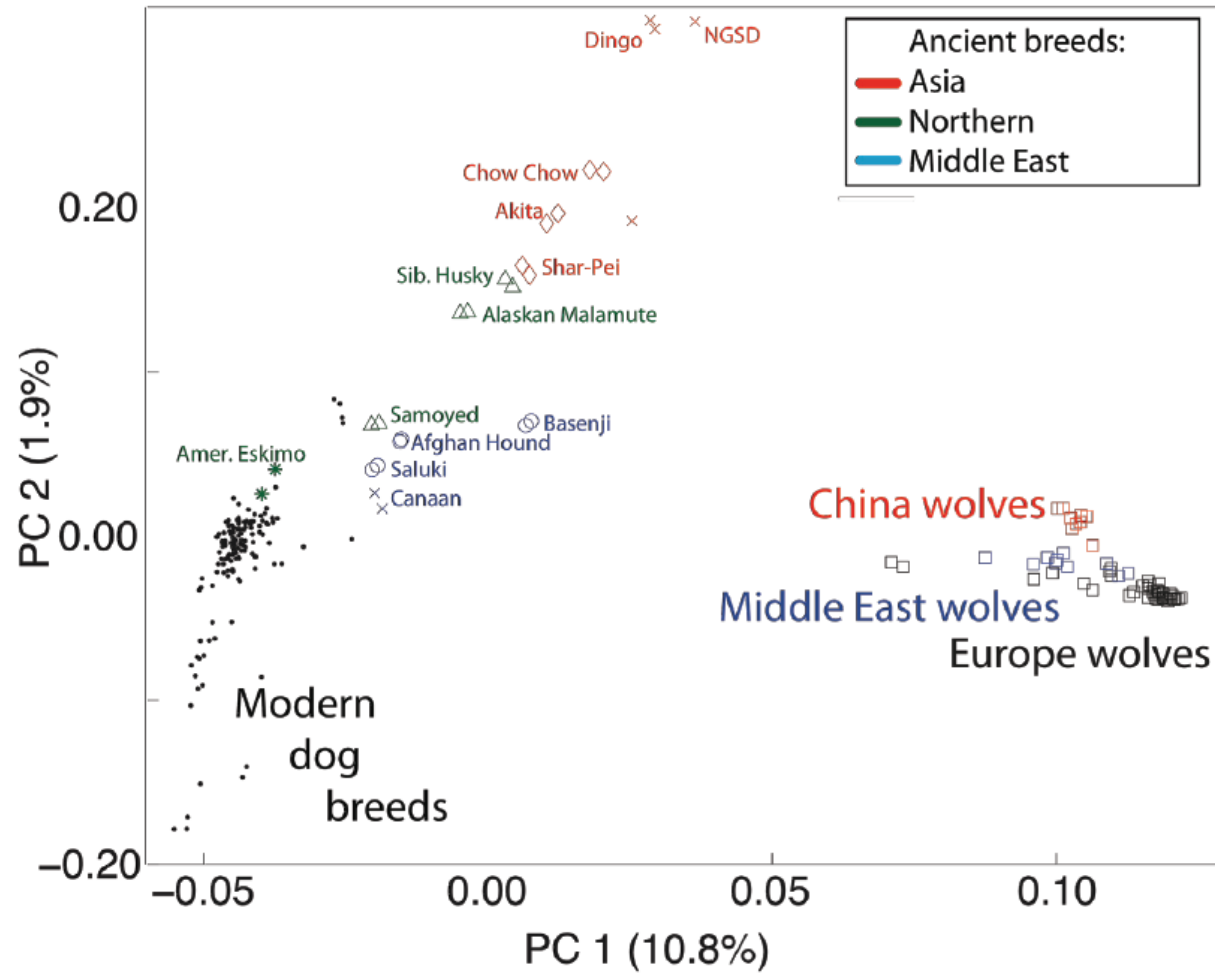
The CTVT genome



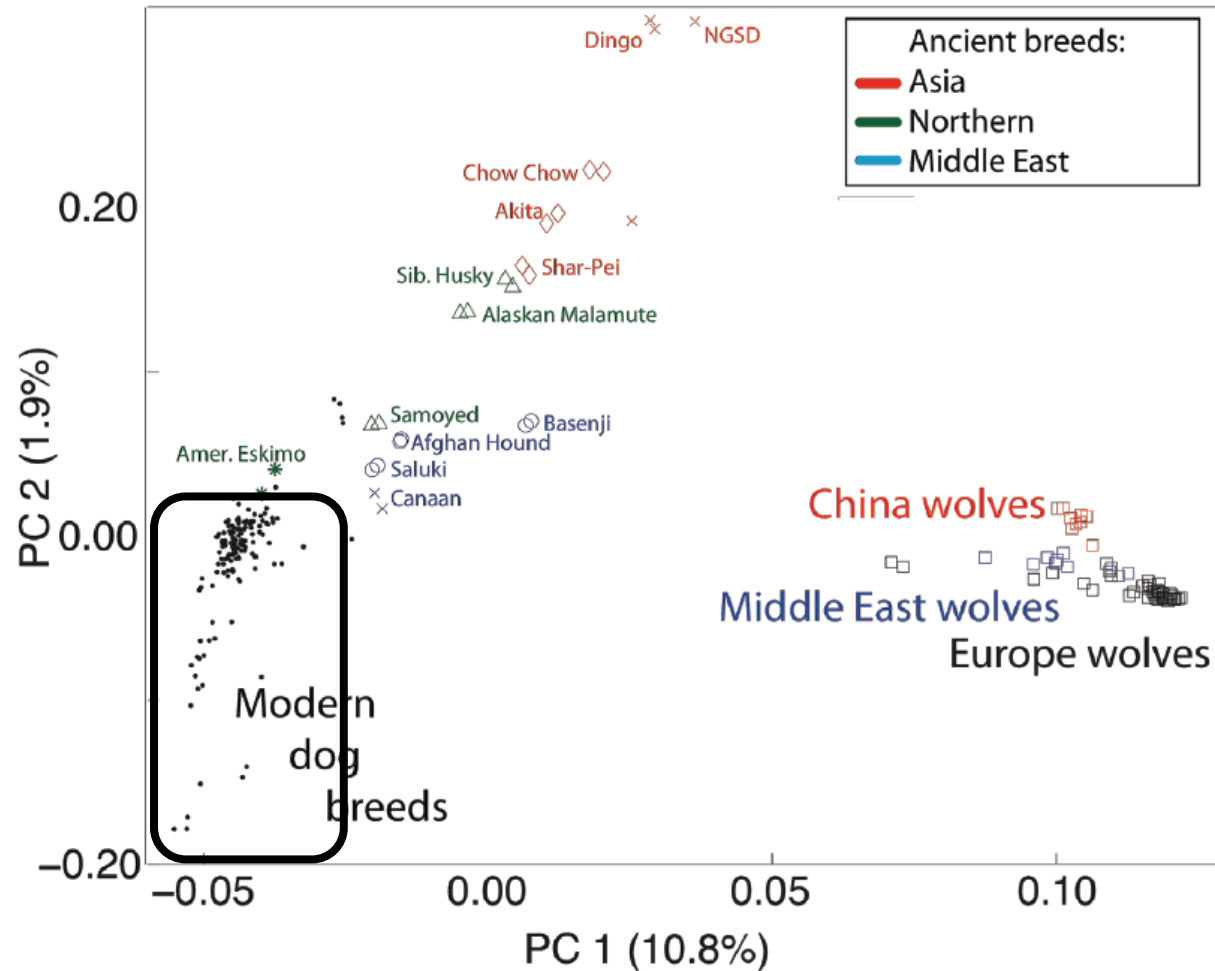
(Image from Albanese et al, *Vet Dermatol*, 13:243 - 246, 2002)

Identity of the CTVT founder
animal

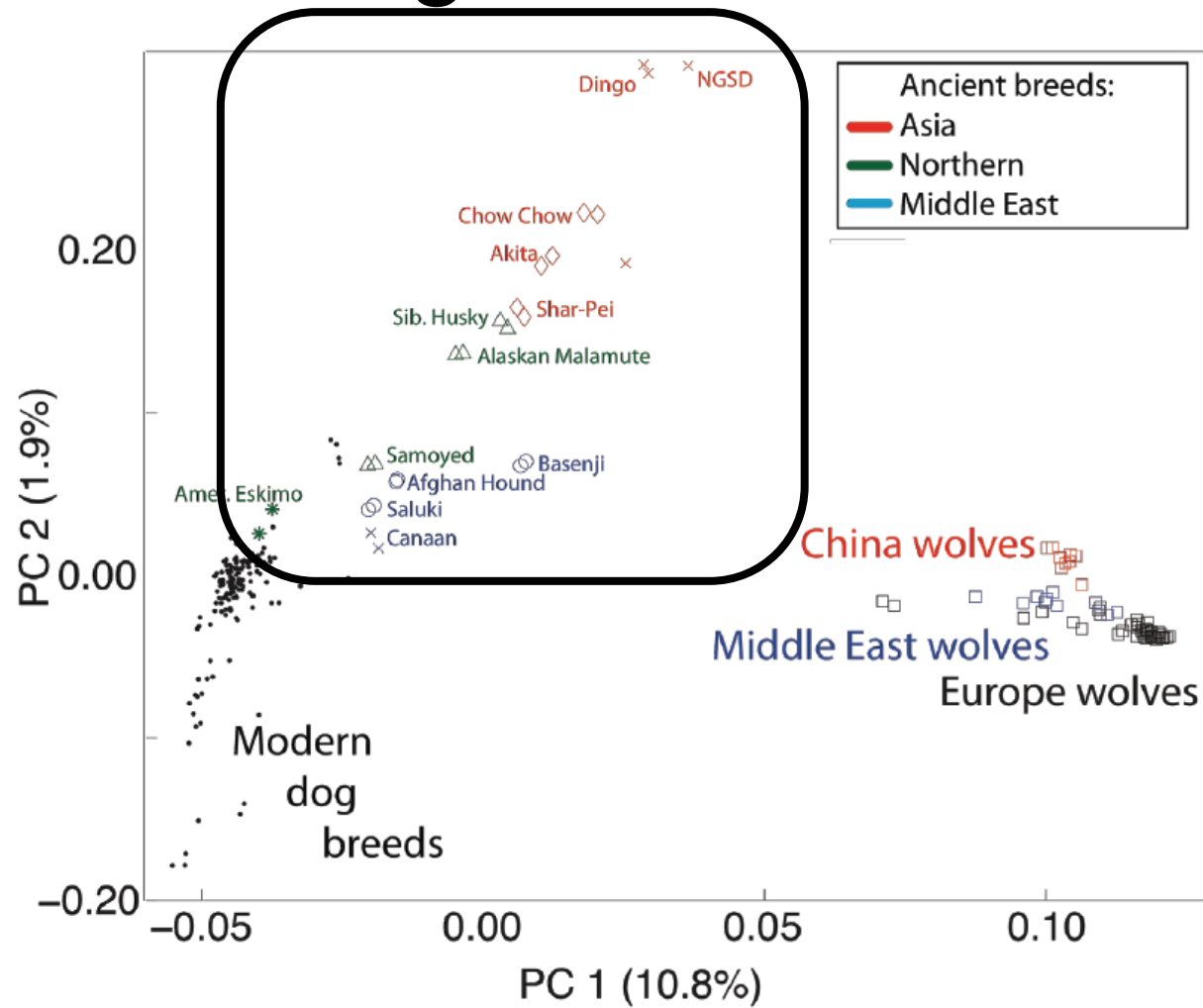
Principal component analysis of dogs and wolves



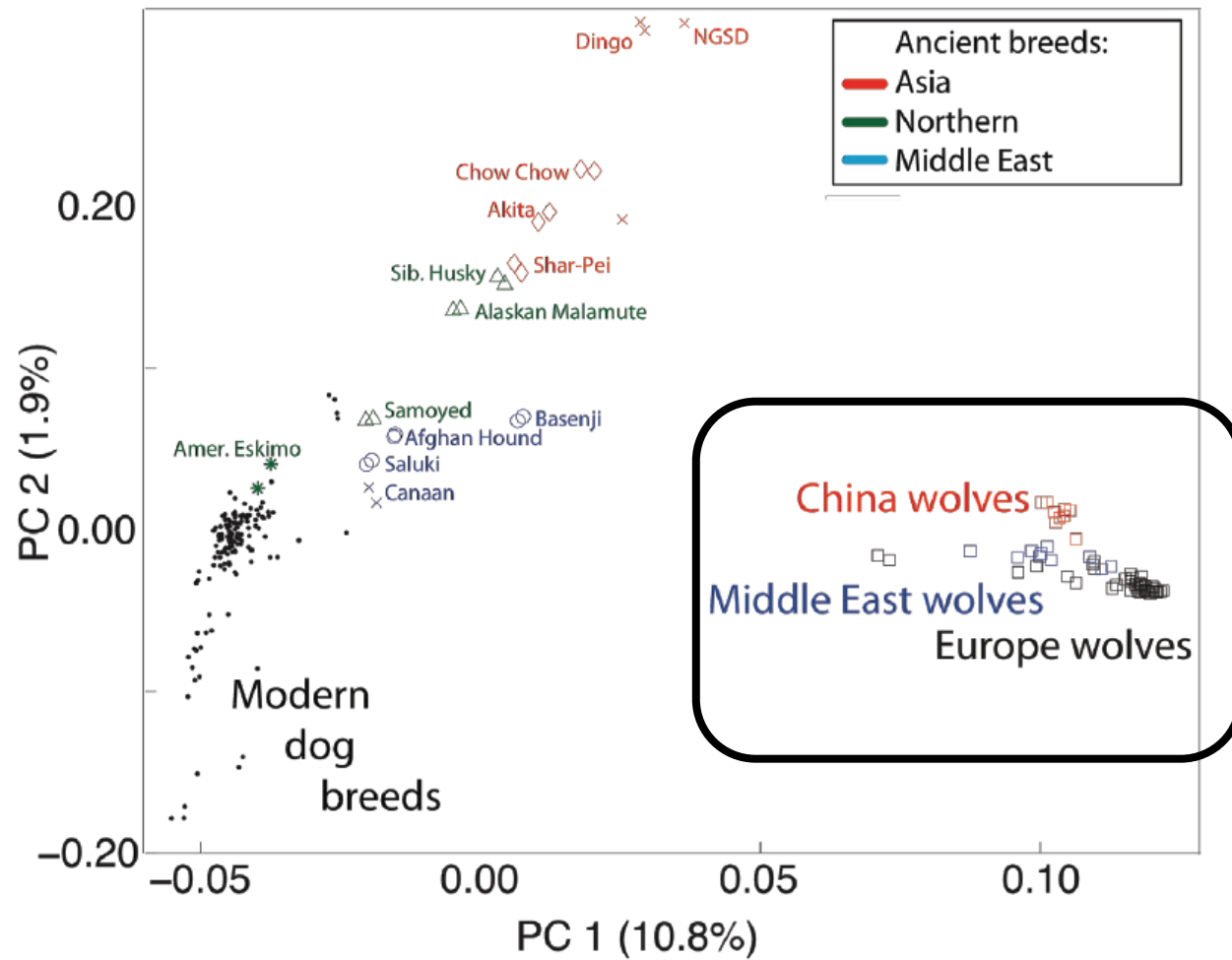
Principal component analysis of dogs and wolves



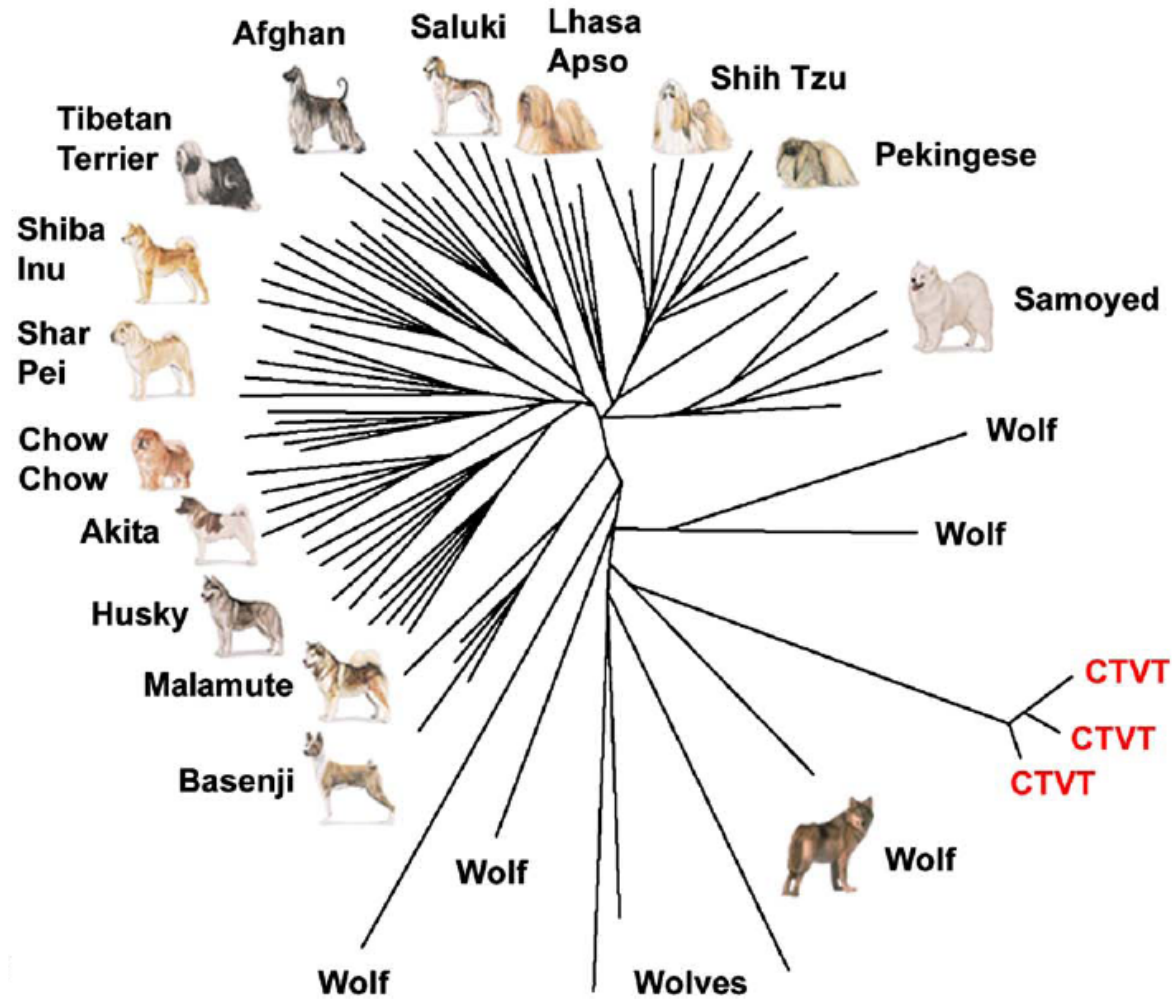
Principal component analysis of dogs and wolves



Principal component analysis of dogs and wolves

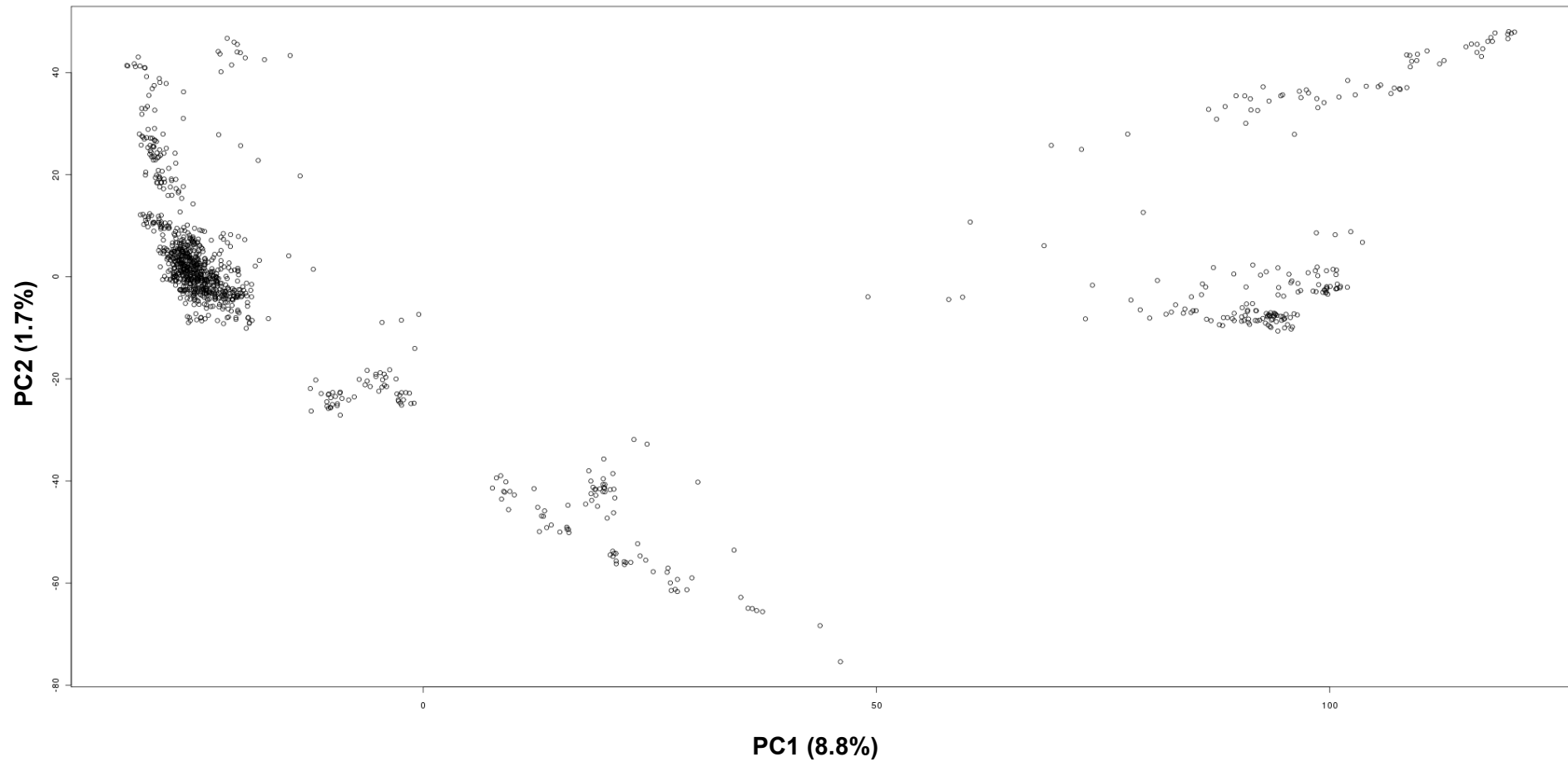


CTVT founder phenotype



Murgia et al, *Cell*, 126:477-87, 2006; Rebbeck et al *Evolution*, 63(9):2340-9, 2009

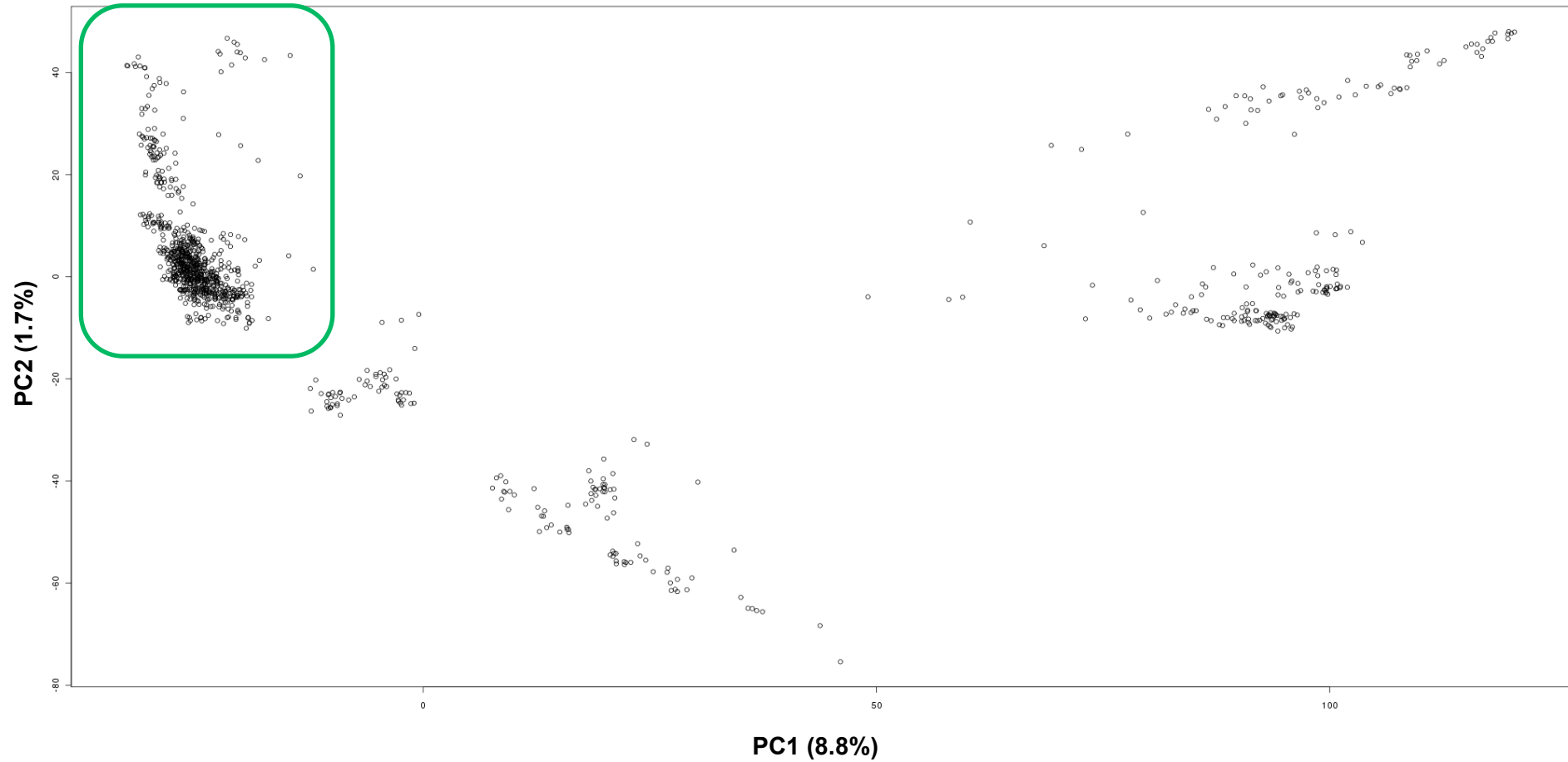
The CTVT founder



With data from vonHoldt et al, *Nature* 364:898 (2010) and vonHoldt et al, *Genome Research* 21:1294 (2011)

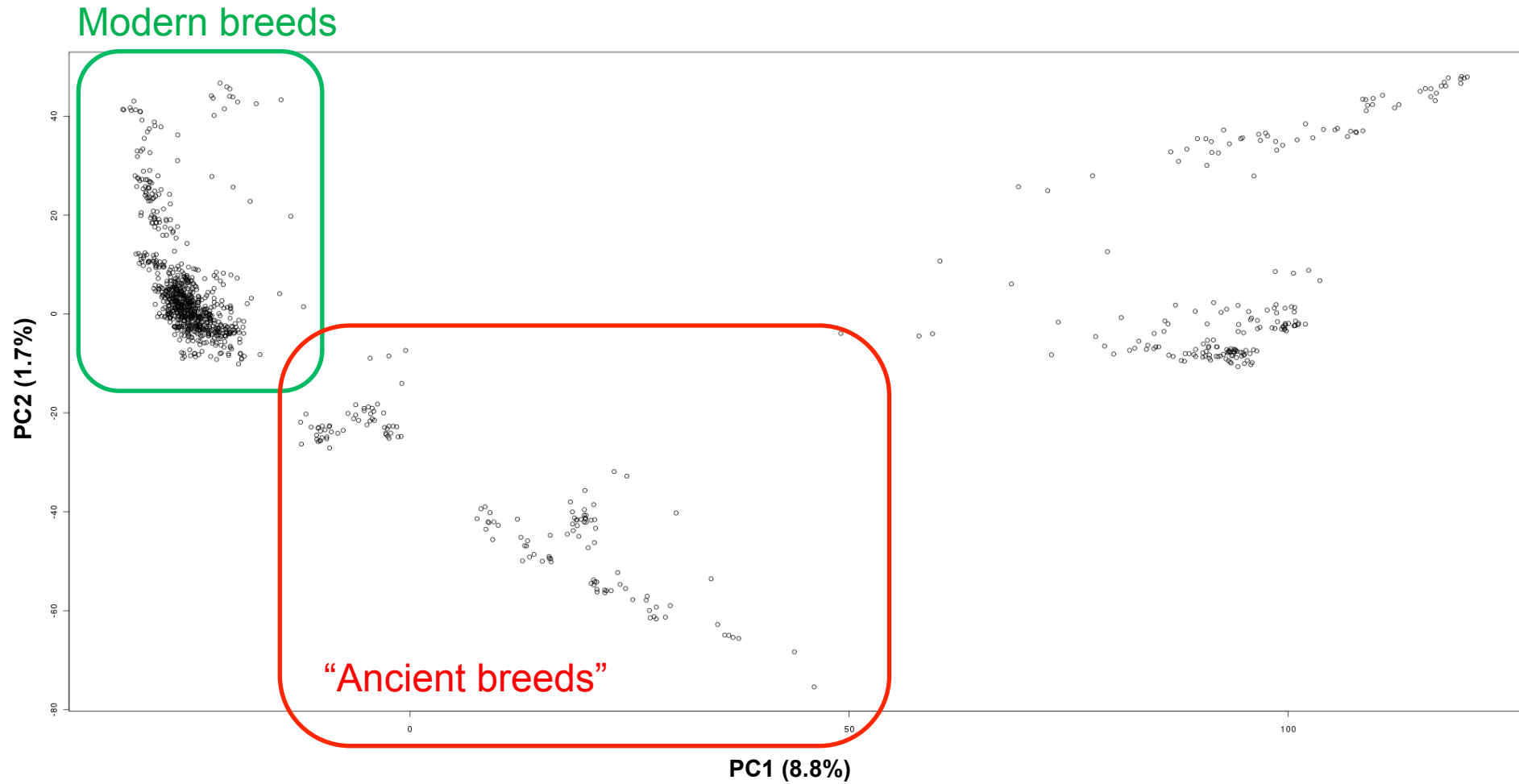
The CTVT founder

Modern breeds



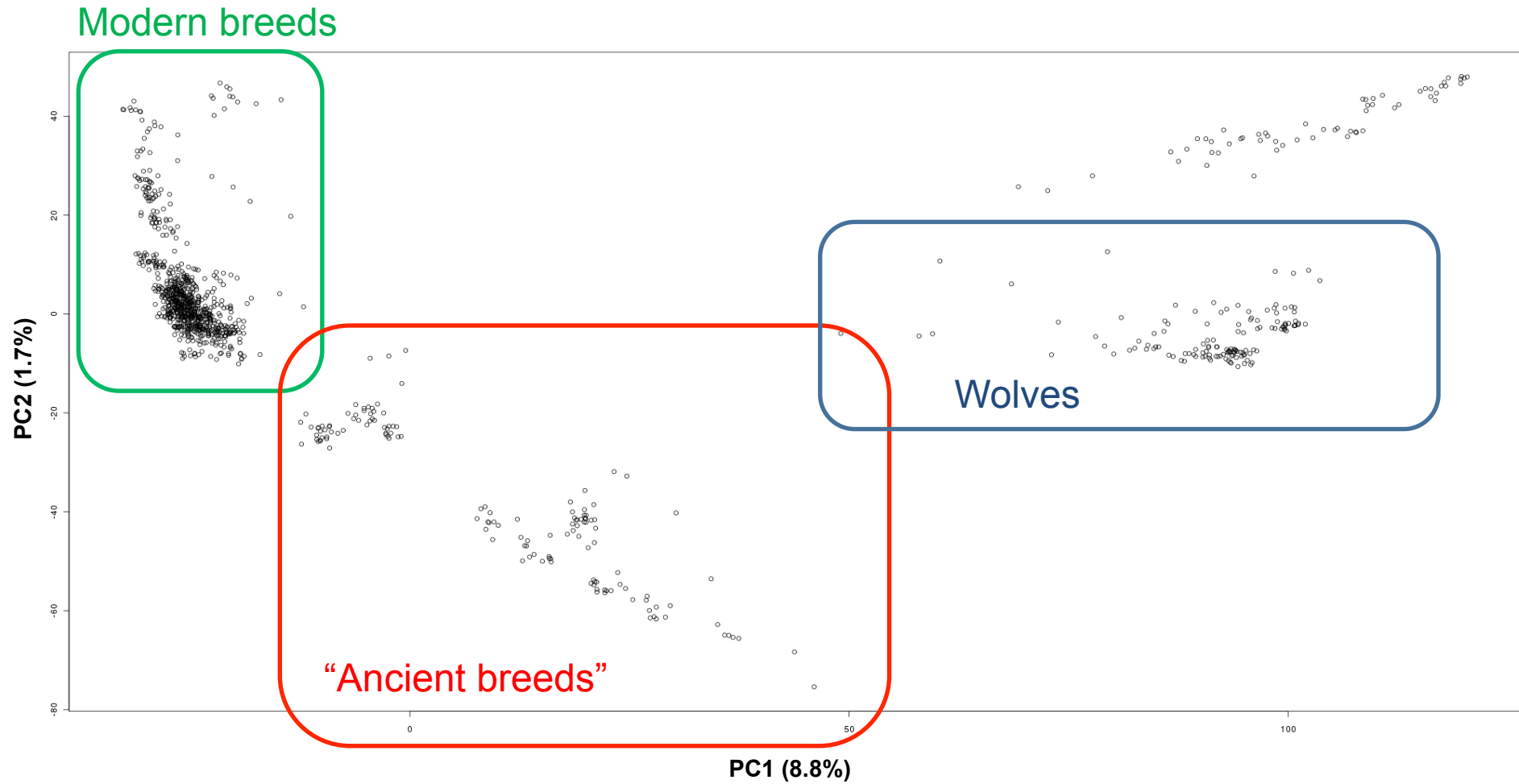
With data from vonHoldt et al, *Nature* 364:898 (2010) and vonHoldt et al, *Genome Research* 21:1294 (2011)

The CTVT founder



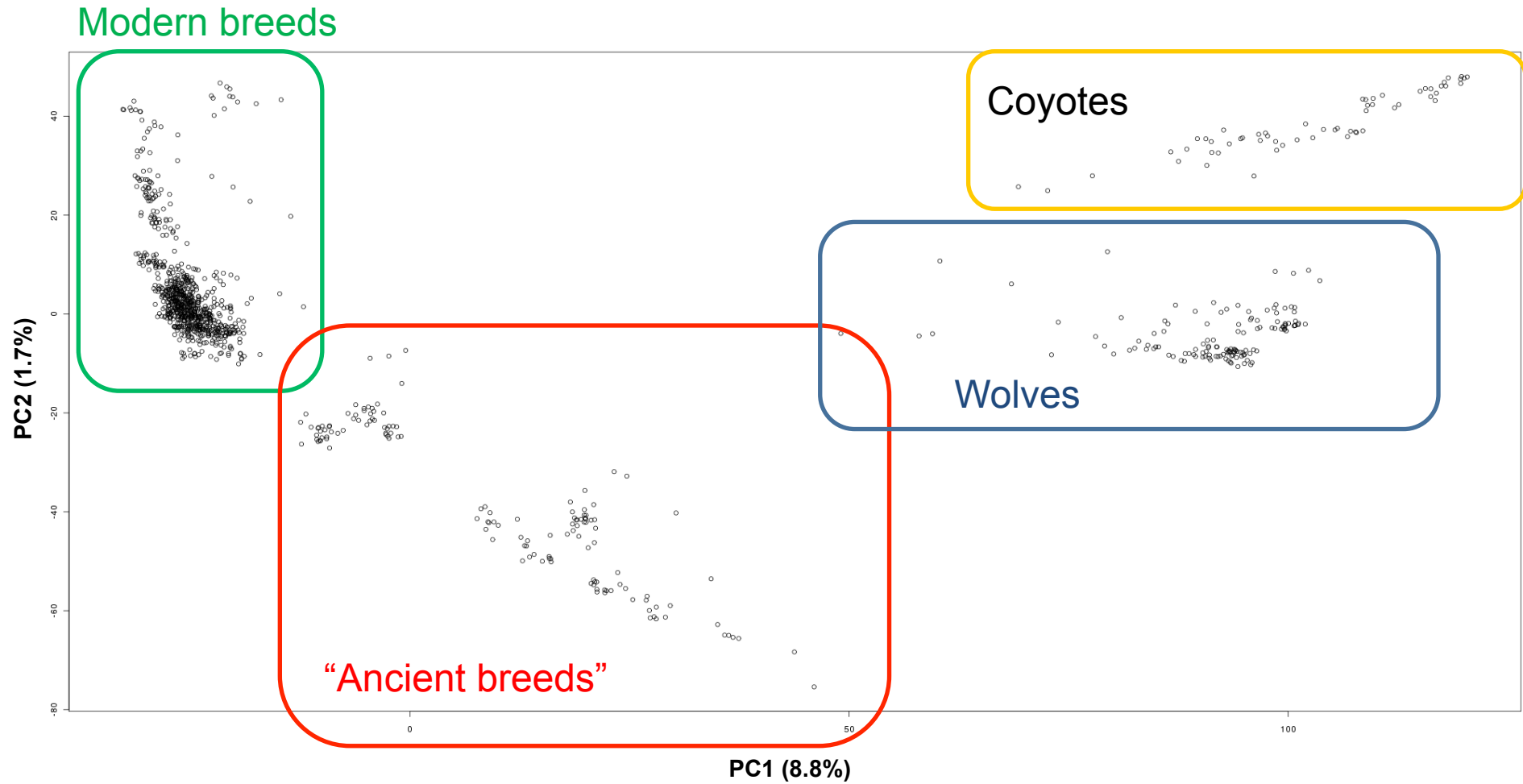
With data from vonHoldt et al, *Nature* 364:898 (2010) and vonHoldt et al, *Genome Research* 21:1294 (2011)

The CTVT founder



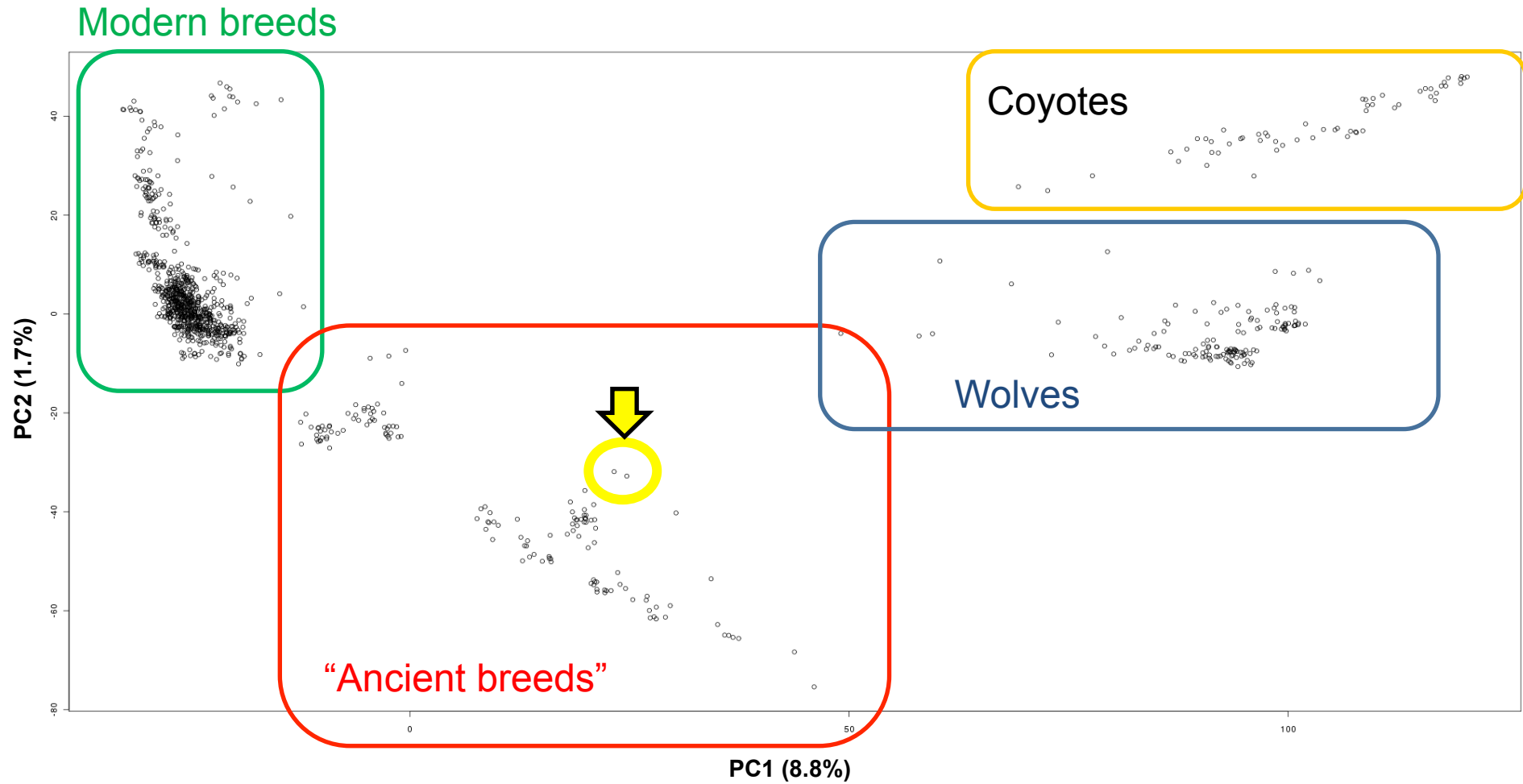
With data from vonHoldt et al, *Nature* 364:898 (2010) and vonHoldt et al, *Genome Research* 21:1294 (2011)

The CTVT founder



With data from vonHoldt et al, *Nature* 364:898 (2010) and vonHoldt et al, *Genome Research* 21:1294 (2011)

The CTVT founder was an “ancient breed” dog



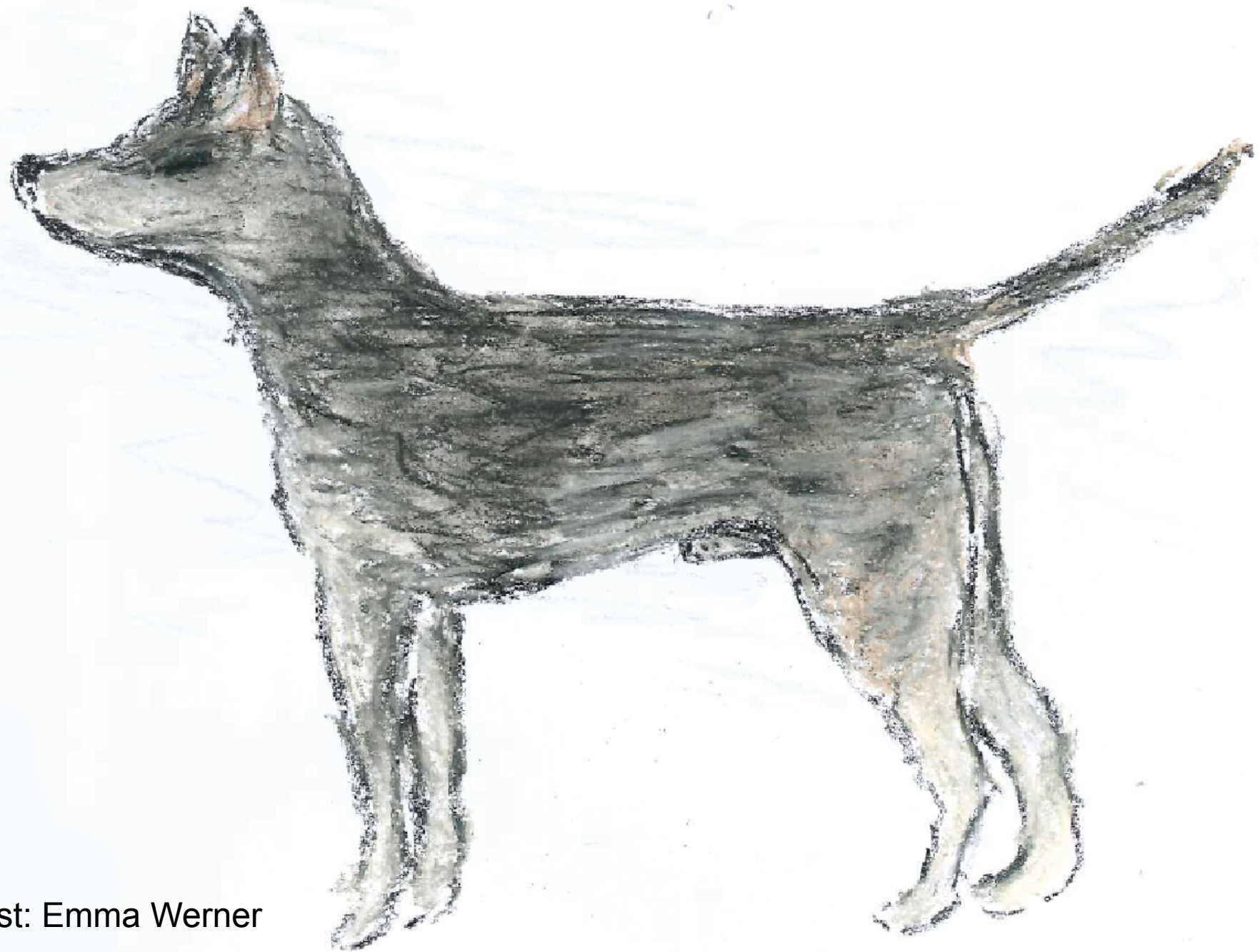
With data from vonHoldt et al, *Nature* 364:898 (2010) and vonHoldt et al, *Genome Research* 21:1294 (2011)



Images downloaded from www.akc.org

Founder dog phenotype

| Gene | Variant | CTVT genotype | CTVT phenotype | Reference |
|---------------|-----------------------|---------------|----------------------|-------------------------|
| <i>ASIP</i> | 24:23,393,510/14TA>GG | GG/GG | Agouti coat | Schmutz et al, 2007 |
| <i>FGF5</i> | 32:7,473,337G>T | G/G | Short hair | Cadieu et al, 2009 |
| <i>KRT71</i> | 27:5,542,806C>T | C/C | Straight hair | Cadieu et al, 2009 |
| - | 10:11,072,007C>T | T/T | Prick ears | Vaysse et al, 2011 |
| - | 10:11,169,956C>T | C/C | Medium to large size | Vaysse et al, 2011 |
| <i>BMP3</i> | 32:8,196,098C>A | C/C | Pointy snout | Schoenebeck et al, 2012 |
| <i>SLC1A2</i> | 18:35,681,332G>A | G/G | Less aggressive | Takeuchi et al, 2009 |
| <i>CDH2</i> | 7:63,867,472 | C/C | Lower risk of OCD | Dodman et al, 2010 |



Artist: Emma Werner

LETTER

doi:10.1038/nature11837

The genomic signature of dog domestication reveals adaptation to a starch-rich diet

Erik Axelsson¹, Abhirami Ratnakumar¹, Maja-Louise Arendt¹, Khurram Maqbool¹, Matthew T. Webster¹, Michele Perloski², Olof Liberg³, Jon M. Arnemo^{4,5}, Åke Hedhammar⁶ & Kerstin Lindblad-Toh^{1,2}

MGAM chr16:10,135,196C>T V1001I

C: Adaptation to starch-rich diet

T: Meat-lover

LETTER

doi:10.1038/nature11837

The genomic signature of dog domestication reveals adaptation to a starch-rich diet

Erik Axelsson¹, Abhirami Ratnakumar¹, Maja-Louise Arendt¹, Khurram Maqbool¹, Matthew T. Webster¹, Michele Perloski², Olof Liberg³, Jon M. Arnemo^{4,5}, Åke Hedhammar⁶ & Kerstin Lindblad-Toh^{1,2}

MGAM chr16:10,135,196C>T V1001I

CTVT: T/T

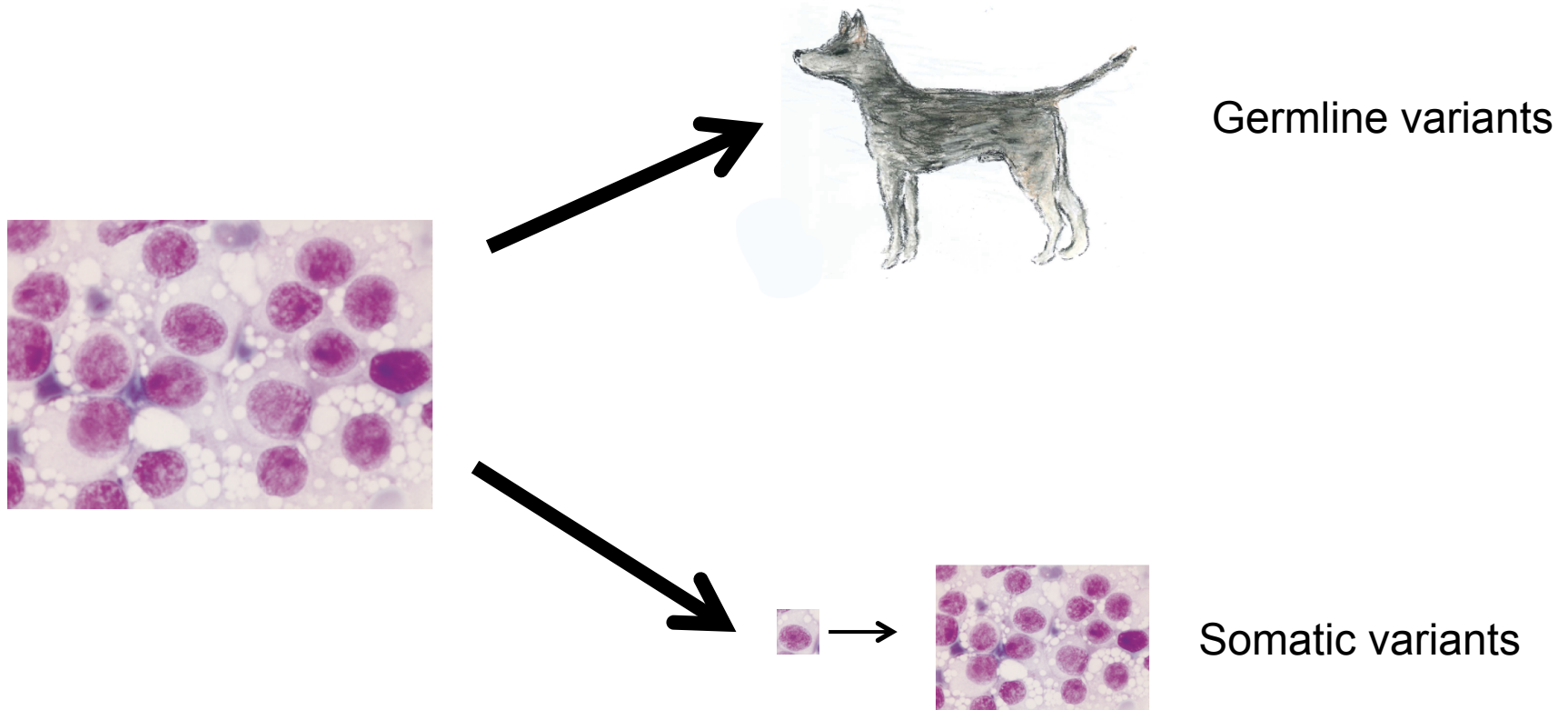
C: Adaptation to starch-rich diet

T: Meat-lover

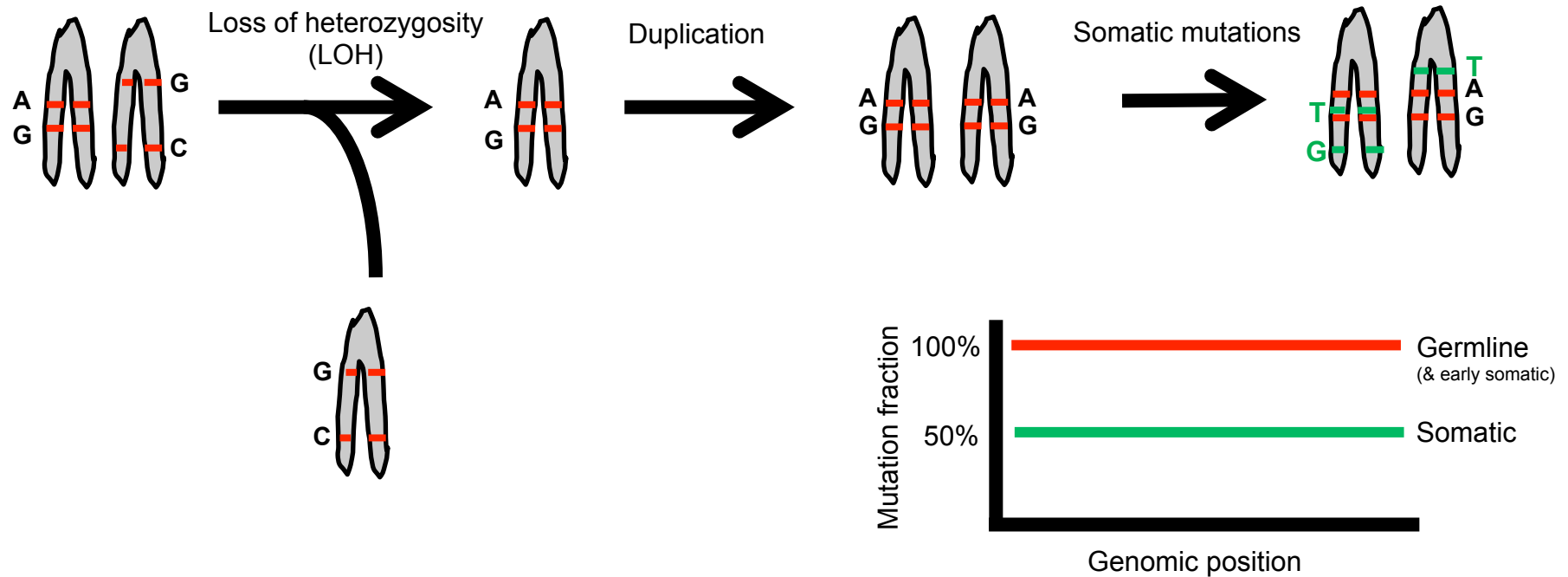


Artist: Emma Werner

The CTVT genome



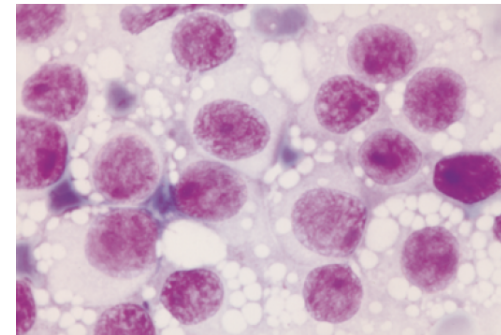
Using Loss of Heterozygosity (LOH) to identify somatic mutations



Number of somatic mutations in CTVT



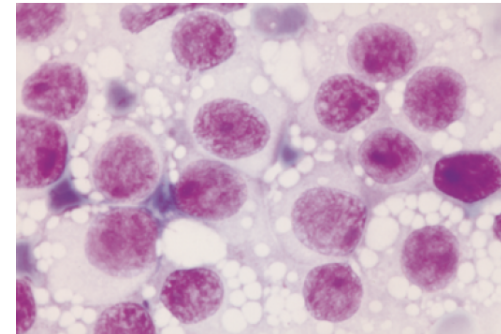
SOMATIC MUTATIONS



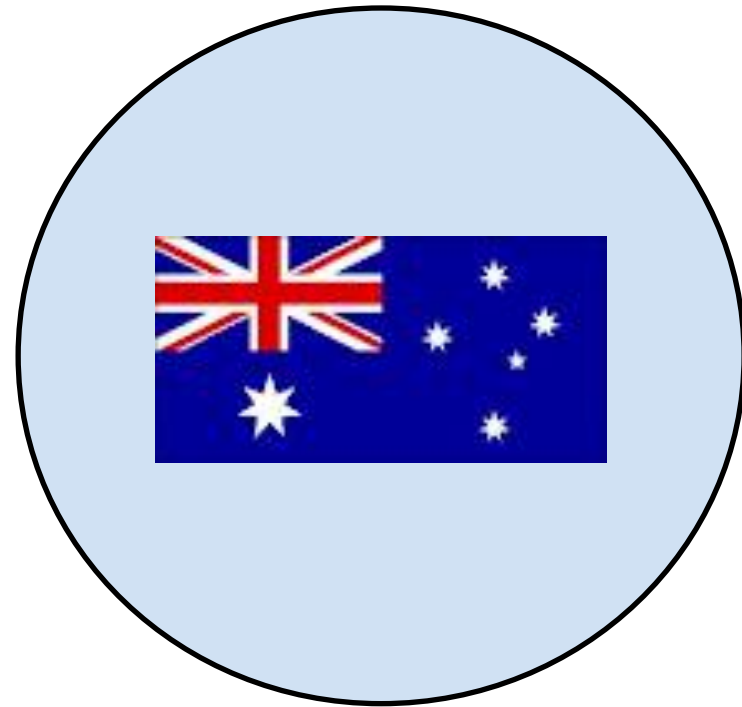
Number of somatic mutations in CTVT



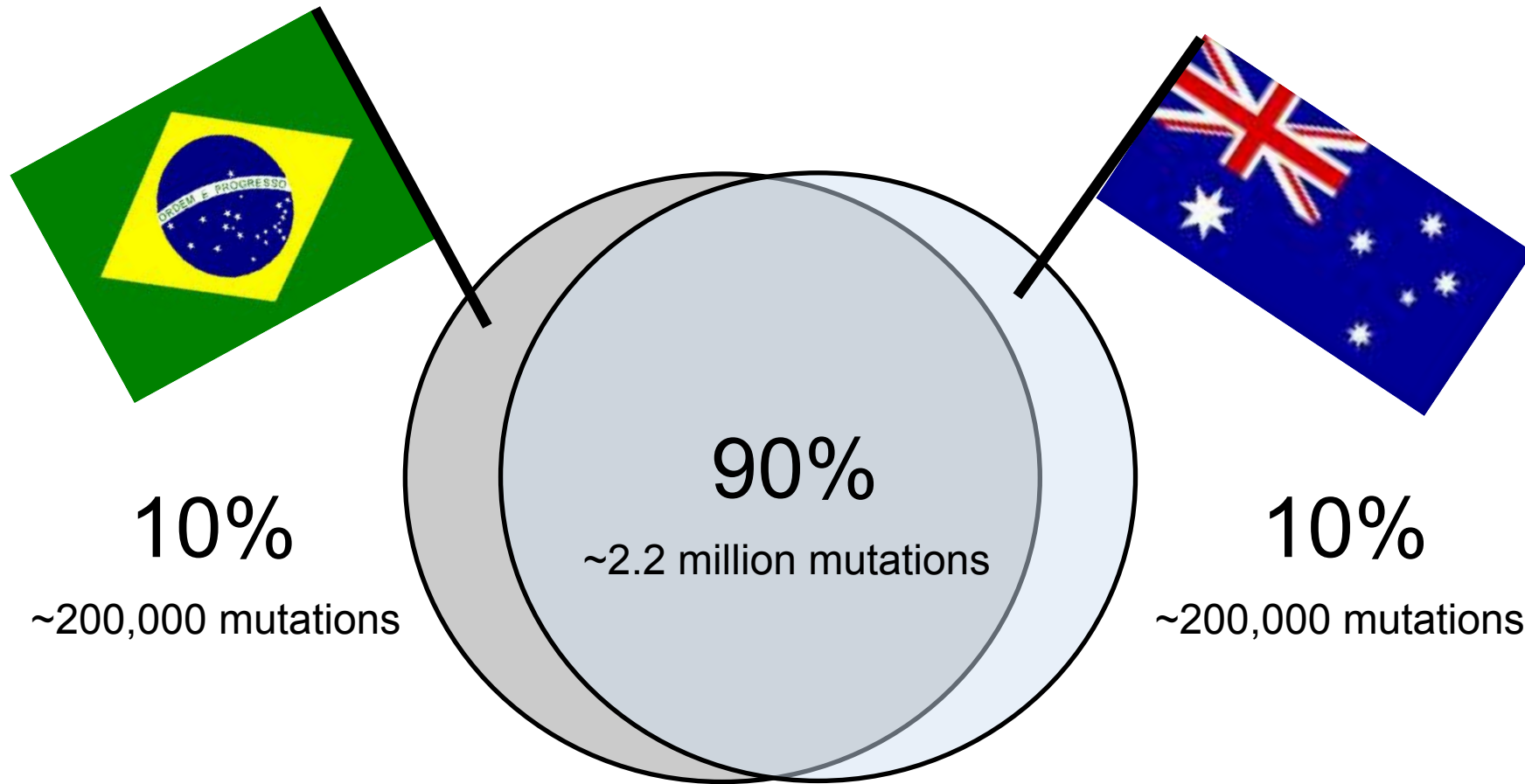
2 – 2.5 million
SOMATIC MUTATIONS



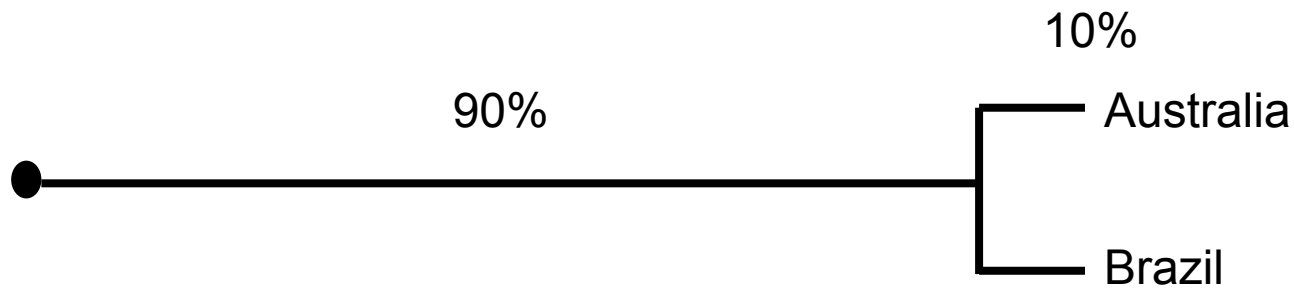
Variants in CTVT genomes



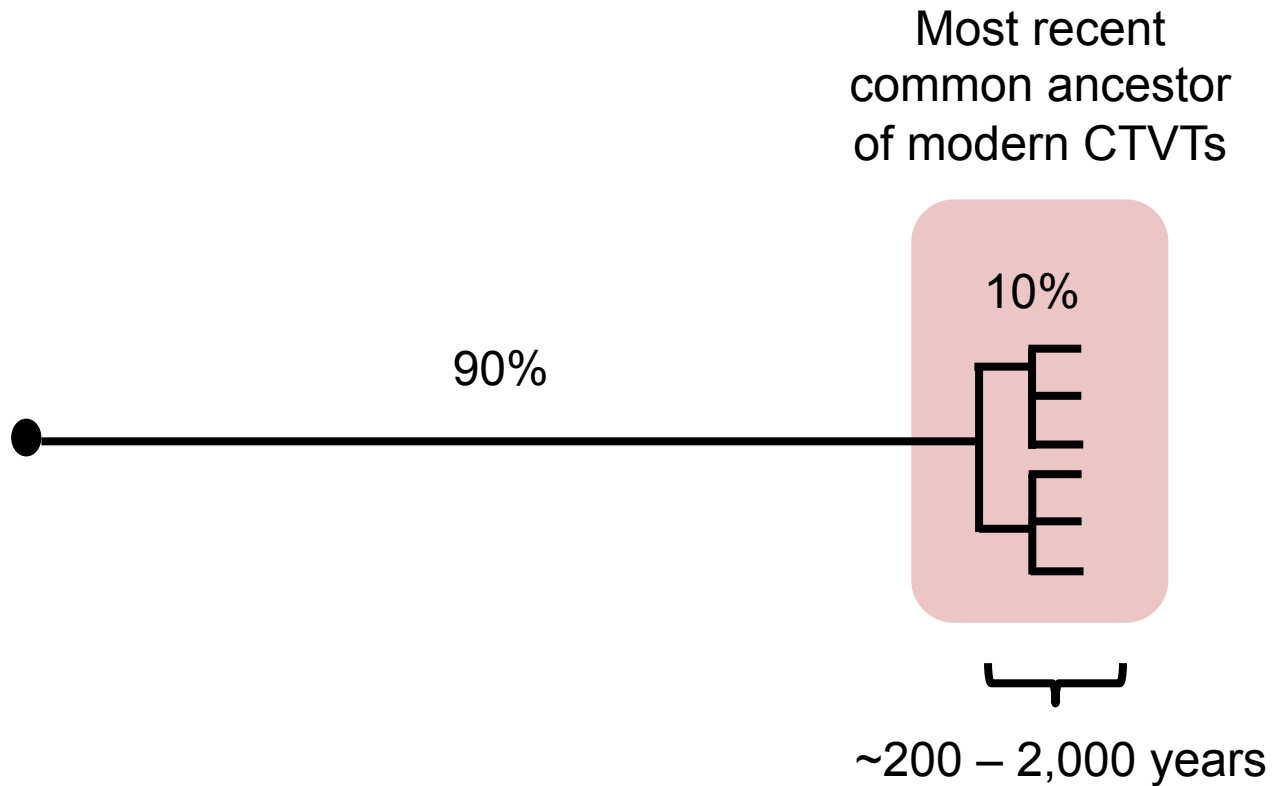
Variants in CTVT genomes



90% of mutations occurred before divergence of modern tumours



Age of CTVT



Age of CTVT

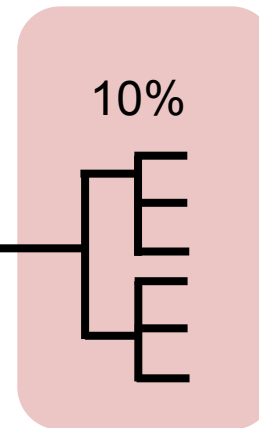


Founder lived
2,000 to 20,000
years ago



90%

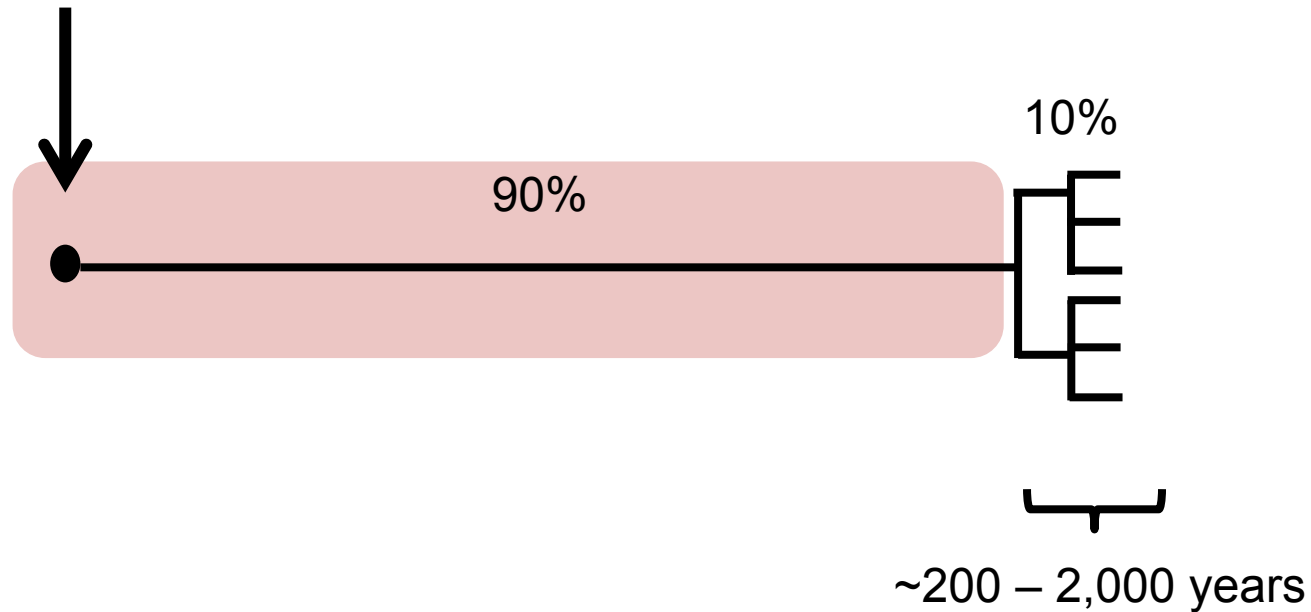
Most recent
common ancestor
of modern CTVTs



~200 – 2,000 years

For most of its history, CTVT was probably confined to an isolated dog population

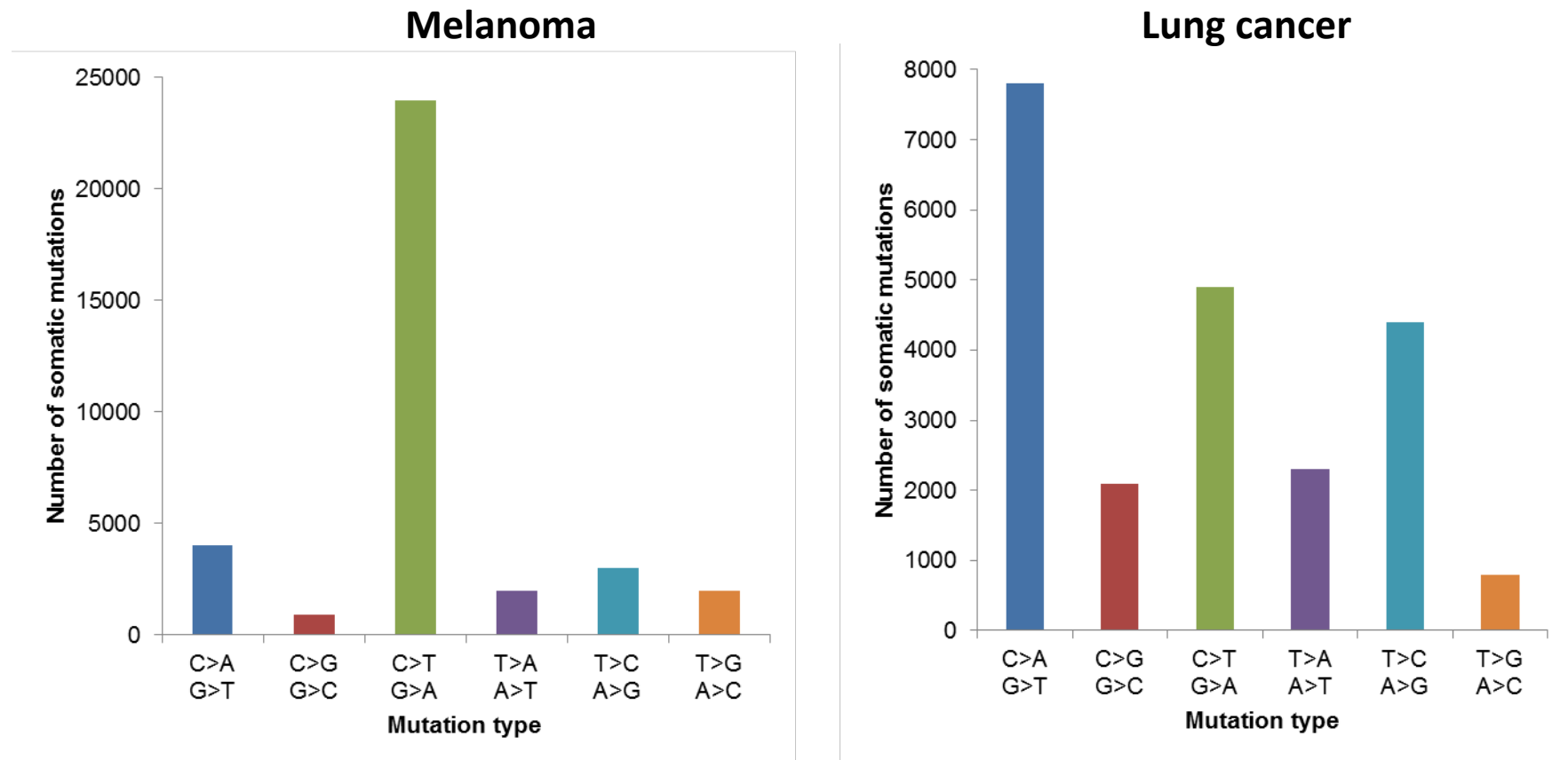
Founder lived
2,000 to 20,000
years ago



What caused the mutations in
CTVT?

Mutation spectrum

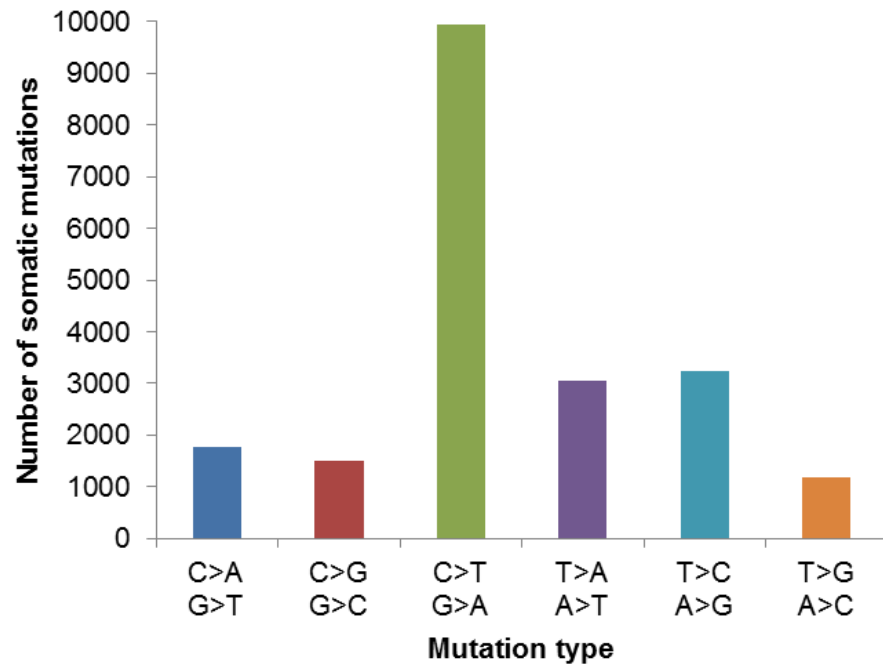
Type of mutation can give clues as to processes that caused cancer



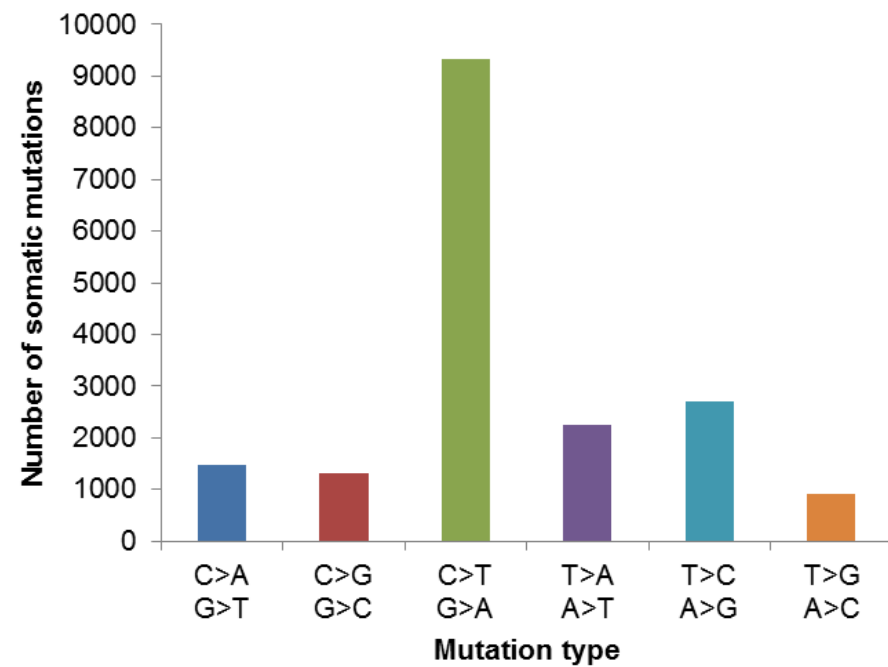
Data adapted from Pleasance et al *Nature* 463:184 (2010) and Pleasance et al *Nature* 463:191 (2010)

Mutation spectrum in CTVT

Australian CTVT



Brazilian CTVT





Artist: Emma Werner

Clonally transmissible cancers

DFTD

- Recent (<20years)
- Metastasis common
- Transmitted by biting
- Facial tumour
- Neural crest origin
- Unresponsive to therapy

CTVT

- Old (>1,000 years)
- Metastasis rare
- Sexually transmitted
- Genital tumour
- Histiocytic origin
- Very sensitive to therapy

Clonally transmissible cancers

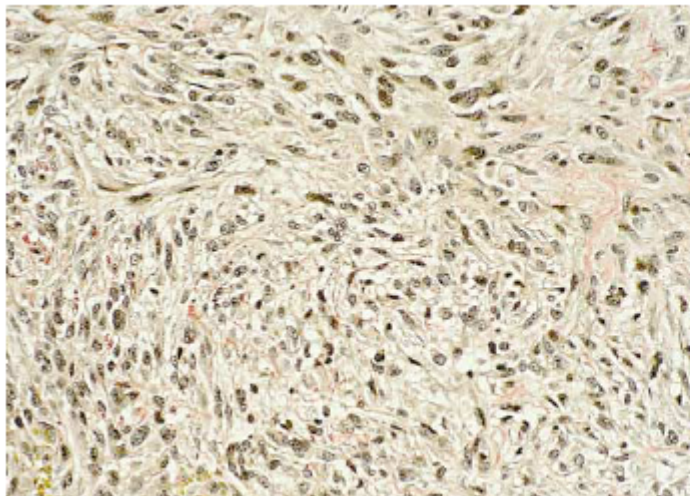


Images:
Tasmanian Devil Conservation Park
www.animalswallpaper.info

Brief Report

GENETIC ANALYSIS OF A SARCOMA
ACCIDENTALLY TRANSPLANTED FROM
A PATIENT TO A SURGEON

HERMINE-VALERIA GÄRTNER, M.D., CHRISTIAN SEIDL, M.D.,
CHRISTINE LUCKENBACH, PH.D., GEORG SCHUMM, M.D.,
ERHARD SEIFRIED, M.D., HORST RITTER, M.D.,
AND BURKHARD BÜLTMANN, M.D.



Patient's Sarcoma

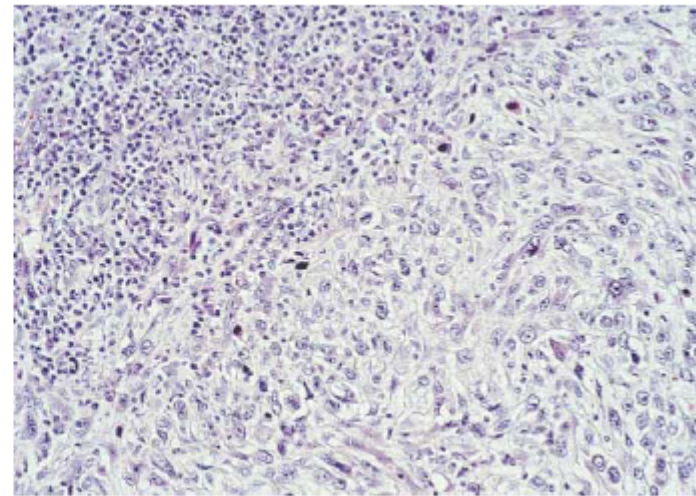
odic acid-Schiff, and van Gieson's stain. Immunostaining was performed with the avidin-biotin-peroxidase method, with antibodies against vimentin, lysozyme, α_1 -antitrypsin, α_1 -antichymotrypsin, keratin, endomysial antigens, S100, actin, and desmin (all antibodies were obtained from Dako, Glestrup, Denmark).

Isolation of DNA

Genomic DNA from peripheral-blood samples was isolated by the "salting-out" method.⁶ DNA from paraffin-embedded tumor and tissue samples was extracted according to a modification⁷ of the method of Goelz et al.⁸

Analysis of Short Tandem-Repeat Polymorphisms

Short tandem-repeat polymorphisms of the loci HUMTH01, HUMCYR04, and HUMACTBP2 were amplified by the polymerase chain reaction (PCR) with fluorescence-labeled primers. Primer sequences were chosen from published sequences.⁹⁻¹¹ The 5' primer for HUMCYR04 was labeled with 5-carboxyfluorescein, whereas



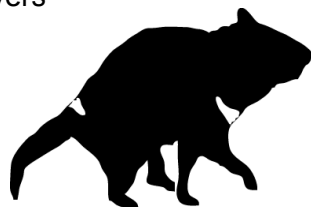
Surgeon's Sarcoma

Wellcome Trust Sanger Institute

Mike Stratton
Zemin Ning
Ludmil Alexandrov
Beiyuan Fu
Zihao Ding
Caitlin Stewart
Bee Ling Ng
Bronwen Aken
Graham Bignell
William Cheng
Tom Connor
Yong Gu
Simon Harris
John Marshall
David Wedge
Keiran Raine
Steve Searle
Simon White
Nigel Carter
Andy Futreal
Peter Campbell
Fengtang Yang
Erin Pleasance
Thierry Voet
Chris Greenman
Stephen Rice
Sue Bumpstead
Emma Werner
Ultan McDermott
Stacey Price

Illumina

Ole Schulz-Trieglaff
Markus Bauer
Matt Hims
Sergii Ivakhno
Wendy Wong
Jennifer Becq
Keira Cheetham
Tony Cox
Russell Grocock
Irina Khrebtukova
Zoya Kingsbury
Shujun Luo
David McBride
Lisa Murray
Isabella Rasolonjatovo
Richard Shaw
Philip Tedder
Carolyn Tregidgo
Niall Gormley
Sean Humphray
Gary Schroth
Geoffrey Smith
Kevin Hall
David Bentley
Dirk Evers



Wellcome Trust Sanger Institute Cancer Genetics and Genomics Group
Wellcome Trust Sanger Institute Sequencing Centre
Wellcome Trust Sanger Institute Central Informatics Team

Australian National University

Amber Alsop
Janine Deakin
Hannah Bender

Walter and Eliza Hall Institute of Medical Research

Zhi-Ping Feng
Mark Kowarsky
Tony Papenfuss

Menzies Research Institute Tasmania

Alex Kreiss
Greg Woods

University of Tasmania

Rodrigo Hamede
Menna Jones

Save the Tasmanian Devil Program

Kate Swift
Bobby Hua
Robyn Taylor
Stephen Pyecroft
Colette Harmsen
Sarah Peck
Anne-Maree Pearse
Chris Boland
Trapping teams and field assistants

University of Cambridge, Department of Veterinary Medicine

Willem Rens
Malcolm Ferguson-Smith

Hamish McCallum (Griffith University)
Kathy Belov (University of Sydney)
David Obendorf (Veterinary pathologist)
Matthew Breen (North Carolina State University)
Albert Vilella (European Bioinformatics Institute)

Wellcome Trust Sanger Institute

Mike Stratton
David Wedge
Emma Werner
Andrea Strakova
Ludmil Alexandrov
Zemin Ning
Keiran Raine
Adam Shlien
Erin Pleasance
Peter Campbell
Fengtang Yang
Bei Yuan Fu
Serena Nik-Zainal
Graham Bignell
Andrew King

Cancer Genetics and Genomics Group
Sequencing Centre
Central Informatics Team



Thank you...

Malcolm Brearley
Jane Dobson
John Cooper
Fernando Constantino-Casas
Michelle Morters
Willem Rens
Malcolm Ferguson-Smith
David Sargan
Inigo Martincorena
Austin Burt
Robin Weiss
Ariberto Fassati
Clare Rebbeck
Claudio Murgia
Bridgett vonHoldt
Bob Wayne
Stephan Beck
Chris Greenman
Mohammed Fazil
Oliver Walkinton
Debra Kamstock
Patrick Kelly
Sally Nixon
Kevin Xie
Michael Meyer
David Walker
Gerry Polton
Branwen Hennig
Christine Kent
Matthew Breen

Frances Nargi
Karina Ferreira de Castro
Saskia Karius
Erika Flores
Hanna Lentz
Kati Loeffler
Ilona Otter
Wendy Edney
Antonio Ortega-Pacheco
Maria Peleteiro
Ceseltina Semedo
Joaquim Henriques
Gabriele Marino
Stephen Cutter
Jude Mulholland
Skye Turner
Anne Corrigan
Ismail Thoya
Hugh Cran
Sophie Constable
Ruth Pye
Jan Allen
Ted Donelan
Andrigo Barboza de Nardi
Karen Allum
Simon Martinez
Oscar Rarieya
Laura Delgadillo
Scott Moroff
Jack Reece

Collaborating Organisations

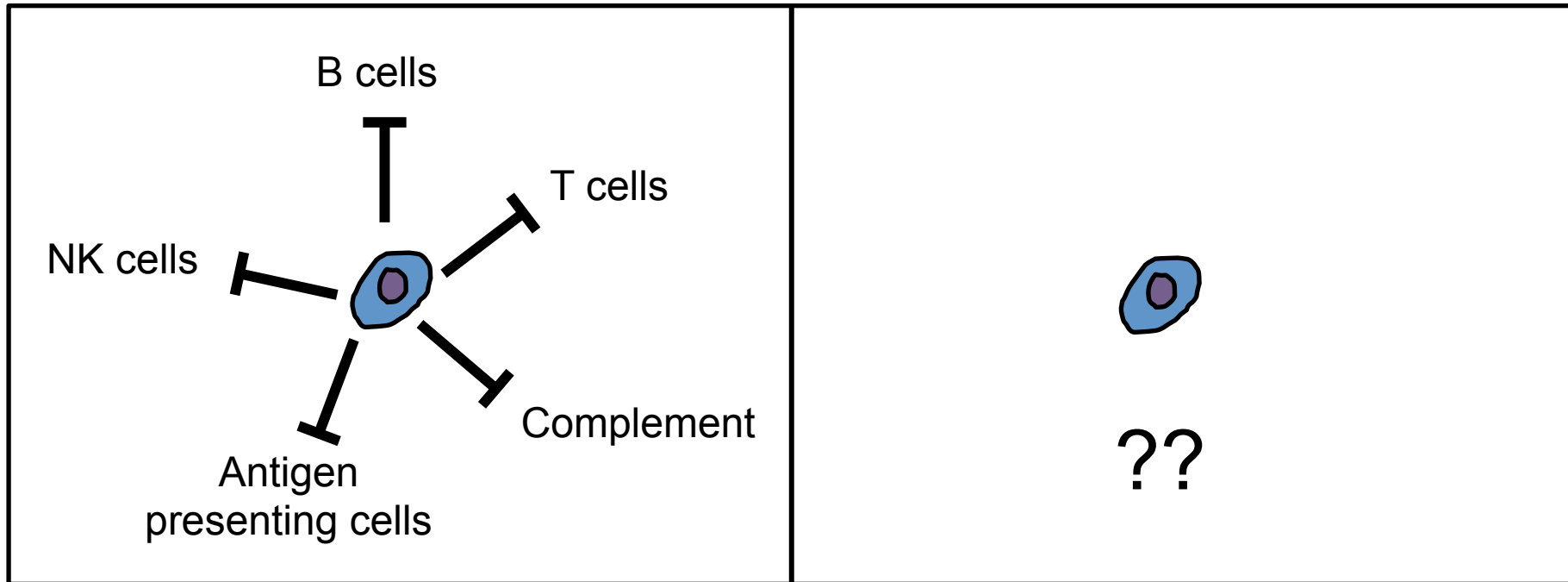
Vets Beyond Borders
World Vets
SCAD Thailand
AMRRIC (Australia)
IFAW
Animal Care Association, The Gambia
St George's University, Grenada
Colorado State University
University of Messina
University of Yucatan
Bons Amigos (Cape Verde)
University of Cambridge, Vet School
European Bioinformatics Institute
Imperial College London
University College London
University of Franca
Sao Paulo State University
Animal Protection Society Samoa
Soi Dog Foundation (Thailand)
Worldwide Veterinary Service
PAWS (Mauritius)
University of Ibadan
University of Nairobi
Usmanu Danfodiyo University
KSPCA (Kenya)
Help in Suffering (India)

Thank you!



Photo: Sarah Peck

Immune evasion



Canine transmissible venereal tumour
(CTVT)

Tasmanian devil facial tumour disease
(DFTD)