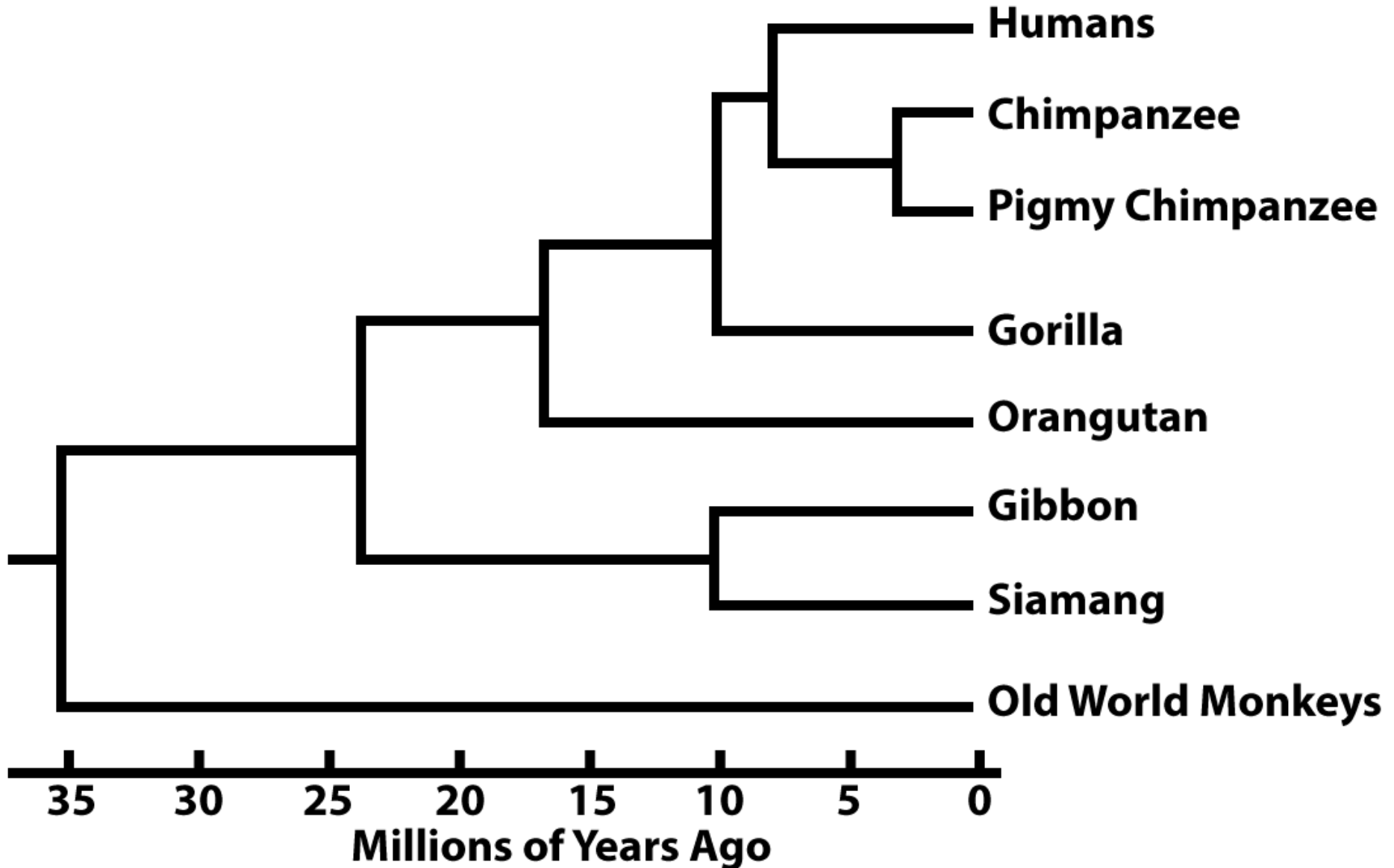


Two Sets of Intertwined Stories



1. EVOLUTIONARY HISTORY

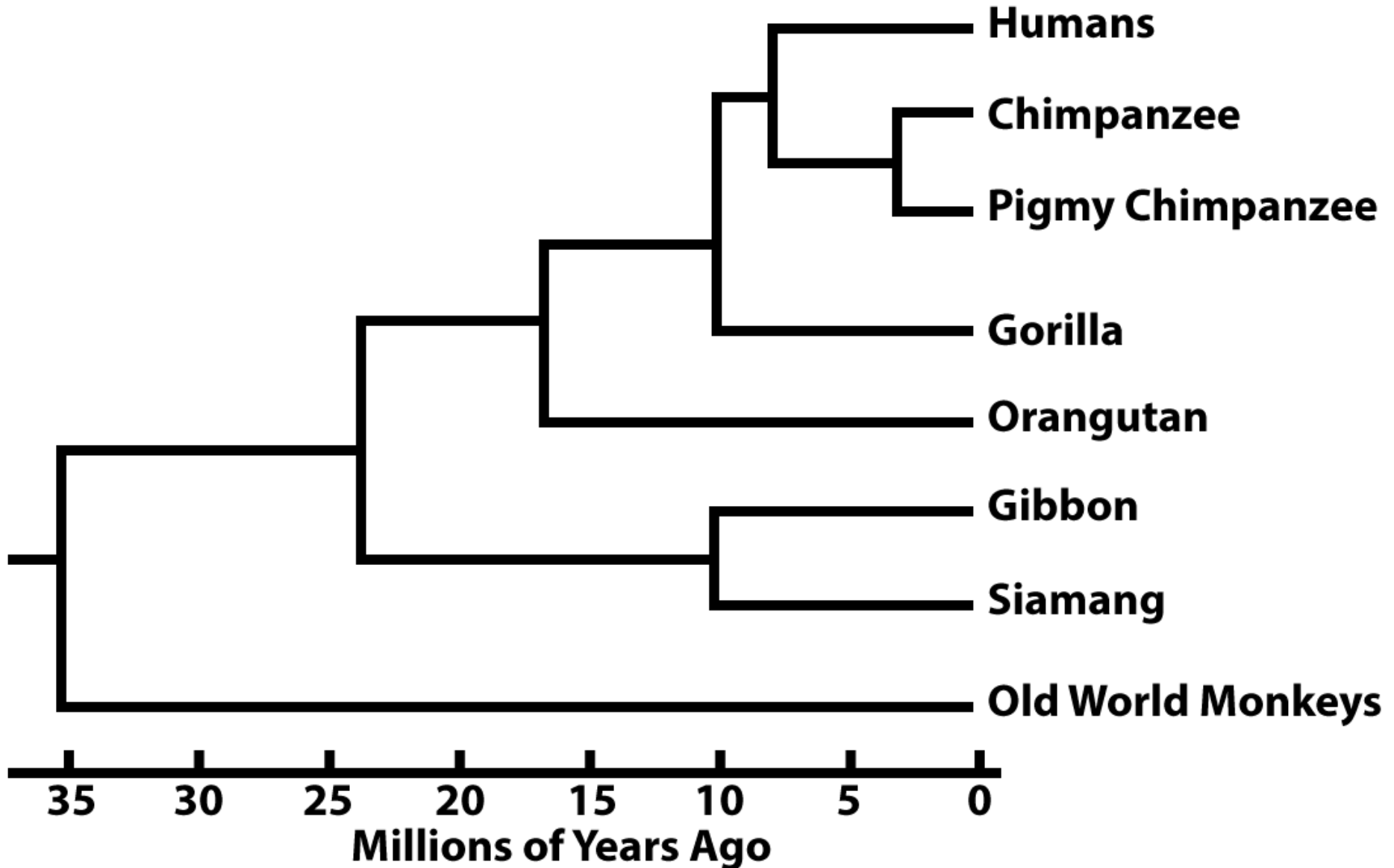


2. BROADER BIODIVERSITY



ALEXIS ROCKMAN, THE ECOTOURIST

1. EVOLUTIONARY HISTORY

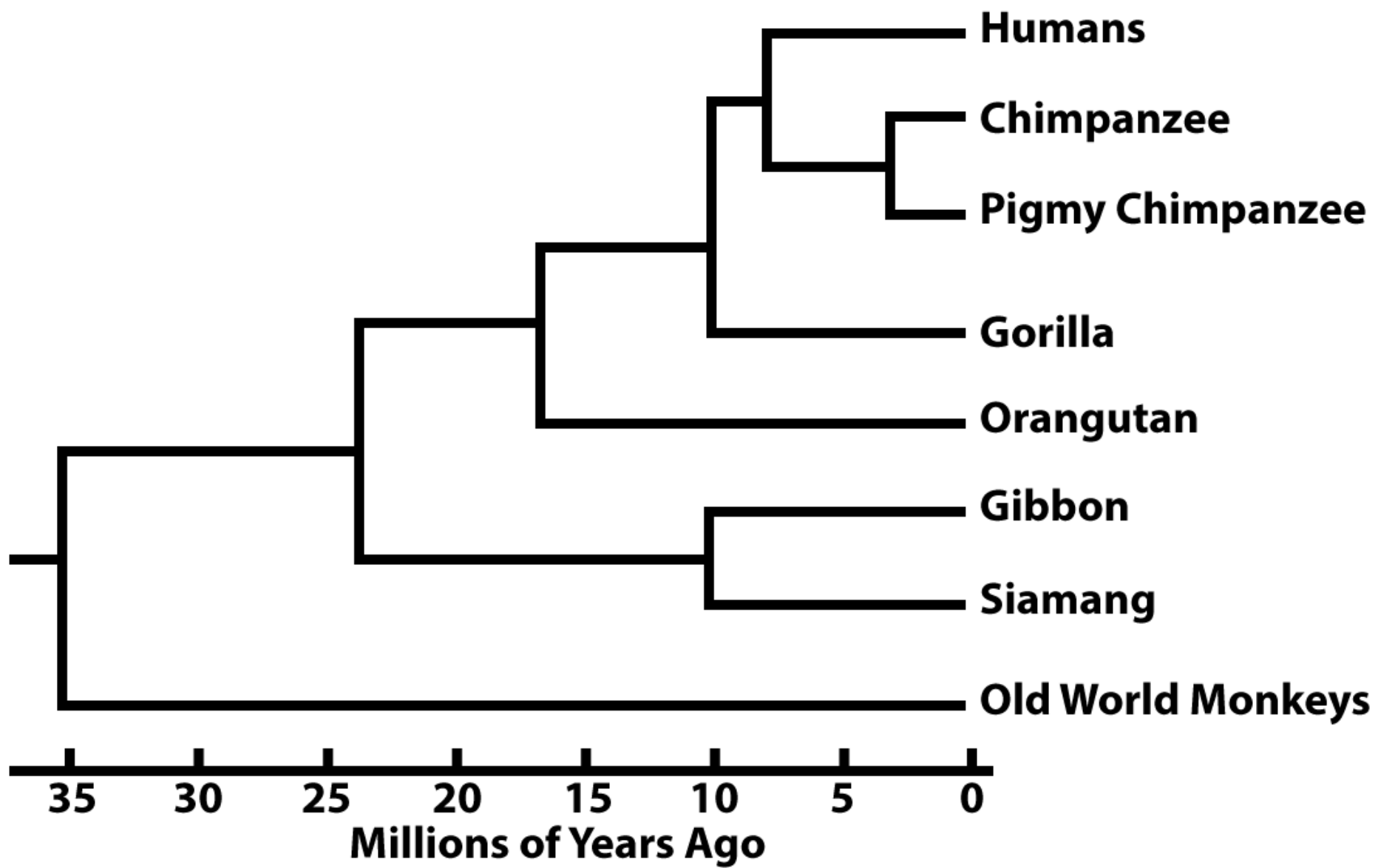


1. EVOLUTIONARY HISTORY (HOUSE)

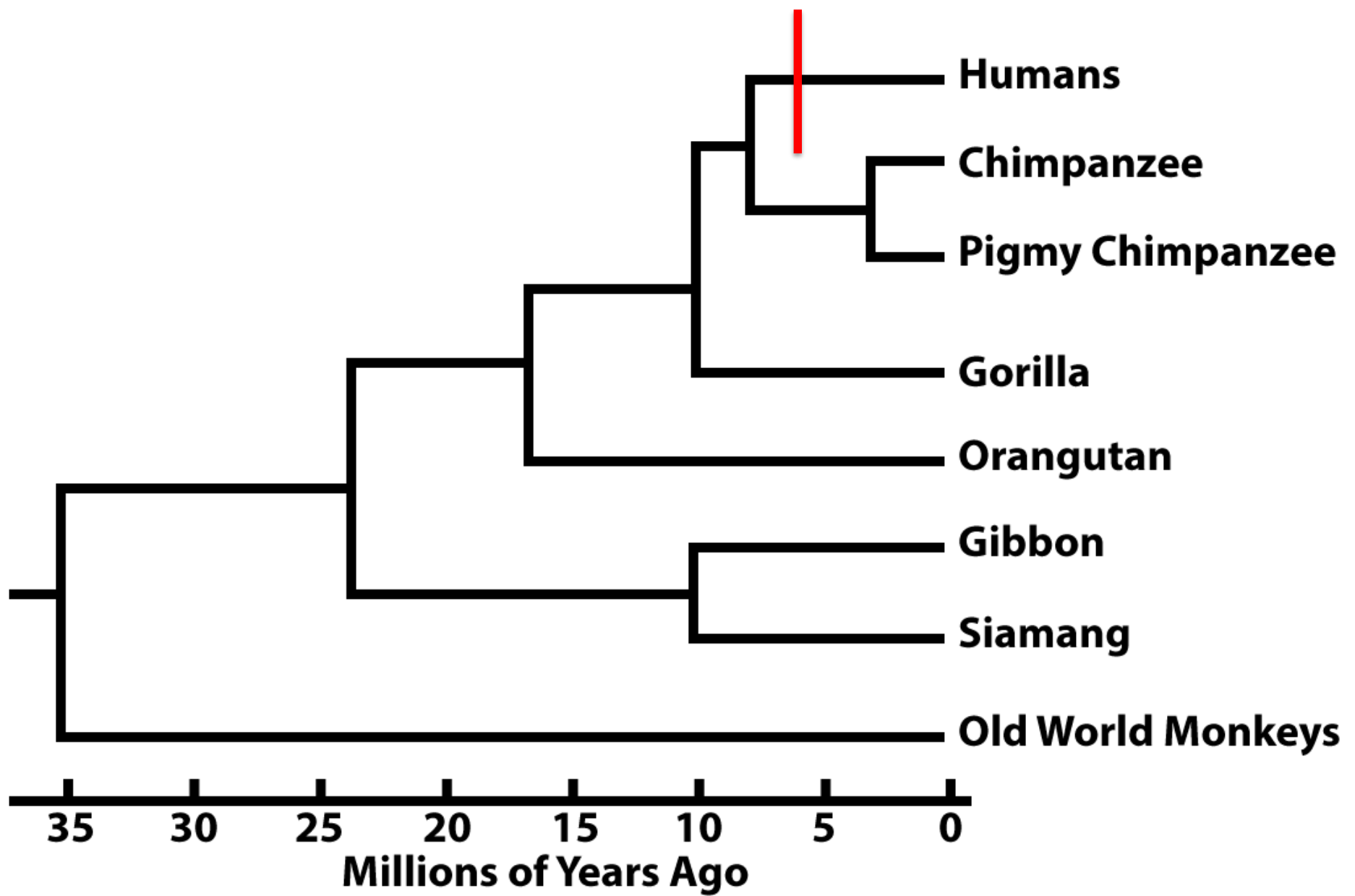


1. EVOLUTIONARY HISTORY (HOUSE MICROBIOME)





Origin of Houses





Western Tanzania



50

Kilometers

Legend

- Ugalla core area
- National Park
- Lake Tanganyika



Research



Check for updates

Cite this article: Thoemmes MS *et al.* 2018

Ecology of sleeping: the microbial and arthropod associates of chimpanzee beds.

R. Soc. open sci. **5**: 180382.

<http://dx.doi.org/10.1098/rsos.180382>

Received: 9 March 2018

Accepted: 13 April 2018

Subject Category:

Biology (whole organism)

Subject Areas:

ecology/evolution

Ecology of sleeping: the microbial and arthropod associates of chimpanzee beds

Megan S. Thoemmes¹, Fiona A. Stewart^{5,6,7},
R. Adriana Hernandez-Aguilar^{5,8}, Matthew A.
Bertone², David A. Baltzegar^{3,4}, Russell J. Borski³,
Naomi Cohen⁵, Kaitlin P. Coyle³, Alexander K. Piel^{5,6}
and Robert R. Dunn^{1,9}

¹Department of Applied Ecology and Keck Center for Behavioral Biology, ²Department of Entomology and Plant Pathology, ³Department of Biological Sciences, and ⁴Genomic Sciences Laboratory, Office of Research, Innovation and Economic Development, North Carolina State University, Raleigh, NC, USA

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⁶School of Natural Sciences and Psychology, Liverpool John Moores University, Liverpool, UK

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⁸Centre for Ecological and Evolutionary Synthesis, Department of Biosciences,







DR. MATT BERTONE, ENTOMOLOGIST AND PHOTOGRAPHER (SELFIE)

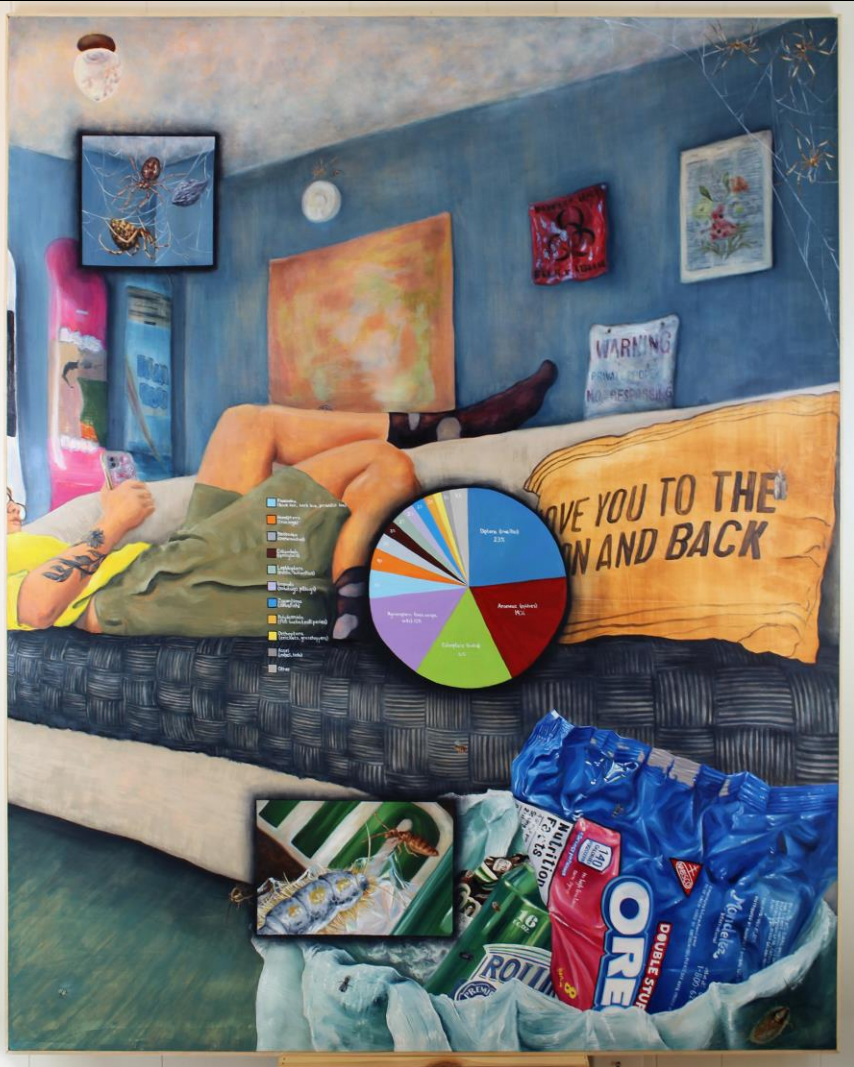




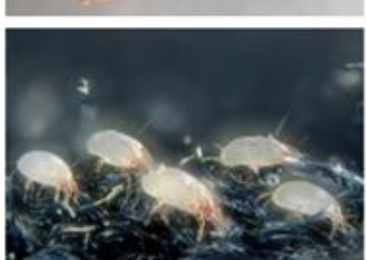
THESE AND OTHER ARTHROPOD IMAGES BY DR. MATT BERTONE.



PAINTING BY ELEANOR Q. C. OLSON



PAINTING BY ELEANOR Q. C. OLSON



More than 1000 *species*



A large proportion of which are house dependent

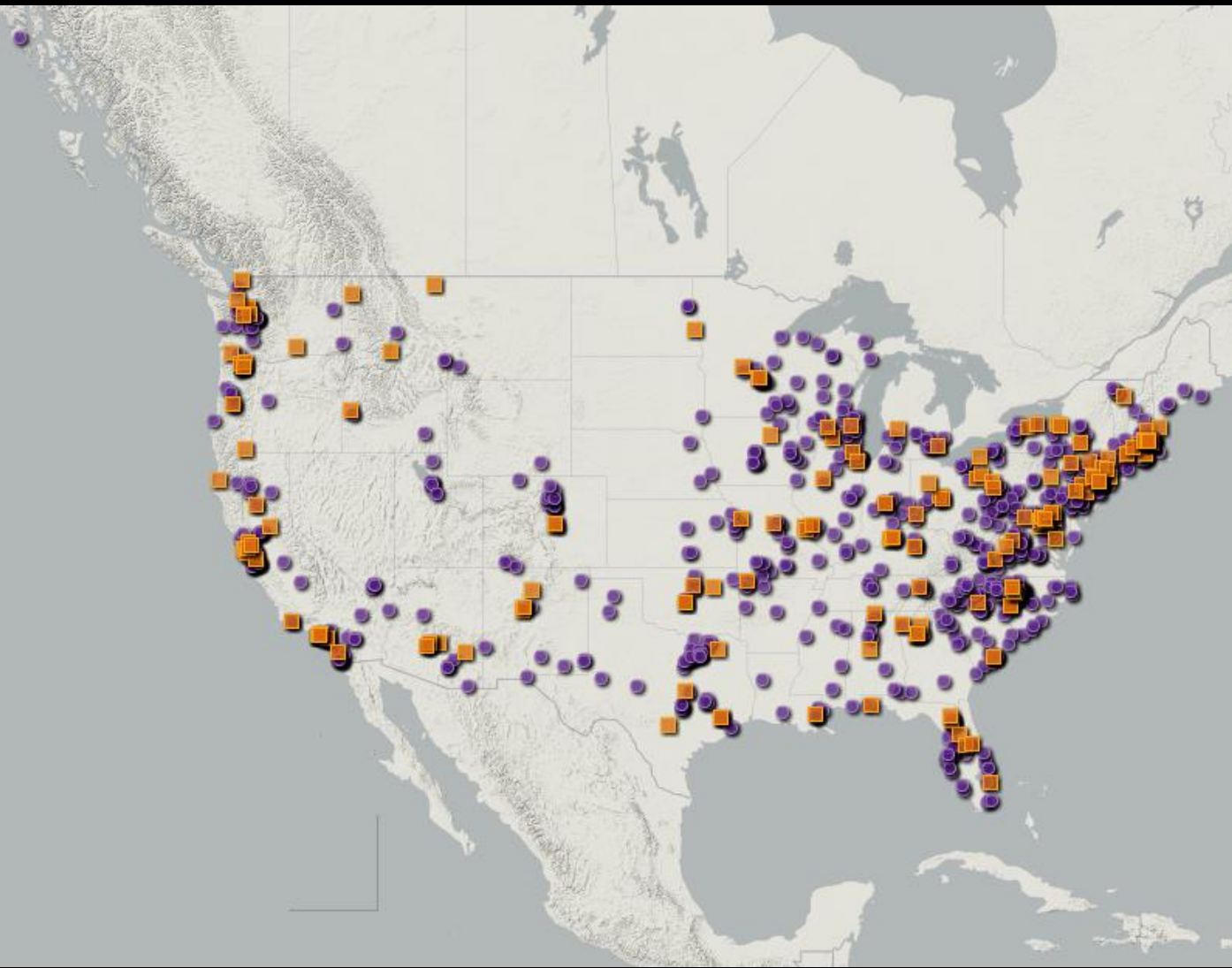
Photos by Matt Bertone



Photos by Matt Bertone



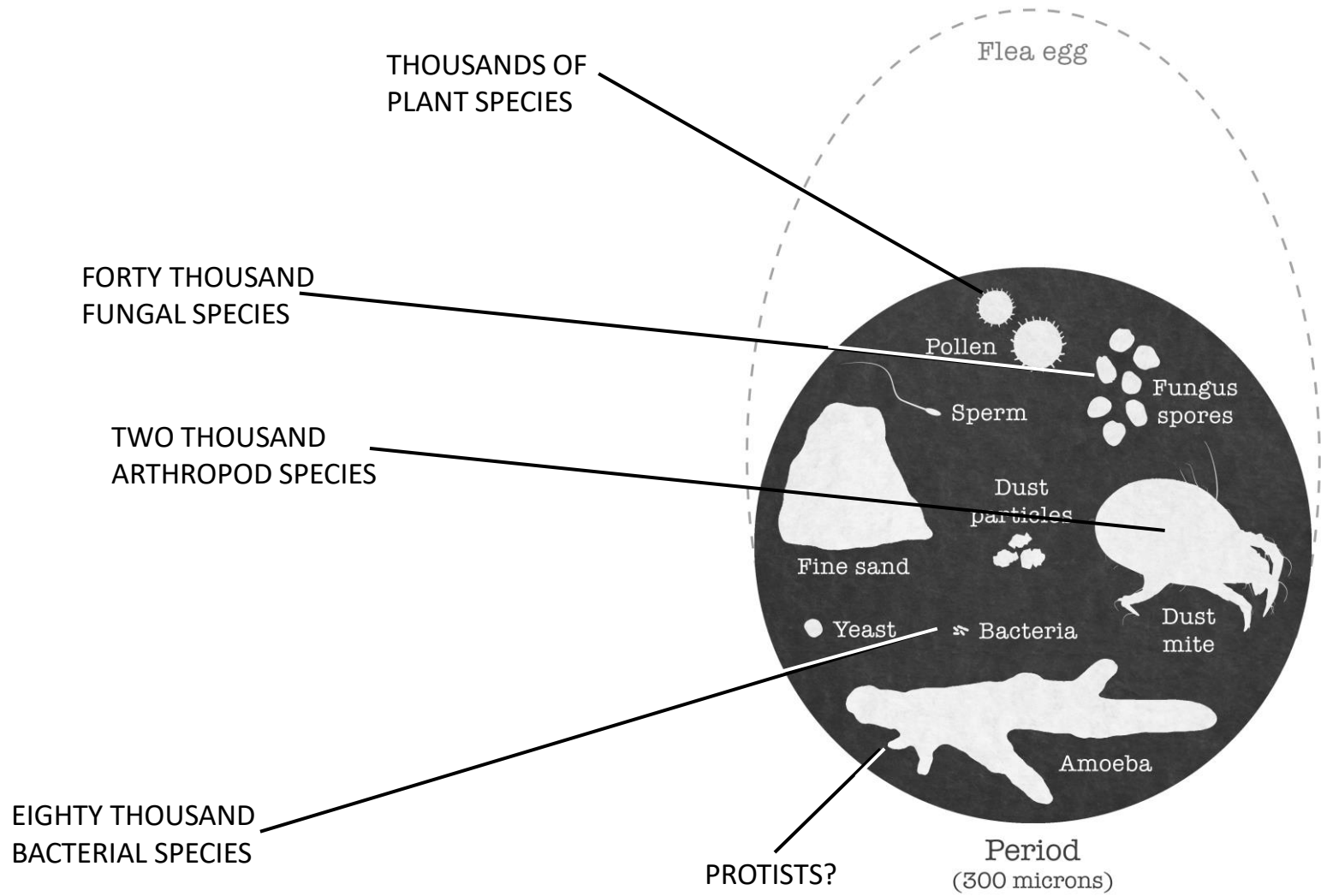




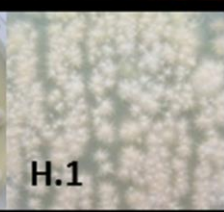
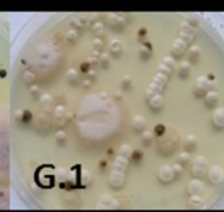
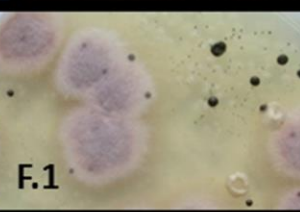
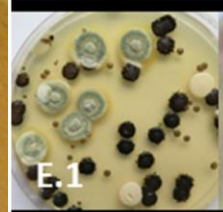
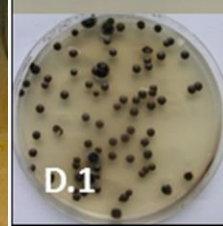
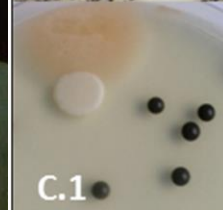
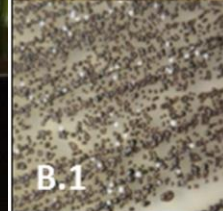
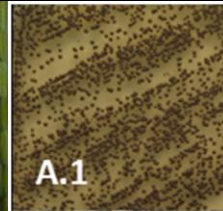


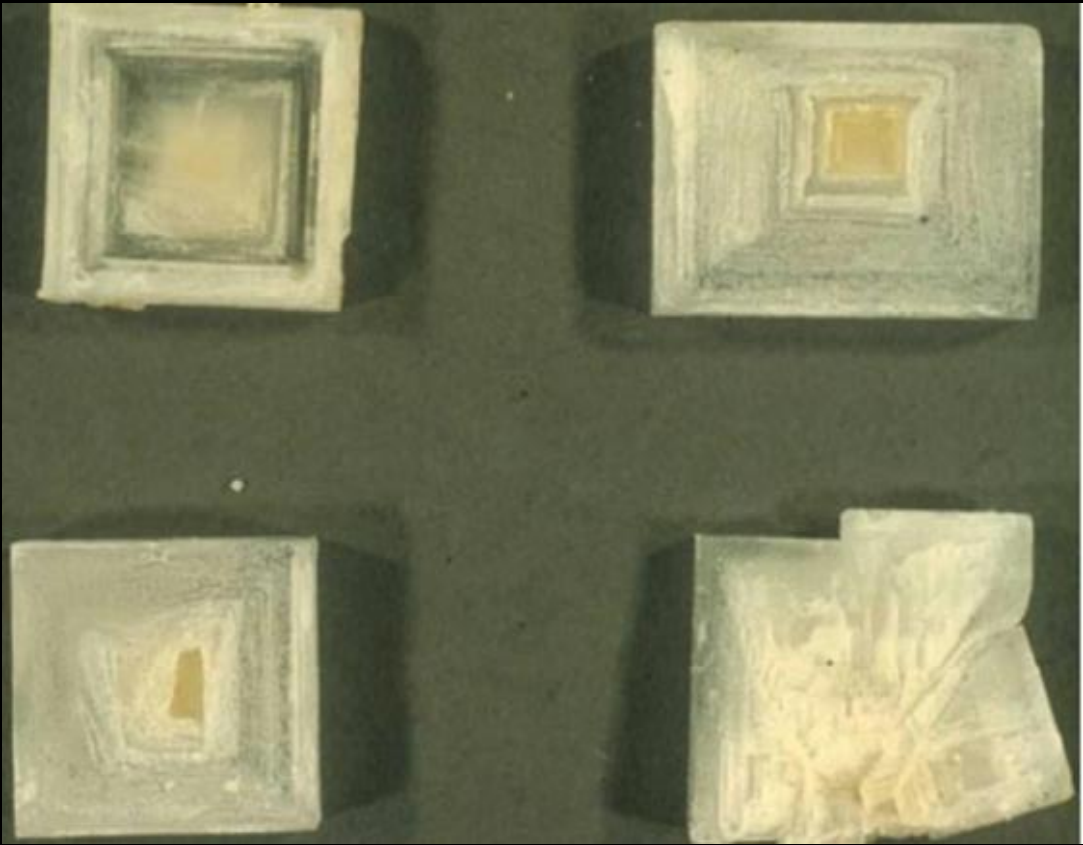


?



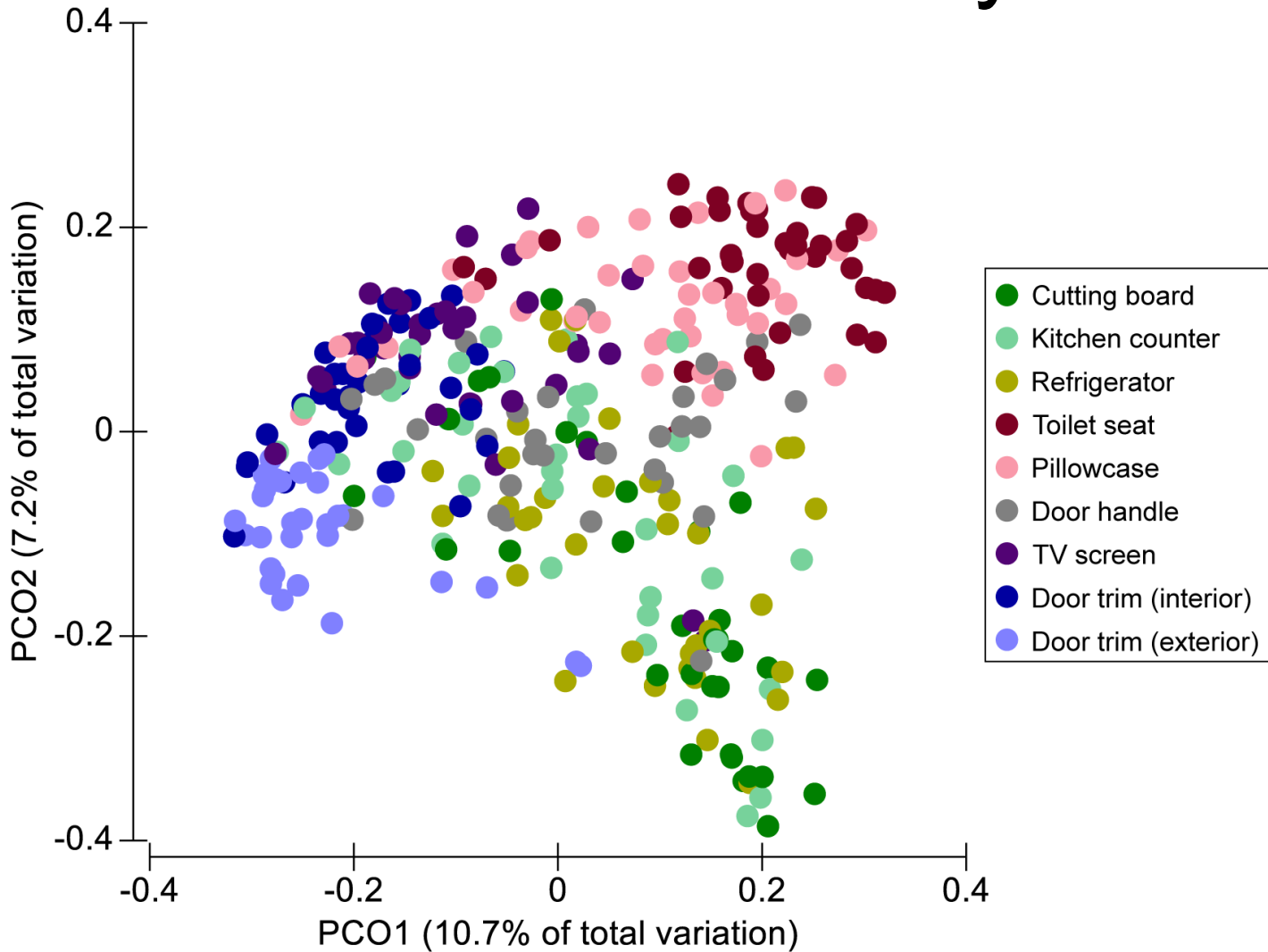








Most “Control” Favors Body Microbes



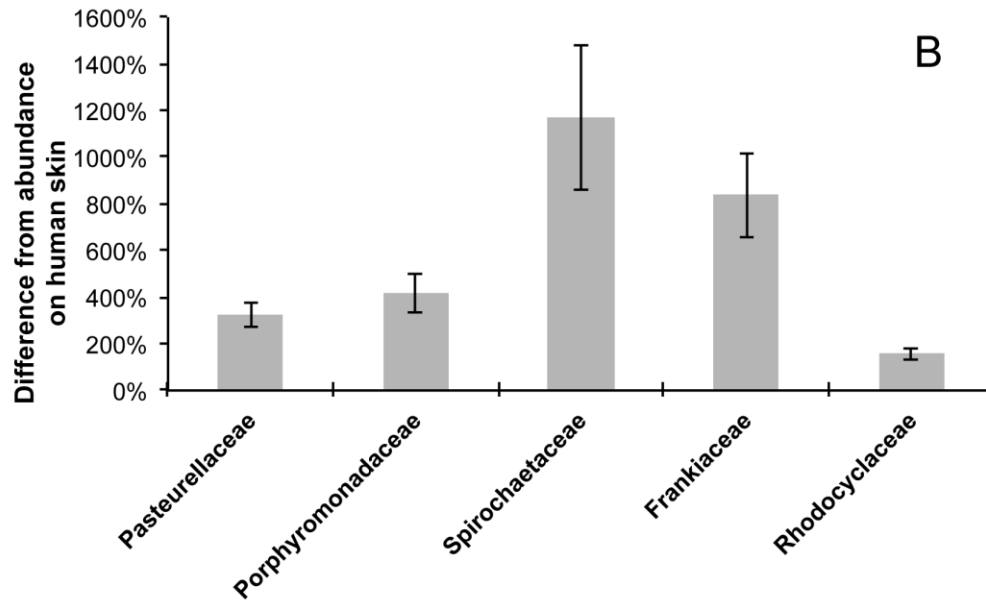
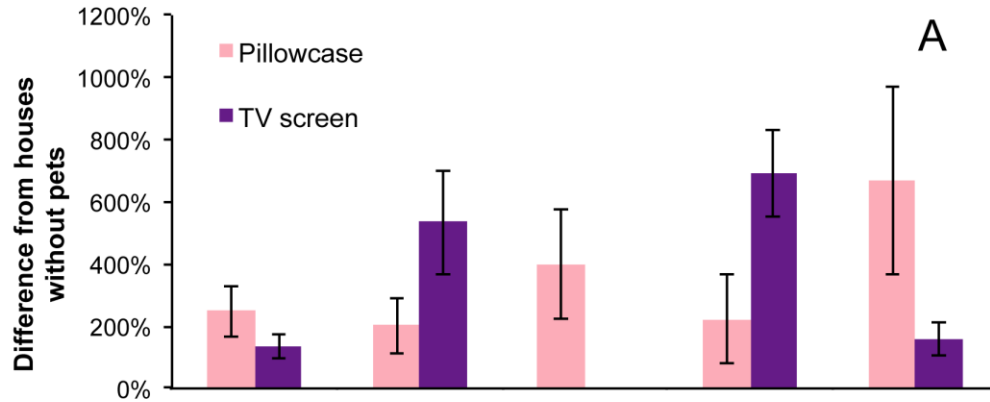
CONTROL OF OCCUPANTS



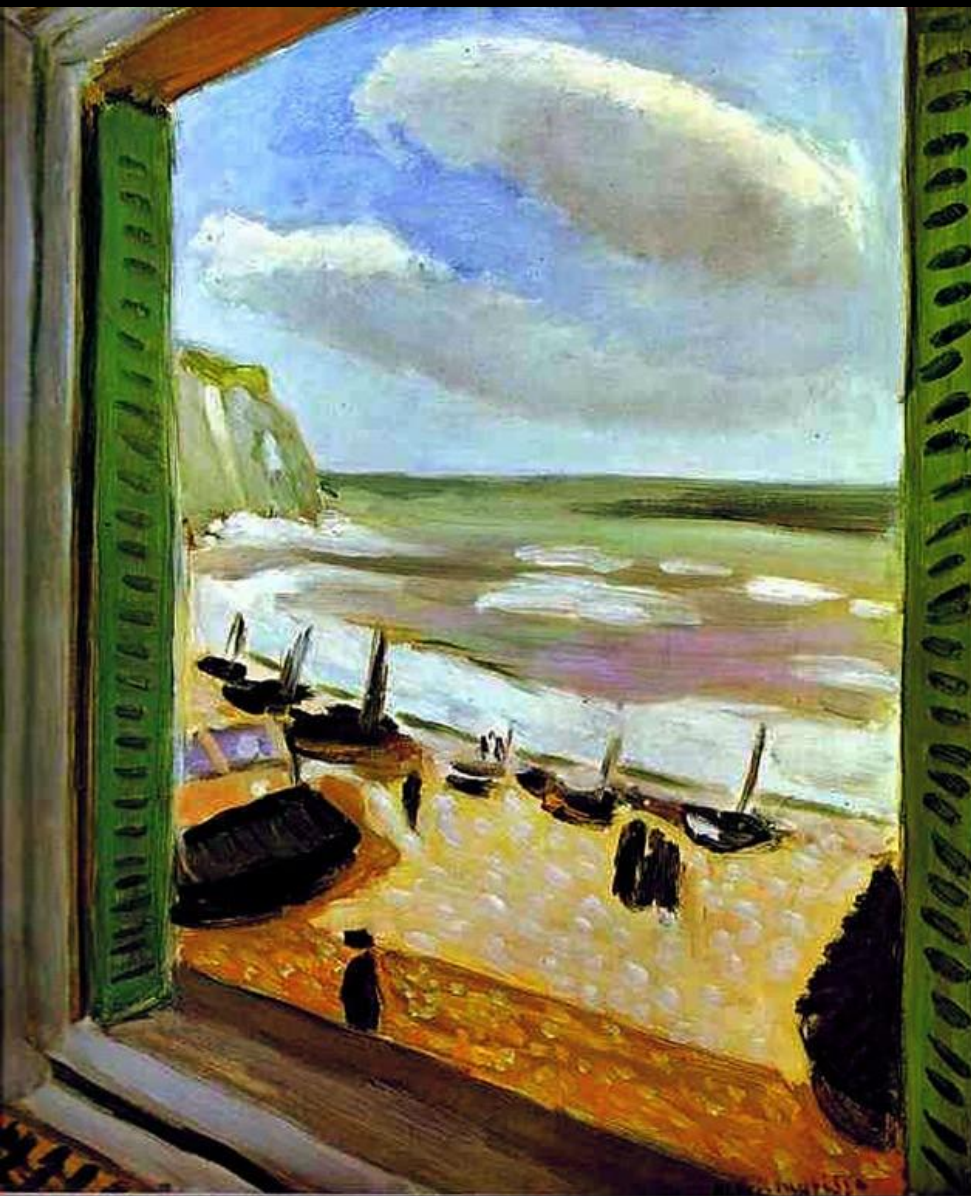


CONTROL VIA NON-HUMAN ANIMALS

Pet Dogs Favor Dog Body Microbes




Closing Windows Favors Body Microbes



Chlorinating Water Favors Chlorine-tolerant Mycobacteria

Ecological Analyses of Mycobacteria in Showerhead Biofilms and Their Relevance to Human Health

Matthew J. Gebert^a, Manuel Delgado-Baquerizo^{a,b}, Angela M. Oliverio^{a,c}, Tara M. Webster^a, Lauren M. Nichols^d, Jennifer R. Honda^e, Edward D. Chan^{f,g,h}, Jennifer Adjemian^{i,j}, Robert R. Dunn^{d,k}, Noah Fierer ^{a,c}

^aCooperative Institute for Research in Environmental Sciences, University of Colorado, Boulder, Colorado, USA

^bDepartamento de Biología y Geología, Física y Química Inorgánica, Escuela Superior de Ciencias Experimentales y Tecnología, Universidad Rey Juan Carlos, Móstoles, Spain

^cDepartment of Ecology and Evolutionary Biology, University of Colorado, Boulder, Colorado, USA

^dDepartment of Applied Ecology, North Carolina State University, Raleigh, North Carolina, USA

^eDepartment of Biomedical Research, Center for Genes, Environment, and Health, National Jewish Health, Denver, Colorado, USA

^fDepartment of Medicine, National Jewish Health, Denver, Colorado, USA

^gDivision of Pulmonary Sciences and Critical Care Medicine, University of Colorado Denver, Aurora, Colorado, USA

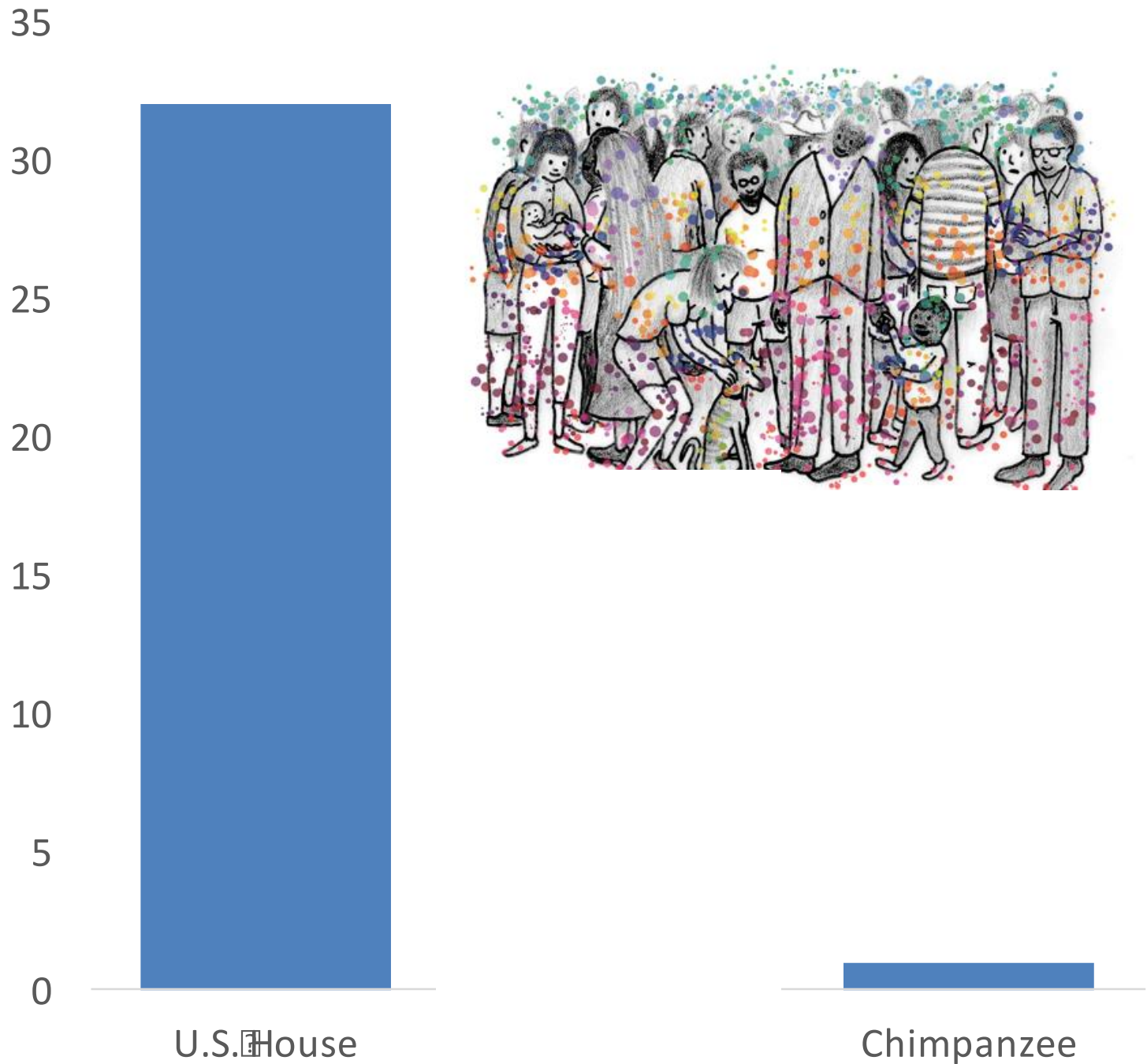
^hDenver Veterans Affairs Medical Center, Denver, Colorado, USA

ⁱNational Institute of Allergy and Infectious Diseases, Bethesda, Maryland, USA

^jUnited States Public Health Service Commissioned Corps, Rockville, Maryland, USA

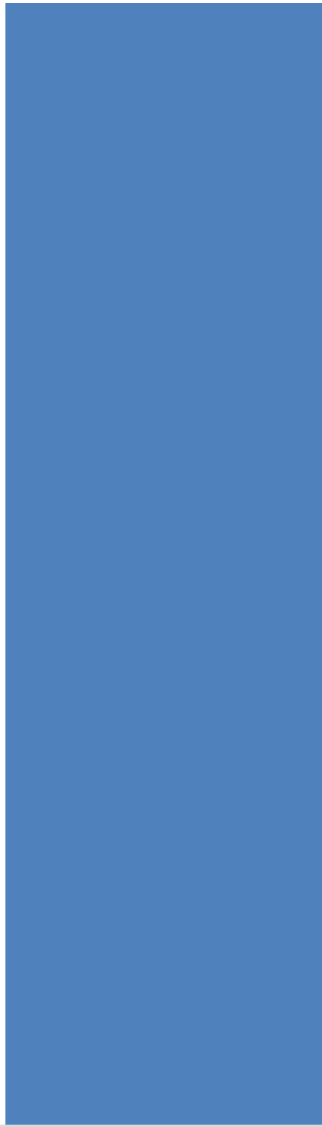
^kNatural History Museum of Denmark, University of Copenhagen, Copenhagen, Denmark

Proportion from bodies

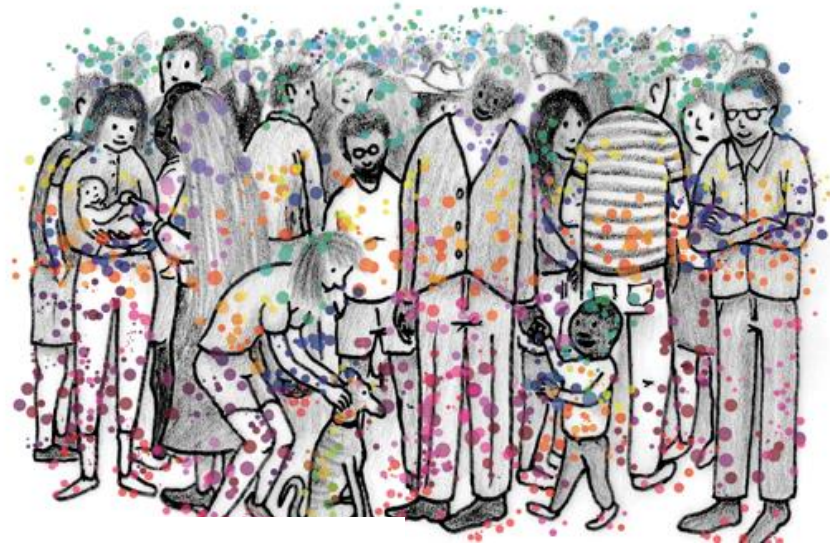


Proportion from bodies

35
30
25
20
15
10
5
0



U.S. House



This is even more extreme for hospitals.



Chimpanzee

~95% of time indoors



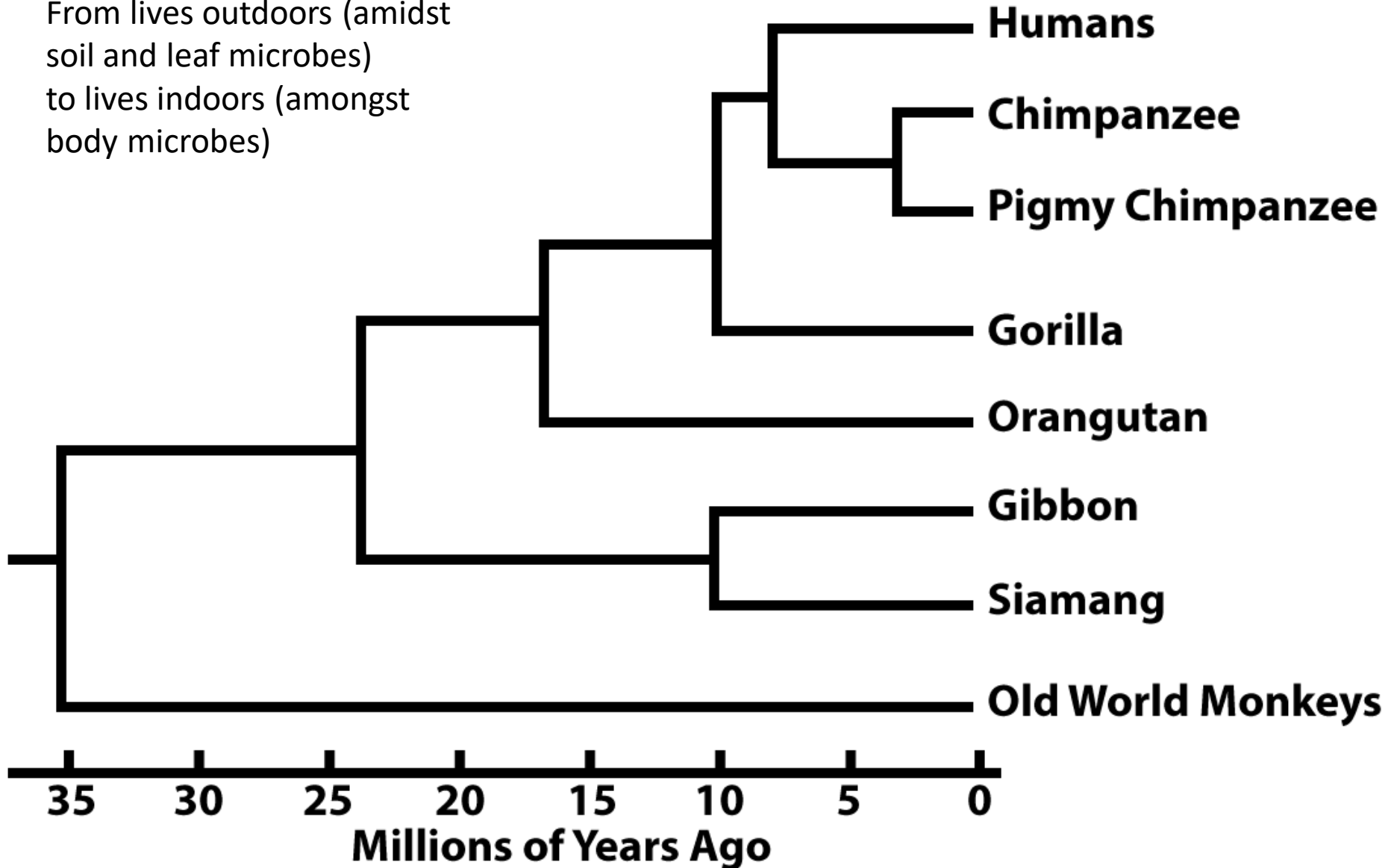
~95% of time indoors



How much time did you spend outdoors today?

1. EVOLUTIONARY HISTORY (HOUSE)

From lives outdoors (amidst soil and leaf microbes)
to lives indoors (amongst body microbes)



Net effects

- Microbial composition that appears to predispose people to allergy, asthma, and other inflammatory diseases (as much via what is absent as what is present)

We don't understand this effect entirely, but it is clear that it is occurring.



Net effects

- Microbial composition that appears to predispose people to allergy, asthma, and other inflammatory diseases (as much via what is absent as what is present)
- **An environment that promotes survival of biocide resistant bacteria and opportunistic pathogens (which are often poor competitors)**

Antibiotic use + bleaching favors resistant organisms (poor competitors)





*Pseudomonas
aeruginosa*
photo of colony by
Scott Chimileski

Structure and Functional Attributes of Bacterial Communities in Premise Plumbing Across the United States

Tara M. Webster, Alexander McFarland, Matthew J. Gebert, Angela M. Oliverio, Lauren M. Nichols, Robert R. Dunn, Erica M. Hartmann, and Noah Fierer*



Cite This: *Environ. Sci. Technol.* 2021, 55, 14105–14114



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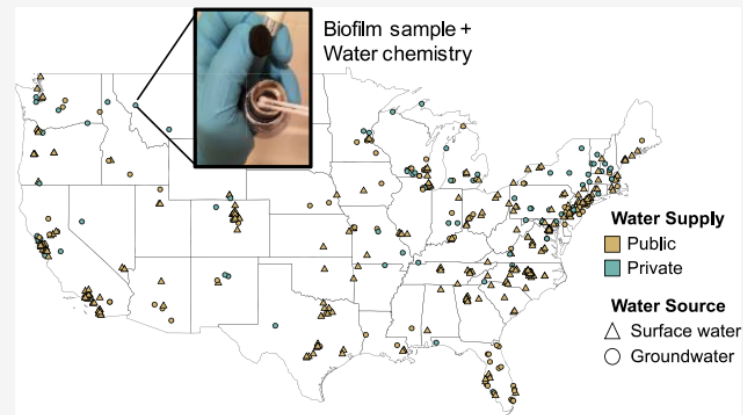


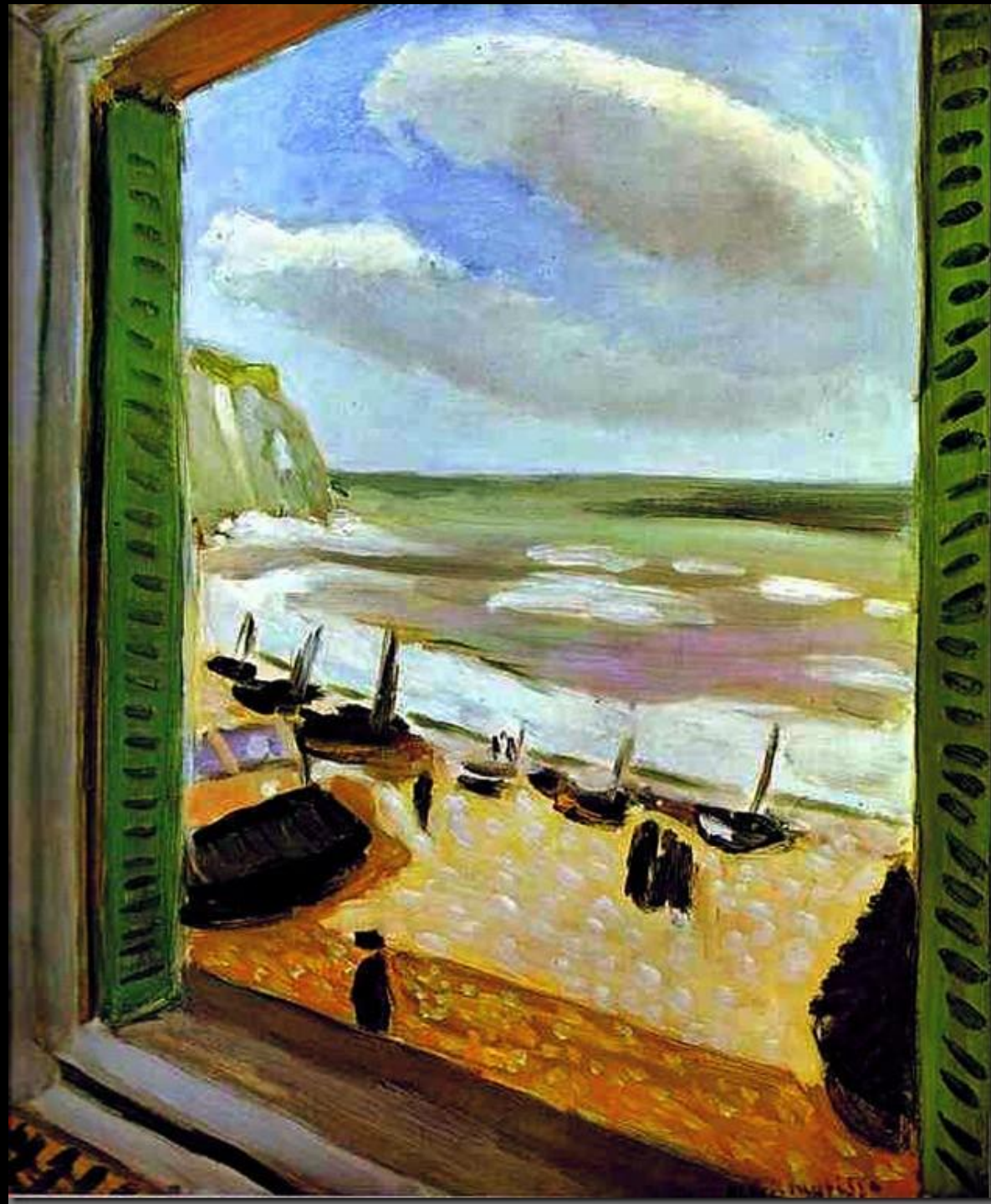
Article Recommendations



Supporting Information

ABSTRACT: Microbes that thrive in premise plumbing can have potentially important effects on human health. Yet, how and why plumbing-associated microbial communities vary across broad spatial scales remain undetermined. We characterized the bacterial communities in 496 showerheads collected from across the continental United States. The overall community structure, determined by 16S rRNA gene amplicon sequencing, revealed high levels of bacterial diversity. Although a large fraction of the observed variation in community composition could not be explained, differences in bacterial community composition were associated with water supply (private well water vs public municipal water), water source (groundwater vs surface water), and associated differences in water chemistry (pH and chlorine). Most notably, showerheads in homes supplied with public water had higher abundances of *Blastomonas*, *Mycobacterium*, and *Porphyrobacter*, while *Pseudorhodoplanes*, *Novosphingobium*, and *Nitrospira* were more abundant in those receiving private well water. We conducted





Net effects

- Microbial composition that appears to predispose people to allergy, asthma, and other inflammatory diseases (as much via what is absent as what is present)
- An environment that promotes survival of antibiotic resistant bacteria and opportunistic pathogens (which are often poor competitors)
- **An environment where beneficial microbes not acquired during birth are very unlikely to be acquired later on.**



Cell Reports Medicine

Volume 1, Issue 9, 22 December 2020, 100156



Recommen

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Cell Host & Mi

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Article

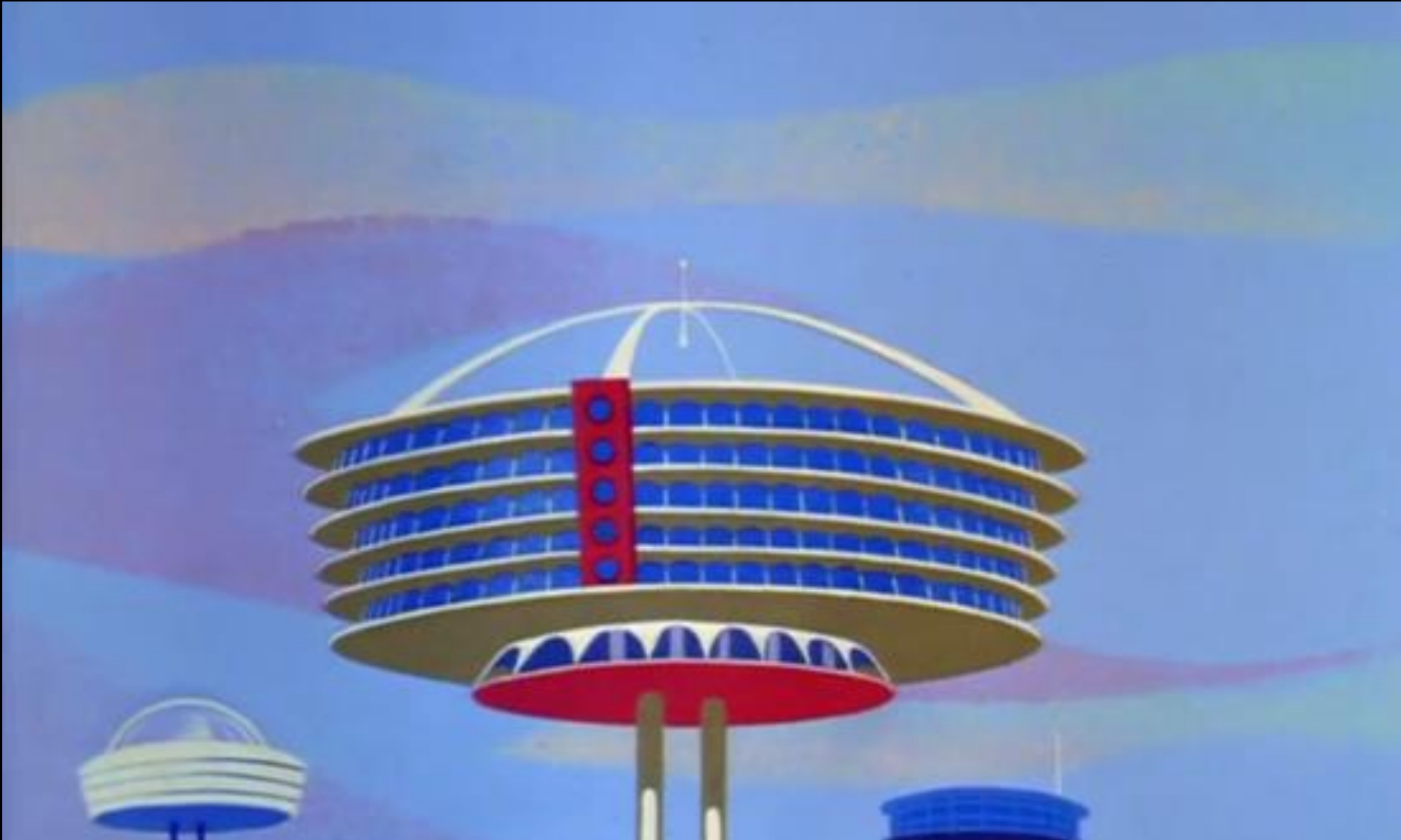
Delivery Mode Affects Stability of Early Infant Gut Microbiota

Caroline M. Mitchell¹, Chiara Mazzoni², Larson Hogstrom³, Allison Bryant¹, Agnes Bergerat¹, Avital Cher²,
Shawna Pochan¹, Penelope Herman¹, Maureen Carrigan¹, Karen Sharp¹, Curtis Huttenhower^{3,4}, Eric S. Lander^{3,5,6},
Hani Yamalik^{3,7}, Demetris Xenarios^{3,7,8}, Manon Yassierli^{2,3,9,10} ✉

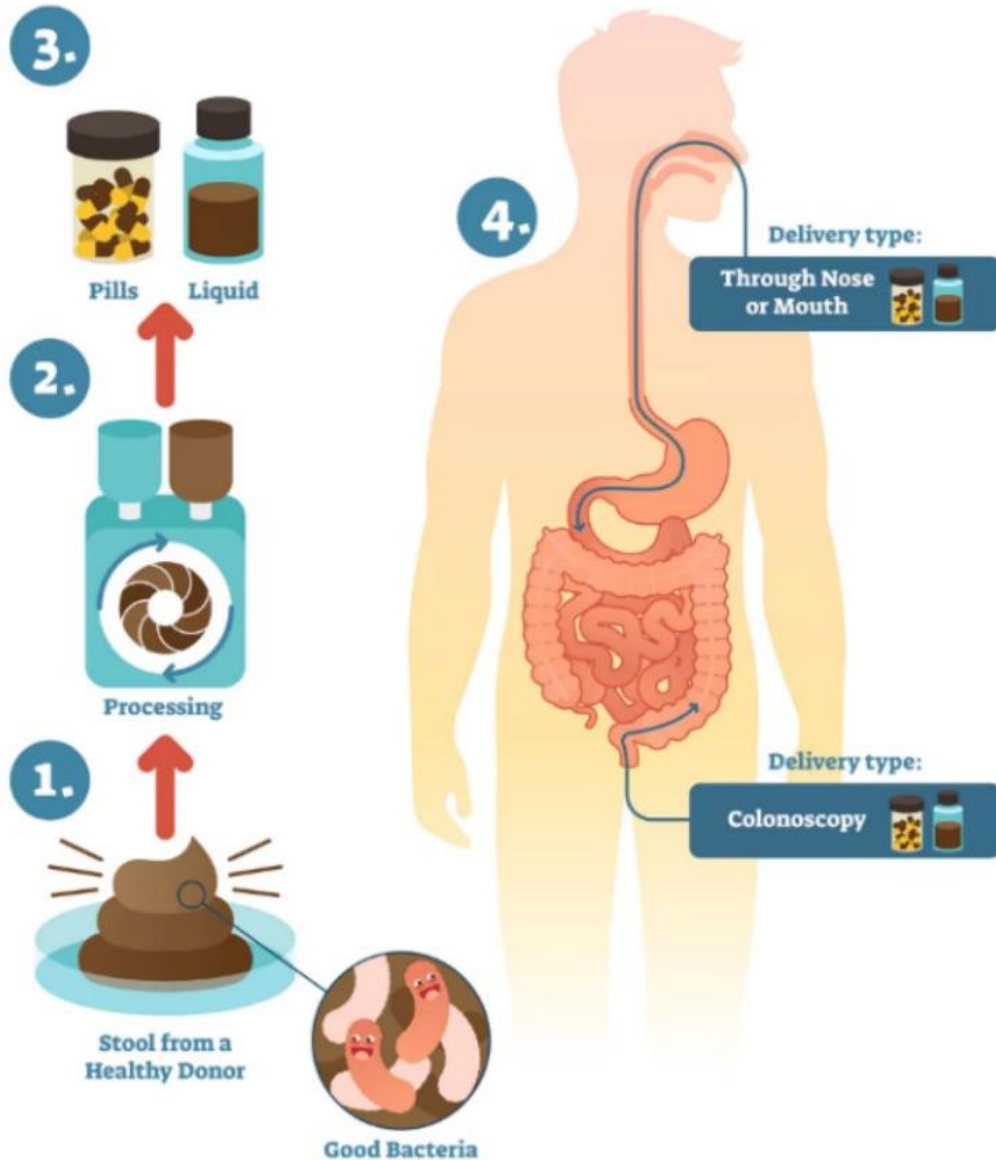
In Trying to Keep Life Out

- We have favored a set of species that have adapted to the indoors (and are not necessarily good for us)
- We have excluded beneficial species (to our own detriment in the context of allergy/asthma/autoimmune disorders)
- We have created homes that are less healthy than those built by our ancestors eight million years ago.

If not this future then what?



FECAL TRANSPLANT THERAPY

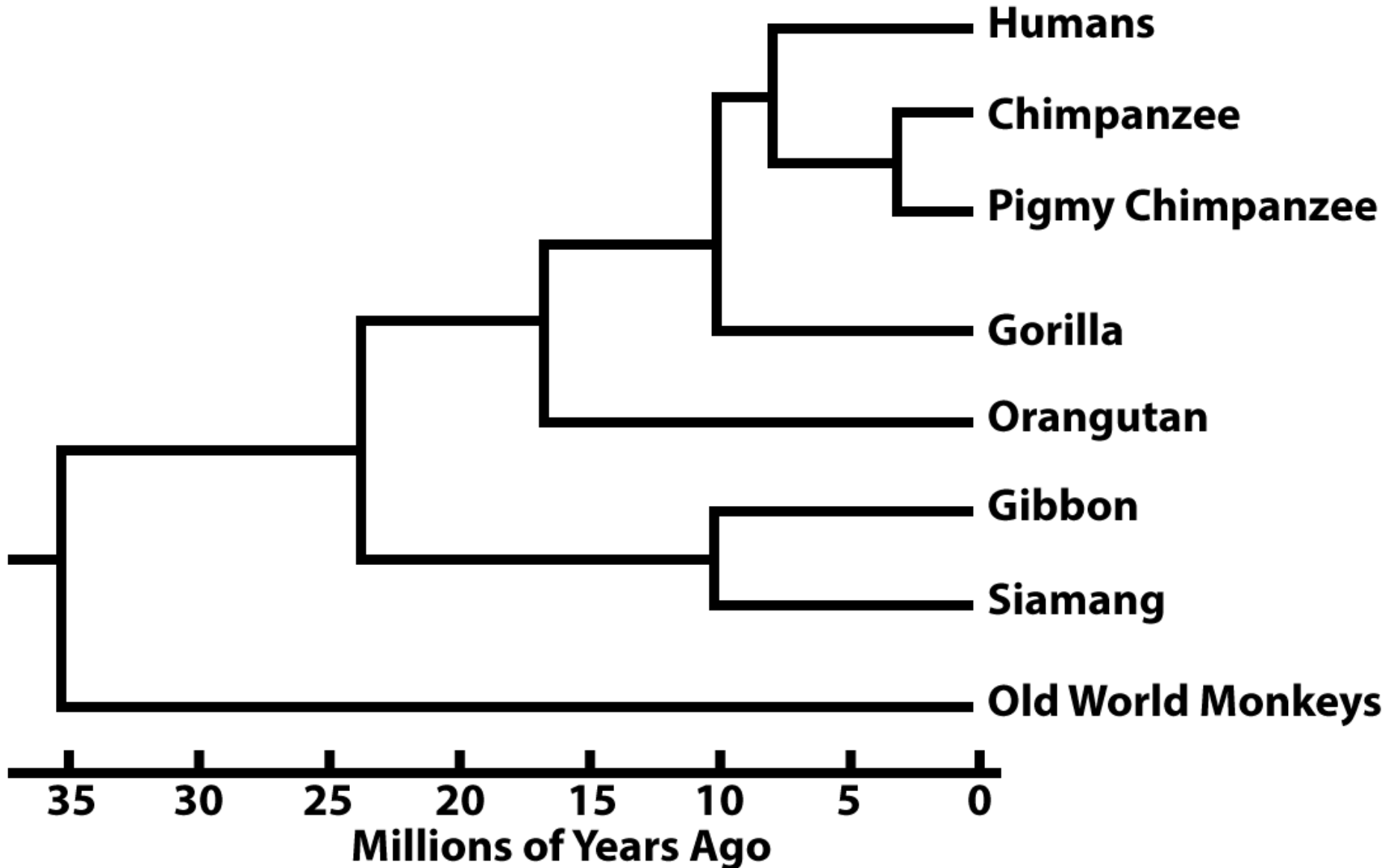


Predictions for the next 5-10 years of medicine (in 2017)

Predictions for the next 5-10 years of medicine (2022)

- An initial focus on re-introduction of outdoor microbes into homes (plants, fermented foods, open windows)
 - This is akin to the earliest fecal transplants
- Initial haphazard results (Which soil? Which plants? Which fermented foods?)
- One thread of future interventions will focus on bringing microbial processes into the home (e.g., fermenters/digesters)
- Another thread will focus on specific microbes used as therapeutics (on surfaces, in the home)
- All of this is already happening, just in remote corners

1. EVOLUTIONARY HISTORY (HEART)

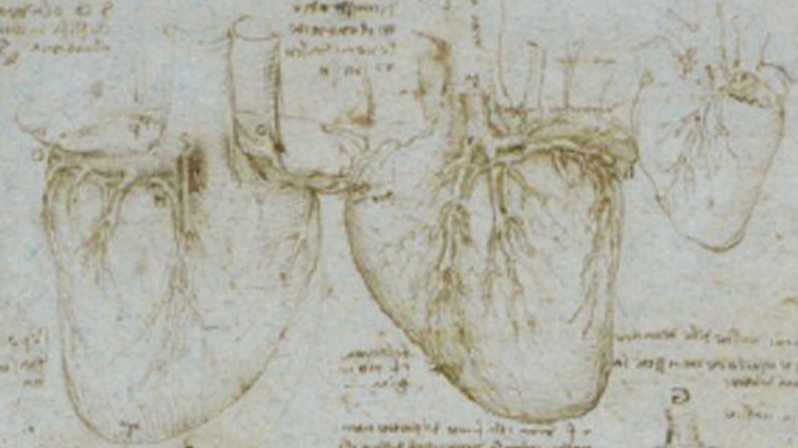




Handwritten text in the top left corner, likely describing anatomical details or experimental procedures.



Handwritten text above the top right anatomical drawings.



Handwritten text between the top right and middle right drawings.



Handwritten text to the right of the four small drawings.



Handwritten text above the two large heart drawings.

Handwritten text between the two large heart drawings.

Handwritten text below the left large heart drawing.

Handwritten text below the right large heart drawing.

Handwritten text in the middle section of the page.



Small handwritten letter 'd' next to the drawing with the tube.



Handwritten text below the drawing with the tube.



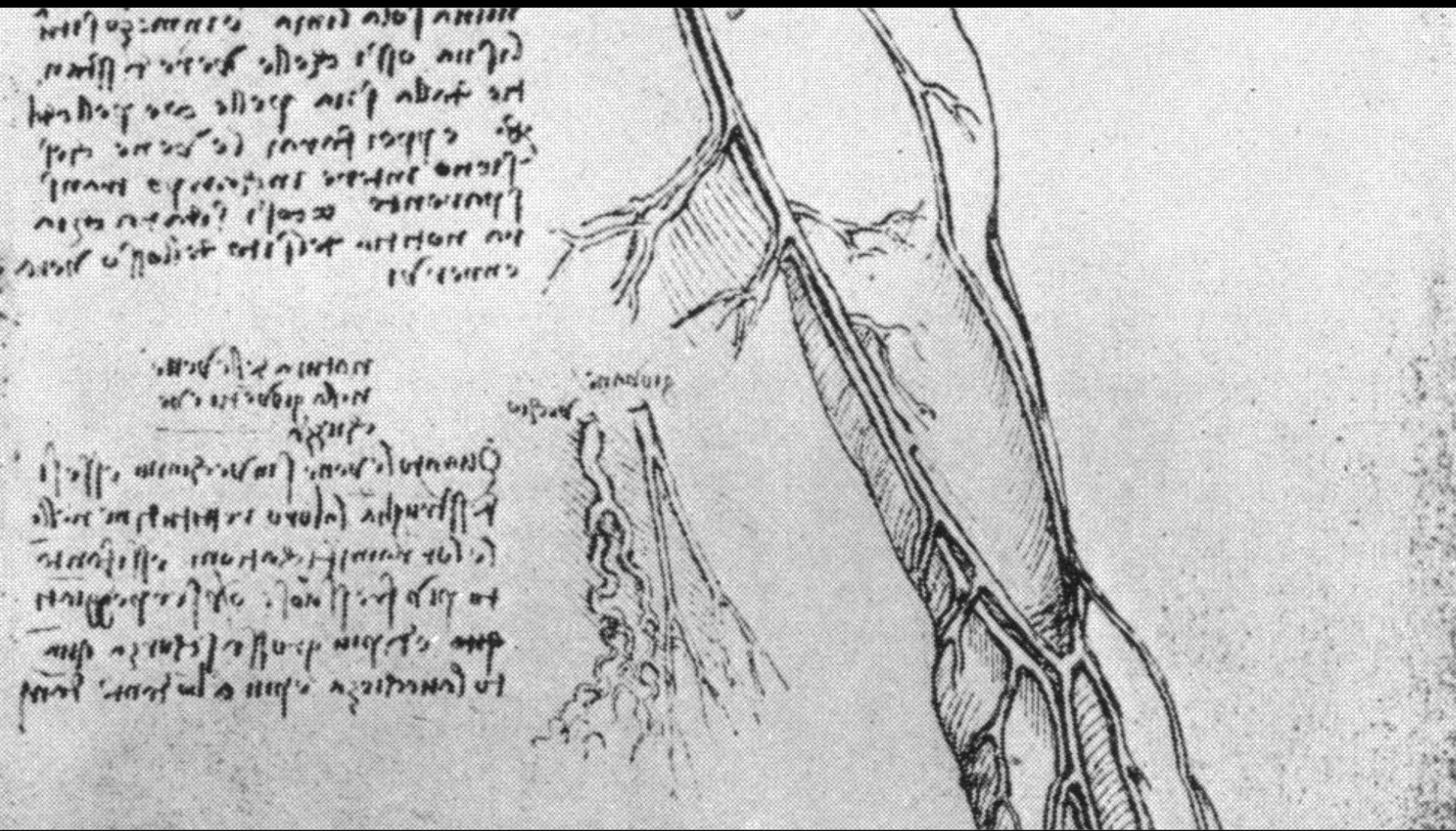
Handwritten text below the bottom right drawing.

The Anatomy of an Old Man (*Il Vecchio*)

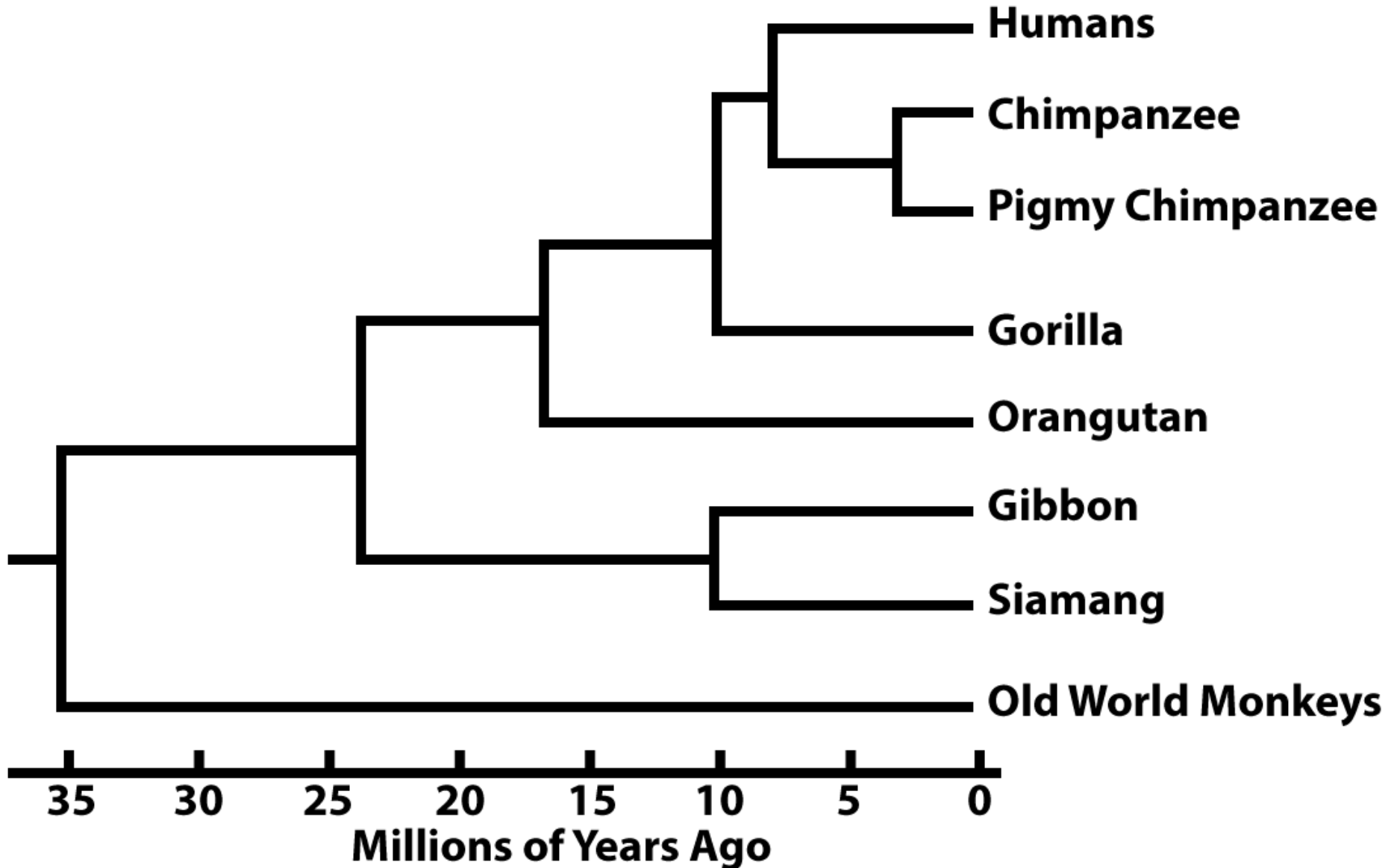


*“As a well-spent day
brings happy sleep, so
life well used brings
happy death”*

Tortuous Arteries

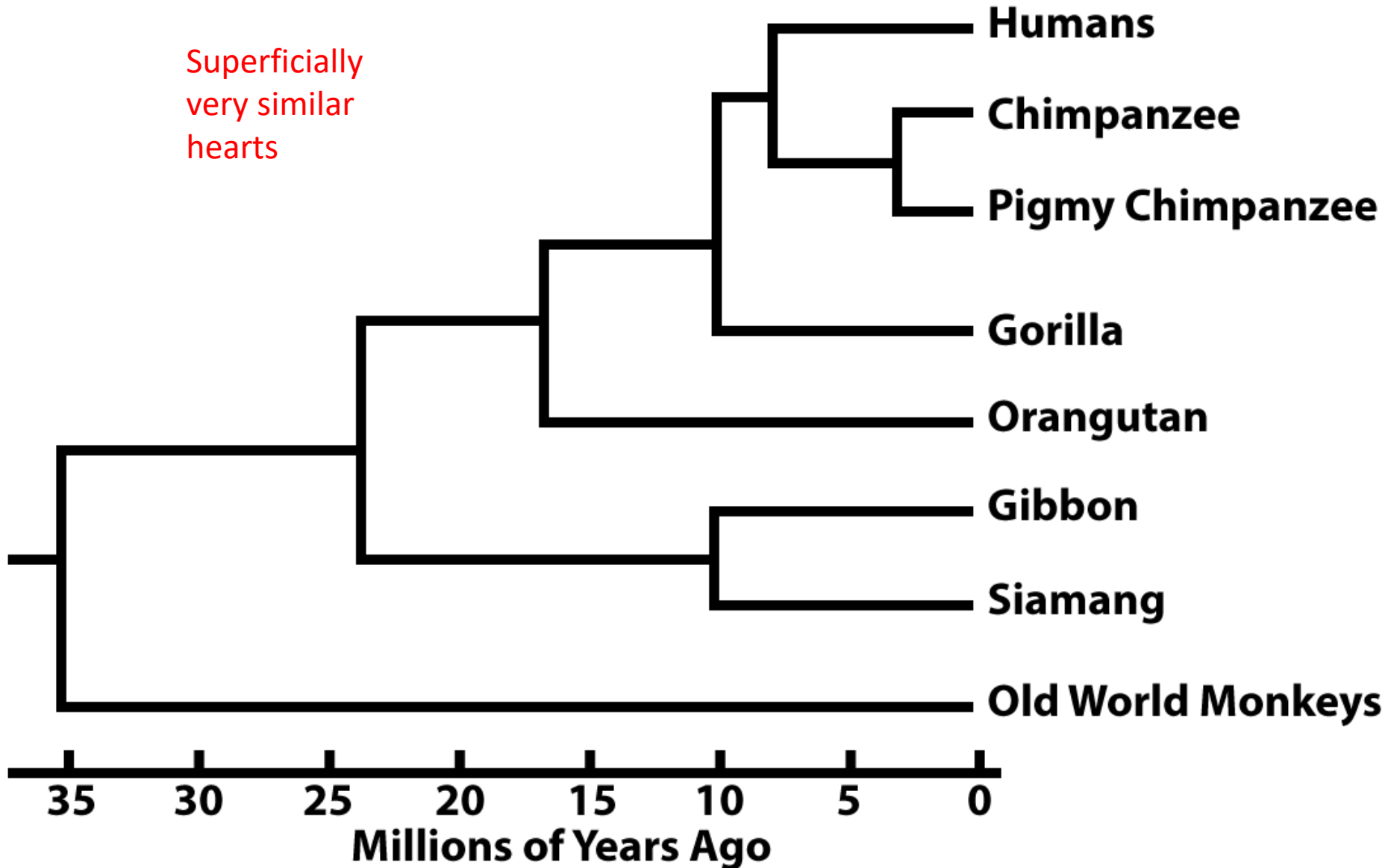


1. EVOLUTIONARY HISTORY (HEART)

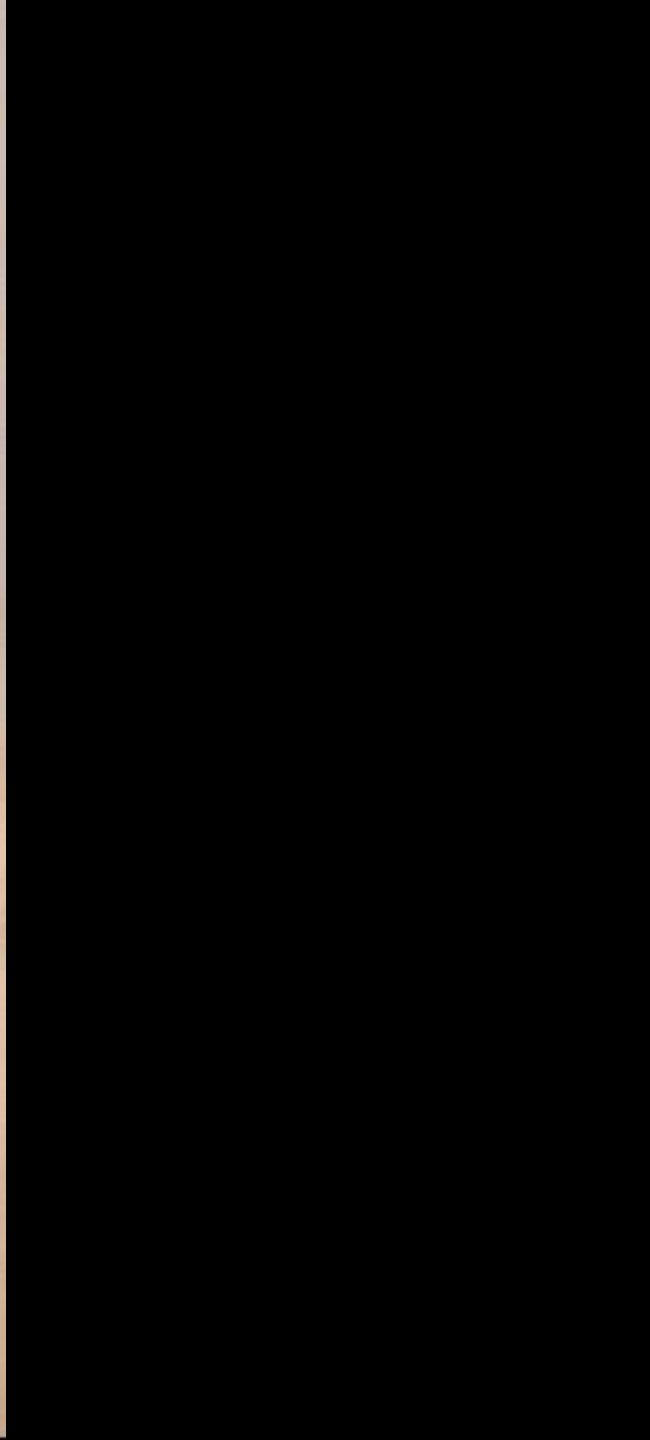
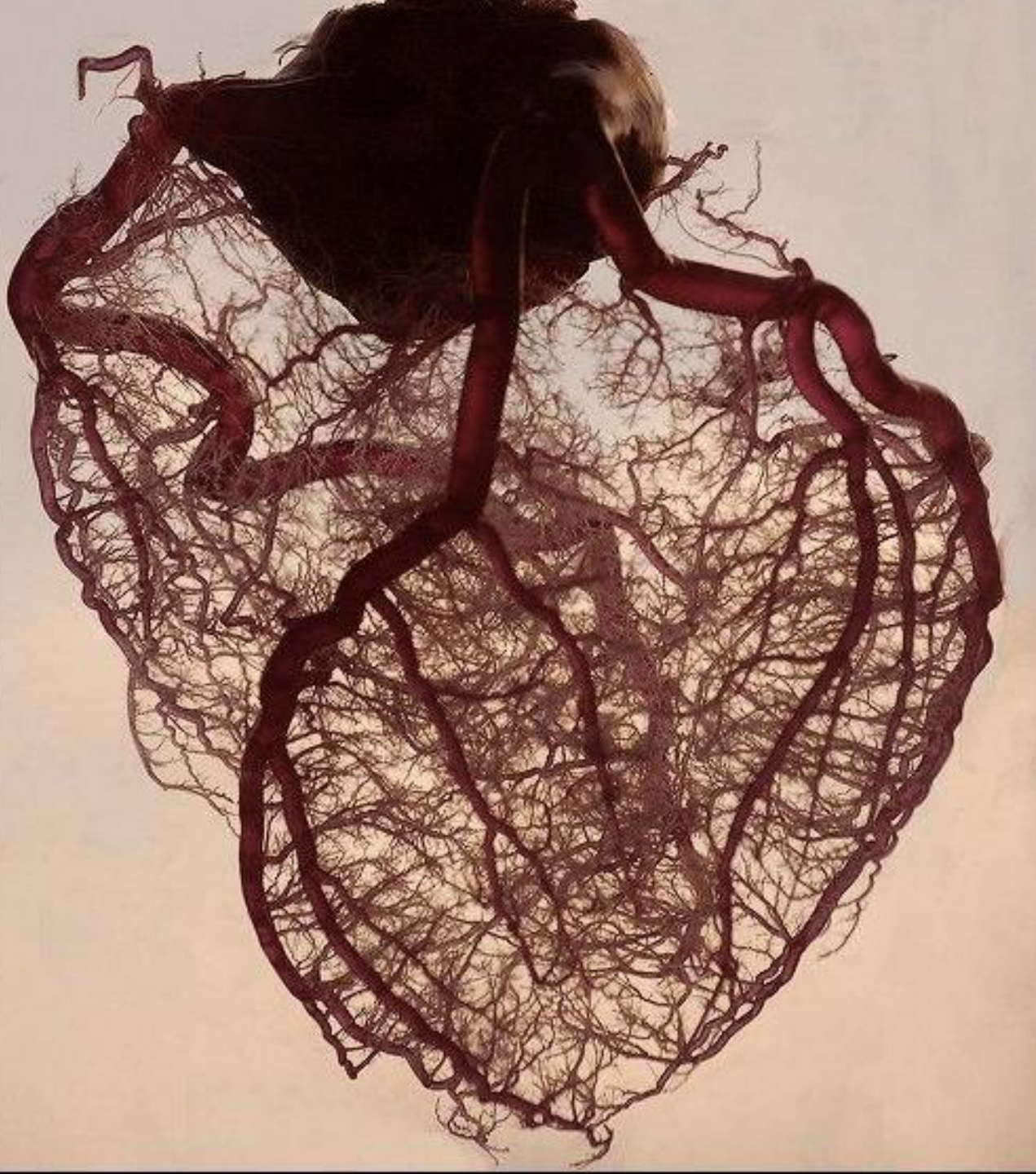


1. EVOLUTIONARY HISTORY (HEART)

Superficially
very similar
hearts









Sanaga-Yong Chimpanzee Rescue Center in Cameroon. Lucy has just died of a heart attack. The other chimpanzees mourn.

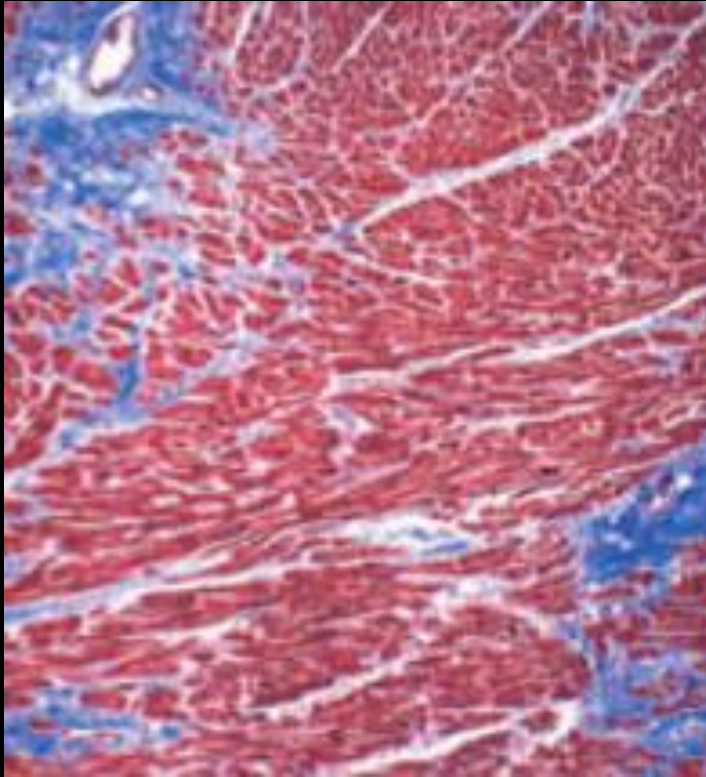


Yerkes National Primate Center, Emory

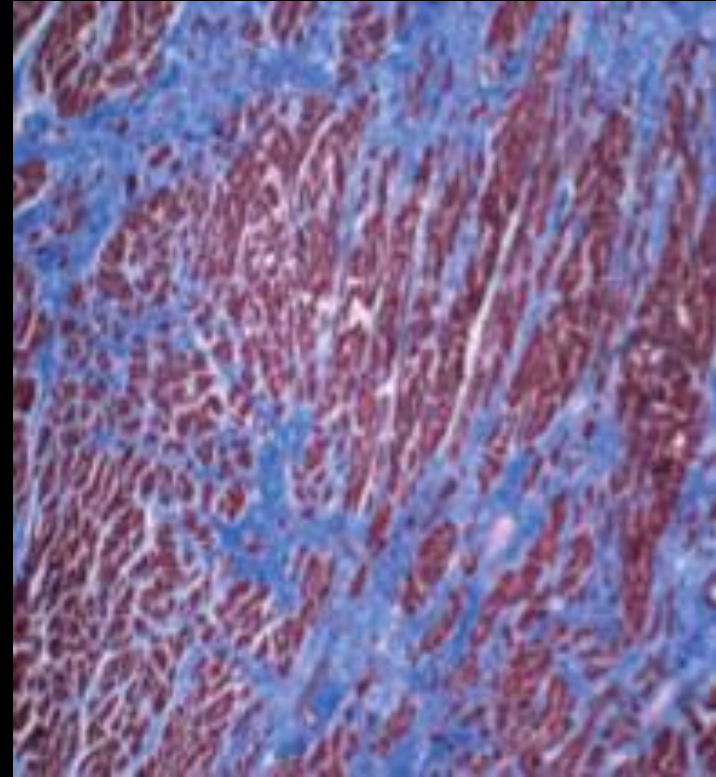


Nissi Varki, UCSD

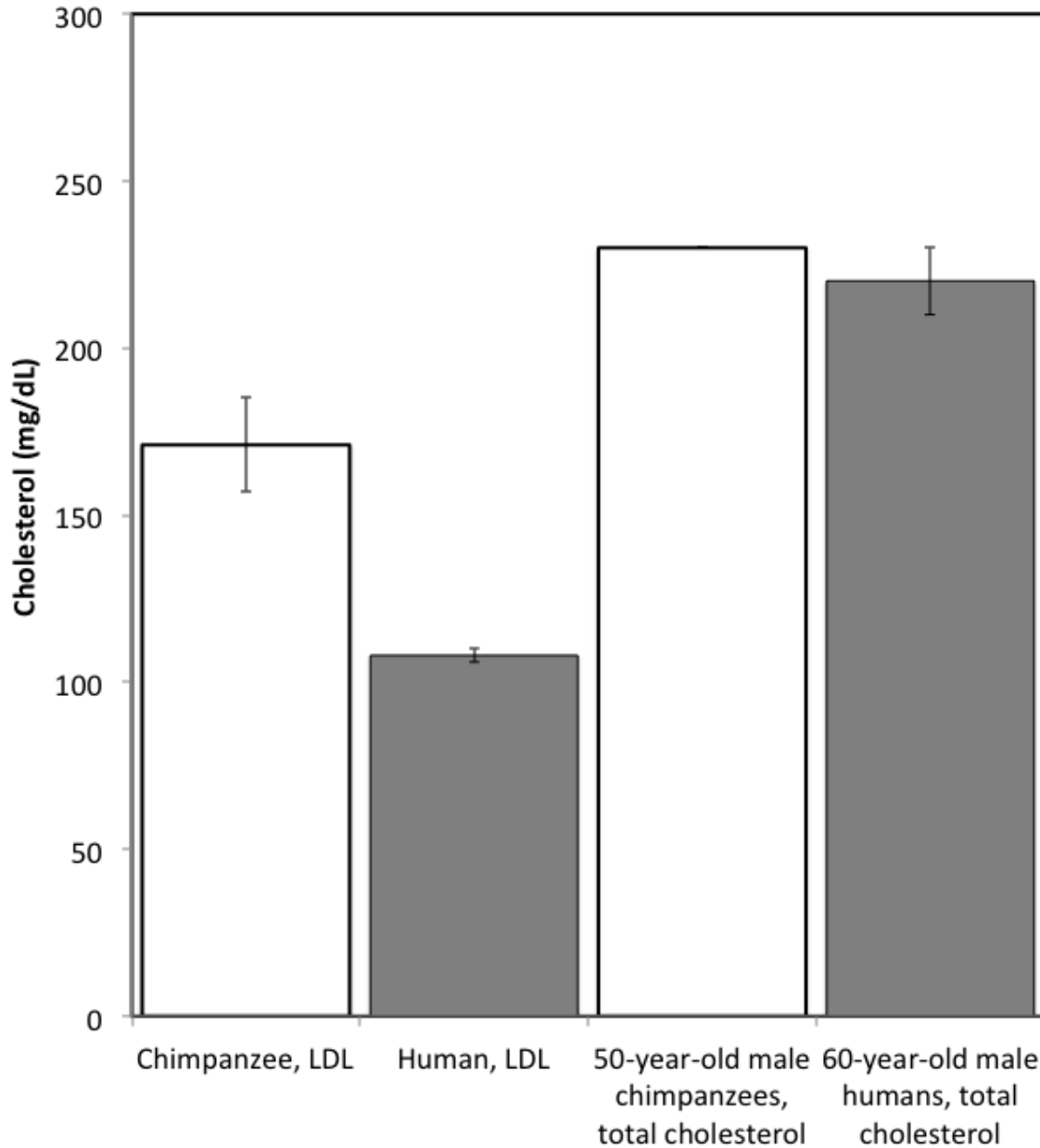
Human



Chimpanzee

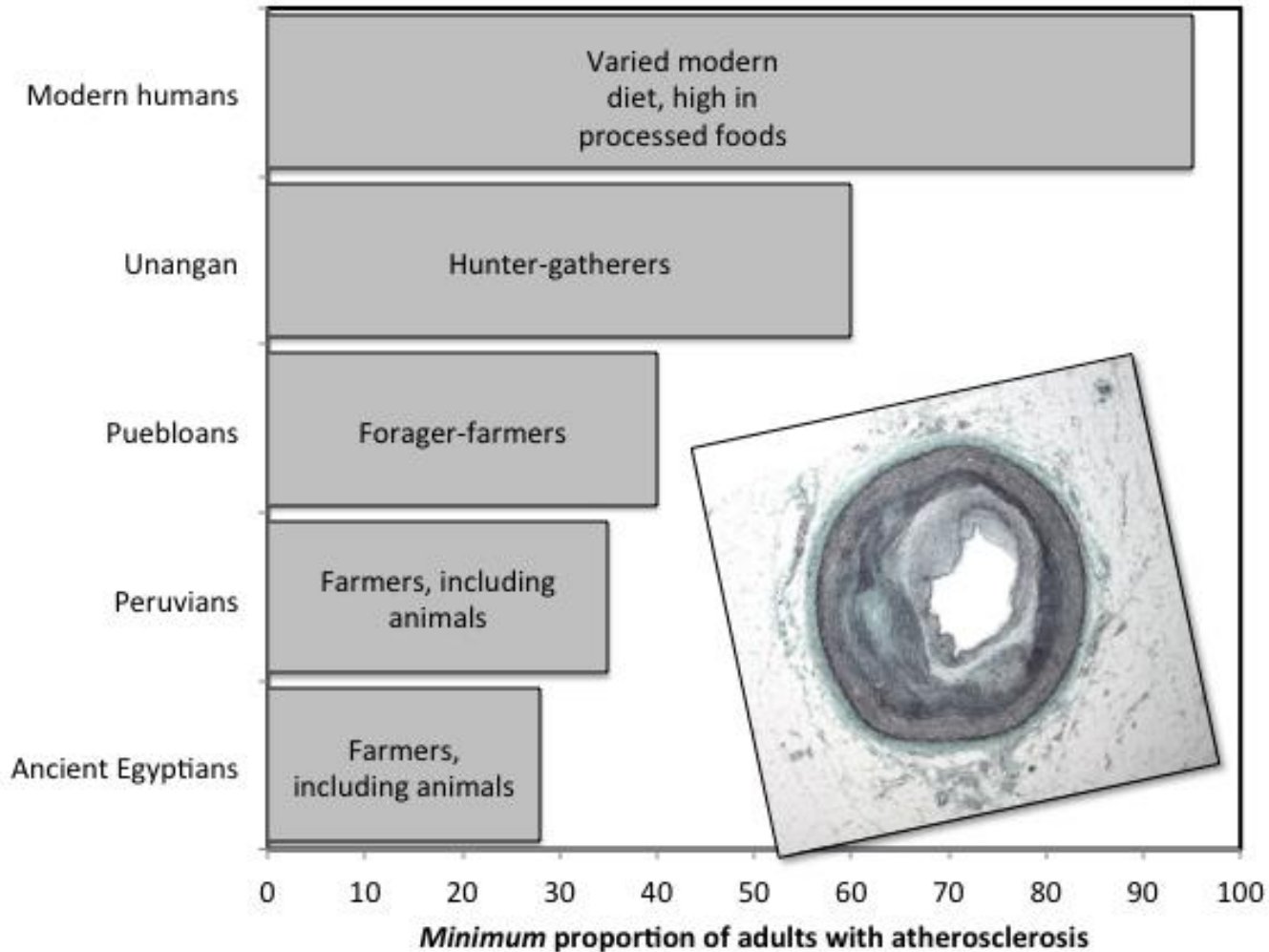


Varki, Nissi, Dan Anderson, James G. Herndon, Tho Pham, Christopher J. Gregg, Monica Cheriyan, James Murphy et al. "Heart disease is common in humans and chimpanzees, but is caused by different pathological processes." *Evolutionary applications* 2, no. 1 (2009): 101-112.

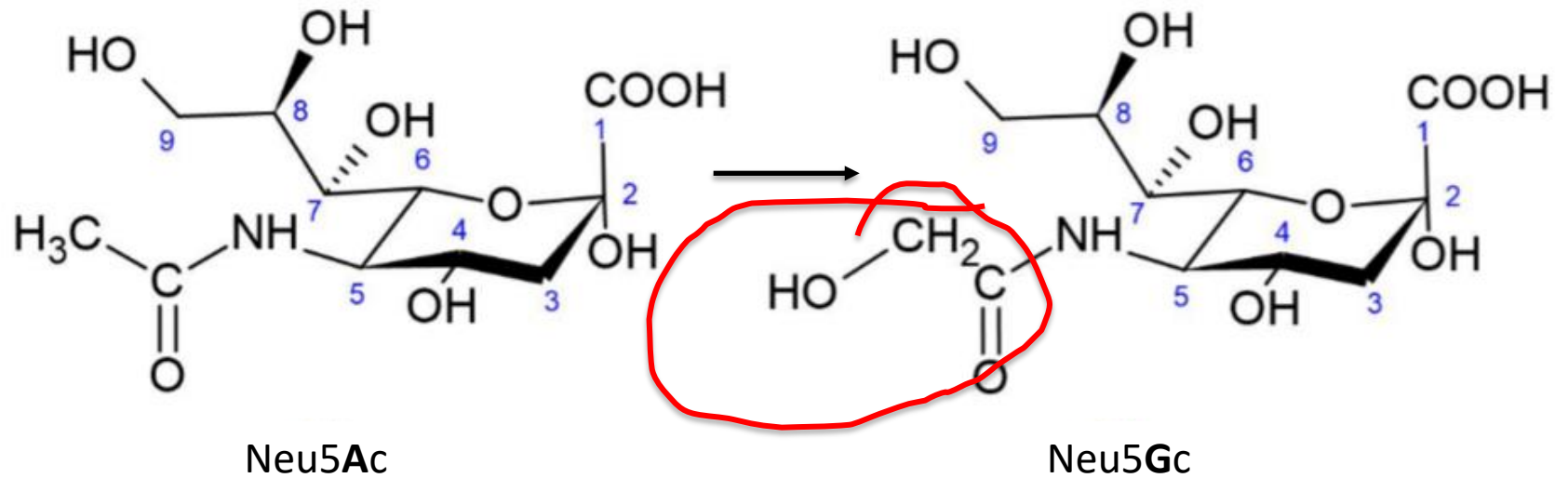


It doesn't appear to be due to differences in cholesterol

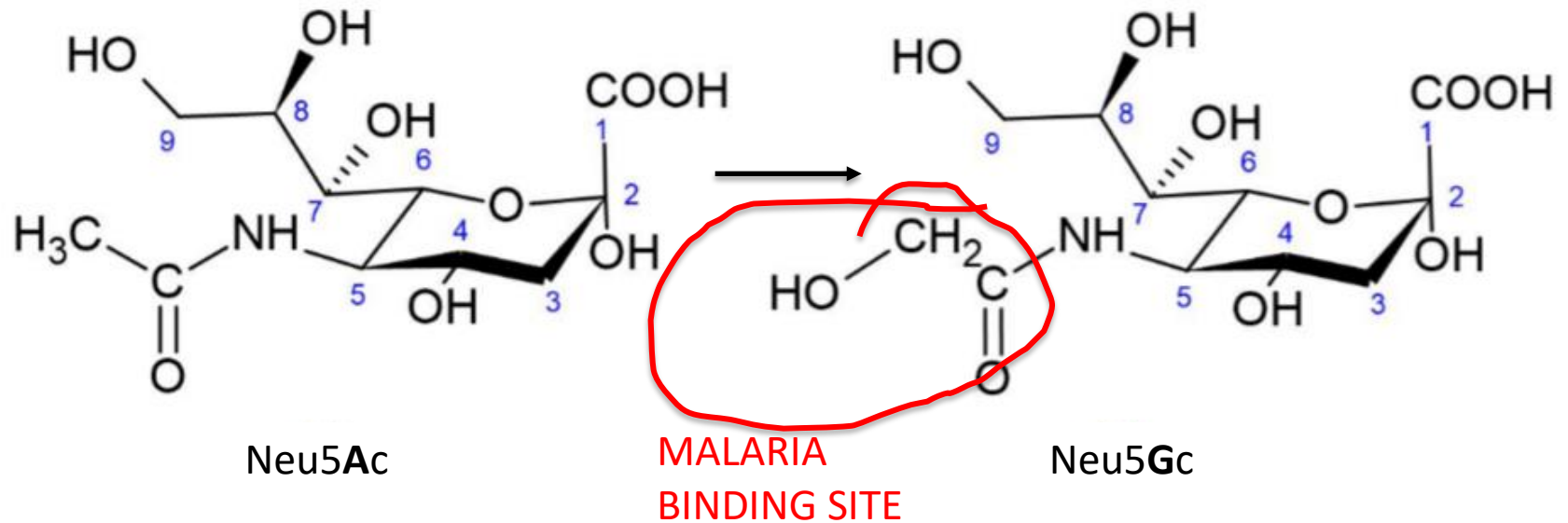
All humans seem to have the propensity for atherosclerosis



Ancient malaria



Ancient malaria

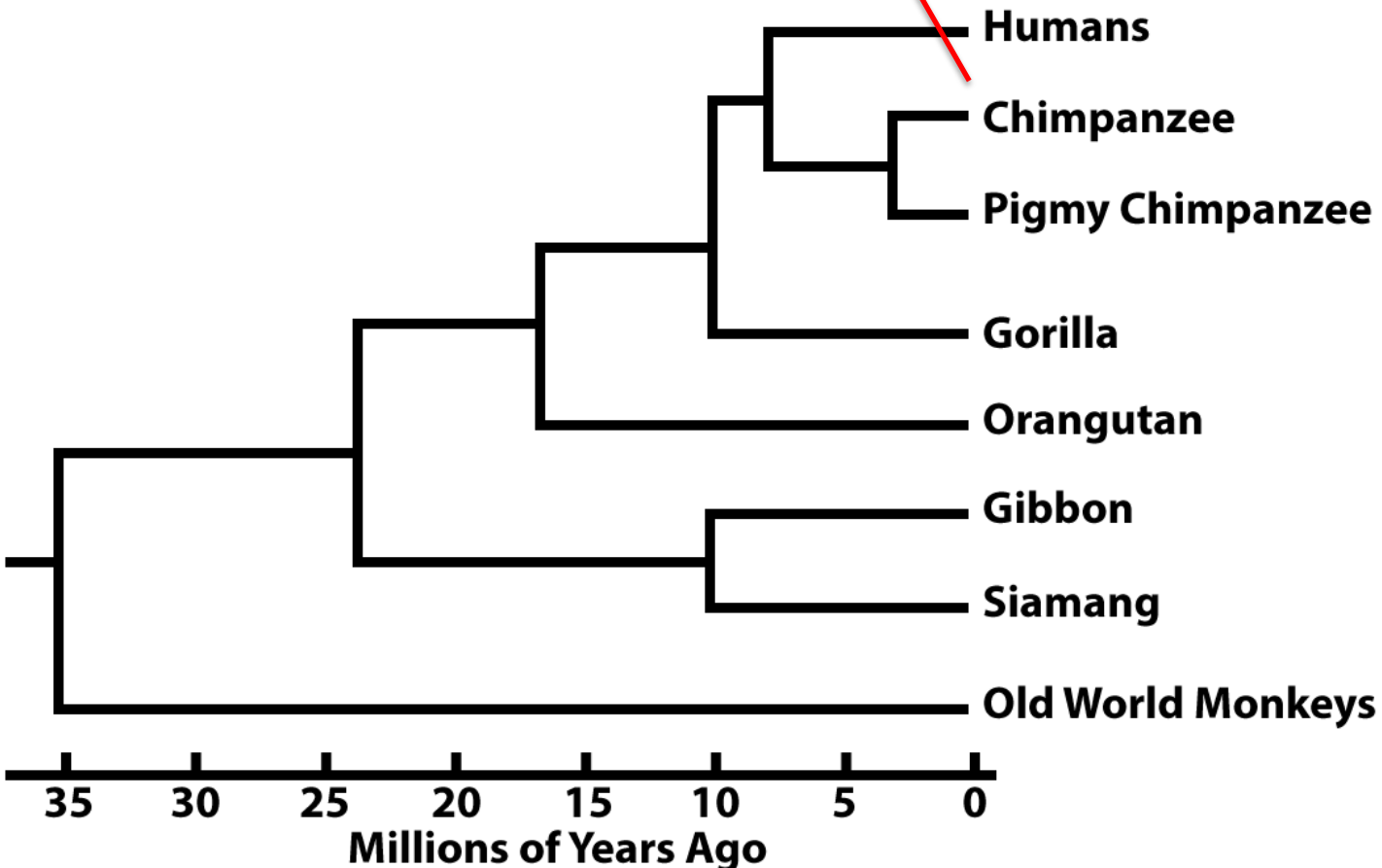


Uniquely human evolution of sialic acid genetics and biology

Ajit Varki  [Authors Info & Affiliations](#)

May 5, 2010 | 107(supplement_2)8939-8946 | <https://doi.org/10.1073/pnas.0914634107>

Broken gene associated with converting Neu5Ac to Neu5Gc



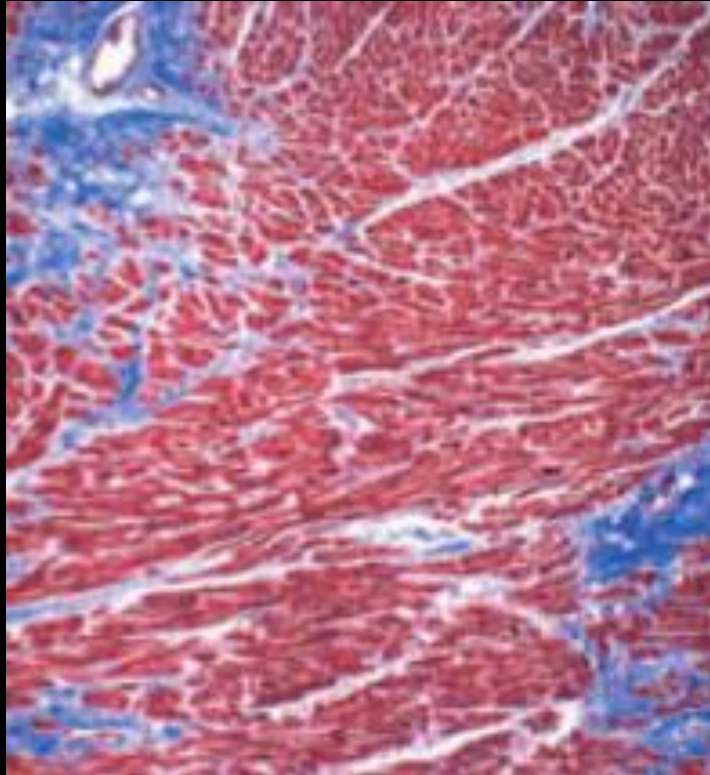




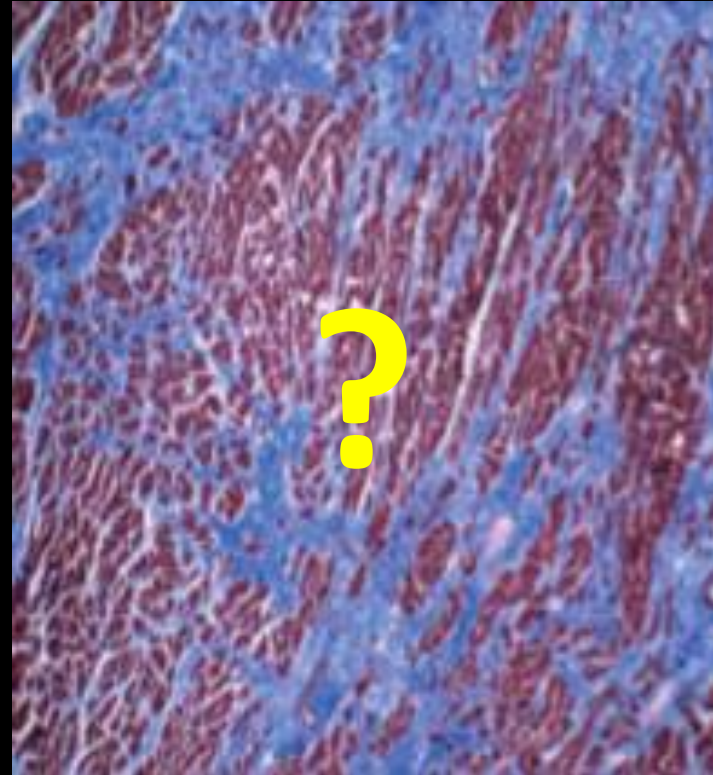
Meat-eating has been shown to accelerate atherosclerosis via sialic acids. Human-like mice develop atherosclerosis more rapidly in general and, especially when fed red meat.

Kawanishi, Kunio, Joanna K. Coker, Kaare V. Grunddal, Chirag Dhar, Jason Hsiao, Karsten Zengler, Nissi Varki, Ajit Varki, and Philip LSM Gordts. "Dietary Neu5Ac Intervention Protects Against Atherosclerosis Associated With Human-Like Neu5Gc Loss—Brief Report." *Arteriosclerosis, thrombosis, and vascular biology* 41, no. 11 (2021): 2730-2739.

Human



Chimpanzee



Varki, Nissi, Dan Anderson, James G. Herndon, Tho Pham, Christopher J. Gregg, Monica Cheriyan, James Murphy et al. "Heart disease is common in humans and chimpanzees, but is caused by different pathological processes." *Evolutionary applications* 2, no. 1 (2009): 101-112.

2. BROADER BIODIVERSITY



ALEXIS ROCKMAN, THE ECOTOURIST

2. BROADER BIODIVERSITY (HOUSES)



ALEXIS ROCKMAN, THE ECOTOURIST



© Piotr Naskrecki

Dr. Stephanie Mathews



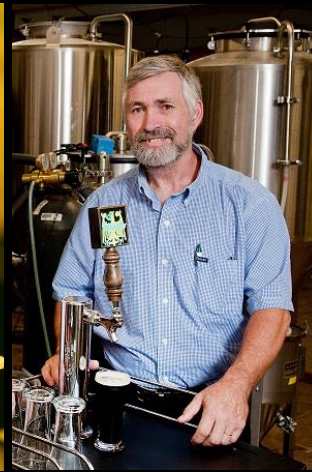
Doubled... the number of bacteria species
on Earth known to be able to break down lignin







The 1st Try Works



Lachancea thermotolerans (YB16), never before used in beer brewing



Led to a patent and a start up (Lachancea LLC, John Sheppard).
Dr. Anne Madden pictured (not here tonight).



RESEARCH ARTICLE

Sugar-seeking insects as a source of diverse bread-making yeasts with enhanced attributes

Anne A. Madden, Caitlin Lahue, Claire L. Gordy, Joy L. Little, Lauren M. Nichols, Martha D. Calvert
... [See all authors](#) ▾

First published: 23 October 2021 | <https://doi.org/10.1002/yea.3676> | Citations: 1

Funding information: North Carolina State University; North Carolina State University Chancellor's Innovation Fund; North Carolina Biotechnology Center, Grant/Award Number: 2019-BIG-6513

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PDF



TOOLS



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Abstract

Insects represent a particularly interesting habitat in which to search for novel yeasts of value to industry. Insect-associated yeasts have the potential to have traits relevant to modern food and beverage production due to insect–yeast interactions, with such traits including diverse carbohydrate metabolisms, high sugar tolerance, and general stress



Otzi, the Iceman
(and his useful yeasts)





Note: Dr. Clint Penick (at far right)
has pillows available
for purchase



© Lauren Nichols

Omar Halawani



© Alex Wild
alexanderwild.com



Almost Dr. Megan Thoemmes



ant and spider by Matt Bertone (Mill moth by anonymous Internet person)





Dr. Carlos Goller



2. BROADER BIODIVERSITY (HEARTS)



ALEXIS ROCKMAN, THE ECOTOURIST

Heart Transplantation in Man

Developmental Studies and Report of a Case

*James D. Hardy, MD, Carlos M. Chavez, MD, Fred D. Kurrus, MD,
William A. Neely, MD, Sadan Eraslan, MD, M. Don Turner, PhD,
Leonard W. Fabian, MD, and Thaddeus D. Labecki, MD, Jackson, Miss*

HEART TRANSPLANTATION has interested many investigators.¹⁻⁵ Studies of related problems were begun in our laboratory in 1956. Webb and his associates studied such factors as practical methods for homologous cardiac transplantation,⁶ cardiopulmonary transplantation,^{7*} restoration of function of the refrigerated heart,⁸ and cardiac metabolism as influenced by ischemia and refrigeration.¹⁰ The operative mortality was high, but extended survival of some dogs with orthotopic homotransplants was achieved. Collateral studies were conducted by other members of the department.¹¹⁻¹⁴ Thus, in the spring of 1963, Webb and the senior author (J.D.H.) considered that the laboratory and clinical heart work justified a planned approach directed toward eventual heart transplantation in man. This objective, a natural outgrowth of transplantation research, was cleared with the administrative officials of the University Medical Center.

It remained for us to evaluate further the methods available for preservation of the heart during its transplantation. While resuscitation of the transplanted heart preserved with cold arrest could be achieved in most animals, it was essential that restoration of a good beat be assured for the first heart transplanted in man. Coronary artery perfusion had previously proved superior to ice slush during aortic valve surgery in animals¹⁵ and in human beings and it was now re-examined. However, the coronary arteries of the dog were often too small and too variable to be perfused consistently with the equipment available, and ventricular ischemia and infarction occurred with some frequency.

Meanwhile, two operating teams had been established, one of which would obtain the heart from

the donor and the other would prepare the recipient. One team continued heart transplantation in dogs (W.R.W.), but the other (J.D.H.) turned to a study of beef hearts and eventually to the use of the hearts of infant calves. It was found that the coronary arteries of even newborn calves were almost equal in size to those of the human adult, and that perfusion of both the right and left coronary arteries with cold oxygenated blood under gravity flow could routinely be achieved within three minutes from the time of excision of the organ. These small calves tolerated the supine position poorly, and ventricular fibrillation frequently occurred even before intrapericardial dissection had been initiated. Even so, the animals proved satisfactory for our purposes, in that coronary artery perfusion was regularly achieved without undue delay, following which the continuously perfused organ was transferred to an adjacent operating table and inserted into the new host who was supported with the pump oxygenator. The two operating teams were combined to complete this final insertion of the transplant. The extracorporeal circuit was primed with "sterile" but unmatched blood obtained from a slaughterhouse two hours previously, and the suture technique described by Lower and associates² was employed. In contrast to experience with heart transplantation in dogs, anastomotic bleeding rarely constituted a problem in calves, whose tissues were tough and held sutures without tearing. Digitalization was found helpful when used cautiously,⁴ but on some occasions the digitalis produced heart block which often responded dramatically to the intravenous infusion of isoproterenol (Isuprel) hydrochloride.

Approximately 50 calves were used in the successful completion of 20 homotransplants.¹⁶ Coronary artery perfusion was found to be satisfactory



In the first transplant the patient lived!



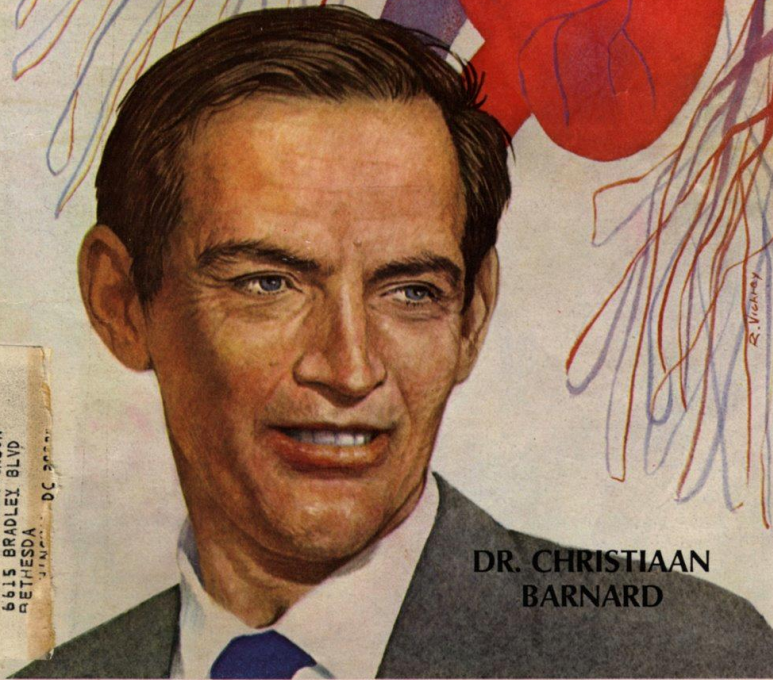
FIFTY CENTS •

DECEMBER 15, 1967

THE TRANSPLANTED HEART

TIME

THE WEEKLY NEWSMAGAZINE

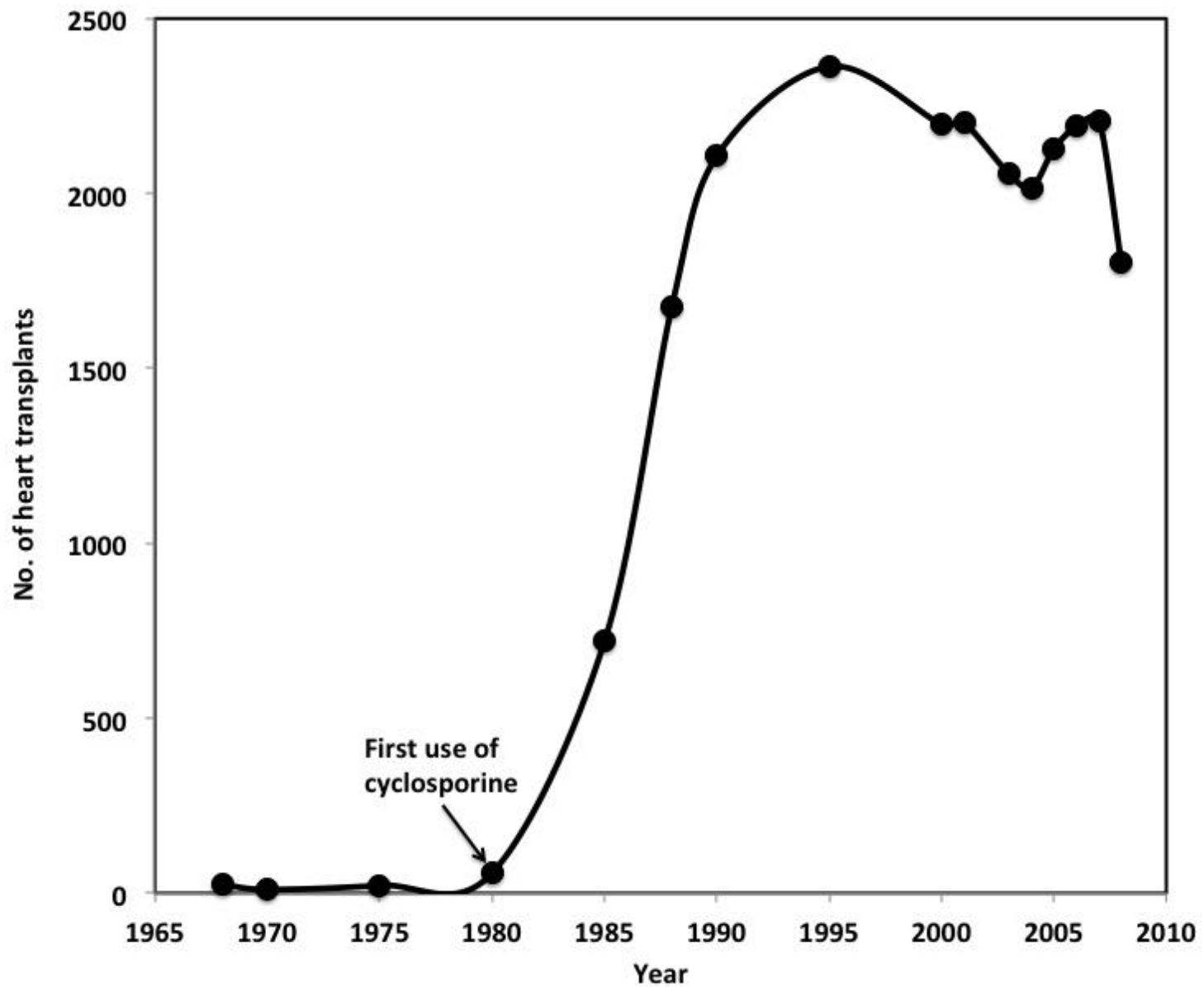


DR. CHRISTIAAN
BARNARD

FEB 71 FED BAS 50 3152 00 34
DR D S FREDRICKSON
6615 BRADLEY BLVD
BETHESDA
M D 20814 DC 30000

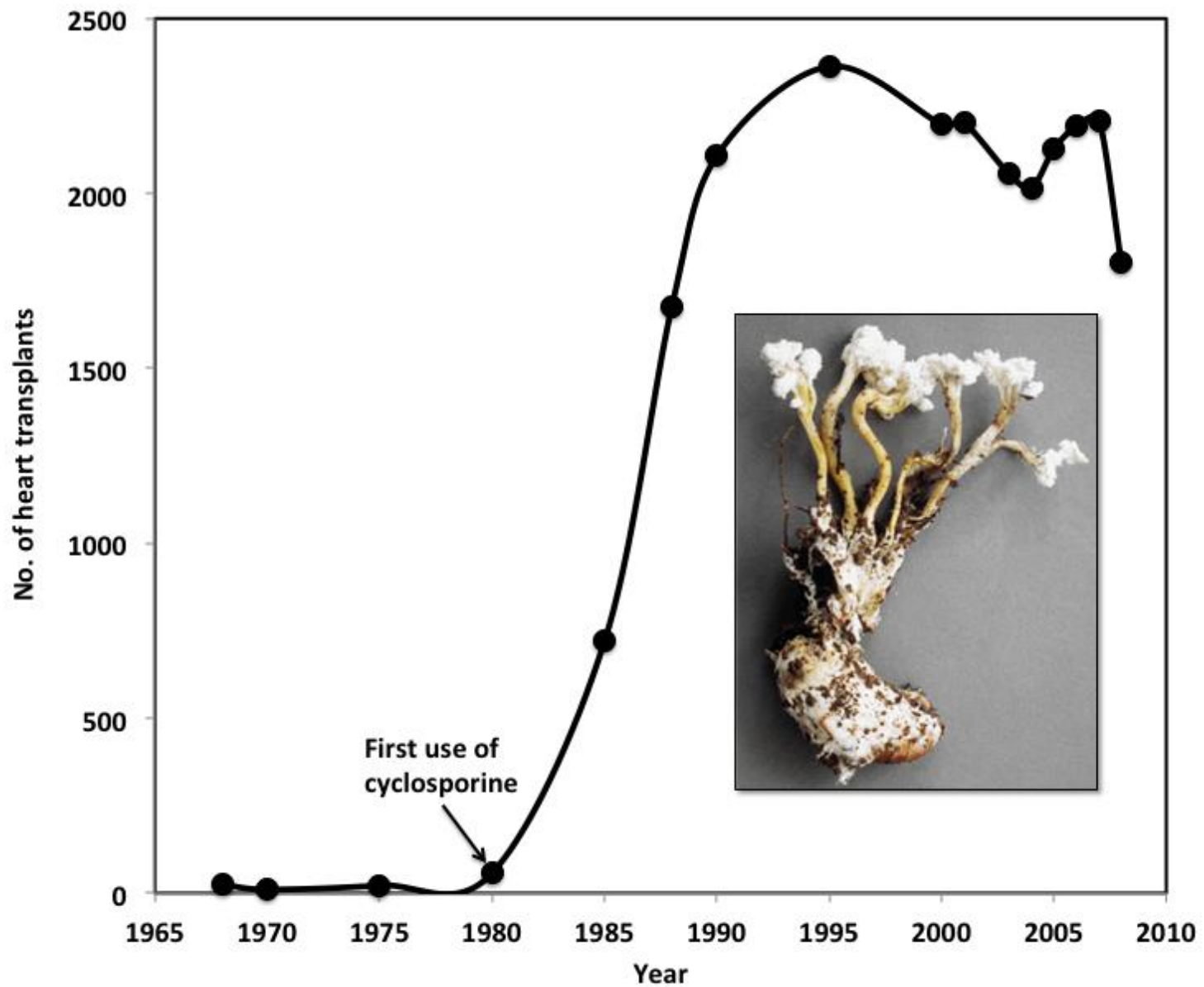
But he lived just 18 days.

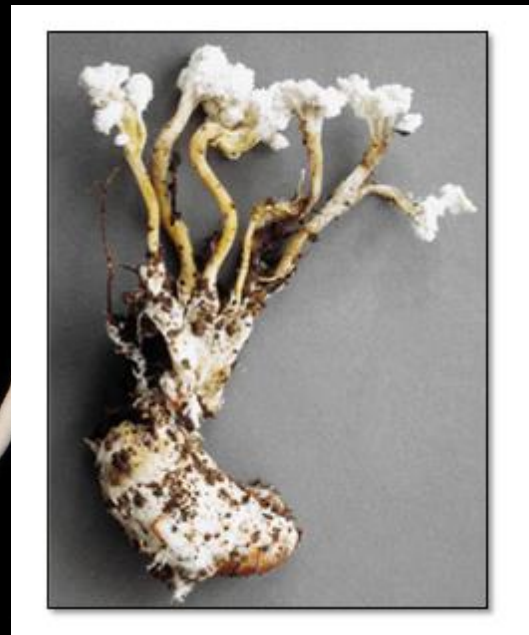




The secret was Cyclosporine.







Kathie Hodge, Cornell

The 'Baby Fae' baboon heart transplant – potential cause of rejection

[David K.C. Cooper](#), [Hidetaka Hara](#), [C. Adam Banks](#), [David Cleveland](#), and [Hayato Iwase](#)

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Baby Fae, received a baboon heart transplant. Rejected the heart after several weeks (1984).

Pig heart transplant

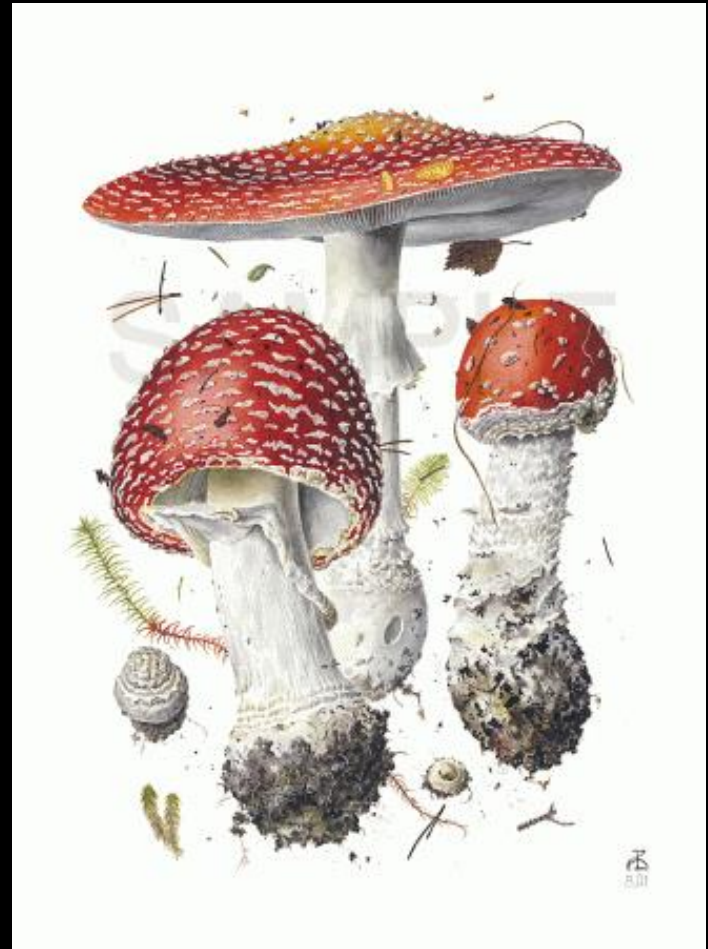


Akira Endo

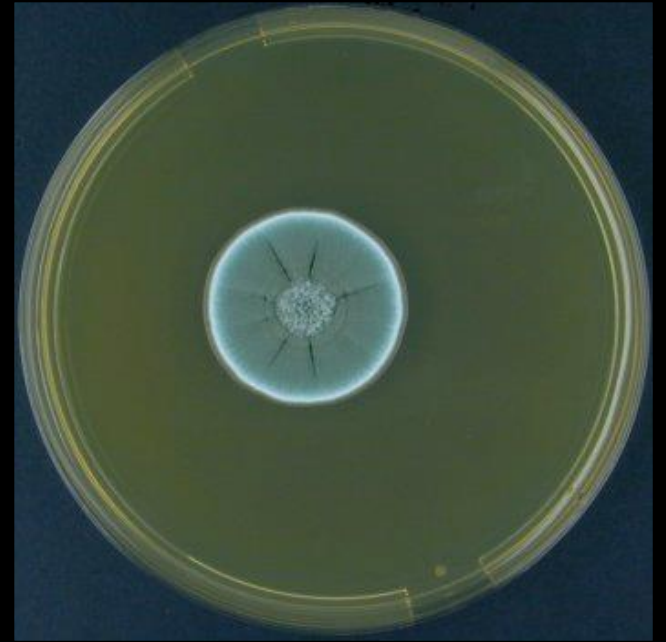


1965





6000 fungal strains



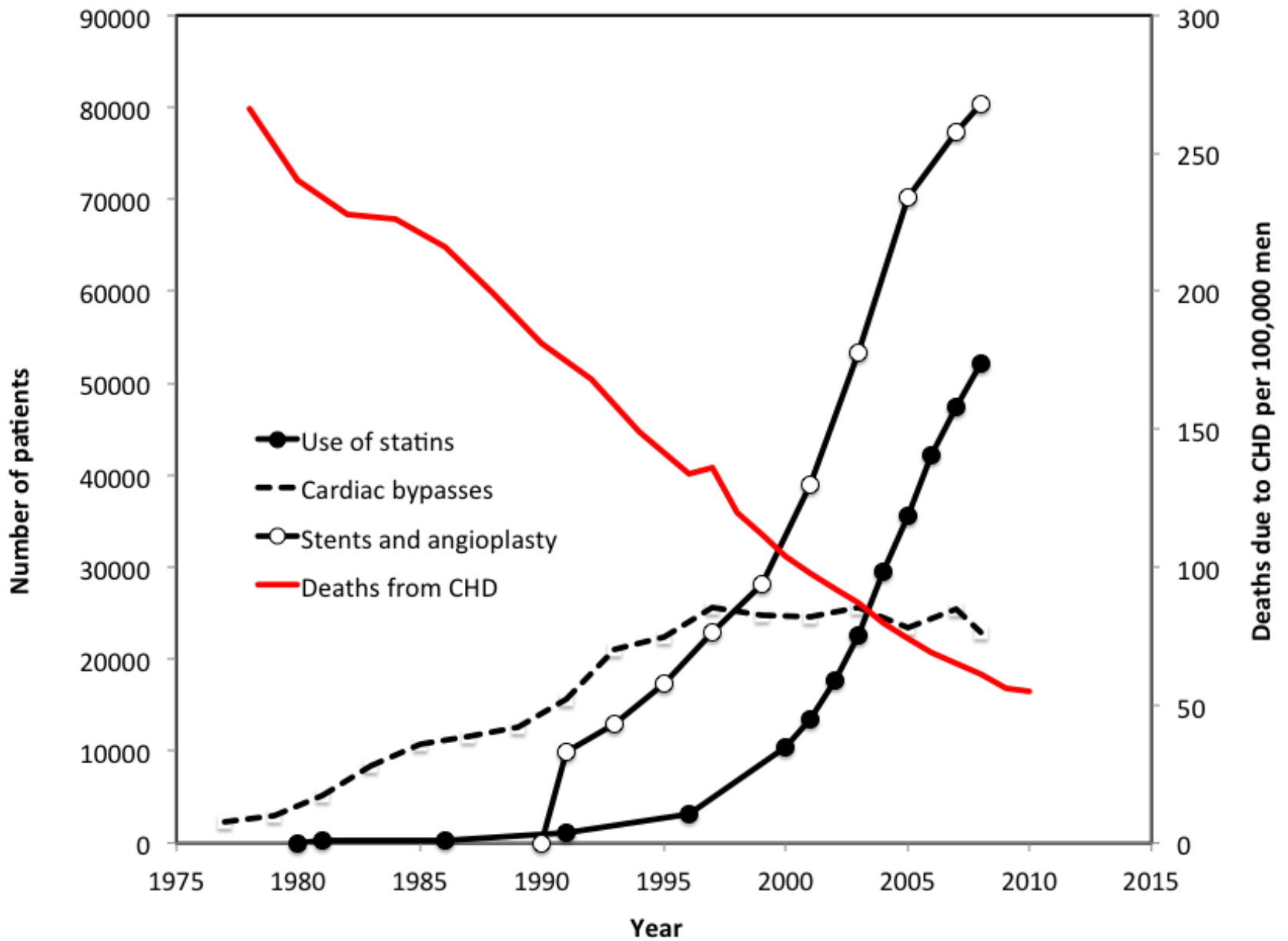
Checked for inhibition of HMG-CoA reductase



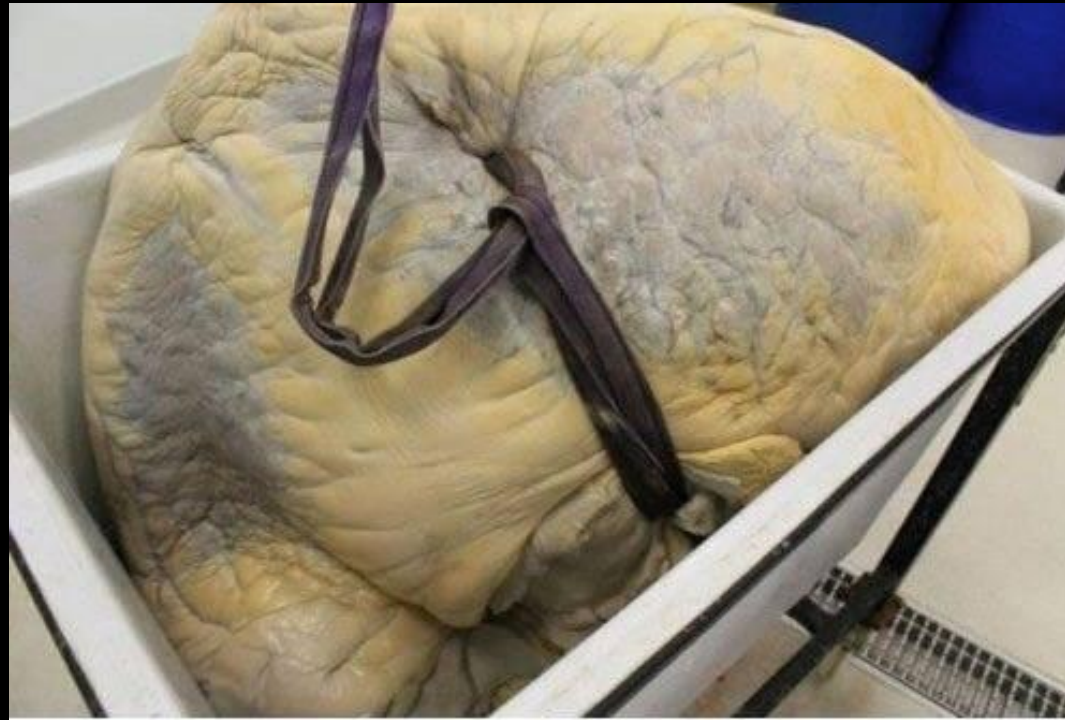
In 1971, citrinin works but causes kidney stones.
In 1973, compactin works in petri dishes, but not in rats

“a wonderful gift from nature”

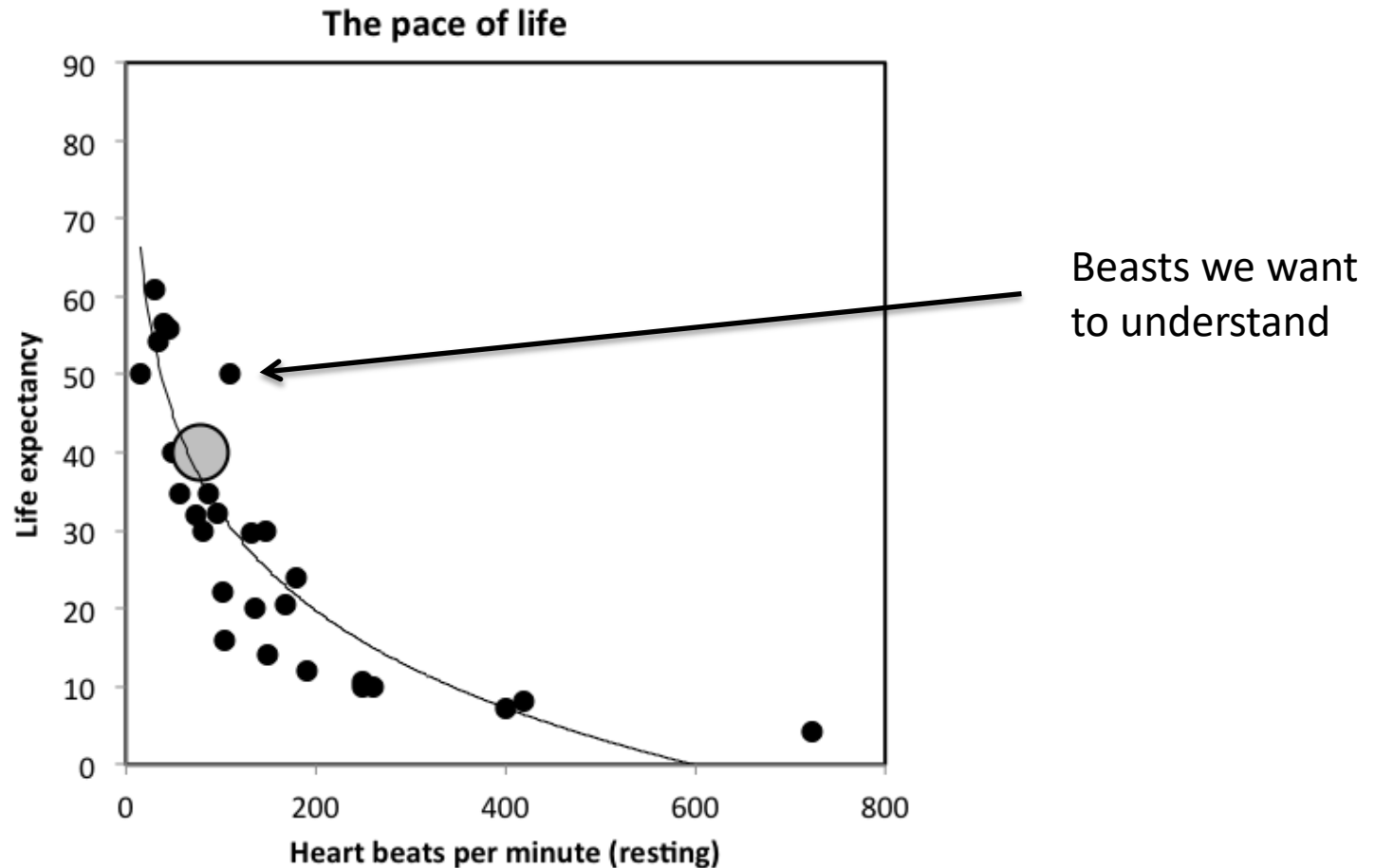




A Baby Blue Whale Heart



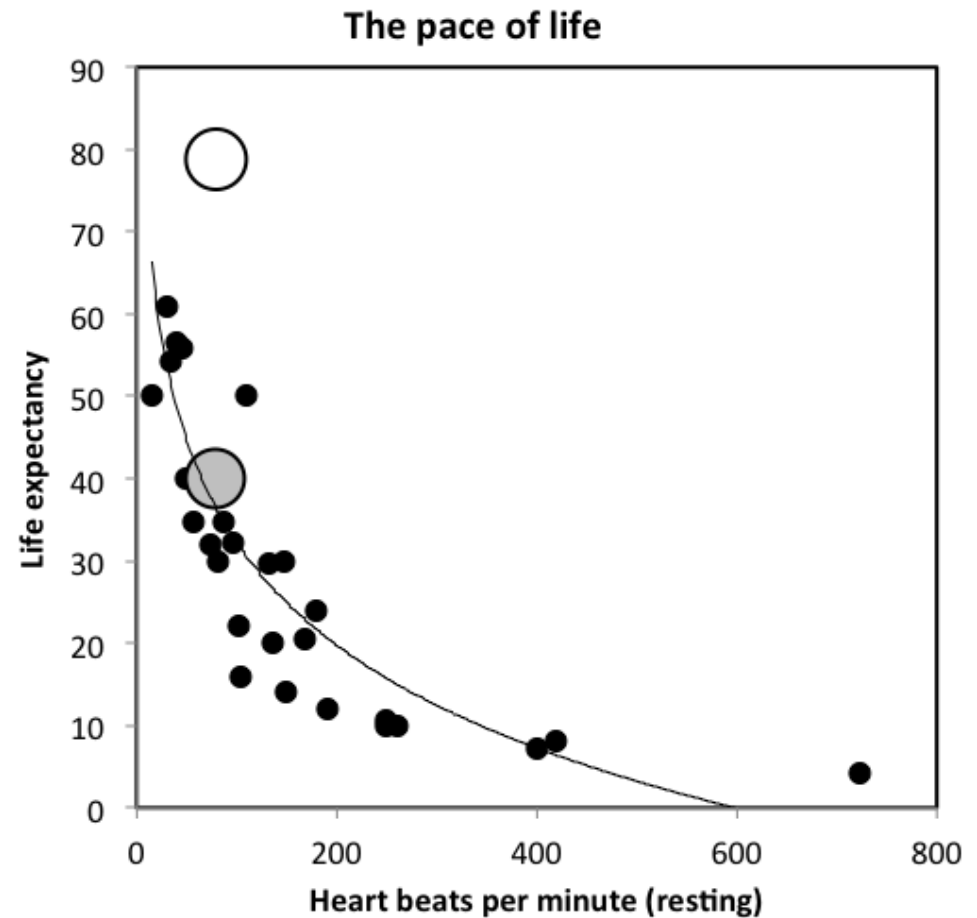
The organisms that get longer lives offer insights into longevity.



Adenosine



Humans get about 1.5 billion extra beats.



WE ARE JUST BEGINNING TO EXPLORE

1.5 MILLION BEETLES
5 MILLION INSECTS
7 MILLION ARTHROPODS
80% UNDISCOVERED





*Pseudomonas
aeruginosa*
photo of colony by
Scott Chimileski



Scaling laws predict global microbial diversity

Kenneth J. Locey^{a,1} and Jay T. Lennon^{a,1}

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Edited by David M. Karl, University of Hawaii, Honolulu, HI, and approved March 30, 2016 (received for review October 27, 2015)

Scaling laws underpin unifying theories of biodiversity and are among the most predictively powerful relationships in biology. However, scaling laws developed for plants and animals often go untested or fail to hold for microorganisms. As a result, it is unclear whether scaling laws of biodiversity will span evolutionarily distant domains of life that encompass all modes of metabolism and scales of abundance. Using a global-scale compilation of ~35,000 sites and ~5.6·10⁶ species, including the largest ever inventory of high-throughput molecular data and one of the largest compilations of plant and animal community data, we show similar rates of scaling in commonness and rarity across microorganisms and macroscopic plants and animals. We document a universal dominance scaling law that holds across 30 orders of magnitude, an unprecedented expanse that predicts the abundance of dominant ocean bacteria. In combining this scaling law with the lognormal model of biodiversity, we predict that Earth is home to upward of 1 trillion (10¹²) microbial species. Microbial biodiversity seems greater than ever anticipated yet predictable from the smallest to the largest microbiome.

biodiversity | microbiology | macroecology | microbiome | rare biosphere

The understanding of microbial biodiversity has rapidly transformed over the past decade. High-throughput sequencing and bioinformatics have expanded the catalog of microbial taxa by orders of magnitude, whereas the unearthing of new phyla is reshaping the tree of life (1–3). At the same time, discoveries of novel forms of metabolism have provided insight into how microbes persist in virtually all aquatic, terrestrial, engineered, and host-associated ecosystems (4, 5). However, this period of discovery has uncovered few,

hold across genomes, cells, organisms, and communities of greatly varying size (13–15). Among the most widely known are the scaling of metabolic rate (B) with body size [M ; $B = B_0 M^{3/4}$ (13)] and the rate at which species richness (i.e., number of species; S) scale with area [A ; $S = cA^z$ (16)]. These scaling laws are predicted by powerful ecological theories, although evidence suggests that they fail for microorganisms (17–19). Beyond area and body size, there is an equally general constraint on biodiversity, that is, the number of individuals in an assemblage (N). Often referred to as total abundance, N can range from less than 10 individuals in a given area to the nearly 10³⁰ cells of bacteria and archaea on Earth (6, 7). This expanse outstrips the 22 orders of magnitude that separate the mass of a *Prochlorococcus* cell (3·10⁻¹⁶ kg) from a blue whale (1.9·10⁵ kg) and the 26 orders of magnitude that result from measuring Earth's surface area at a spatial grain equivalent to bacteria (5.1·10²⁶ μm²).

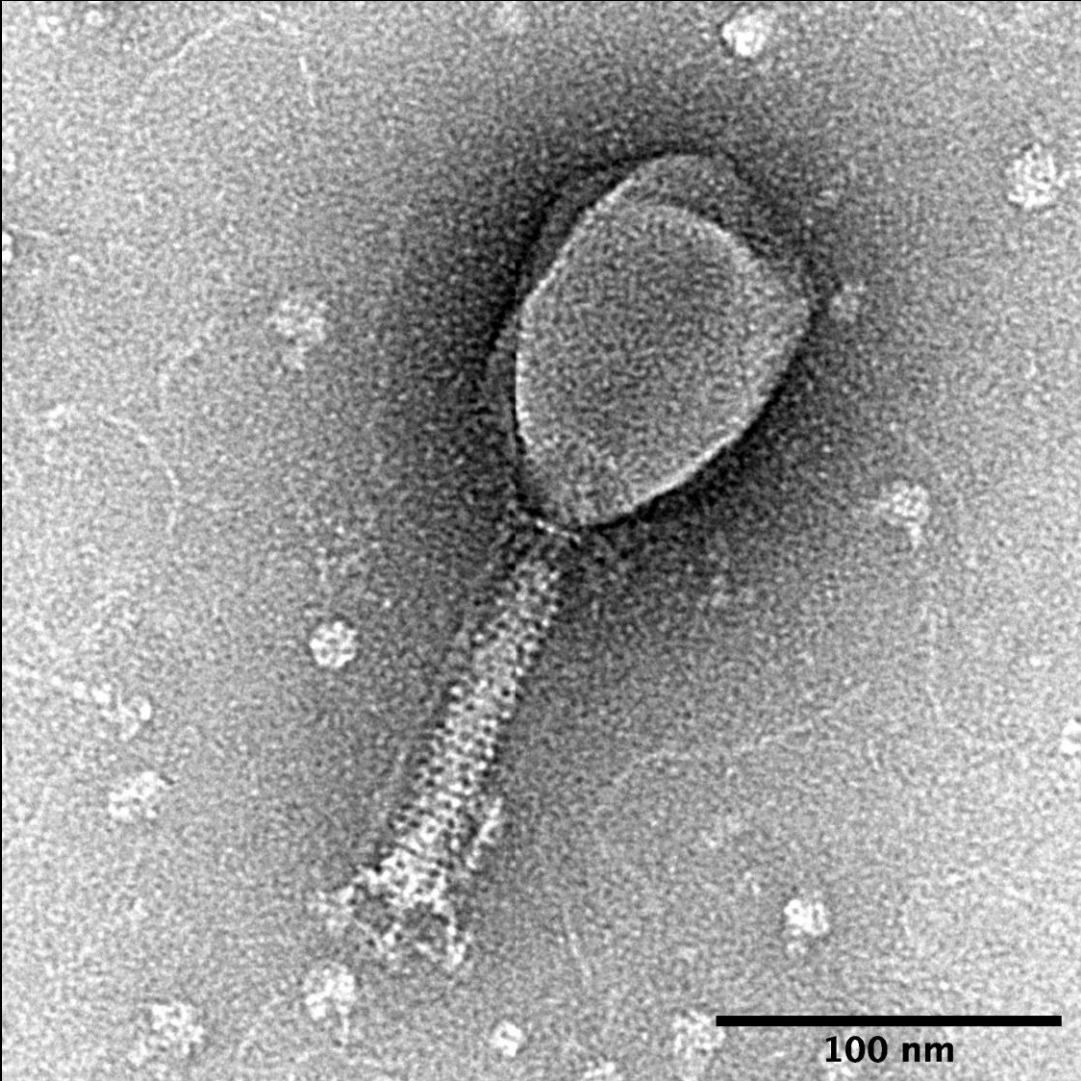
Here, we consider whether N may be one of the most powerful constraints on commonness and rarity and one of the most expansive variables across which aspects of biodiversity could scale. Although N imposes an obvious constraint on the number of species (i.e., $S \leq N$), empirical and theoretical studies suggest that S scales with N at a rate of 0.25–0.5 (i.e., $S \sim N^z$ and $0.25 \leq z \leq 0.5$) (20–22). Importantly, this relationship applies to samples from different systems and does not pertain to cumulative patterns (e.g., collector's curves), which are based on resampling (20–22). Recent studies have also shown that N constrains universal patterns of commonness and rarity by imposing a numerical constraint on how abundance varies among species, across space, and through time (23, 24). Most notably, greater N leads to increasingly uneven distributions

1,000,000,000,000 SPECIES

1,000,000,000,000 SPECIES
(IF OFF BY 3 ZEROS, STILL A
BILLION)

1,000,000,000,000 SPECIES
(IF OFF BY 3 ZEROS, STILL A
BILLION)

JUST 30,000 ARE NAMED



Electron micrograph of T2 bacteriophage

Two Sets of Intertwined Stories (of our relationship to the rest of life)

