

Microbes on fish skin display distinct evolutionary patterns

Michael Doane Ph.D
Megan M. Morris
Bhavya Papudeshi
Lauren Allen
Dnyanada Pande
John M. Haggerty
Shaili Johri
Abby Turnland
Meredith Peterson
Dovi Kacev
Andy Nosal
Deni Ramirez
Kevin Hovel
Julia Ledbetter
Amanda Alker
Jackeline Avalos
Kristi Baker
Shruti Bhide
Emma Billings
Steven Byrum
Molly Clemens
Amelia Juliette Demery
Lais Farias Oliveira Lima
Oscar Gomez
Omar Gutierrez
Selena Hinton
Donald Kieu
Angie Kim
Rebeca Loaiza
Alexander Martinez
Jordan McGhee
Kristine Nguyen
Sabrina Parlan
Amanda Pham
Rosalyn Price-Waldman

Robert A. Edwards
Elizabeth A. Dinsdale

Video Byte

Keywords: Microbiome, phyllosymbiosis, metagenomics, elasmobranch, skin, teleost, vertebrate fishes, microbial community, community ecology, Chondrichthyes, Osteichthyes, shotgun metagenomics, clade, host-microbiome similarity, functional composition, assemblage, holobiont, cartilaginous, bony fish, denticle

Posted Date: October 28th, 2020

DOI: <https://doi.org/10.21203/rs.3.rs-99202/v1>

License:   This work is licensed under a Creative Commons Attribution 4.0 International License.
[Read Full License](#)

Abstract

Microbiomes share an intimate relationship with the organisms they colonize, even across evolutionary timescales. That's the basis of a theory called phylosymbiosis. Phylosymbiosis holds that microbial communities evolve as their host evolves and has been confirmed to exist for certain insects and mammals. Researchers recently tested whether that relationship holds among fish. Approximately 420 million years ago, fish made an epic evolutionary split into elasmobranchs – creatures with all-cartilage skeletons – and bony fish. Since then, the two have accumulated vast differences in anatomy and physiology, most notably in their skin. That's where the researchers zeroed in. For a small sample of fish, they used metagenomics to compare the makeup of microbial communities living on fish skin. Between fishes considered closely or distantly related in evolutionary terms, findings revealed that elasmobranchs displayed patterns of phylosymbiosis, while bony fish did not. A difference that might be linked to alternative processes of microbiome assembly. Scaling up to a larger sample size could offer researchers further insight and help explain the skin microbiome dynamics of marine fishes.