



# A microbial consortia approach is crucial for understanding human health

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# African Microbiomes hold the key to the future of microbiome research!

### Roadmap to expanding microbiome research in Africa

#### nature reviews microbiology

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Comment | Published: 16 June 2023

#### African microbiomes matter

Thulani P. Makhalanyane , Oliver K. I. Bezuidt, Rian E. Pierneef, Eshchar Mizrachi, Adolphe Zeze,

Romain K. Fossou, Claude Ghislaine Kouadjo, Samuel Duodu, Chioma B. Chikere, Olubukola O. Babalola,

Ashwil Klein, Marshall Keyster, Mornè du Plessis, Nourou S. Yorou, Mohamed Hijri, Theresa Rossouw,

Casper N. Kamutando, Stephanus Venter, Lucy N. Moleleki & Colin Murrell

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430 Accesses | 182 Altmetric | Metrics

#### Coordinated research networks

Continental microbiome research networks



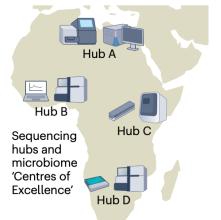
#### **Policy interventions**

Coordinated continental science plans to advance microbiome research



#### Infrastructure development

Urgent need to establish the sequencing capacity and research infrastructure



#### **Comment**

https://doi.org/10.1038/s41591-024-03026-2

# Microbiome research in Africa must be based on equitable partnerships

Ovokeraye H. Oduaran, Moréniké Oluwátóyìn Foláyan, Arox W. Kamng'ona, Annettee Nakimuli, Lamech M. Mwapagha, Mathabatha E. Setati, Michael Owusu, Nicola Mulder. Thulani P. Makhalanyane & Soumaya Kouidhi

Check for updates

#### Table 1 | Pillars of an implementation framework for African microbiome research

Pillar	Rationale	
Local research leadership	Empowering local scientists and scholars ensures consideration of cultural and contextual relevance in research efforts, fostering community-engaged research and network strengthening.	
Contextualized global research	Addressing locally relevant public health priorities and public health research priorities, while aiming for globally applicable solutions, is crucial to the advancement of microbiome research on the continent.	
Ethical and equitable partnerships	Fair engagement practices with shared goals and objectives, clear guidelines on samples and data ownership, along with future-minded initiatives bode well for long-lasting, mutually beneficial collaborations.	
Standardized microbiome protocols	Standardized procedures suitable for the African setting, ranging from sample collection and storage practices to laboratory and analytical approaches, will facilitate reproducibility and consistency in research efforts.	
Governmental involvement	The role of national governments is vital for policy development, enabling conducive regulatory research environments, and can lead to coordinated efforts to tackle public health priorities.	

#### Research focus

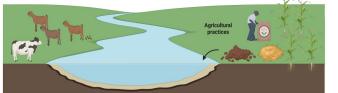
Connection between plants, animals and humans

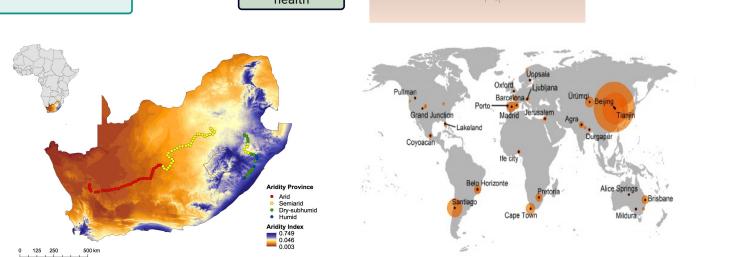
Microbial diversity

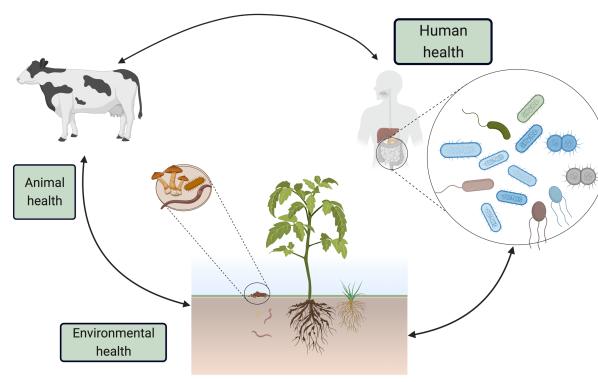
Evolutionary adaptation

Microbial interactions

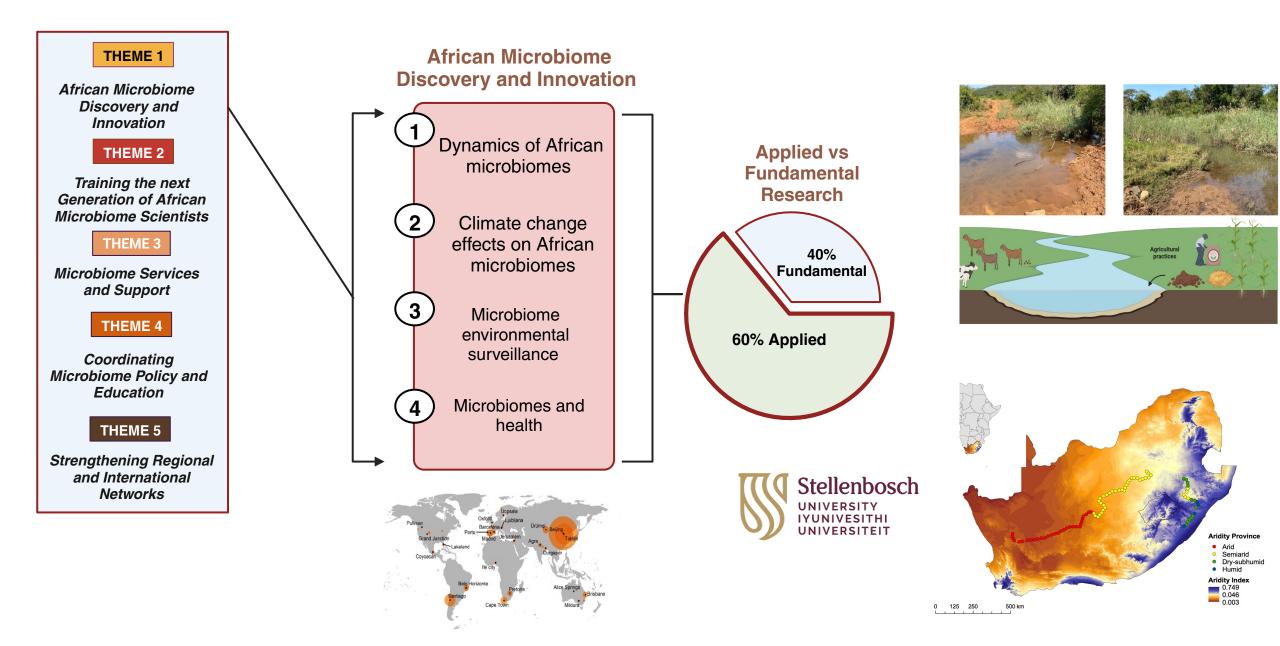








# African Microbiome Group@SUN thematic research areas

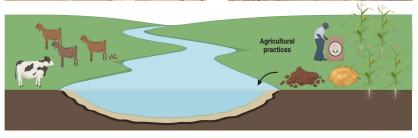


# **Current funded projects**

- 1. Establishing genetic, phylogenetic and functional Mechanisms that shape microbiomes (NRF/NSF)
- 2. Microbiomes as sentinels of change in coastal and marine environments (NRF)
- 3. Leapfrogging the discovery of microbial bioproducts using machine learning
- 4. Understanding the persistence and transmission of antimicrobial resistance genes in African wastewater and freshwater environments.









# **African Microbiome Group Members**













**Postdoctoral Fellows** 









Keith Dube Nombuso Sithole

Runesu Bakasa

SU New PhD students









Mayibongwe Buthelezi Phillip Mawire

Benjamin Abraham Johan Paul Makumbi

UP registered students committed to their degrees









Michelle Bekker Mancha Mabaso

UP registered students who wouldn't leave their moms and boyfriends



# New Honours students and projects



Nobuhle Kambunga

Antimicrobial resistance in children living in informal housing



**Louis Burger** 

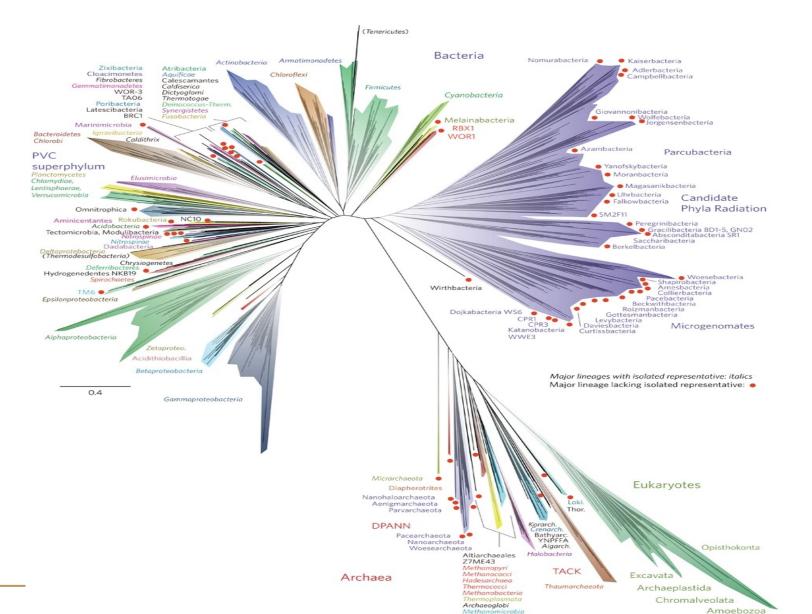
The role of the gut microbiome in long covid



Isabella Blumeris

Childhood stunting in children born from HIV positive mothers

### A new view of microbial diversity





- 1. Role in human **health**
- 2. Role in **biochemical** cycling
- 3. Impact on **climate change**

Hug et al. "A new view of the tree of life." *Nature microbiology* 1.5 (2016): 1-6.

# The complexity of studying microbiomes: Polar and Marine environments as model ecosystems



1

Microbial communities are **numerous** 

Marine sediment

7.85 × 10<sup>3</sup> to 6.10× 10<sup>5</sup> (bacteria)

and
3.28 × 10<sup>4</sup> to 2.46 × 10<sup>6</sup> (archaea)

Widespread horizontal gene transfer, gene loss and convergent evolution may confound the relationship between traits and phylogeny

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REVIEWS

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Examining horizontal gene transfer in microbial communities

Ilana Lauren Brito

Environmental
conditions
disproportionally shape
phylogeny

The ISME Journal (2018) 12:2470–2478 https://doi.org/10.1038/s41396-018-0158-1

**ISME** 

ARTICLE



Microbes in polar and marine environments are excellent models for understanding the effects of change

A strong link between marine microbial community composition and function challenges the idea of functional redundancy

Pierre E. Galand 601 · Olivier Pereira1 · Corentin Hochart2 · Jean Christophe Auguet3 · Didier Debroas2

#### Five lessons from studying environmental microbiomes

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- 1. Ecological context matters
- 2. Microbial interactions are consequential than single microbes
- 3. High diversity leads to community stability
- 4. Microbial communities are subject to succession
- 5. Changes environmental conditions influence microbial evolution and adaptation

#### nature climate change

#### Biogenic factors explain soil carbon in paired urban and natural ecosystems worldwide

Received: 11 July 2022 Accepted: 27 February 2023

Manuel Delgado-Baquerizo ® 1,2 ≥ , Pablo Mark A. Bradford 95, David J. Eldridge 96 Tadeo Sáez-Sandino ®8, Yu-Rong Liu ®9, Sebastian Abades © 10. Adebola R. Bamig José L. Blanco-Pastor14, Jorge Duran @ 15,1 Tine Grebenc<sup>19</sup>, Thulani P. Makhalanyane Tina U. Nahberger<sup>19</sup>, Gabriel F. Peñaloza-I Alexandra Rodríguez 6 15,16, Christina Siek Pankaj Trivedi @ 27, Jay Prakash Verma 21, L Tianxue Yang<sup>26</sup>, Eli Zaady <sup>© 28</sup>, Xiaobing Z



RESEARCH ARTICLE

https://doi.org/10.1038/s41564-022-01266-x

**Bacteria and Archaea Regulate Particulate Organic Matter Export in Suspended and Sinking Marine Particle Fractions** 

Choaro D. Dithugoe, Ab, Coliver K. I. Bezuidt, Emma L. Cavan, William P. Froneman, Sandy J. Thomalla,

#### Global hotspots for soil nature conservation

Published online: 12 October 2022 Check for updates

https://doi.org/10.1038/s41586-022-05292-x Carlos A. Guerra<sup>12,3</sup> , Miguel Berdugo<sup>4</sup>, David J. Eldridae<sup>5</sup>. Nico Eisenhauet<sup>1</sup> Brajesh K. Singh<sup>6,7</sup>, Haiying Cui<sup>6,9</sup>, Sebastian Ab Adebola R. Bamigboye<sup>12</sup>, Felipe Bastida<sup>13</sup>, Jose Jorge Durán 16.77, Tine Grebenc 18, Javier G. Illán 19. nature microbiology even Mamet<sup>22</sup>, Marco A. Molina-Montenegro Jay Prakash Verma<sup>25</sup>, Ana Rey<sup>15</sup>, Alexandra Roc Alberto L. Teixido<sup>31</sup>, Cristian Torres-Díaz<sup>32</sup>, Pank Ling Wang<sup>a</sup>, Jianyong Wang<sup>a</sup>, Eli Zaady<sup>34</sup>, Xiaob

Standardized multi-omics of Earth's microbiomes reveals microbial and metabolite diversity

Van Goethem et al. Microbiome (2018) 6:40 https://doi.org/10.1186/s40168-018-0424-5

Microbiome

#### RESEARCH

**Open Access** 

A reservoir of 'historical' antibiotic resistance genes in remote pristine

Antarctic soils

( CrossMark

Marc W. Van Goethem<sup>1†</sup>, Rian Pierneef<sup>2†</sup>, Oliver K. I. Bezuidt<sup>1</sup>, Yves Van De Peer<sup>1,3,4,5</sup>, Don A. Cowan<sup>1</sup> and Thulani P. Makhalanyane 1\*00



# How do the gut microbiomes of healthy individuals in urban and rural areas of South Africa differ, and what are the functional implications of these differences?



## Studies on African gut microbiomes

**ARTICLE** 

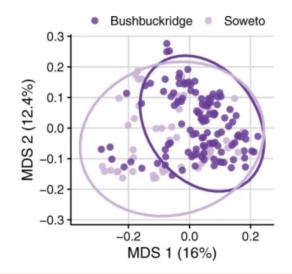


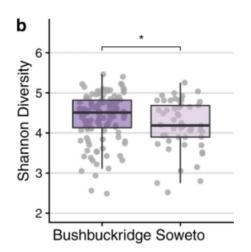
https://doi.org/10.1038/s41467-021-27917-x

OPEN

Short- and long-read metagenomics of urban and rural South African gut microbiomes reveal a transitional composition and undescribed taxa

Fiona B. Tamburini <sup>1</sup>, Dylan Maghini <sup>1</sup>, Ovokeraye H. Oduaran <sup>2</sup>, Ryan Brewster <sup>3</sup>, Michaella R. Hulley <sup>2,4</sup>, Venesa Sahibdeen <sup>4</sup>, Shane A. Norris <sup>5,6</sup>, Stephen Tollman <sup>7,8</sup>, Kathleen Kahn <sup>7,8</sup>, Ryan G. Wagner <sup>7,8</sup>, Alisha N. Wade <sup>7</sup>, Floidy Wafawanaka <sup>7</sup>, F. Xavier Gómez-Olivé <sup>7,8</sup>, Rhian Twine <sup>7</sup>, Zané Lombard <sup>4</sup>, H3Africa AWI-Gen Collaborative Centre, Scott Hazelhurst <sup>2,9,11</sup> & Ami S. Bhatt <sup>1,3,10,11</sup>





Comparatively **few studies** on African gut
 microbiomes

Stellenbosch

- Some evidence showing that the gut microbiome of South Africans does not conform to a simple "western-nonwestern" axis and contains undescribed microbial diversity.
- Due to differences in diets, vital to expand these studies including different regions

# Studies on African gut microbiomes

**SA Gut team (past and present)** 









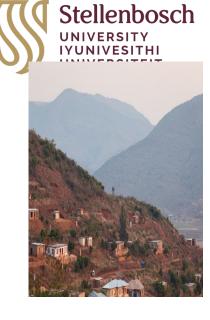


- 1. Elucidate gut microbiomes of healthy **SA individuals**
- 2. We collected stool samples from **Venda** and **Pretoria** (n=100)
- Amplicon sequencing using
   16S and ITS rRNA gene
   markers
- **4. Shotgun** analysis

metagenomic



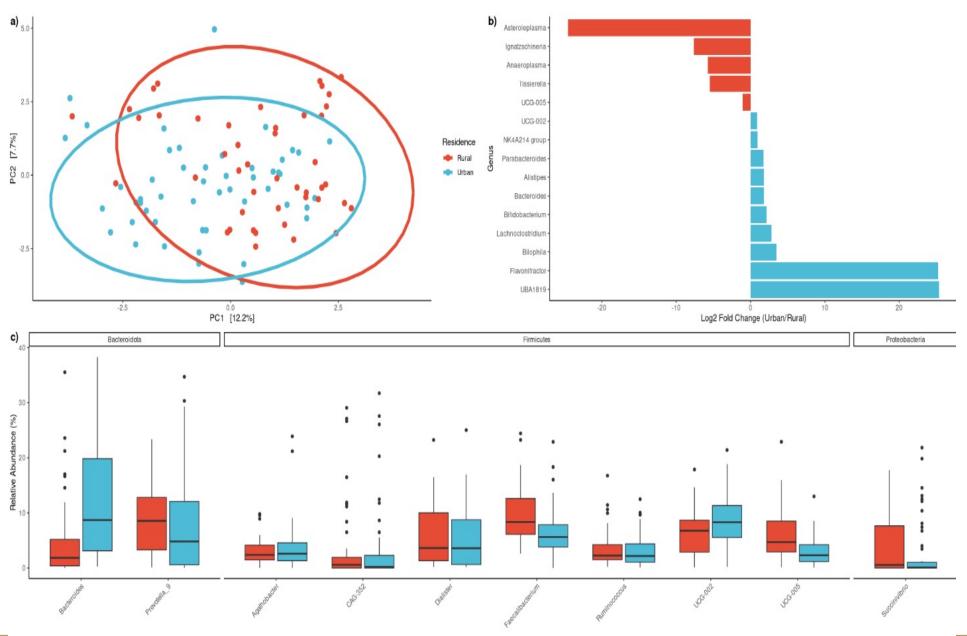




Vhembe district, Limpopo



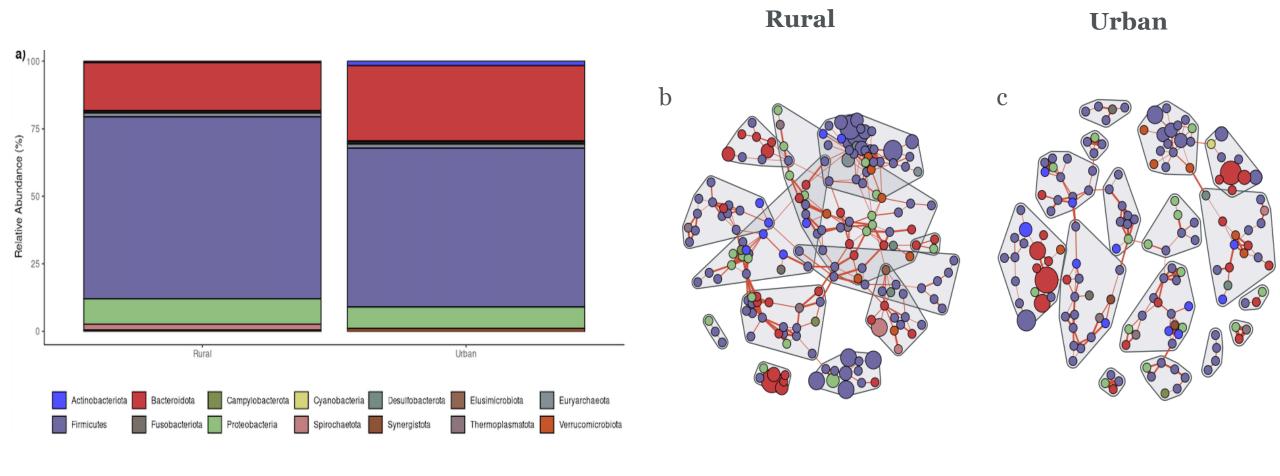
## Structural patterns of bacterial communities



Stellenbosch
UNIVERSITY
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Bars in red and blue
indicate the urban
and rural
populations,
respectively.

Science · EyeNzululwa Thee 100 mostwabundant genera in rural and urban samples

#### Gut microbiomes of rural South Africans more diverse and connect



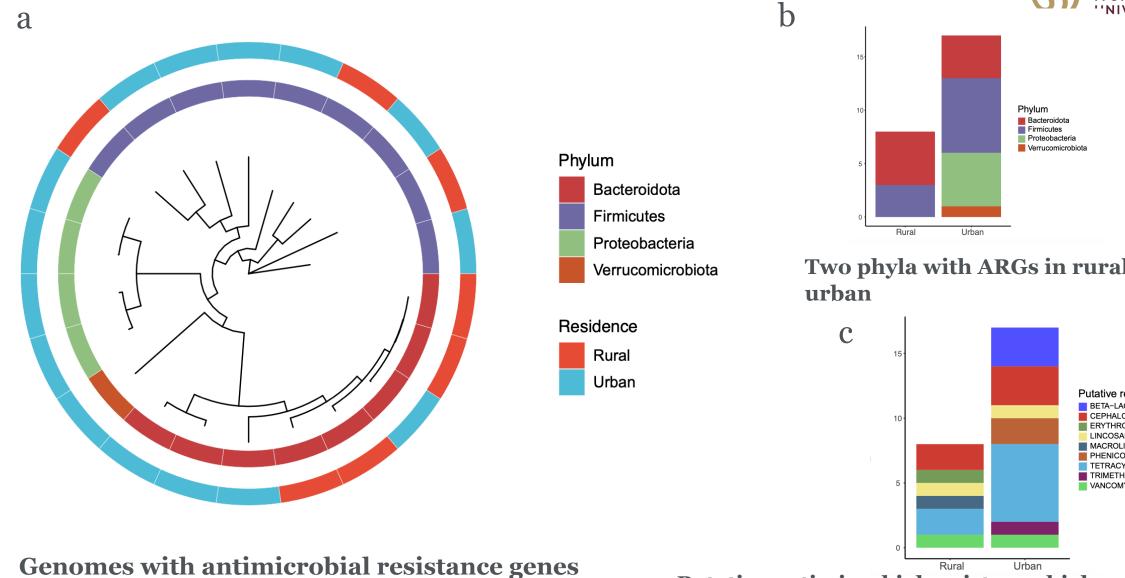
Alpha diversity shows that rural microbiomes more diverse

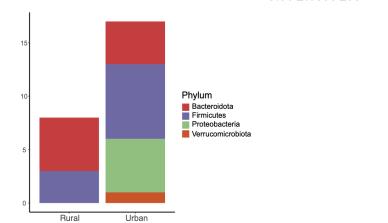
Rural networks are more connected suggesting increased cooperativity in these microbiota

# Metagenomic analysis reveals over 380 diverse high to medium Stellenbosch quality genomes

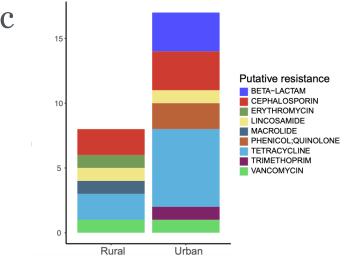
Phylum	Rural	Urban	Total
Actinobacteriota	1	8	9
Bacteroidota	56	41	97
Desulfobacterota	0	1	1
Elusimicrobiota	0	2	2
Firmicutes	125	125	250
Methanobacteriota	0	2	2
Proteobacteria	10	9	19
Spirochaetota	2	0	2
Verrucomicrobiota	0	2	2

South African gene catalogue shows high diversity and ARGs Stellenbosch abundant in urban individuals UNIVERSITY





Two phyla with ARGs in rural vs four in



Putative antimicrobial resistance higher in urban

## **Summary**



- While previous studies have shown the difference between high-income countries and traditional communities, our data suggests some clear differences in urban and rural Africans
- Urban and rural South Africans harbor highly similar gut microbiota
- Results from co-occurence networks suggest clear differences in the **stability** of **rural and urban microbiota**
- These differences may explain the considerably varied functional profiles (based on ARGs)

## Conclusion and future directions



- 1. Studies on environmental ecosystems have provided remarkable insights on microbial evolution
- 2. There are significant gaps in microbiome studies
- 3. A need to build collaborations and partnerships



# Acknowledgements

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School for Data Science and
Computational Thinking



Siyakhula project team

- 1. Theresa Rossouw
- 2. Louisa du Toit



Dr. Rian Pierneef



Dr. Oliver Bezuidt



Dr. Girish Nair











#### **Housing study Core team**

- 1. Ronelle Burger (SU)
- 2. Marisa von Fintel (SU)
- 3. Dumisani Hompashe (UFS)
- 4. Vuyiswa Dlamini (SU)
- 5. Kuhle Ndyonda (SU)
- 6. Lauren Tavener-Smith (SU)
- 7. Renate Campher (SU)
- 8. Eigelaar-Meets (SU)







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