

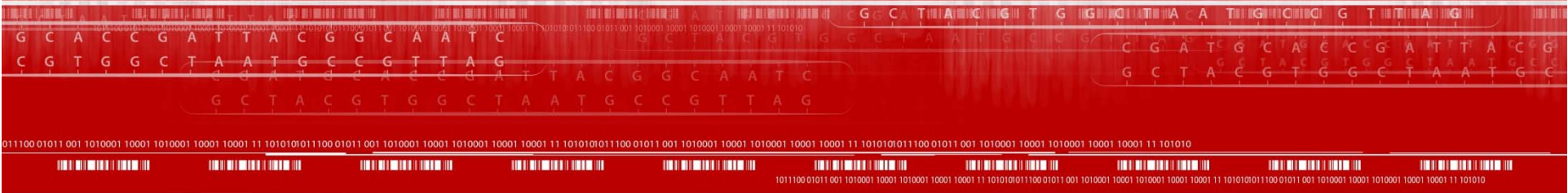
Global One Health and Role of –Omics in Strengthening Integrated Capacity to Address Food- and Waterborne Diseases

Wondwossen A. Gebreyes DVM, PhD, DACVPM
Professor and Exec. Director, Global One Health initiative (GOHi)

Global Water Food Safety Summit
November 19-21, 2019 (College Park, MD)



THE OHIO STATE UNIVERSITY
COLLEGE OF VETERINARY MEDICINE



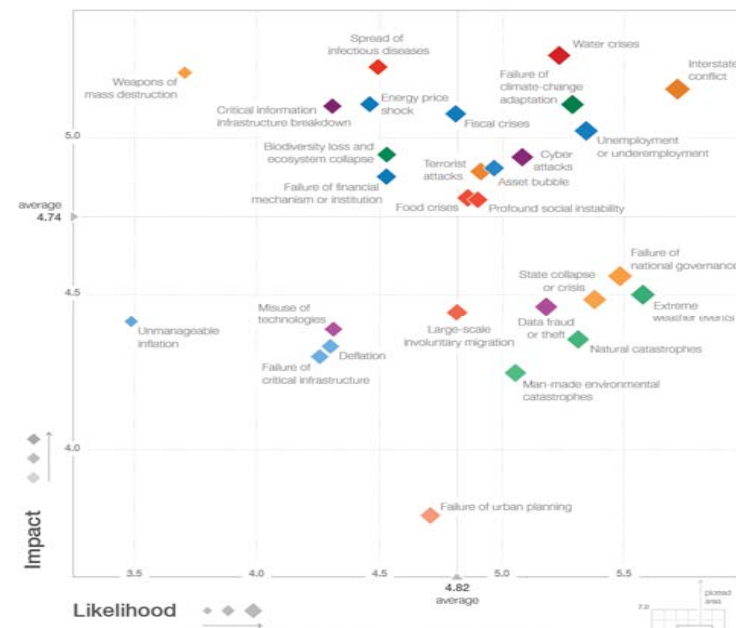


Role in the changing global dynamics.

The Global Risk

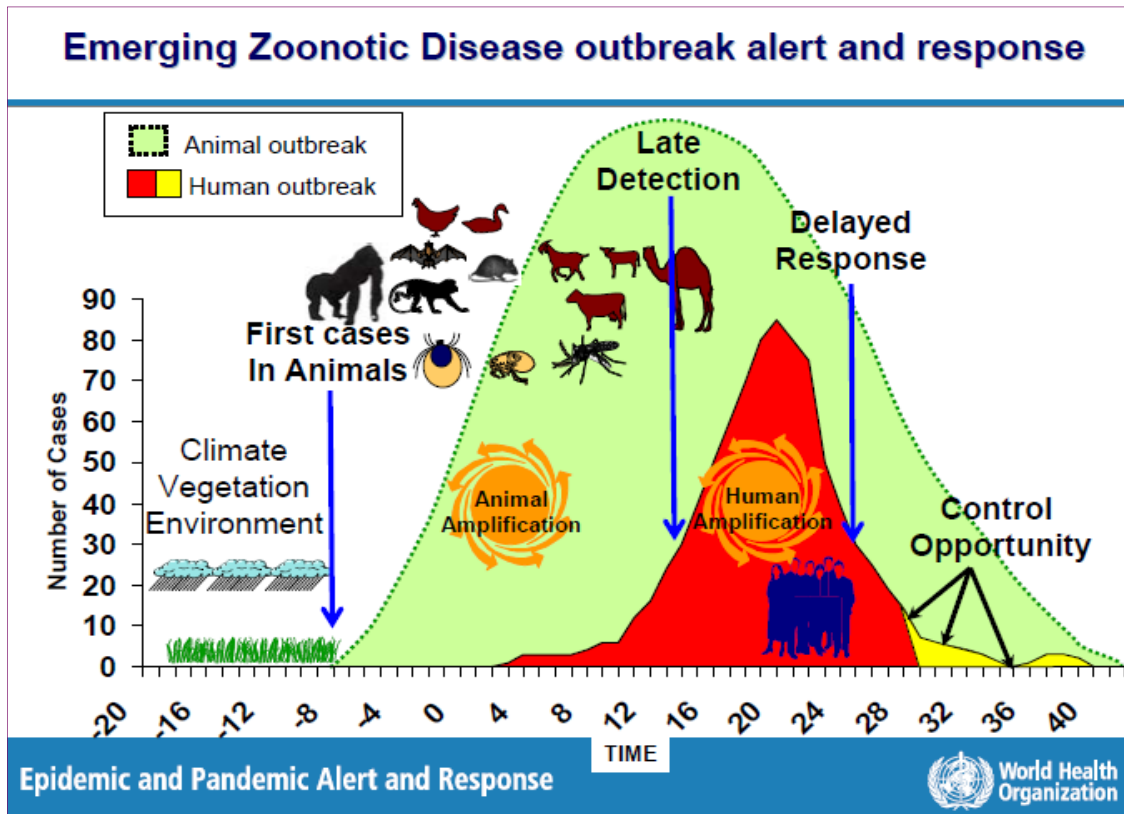
- Spread of infectious diseases;
- **Water crises**;
- Failure of climate change adaptation;
- Global travel;
- Man-made environmental catastrophes;
- Biodiversity loss and ecosystem collapse;
- Large-scale migration, conflict, refugees

Consequences – Very complex, wider in spectrum.



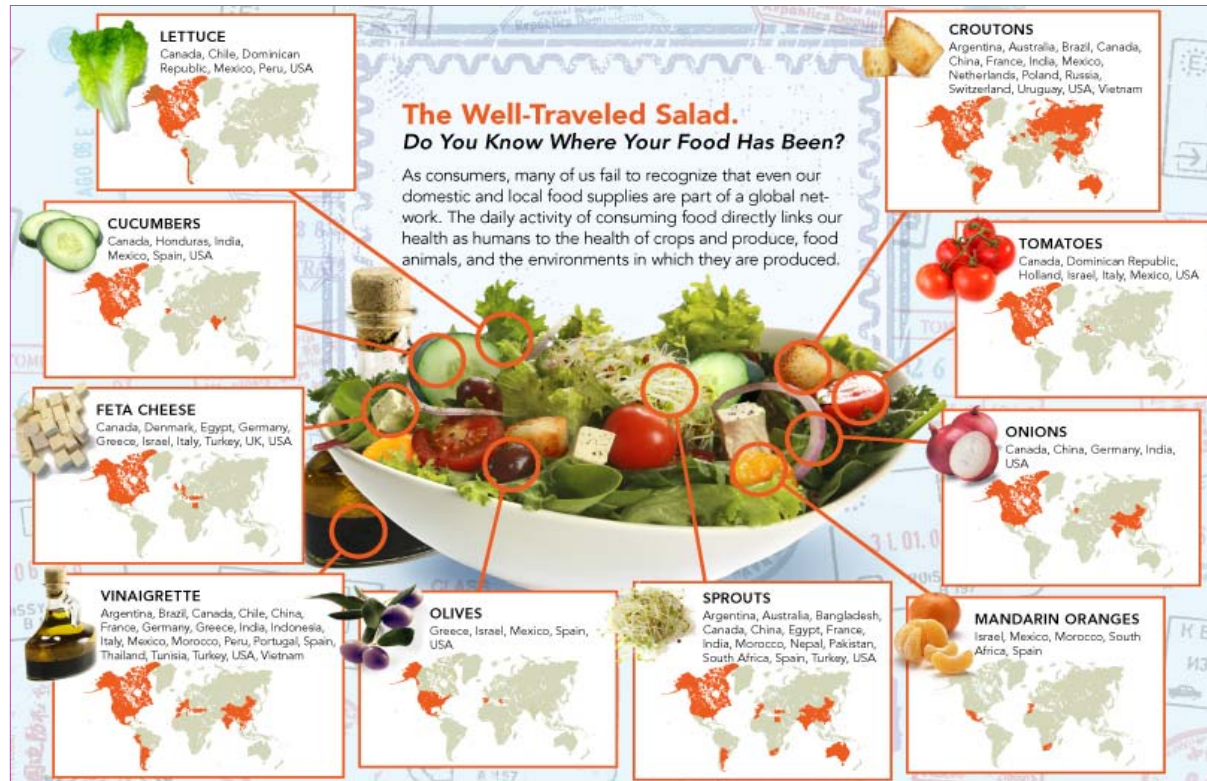
Global Risk Forum, 2016







An expanded paradigm



<http://resources.nationalacademies.org/widgets/Food-origins/flashfile/Food%20origins.html?keepThis=true&>



Climate change and water-borne diseases

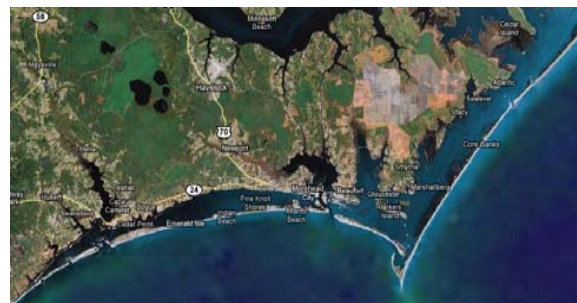
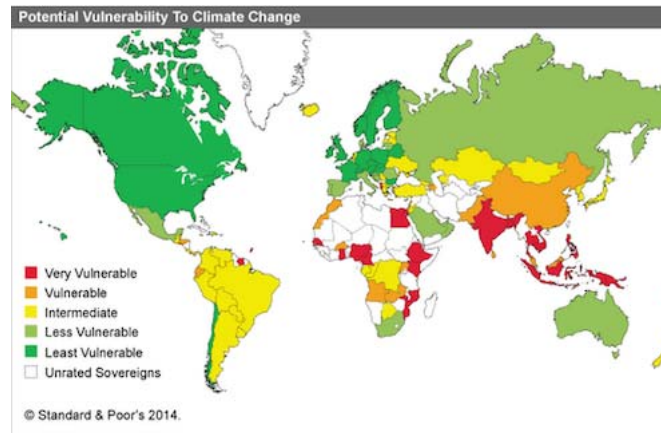
#ClimateChange
WHETHER YOU LIVE IN A...

Rural village Small island or coastal town Big city

CLIMATE CHANGE THREATENS YOUR HEALTH

Drought, floods and heat waves will increase. Vector-borne diseases, like malaria and dengue fever will increase with more humidity and heat.

Between 2030 and 2050 climate change is expected to cause **250 000 ADDITIONAL DEATHS PER YEAR** due to malaria, malnutrition, diarrhoea and heat stress.

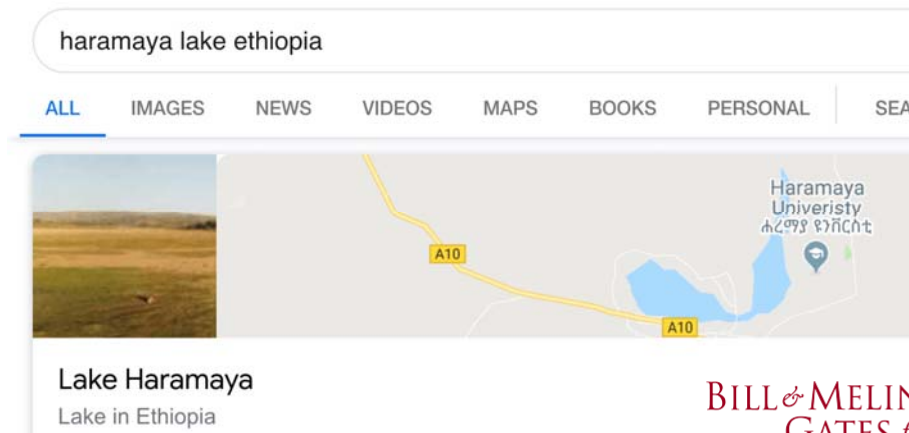




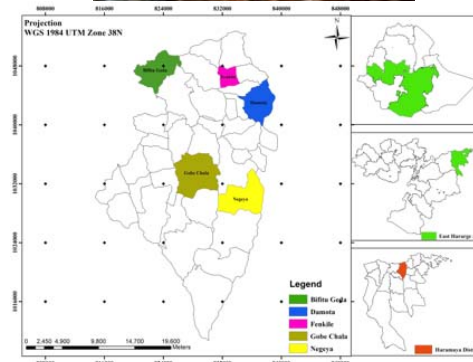
Environmental Enteric Dysfunction (EED)

Impact of water availability on sanitation- Foodborne Pathogens

- Reduced colonization by *Campylobacter*, in children, improve sanitation and hygiene to reduce the prevalence of Environmental Enteric Dysfunction (EED) and thus childhood stunting.
- Haramaya University, Eastern Ethiopia and UFL (USA)

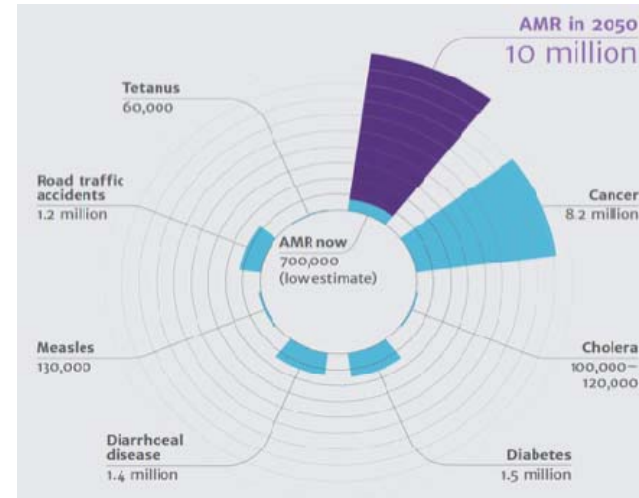
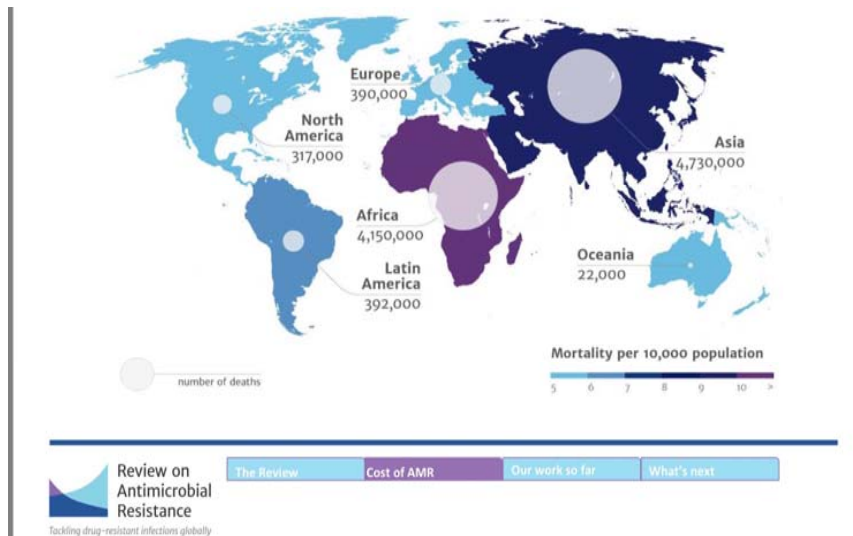


BILL & MELINDA
GATES foundation





The Added Burden of Antimicrobial Resistance (AMR) Global Public Health priority-Crisis

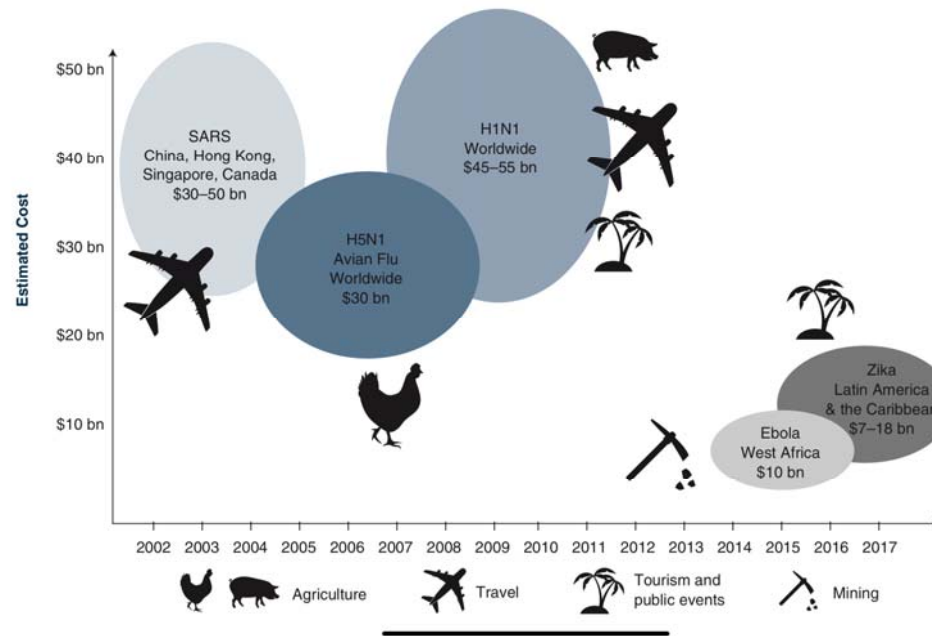


Cumulative economic impact of AMR, > \$100 Trillion by 2050.





Economic Impact of One Health issues- Zoonotic Food- Water- and vector-borne Diseases



“As a physician, the issue of antimicrobial resistance—or AMR—is very familiar to me. ...The problem goes beyond hospitals. Antimicrobial resistance crosses boundaries of nations, sectors, and even species— affecting livestock, crops, and wildlife...

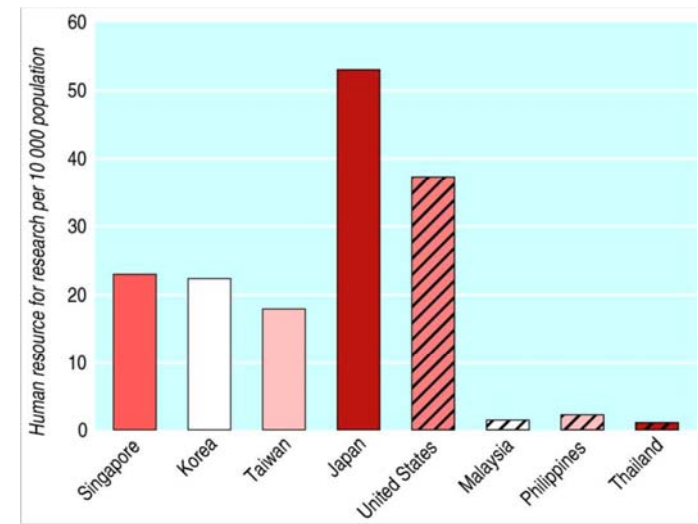
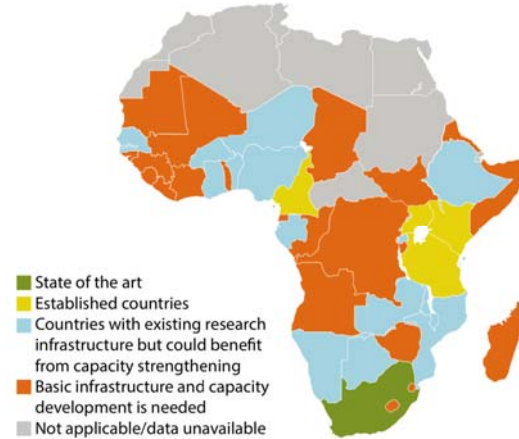
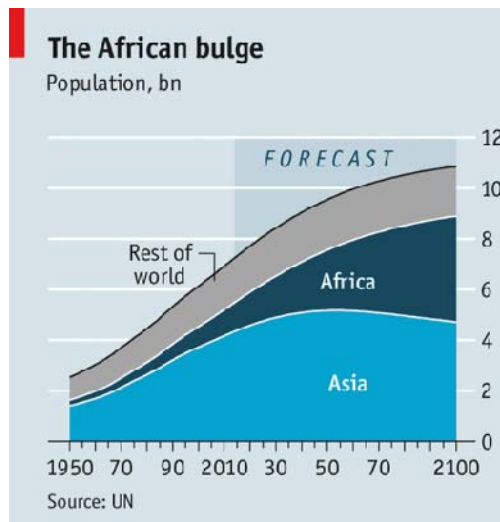
The World Bank is coordinating efforts across agricultural, environmental, and health sectors under the umbrella of One Health—”

Dr. Jim Yong kim, 2014
President (Former)





Population, Global Preparedness on Research Capacity



Sitti-Amorn et al., BMJ (2000)

Demand for Animal Source Food (ASF) and water is increasing at high rate.



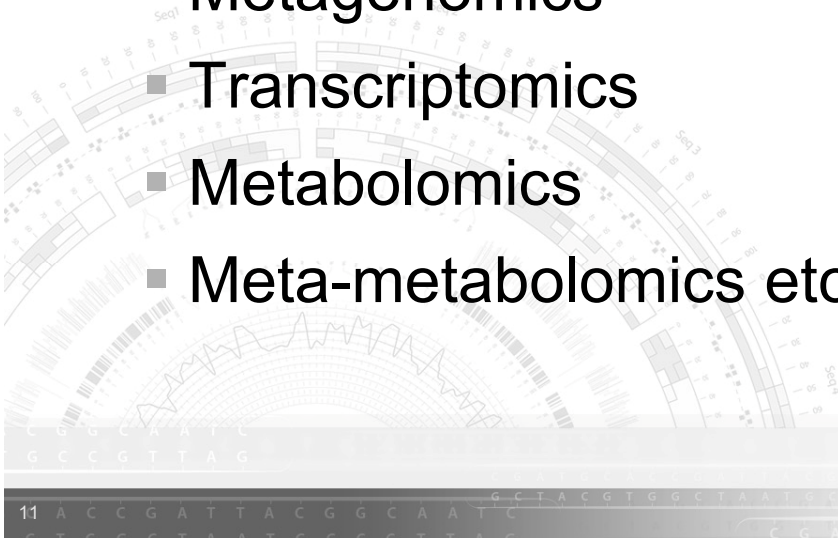


Global investment on Health Research

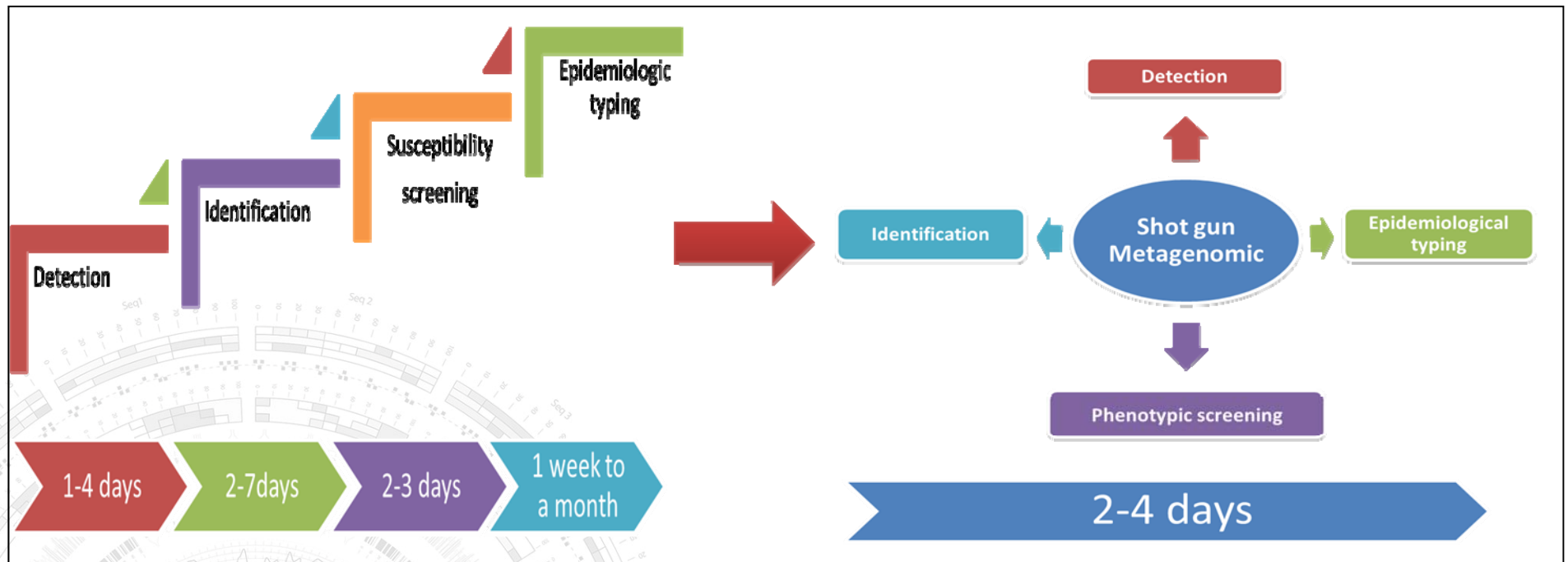
Name of funder	Type of funder	Total Amount Invested (US\$)	Disease area	Total Amount Invested (US\$)	% of total funding allocated
US National Institutes of Health (NIH)	Government	5,795,734,228	HIV/AIDS	5,488,168,543	43.5%
Bill & Melinda Gates Foundation	Private foundation	2,530,342,885	Malaria	2,709,915,003	21.5%
Aggregate Pharmaceutical and Biotechnology Company Respondents	Private	2,037,077,360	TB	2,508,349,300	19.9%
European Commission: Directorate-General for Research and Innovation*	Government	567,311,143	Dengue	781,191,533	6.2%
United States Agency for International Development (USAID)	Government	416,278,263	Kinetoplastids*	706,168,846	5.6%
US Department of Defense (DOD) including DOD Defense Advanced Research Projects Agency (DARPA)	Government	403,029,646	Helminths (Worms & Flukes)	352,659,673	2.8%
The Wellcome Trust	Charity	361,225,501	Leprosy	42,627,803	0.3%
UK Department for International Development (DFID)	Government	348,154,635	Trachoma	19,653,909	0.2%
UK Medical Research Council (MRC)	Government	270,742,509	Buruli Ulcer	17,429,734	0.1%
Institut Pasteur	Private foundation	161,012,834	Grand Total	12,626,164,344	100%
Dutch Directorate General of International Cooperation	Government	128,593,178			
Australian National Health and Medical Research Council	Government	100,613,706			
Grand total		13,120,115,888			

-Omics and associated technologies provides efficient, high-throughput solutions

- Genomics
- Proteomics
- Metagenomics
- Transcriptomics
- Metabolomics
- Meta-metabolomics etc.



Need for innovative and cost-effective systems



- Many tests for one pathogen
- One test for all pathogens

GenomeTrakr



Basic Data Flow for Global WGS Public Access Databases

DATA ACQUISITION

Sequence and upload genomic and geographic data



Other distributed sequencing networks



DATA ASSEMBLY, ANALYSIS, AND STORAGE

International Nucleotide Sequence Database Collaboration (INSDC)
Shared Public Access Databases

- NCBI – National Center for Biotechnology Information
- EMBL – European Molecular Biology Laboratory
- DDBJ – DNA Databank of Japan



PUBLIC HEALTH APPLICATION AND INTERPRETATION OF DATA

- Find clinical links
- Identify clusters
- Conduct traceback
- Develop rapid methods
- Develop culture independent tests
- Develop new analytical software

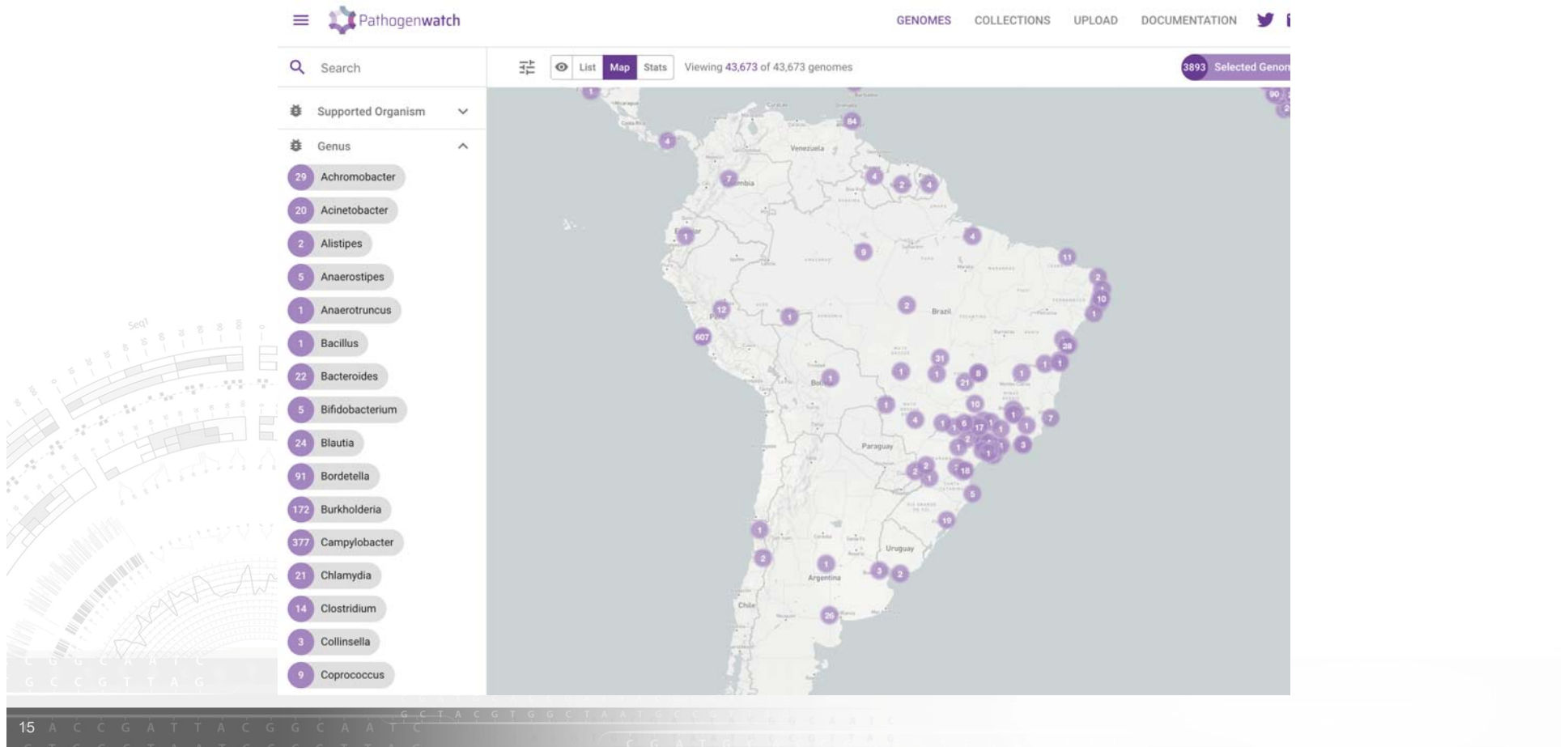


11/2014

State, Local, Federal, and Foreign Public Health Agencies

Academia/Industry

Portable data and visualization





Vision

Capable professionals and institutional systems that support and advance a healthy, enduring global community





Compelling Global Challenge



*Integrated approach to build capacity, to mitigate diseases and associated hazards; influence science & policy at the interface between & among humans, animals, plants and the environment- **Global One Health.***

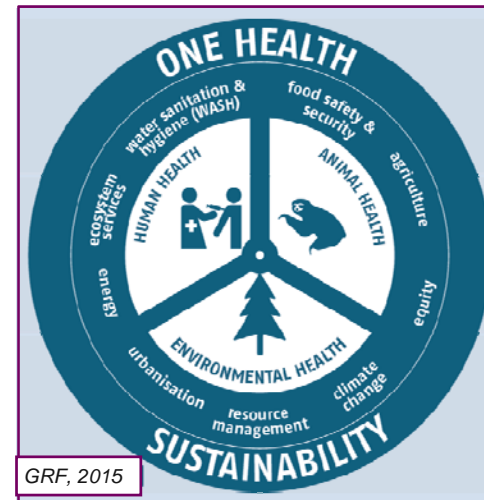
Integrated approach is essential for the sustainability of our planet.

Challenge- How do you build capable professions; influence policy and sustain impact?



Global One Health- Capacity Needs

- **Food and water safety**
- Re-emergence: multi-drug resistance (MDR), hypervirulent,
- Vector-borne Diseases
- Chemical hazards: antibiotics, pesticides...
- Biodefense and biosecurity
- Malnutrition and environmental enteric dysfunction (EED)
- Interaction and chronic outcomes (cancer)
- Lives and livelihood
- International trade
- Sustainability





ICOPHAI GenomeTrakr Partnership

Home | ICOPHAI | Salmonella enterica (ID 27: x)

www.ncbi.nlm.nih.gov/bioproject/275961

NCBI Resources How To Sign in to NCBI

BioProject BioProject Search Advanced Help

Display Settings: Send to:

Salmonella enterica Accession: PRJNA275961 ID: 275961

GenomeTrakr: OSU-ICOPHAI project

Whole genome sequencing of cultured *Salmonella enterica* as part of the US Food and Drug Administration's WGS surveillance effort for the rapid detection of foodborne illness outbreaks.

Related Resources:

- International Congress on Pathogens at the Human-Animal Interface (ICOPHAI)

Project Type: Genome sequencing and assembly

Attributes: Scope: Multisolate; Material: Genome; Capture: Whole; Method type: Sequencing

Relevance: Medical

Project Data:

Resource Name	Number of Links
BioSample	924

SRA Data Details

Parameter	Value
Data volume, Gbases	4
Data volume, Mbytes	2749

Lineage: Bacteria; Proteobacteria; Gammaproteobacteria; Enterobacteriales; Enterobacteriaceae; Salmonella; Salmonella enterica [Taxonomy ID: 28901]

Submission: Registration date: 20-Feb-2015
US Food and Drug Administration

Related information: BioSample, Taxonomy

Related Resources: International Congress on Pathogens at the Human-Animal Interface (ICOPHAI)

Recent activity: Turn Off Clear

- Salmonella enterica BioProject
- icophai (1) BioProject
- horizontal transfer of resistance (1834) PubMed
- Minimization of bovine control costs in US d

See Genome Information for Salmonella enterica

NAVIGATE ACROSS: 2808 additional projects are related by organism.



1000+ MDR *Salmonella* submitted to FDA CFSAN

- Brazil (104)
- Ethiopia (401)
- Kenya (86)
- Mexico (63)
- Tanzania (64)
- Thailand (60)
- U.S. –OSU (247)

E. Coli

- Eastern Africa

Campylobacter

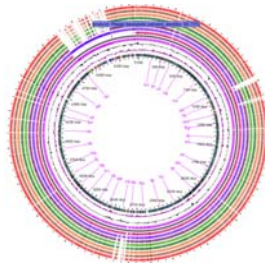
- Ethiopia



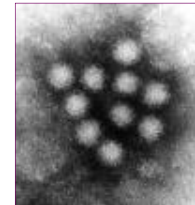


Building Capacity- NIH-GID/ Research Training Program (OHEART)

Molecular Epidemiology of MDR *Salmonella* in Tanzania (Julius Medardus)



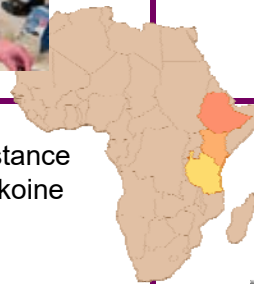
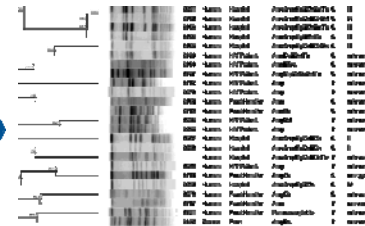
Enteric caliciviruses in pigs and cattle (Zufan Sisay – Addis Ababa University, Ethiopia)



Molecular Typing and Antimicrobial Resistance of *Campylobacter* (Isaac Kashoma – Sokoine University of Agriculture, Tanzania)

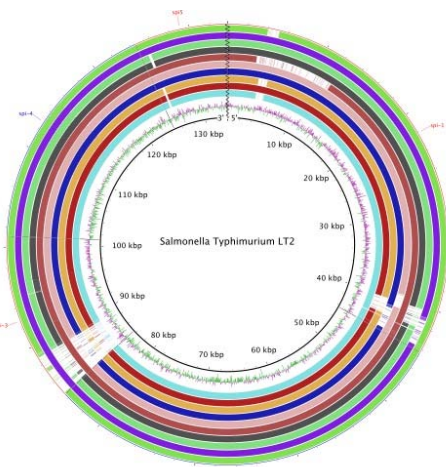


MRSA among HIV, Hospital and Food handler cohorts (Benear Obanda – Kenya Medical Research Institute, KEMRI)

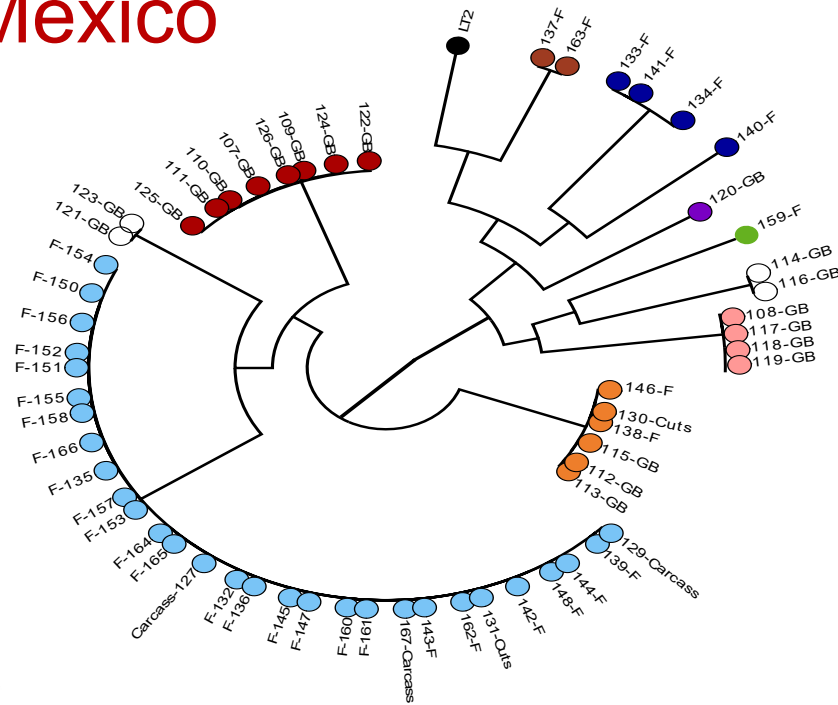


Salmonella WGS- Mexico

- Blast Atlas
- SNP- based phylogeny



CC Skew
 CC (Salmonella)
 Montevideo
 GIVE
 Muenster
 Newport
 Derby
 Reading
 Roodepoort
 Senftenberg
 London
 Bergen



Serovar
 Bergen
 Derby
 Give
 London
 Montevideo
 Muenster
 Newport
 Reading
 Roodepoort
 Senftenberg
 Typhimurium LT2

Distribution of strains per isolation site:
 Mexico City: samples 107-119
 Guadalajara City: samples 120-126
 Mexicali City: samples 127-167

Antimicrobial Resistance and Virulence

- Increasing prevalence of antibiotic resistant strains
- Special case of directed evolution
- Acquire AMR genes or virulence-associated genes from the environment.
- Often independent of serotype or gene markers
- NGS comprehensively and unambiguously track AMR and virulence-associated genes



Culture of a multiresistant bacterium

Identify and correlate- AMR phenotype/ genes

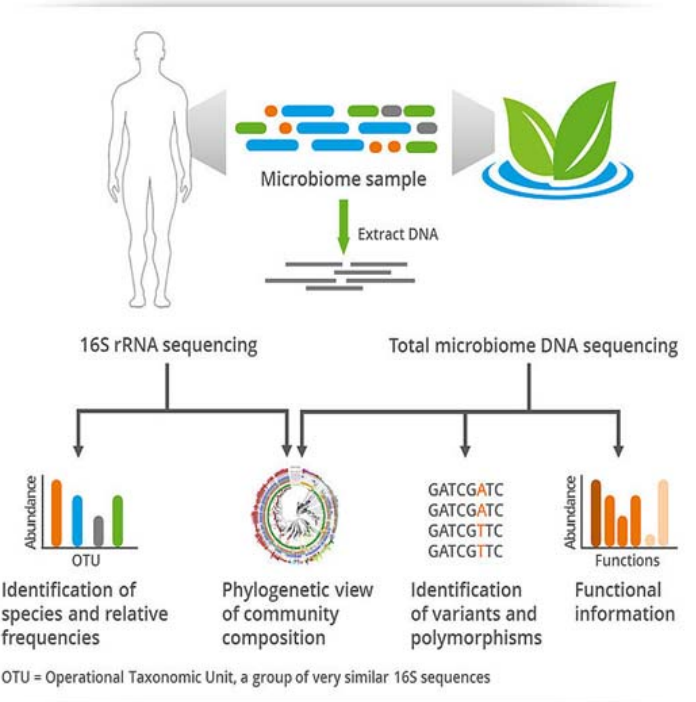
AMR phenotype	AMR genotype																																		
	β-lactams		Quinolones				Aminoglycosides						Sulfonamides			Amphenicols		Tetracyclines			Multidrug efflux pumps families														
	blaTEM-1	blaCARB-2	qnrA1	qnrB	qnrB19	oqxAB	aac(3)-IId	aadA	aadA2	aadA5	aadA22	aph(3'')-Ib	aph(3')-Ia	aph(6)-Id	sul1	sul2	dfrA1	dfrA12	dfrA17	floR	catA1	tetA	tetB	qacEDelta1	mph(A)	bleO	fosA	mdtABCD	MATE	MFS	RND	ATP-depdent ABC			
Amp																																			
Tet																																			
Car																																			
Amp/Cef																																			
Amp/Car																																			
Amp/Tet																																			
Amp/Car/Tet																																			
Amp/Car/Sxt/Tet																																			
Amp/Car/Sxt/Tet																																			
Chl/Str/Sxt/Tet/Kan																																			
Amp/Car/Cef/Chl/Tet																																			
Amp/Car/Sxt/Gen/Tet																																			
Amp/Car/Sxt/Chl/Tet																																			
Amp/Car/Cef/Sxt/Tet																																			
Chl/Str/Sxt/Tet/Kan																																			
Amp/Car/Cef/Tet/Cro/Net																																			
Amp/Car/Cef/Sxt/Chl/Gen/Tet																																			
Amp/Car/Cef/Sxt/Gen/Net/Tet																																			

Amp, ampicillin; **Amx**, amoxicillin; **Amc**, amoxicillin clavulanic acid; **Car**, carbenicillin; **Cro**, ceftriaxone; **Cef**, cephalotin; **Ctx**, cephotaxime; **Cip**, ciprofloxacin; **Pef**, pefloxacin;
Amk, amikacin; **Kan**, kanamycin; **Gen**, gentamicin; **Str**, streptomycin; **Net**, netilmicin; **Sxt**, trimethoprim-sulfamethoxazole; **Chl**, chloramphenicol; **Nit**, nitrofurantoin; **Tet**, tetracycline

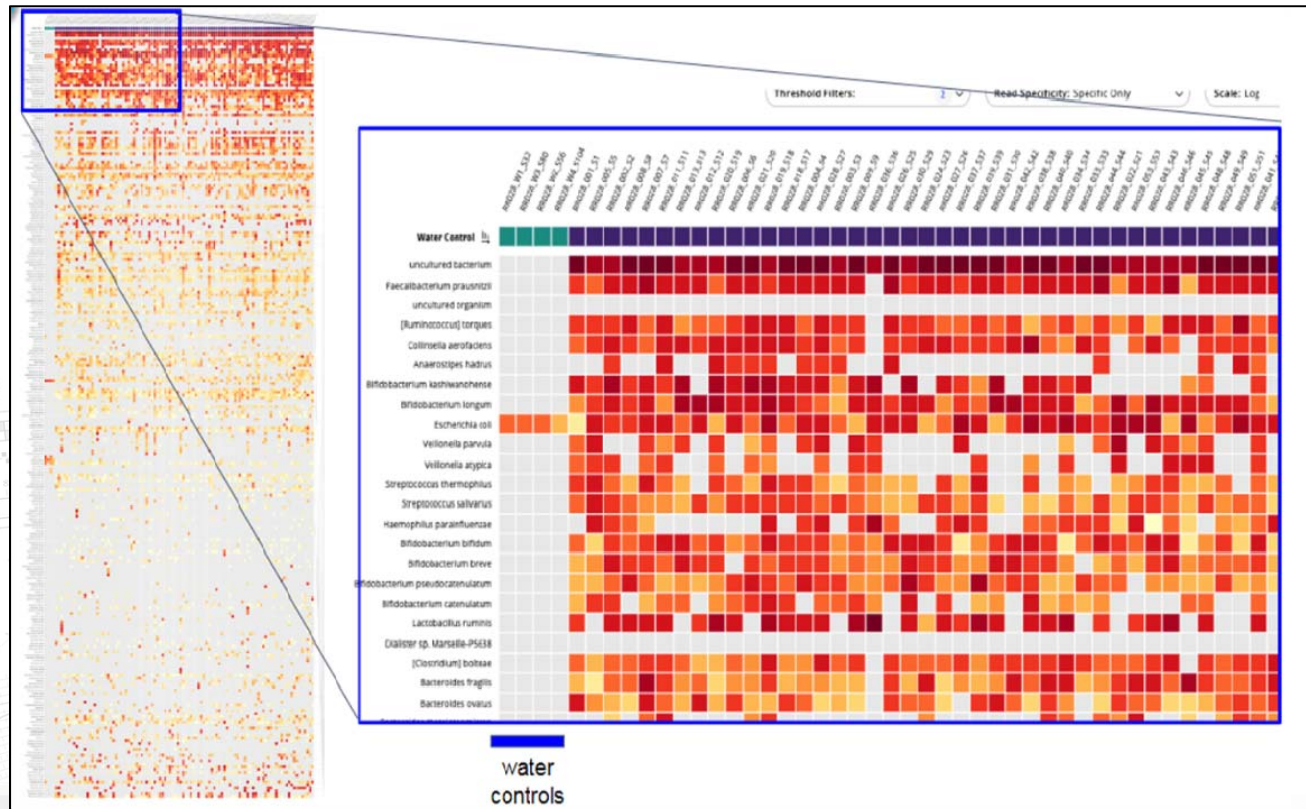
■ Gene present □ Gene absent

Shotgun Metagenomics

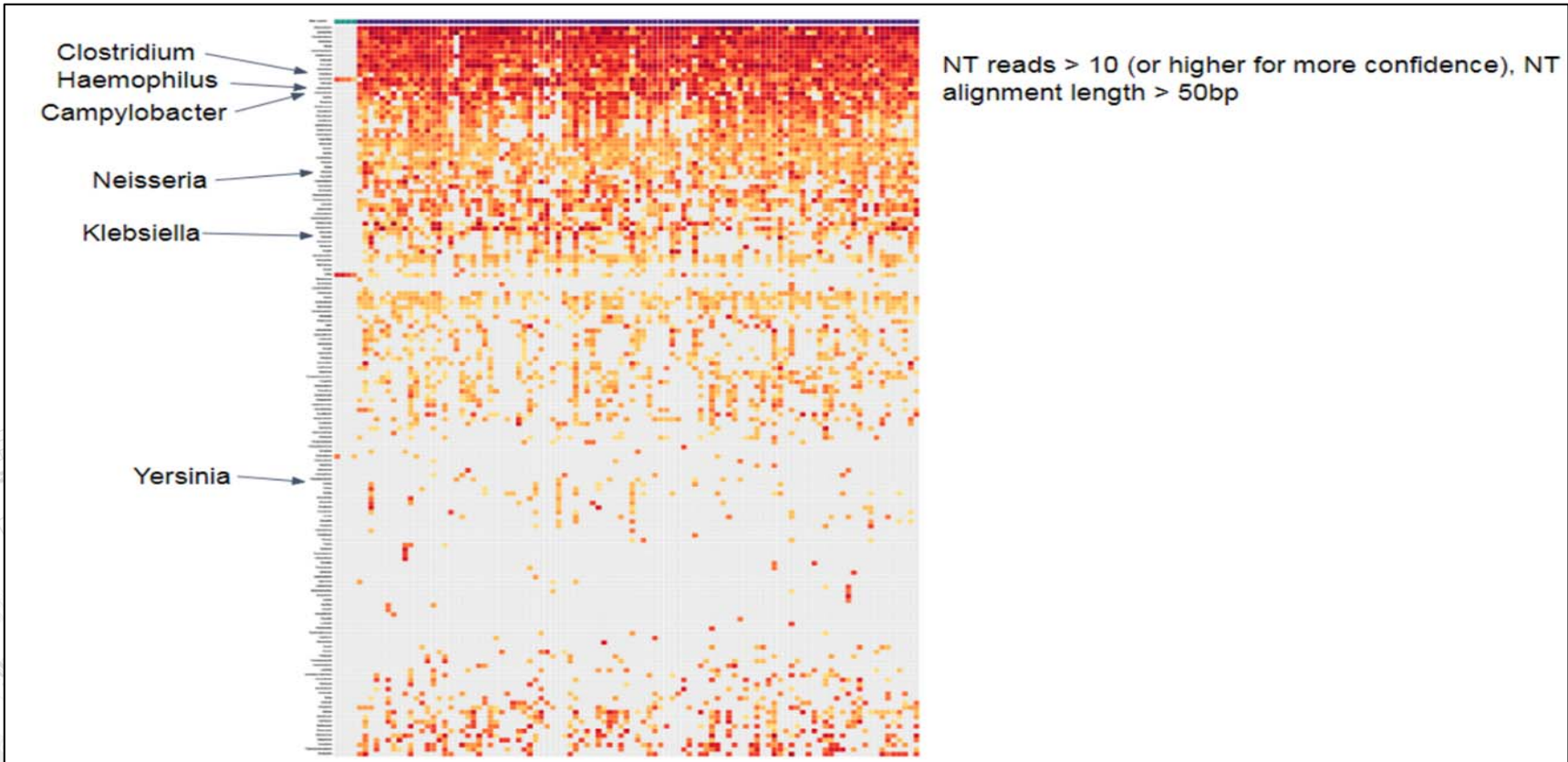
- Comprehensive sampling all genes in all organisms in a complex sample
- Metabolic pathways and gene function
- Evaluation of bacterial diversity
- Detect the abundance of microbes in various environments
- Studying unculturable microorganisms



Campylobacter Genomics and Enteric Dysfunction (CAGED)



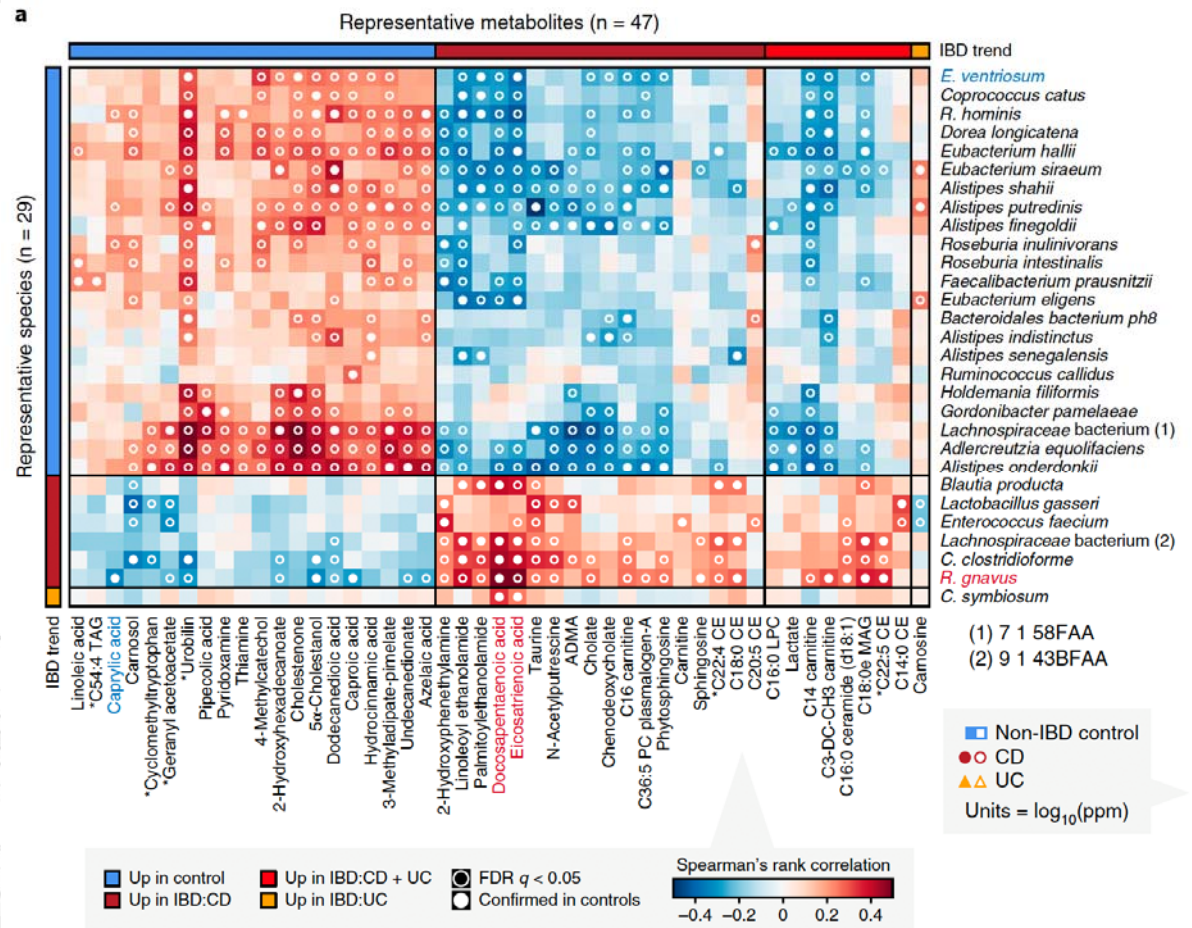
CAGED- Identification of multiple bacteria



Metabolomics

IBD associated Microbes and Metabolic profiles

Fransoza et al., 2019



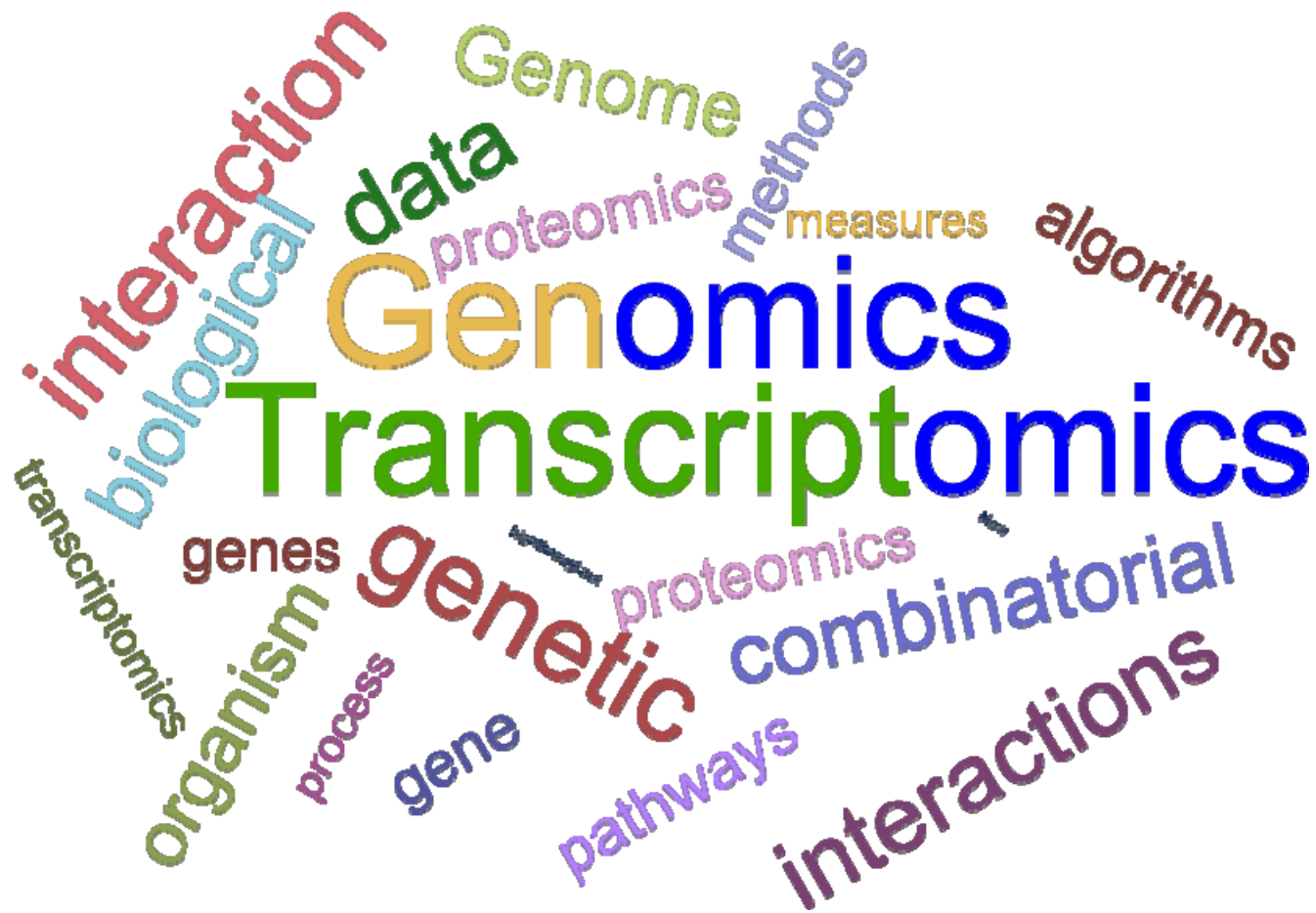
Global context

- Oxford Nanopore (MinION)
- Portable
- Very long read length
 - Easier bioinformatics
 - Quality? V. short reads
- Complement with Illumina
 - No capital cost
 - Catch?



Why MinION?

Long reads	Real time	Easy, rapid prep	On demand	Accessible
 Choose your read length: 5kb? >200kb? Longer?	 Immediate access to data	 1D library prep: <10mins, 2 pots	 Run different experiments in sequence on one flow cell	 No capital cost
 Easier assembly, phasing	 Rapid time to result – move on	 Low cost of materials	 Barcode for even more samples	 Easy install
 Covering repetitive regions	 Rapid insight into whether status of sample	 De-skill prep	 Run many experiments on one device	 No additional lab infrastructure requirements





Training for Transformative Impact

GLOBAL ONE HEALTH

Molecular
Epid.

GLOBAL ONE HEALTH

Environmen
tal Health

GLOBAL ONE HEALTH

Food
Safety

GLOBAL ONE HEALTH

Data
Analytics

GLOBAL ONE HEALTH

Geospat.
Epid.



FOGARTY



THE OHIO STATE UNIVERSITY

Global One Health initiative (GOHi)

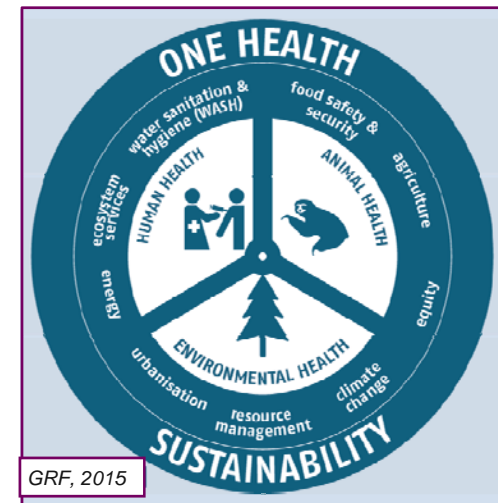
Bringing multi-disciplinary teams/ Global One Health knowledge-sharing





Take Home Message

- Water/ Food- Key conduits for One Health;
- Integrated approaches- addressing biological and chemical hazards
- Lack of capacity and harmonization of research and surveillance is critical;
- Genomics+ play key role;
- Resource mobilization to address Global Water and Food Safety is key for success;
- FDA and global partners leadership is needed.





THE OHIO STATE UNIVERSITY

Global One Health initiative (GOHi)

Thank You!

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Muchas Gracias!

谢谢

Asante Sana!

شكرا جزيلًا

Merci Beaucoup!

Благодаря ти

Terima Kasih

நன்றி

Muito Obrigado!

תודה רבה

Vielen Dank!

Murakoze!

ขอบคุณมาก



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