

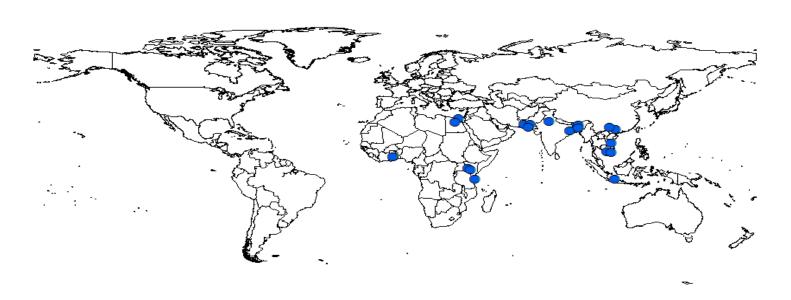
The Typhoid Fever Surveillance in Africa Program (TSAP)



Introduction

Present global disease burden model for typhoid fever (TF)

- Studies conducted in the period of 1990-2011 covered
- Data available from the below sites:



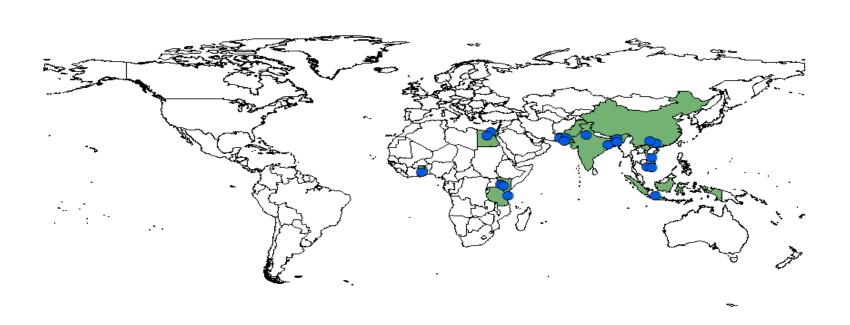
Central America: none South America: none North Africa: 2 West Africa: 1 East Africa: 3

Central Africa: none

East Asia: 2 South Asia: 7 Southeast Asia: 4 Central Asia: none

Introduction

Extrapolation of available data to a wider region





Introduction – Kilifi 2009

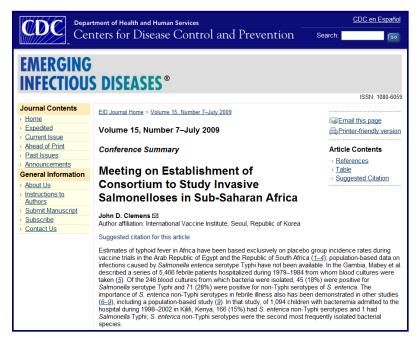
Meeting on invasive Salmonelloses convened by the IVI in Kilifi, Kenya in January 2009

Question:

Is there typhoid fever in Africa? If so, how much?

Outcome:

Recommendation that a sentinel network of field sites be created across Africa that follows similar standards and study procedures.



http://www.cdc.gov/EID/content/15/7/e2.htm



TSAP – Objective, procedures and study sites

Objective:

to generate standardized data on incidence of invasive *Salmonella* bloodstream infections in sub-Saharan Africa

Standardized procedures:

Protocol

- 1) Inclusion criteria
- 2) Clinical procedures
- 3) Laboratory procedures
- Logistics incl. sample shipment
- Health care utilization survey (HCUS)
- Database/Data management

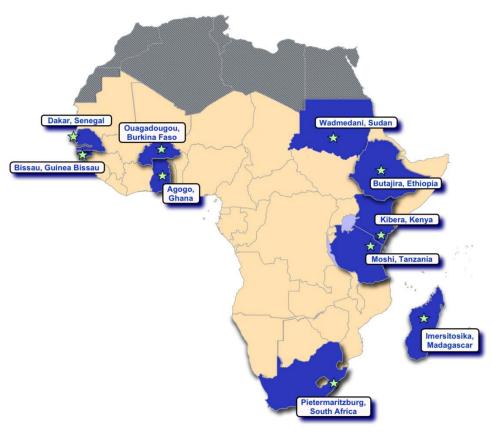


Fig. Location of TSAP study sites

TSAP – Outcomes

Research Topics	Expected outcomes
Incidences for typhoid/paratyphoid fever and NTS infections	Identification of risk groups/target groups for preventive interventions
Antimicrobial susceptibility patterns	Adaption of treatment with adequate antibiotics/Re-evaluation of national guidelines/Awareness raising
Other organisms identified	Information on further causes of febrile diseases (i.e. incidence of viral infections in particular risk groups/per season)
Co-infection with malaria	Occurrence of <i>P. falciparum</i> on invasive bacterial infections
Clinical data/risk factor assessment	Determination of risk factors associated with enteric fever
Information on Salmonella carrier status	Proportion of Salmonella carriers in population



TSAP – Extrapolation model

To develop and validate a disease burden prediction model using standardized data

Improvement of the current disease burden model to

- Predict burden of disease at population level and
- Identify populations at higher risk of Salmonella infections, using standardized data incl. risk factors collected at TSAP sites.

Step 1: Decision on input variables that are transferable and available/or easily determinable

Step 2: model to be calibrated against known output variables such as current disease burden estimates and validated in at least three TSAP sites on its ability to predict what it projected.



This model intends to enable the global community to predict the disease burden based on input variables in differing countries or geographical locations and help inform GAVI/country-level decision-making for vaccine introduction.



Active, household-level surveillance

Active household-level surveillance of fever cases

Rationale:

- Avoid self-treatment of fever with antibiotics (leading to reduced microorganism detection rates in blood culture diagnostics)
- Capture patients and perform diagnostics early in the course of the disease (when bacteremia is high)
- Adjust for the differing health systems/health-seeking behaviors in TSAP countries, which might lead to incomparable incidence calculations



Active household-level surveillance of febrile cases in selected TSAP countries where health-care seeking attendance is hampered due to lack of health-are systems and at sites with considerable antibiotic self-treatment prior to fever diagnostics



Summary

- Standardized surveillance established and running in 10 sites
- Initial data indicate S. Typhi and NTS disease burden in study sites
- One year of surveillance data insufficient
- Design/validation of extrapolation model where data feed into



Acknowledgments

- Peter Aaby
- Mohammad Ali
- Yaw Adu-Sarkodie
- Abraham Aseffa
- Baya Banza
- Holly Biggs
- Morten Bjerregaard-Andersen
- Robert F. Breiman
- John A. Crump
- Ligia Maria Cruz Espinoza
- Jessice Fung Deerin
- Denise Dekker
- Nagla Gasmelseed
- Amy Gassama
- Justin Im
- Hyon-Jin Jeon
- Vera von Kalckreuth
- Karen H. Keddy
- Frank Konings
- Ralf Krumkamp
- Sooyoung Kwon

- Benedikt Ley
- Jürgen May
- Christian G. Meyer
- Eric Mintz
- Vittal Mogasale
- Joel Montgomery
- Pem Namgyal
- Ursula Panzner
- Gi Deok Pak
- Se Eun Park
- Raphael Rakotozaindrindrainy
- Nimako Sarpong
- Heidi Schütt-Gerowitt
- Norbert G. Schwarz
- Arvinda Sooka
- Abdramane Soura
- Adama Tall
- Muna El-Tayeb
- Mekonnen Teferi
- Kamala Thriemer
- Michelle Warren
- Thomas F. Wierzba

THANK YOU



Published *S*. Typhi incidence: Asia*

Site	Hechi, China	Kolkata, India	North Jakarta, Indonesia	Karachi, Pakistan	Hue, Viet Nam	Total	<i>P</i> -value ^a
Site setting	Urban / rural	Urban slums	Urban slums	Urban slums	Urban	-	-
Total population from the study census [year of census]	112 889 [2001]	56 946 [2003]	160 261 [2002]	101 937 [2001, 2003] ^b	281 781 [2002]	713814	-
Surveillance period	1 Aug 2001 to 31 Jul 2002	1 Nov 2003 to 30 Oct 2004	1 Aug 2002 to 31 Jul 2003	1 Aug 2002 to 31 Jul 2004°	1 Jun 2002 to 30 Jun 2003	-	-
Surveillance target population (age in years)	5–60	All ages	All ages	2–15	5–18	-	-
Typhoid Incidence (per 100 000/year) By age	15.3	214.2	81.7	451.7	21.3	107.6	< 0.0001
0-1 years	NA	89.2	0.0	NA	NA	21.6	0.0777
2-4 years	NA	340.1	148.7	573.2	NA	364.8	< 0.0001
5–15 years	29.3	493.5	180.3	412.9	24.2	170.8	< 0.0001
≥ 16 years	12.4	119.7	51.2	NA	10.9	46.8	< 0.0001

• Ochiai et al., Bull WHO 2008