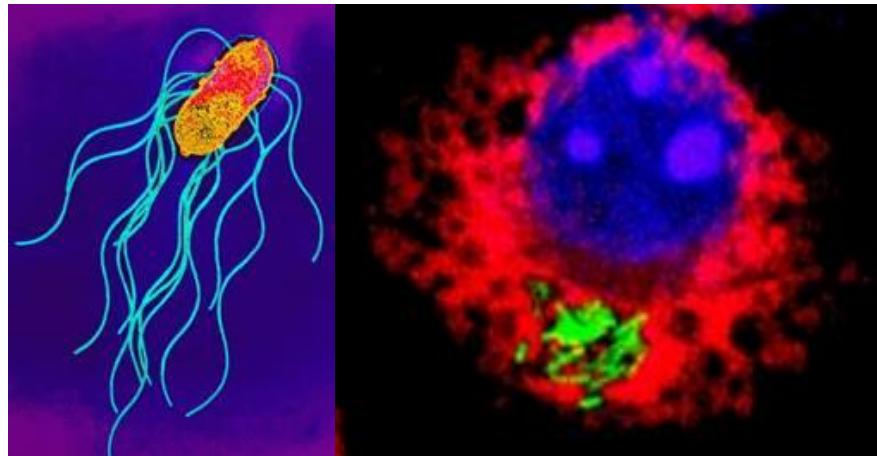




## *Salmonella* – Taxonomie, Vorkommen, Serovare und Pathogenese



Workshop "Salmonellen – ein komplexes Thema für Human- und Veterinärmedizin"  
National Forschungsplattform für Zoonosen

Dr. Karsten Tedin

Institut für Mikrobiologie und Tierseuchen, Freie Universität Berlin



## Introduction

- Salmonellose - Facts and Figures
- *Salmonella* - Taxonomy, Characteristics and Nomenclature
- Recent statistics human cases (RKI), monitoring studies, food screening (BfR)
- *Salmonella* pathogenesis and virulence Mechanisms
- Future concerns - Multi-Drug-Resistant (MDR) *Salmonella* serovars and changes in the pathogenesis of non-Typhoidal *Salmonella* infections



## The Importance of *Salmonella enterica* Serovars in Infections and Disease

- According to the World Health Organization\*, there are an estimated 20 million cases of **Typhoid** each year in the world (*S. Typhi*), resulting in approx. 600 000 deaths
- A further 1,3 billion (Mrd.) **non-Typhoid** human cases of Salmonellosis (*S. Typhimurium*, *S. Enteriditis*, *S. Choleraesuis*, etc.) are estimated to result in an additional 3 million deaths
- *Salmonella* infections account for  $\geq 30\%$  of all food-related deaths
- *Salmonella* infections can range from a self-limiting gastroenteritis to systemic infections
- Alone in the USA, nearly 7 billion (Mrd.) US-Dollar in economic losses yearly are attributed to contaminated food-related infections caused by *Salmonella* spp.
- **Non-symptomatic** infections and intermittent shedding in domestic animals provide a constant reservoir for infection and contamination of food

\* Pang *et al.* (1995) Trends Microbiol. 3:253-255; Pang *et al.* (1998) Trends Microbiol. 6:131-133



## *Salmonella* Taxonomy

- *Salmonella* spp. are members of the *Enterobacteriaceae*, which includes other Gram-negative bacterial genera such as *Enterobacter*, *Escherichia coli*, *Klebsiella*, *Shigella*, *Yersinia*, etc.
- These are all members of the "Proteobacteria", characterized by a high degree of metabolic diversity or "heterotrophy", i.e. aerobic and anaerobic respiration, fermentation, etc.
- High degree of metabolic flexibility
- Principal habitat of the *Salmonellae* is the intestinal tract of humans and animals (reservoir), but disseminate in the environment (water, soil, plants and food) through contamination with human or animal waste (faeces)

## Salmonella Taxonomy

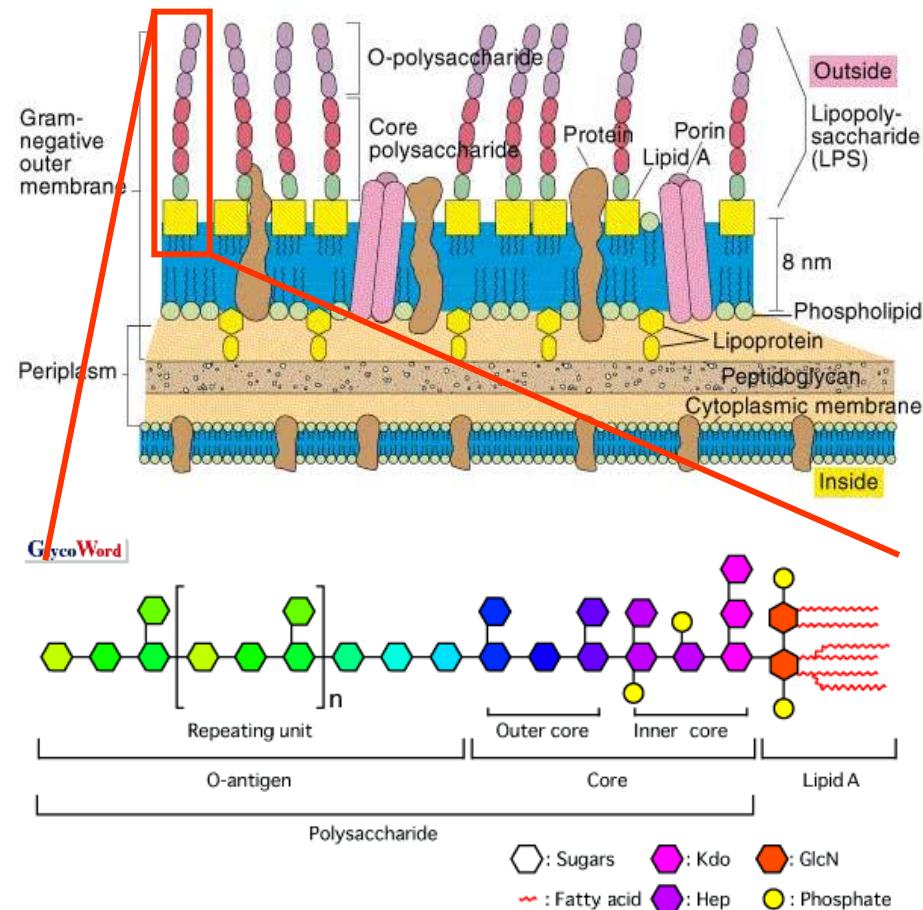
*Salmonella* spp. are identified through the presence of combinations of antigenic determinants, the **Somatic**, **Surface** and **Flagellar** antigens which are used to establish the serovar. Currently, there are > **2500** serovars

### Somatic (O) Cell Wall Antigens

- Heat stable, alcohol-resistant cell wall **lipopolysaccharide (LPS)** antigens
- ca. 70 different antisera used for serology
- the LPS component is the so-called "endotoxin"

### Surface (Capsular) Vi Antigens

- Occurs in only a few *Salmonella* serovars (common in *E. coli* and *Klebsiella* = K antigen)
- May mask O antigens
- *Salmonella Typhi*, *Paratyphi C* and *Dublin* are the only serovars with the Vi antigen





## *Salmonella* Taxonomy

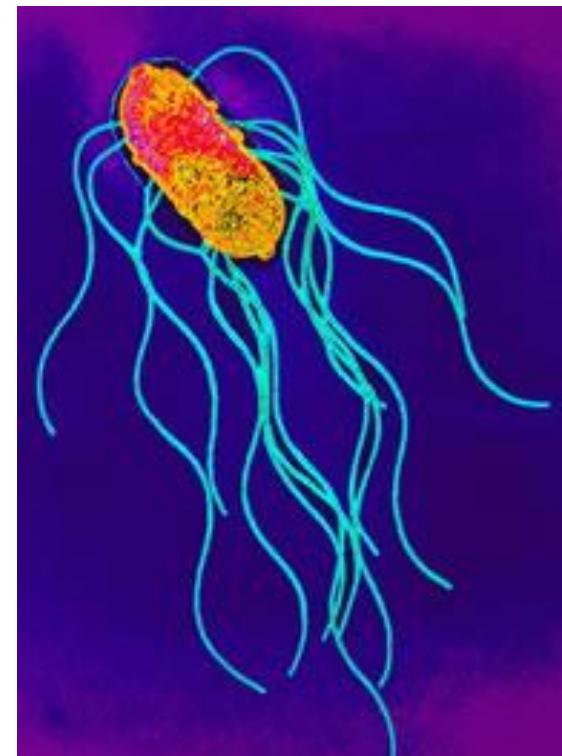
*Salmonella* spp. are identified through the presence of combinations of antigenic determinants, the **Somatic**, **Surface** and **Flagellar** antigens which are used to establish the serovar. Currently, there are > **2500** serovars

### Flagellar (H) Antigens

- Some serovars have a constant (**monophasic**) H antigen (*S. Enteriditis*, *S. Typhi*), however most *Salmonella* serovars have two, different H antigens, 1 or 2 (**diphasic**)
- Serovars with one H antigen may acquire the other at a rate of  $10^{-3}$  to  $10^{-5}$  (**phase variation**)
- Example of a serological formula:

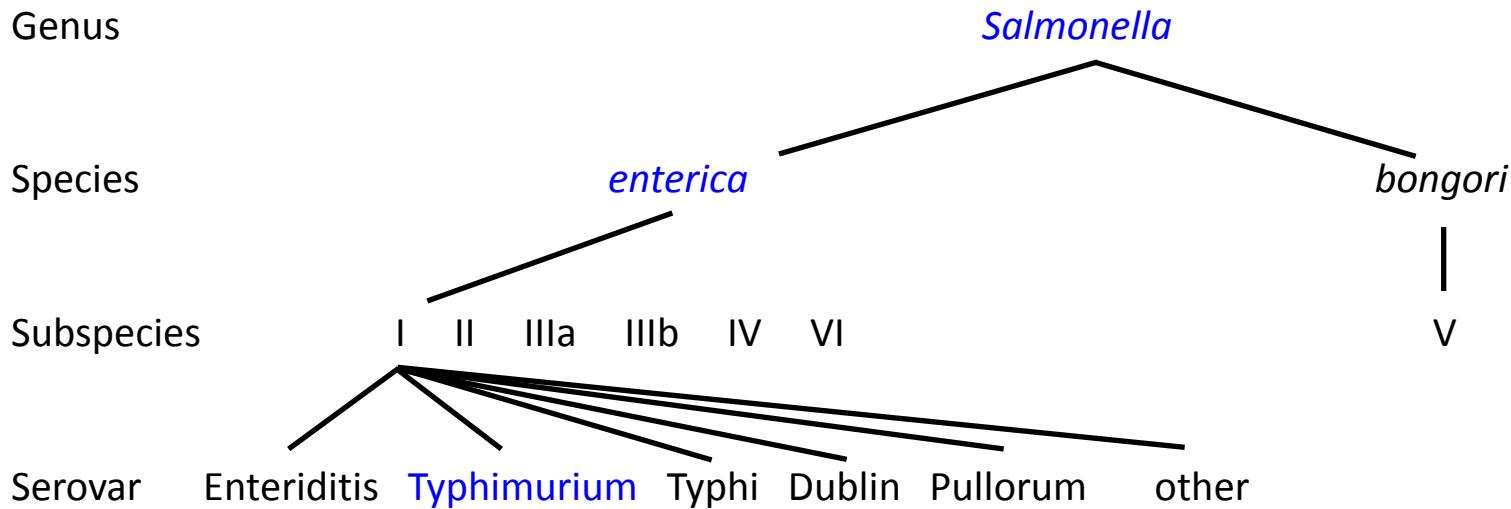
***S. ser. Typhimurium – 1, 4, 5, 12 : i : 1, 2***

O antigens detected: 1, 4, 5 and 12  
H1 antigens detected – phase 1: i  
H2 antigens detected – phase 2: 1 and 2



## *Salmonella* Taxonomy

Due to the close genetic (DNA) relatedness of *Salmonella* strains, a new taxonomic system was proposed by Popoff and Le Minor in 1987 in which only **two species** are recognized:



The full taxonomic name for *Salmonella typhimurium* following this system is:

## *Salmonella enterica* subspecies *enterica* serovar Typhimurium



## *Salmonella* Taxonomy

### *Salmonella enterica* Subspecies Group Designations:

Subspecies*	Group	Mal.	ONPG	Dulcit	Salicin	Galakt.	Gel.
<i>S. enterica</i> ssp. <i>enterica</i>	I	neg.	neg.	pos.	neg.	neg.	neg.
<i>S. enterica</i> ssp. <i>salamae</i>	II	pos.	neg.	pos.	neg.	pos.	pos.
<i>S. enterica</i> ssp. <i>arizonae</i>	IIIa	pos.	pos.	neg.	neg.	(±)	pos.
<i>S. enterica</i> ssp. <i>diarizonae</i>	IIIb	pos.	pos.	neg.	neg.	(±)	pos.
<i>S. enterica</i> ssp. <i>houtenae</i>	IV	neg.	neg.	neg.	pos.	pos.	pos.
<i>S. enterica</i> ssp. <i>indica</i>	VI	neg.	var.	var.	neg.	pos.	pos.
<i>S. bongori</i> *	V	neg.	pos.	pos.	neg.	pos.	neg.

Abbreviations: (±), schwach; var., variabel; Mal., Malonat; ONPG, ; Galakt., Galakturonsäure; Gel., Gelatin.

\* *S. bongori* eigene Spezies, zum Vergleich mit *S. enterica* ssp. aufgeführt.



## *Salmonella* Taxonomy

### *Salmonella* Subspecies Designations:

Group	Species/Subsp.	Comments*
I	<i>enterica</i>	99.5% of all <i>Salmonella</i> isolates in mammals/birds
II	<i>salamae</i>	
IIIa	<i>arizonae</i>	
IIIb	<i>diarizonae</i>	largely cold-blooded animal isolates
IV	<i>houtenae</i>	
V	<i>bongori</i>	<i>S. bongori</i> recognized as separate species
VI	<i>indica</i>	

\* The pathogenesis and virulence of groups II – VI has not been well studied, and human infections are rare



## *Salmonella enterica* subspecies *enterica* Serovars

Serovar	Host specificity / Disease
<i>S. Typhi</i>	Humans / Typhoid, Septicemia
<i>S. Typhimurium</i>	Broad Host Range Humans / Gastroenteritis, Diarrhea, Enteric Fever
<i>S. Abortusovis</i>	Mice / Typhoid, Septicemia
<i>S. Arizonae</i>	Sheep / Goats Diarrhea, Septicemia
<i>S. Choleraesuis</i>	Reptiles Swine / Swine Paratyphoid, Septicemia
<i>S. Dublin</i>	Cattle / Diarrhea, Septicemia
<i>S. Enteritidis</i>	Broad host range / Gastroenteritis, Diarrhea
<i>S. Gallinarum</i>	Poultry / Fowl Typhoid, Diarrhea, Septicemia
<i>S. Paratyphi A,B,C</i>	Humans / Paratyphoid, Septicemia
<i>S. Pullorum</i>	Poultry / Pullorum Disease (young birds < 3 weeks old)
<i>S. Typhisuis</i>	Swine / Chronic Paratyphoid, Diarrhea

# *Salmonella* Nachweis

(Anlehnung an DIN EN ISO 6579:2003; 6579-1:2014)

## Lebensmittel / Futtermittel

■ Enrichment (Voranreicherung)	Buffered Peptone Water Nährbouillon	16 - 20 h, 37°C
■ Selective Enrichment (selektive Anreicherung)	Rappaport-Vassiliadis-Bouillon bzw. halbfesten RV Agars	0,1 ml + 10 ml "Rappaport" ≥24 h, 42°C (2. enrichment ≥24 h, 42°C)
■ Subculture (Subkultivierung)	Xylose-Lysin-Desoxycholat-Agar (XLD) Gassner Chromagar Nähragar	≥24 h, 37°C
■ Serology/Biochemistry/PCR (serologische u. biochemische Bestätigung)	meist Gruppen-spezifisch Antisera* Weiterleitung NRZ- <i>Salmonella</i> (RKI)	

\*Serogroup

- A: Paratyphi A, u.a.
- B: Typhimurium, Agona, Heidelberg, u.a
- C1: Choleraesuis, Infantis, u.a.
- C2: Newport, Hadar, Bovismorbificans, u.a.
- D: Enteriditis, Dublin, Typhi
- E: Anatum, u.a.
- G: Poona, Havana, u.a.





## *Salmonella* Serovars in Selected Animal Species/Food Products

Tierart	<i>Salmonella</i> -Nachweisrate (%)		
	2012	2013	Lebensmittel-2013
Hühner	2,6	2,1	4 - 12%
Rinder	2,8	3,0	0,09%
Schweine	13,5	17,4	2,7%
Schafherden	5,2	7,3	
Ziegenherden	1,3	1,3	
Pferde	3,3	1,1	
Hunden	2,3	2,4	
Katzen	1,2	1,7	

Bundesinstitut für Risikobewertung - Erreger von Zoonosen in Deutschland im Jahr 2013



## *Salmonella* Serovars in Selected Animal Species

*Salmonella*-Nachweisrate in Tierkörpern (%)

Tierart	2009	2010	2011
Kälber	0/80 (0)	0/55 (0)	3/101 (2,9)
Kühe	1/36 (2,8)	2/96 (2,1)	0/26 (0)
Mastschweine	13/504 (2,6)	4/209 (1,9)	15/166 (9,0)
Schaf	1/52 (1,9)	0/43 (0)	0/38 (0)
Ziegen	0/21 (0)	0/12 (0)	0/9 (0)
Pferde	0/39 (0)	0/22 (0)	0/20 (0)
Hunden	0/149 (0)	0/66 (0)	6/58 ( <b>10,3</b> )
Katzen	1/155 (0,6)	0/60 (0)	3/88 (3,4)
Reptilien	5/21 ( <b>23,8</b> )	6/23 ( <b>26,1</b> )	1/9 ( <b>11,1</b> )
Tauben	10/48 ( <b>20,8</b> )	6/53 ( <b>11,3</b> )	12/56 ( <b>21,4</b> )

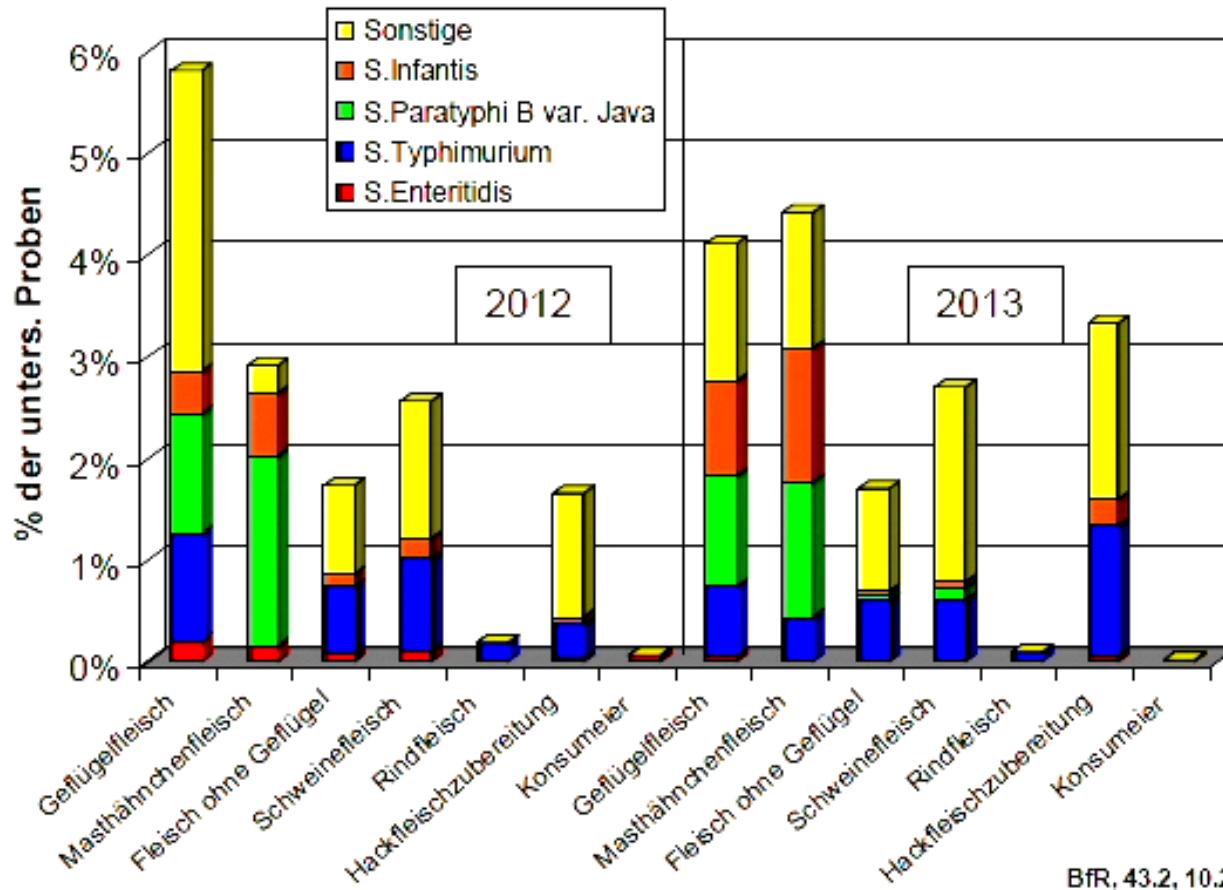


## *Salmonella* Serovars in Selected Animal Species

*Salmonella*-Nachweisrate in Kotproben (%)

Tierart	2009	2010	2011
Kälber	9/267 (3,4)	10/283 (3,5)	19/302 (6,3)
Kühe	1/882 (0,1)	3/416 (0,7)	554/5266 ( <b>10,5</b> )
Mastschweine	3/135 (2,2)	15/220 (6,8)	12/76 ( <b>18,8</b> )
Schaf	0/18 (0)	0/5 (0)	0/4 (0)
Ziegen	0/17 (0)	0/4 (0)	0/6 (0)
Pferde	0/14 (0)	0/21 (0)	1/22 (4,5)
Hunden	4/187 (2,1)	3/188 (1,6)	13/113 ( <b>11,5</b> )
Katzen	0/97 (0)	1/63 (1,6)	0/45 (0)
Reptilien	10/43 ( <b>23,3</b> )	13/51 ( <b>25,5</b> )	5/13 ( <b>38,5</b> )
Tauben	17/267 (6,4)	10/229 (4,4)	5/330 (1,5)

## Salmonella Serovars in Selected Animal Species/Food Products



BfR, 43.2, 10.2014



## *Salmonella* Serovars in Food Samples of Animal and Plant Origins

Anzahl der zehn häufigsten Serovare aus Einsendungen von  
Lebensmitteln in den Jahren 2004 – 2008  
(NRL BfR, 2010)

<i>S. enterica</i> Serovar	Anzahl der Isolate aus Lebensmitteln insgesamt	Anzahl der Isolate aus Lebensmitteln tierischer Herkunft	Aus pflanzlichen Lebensmitt.
<i>Salmonella</i> gesamt	6241 (100%)	5780	265
Typhimurium	1939 (31,1%)	1908 (33,0%)	10
Enteritidis	950 (15,2%)	805 (13,9%)	12
1,4,[5],12:l:-	373 (6,0%)	373 (6,5%)	0
Derby	323 (5,2%)	308 (5,3%)	12
Infantis	244 (3,9%)	239 (4,1%)	4
Paratyphi B (dT+)	229 (3,7%)	227 (3,9%)	1
S. Subspezies I, rau	177 (2,8%)	170 (2,9%)	3
Saintpaul	155 (2,5%)	150 (2,6%)	5
Bovismorbificans	135 (2,2%)	133 (2,3%)	2
Hadar	135 (2,2%)	133 (2,3%)	0

## Reported Cases of *Salmonella* Infections of Humans 2001 - 2013

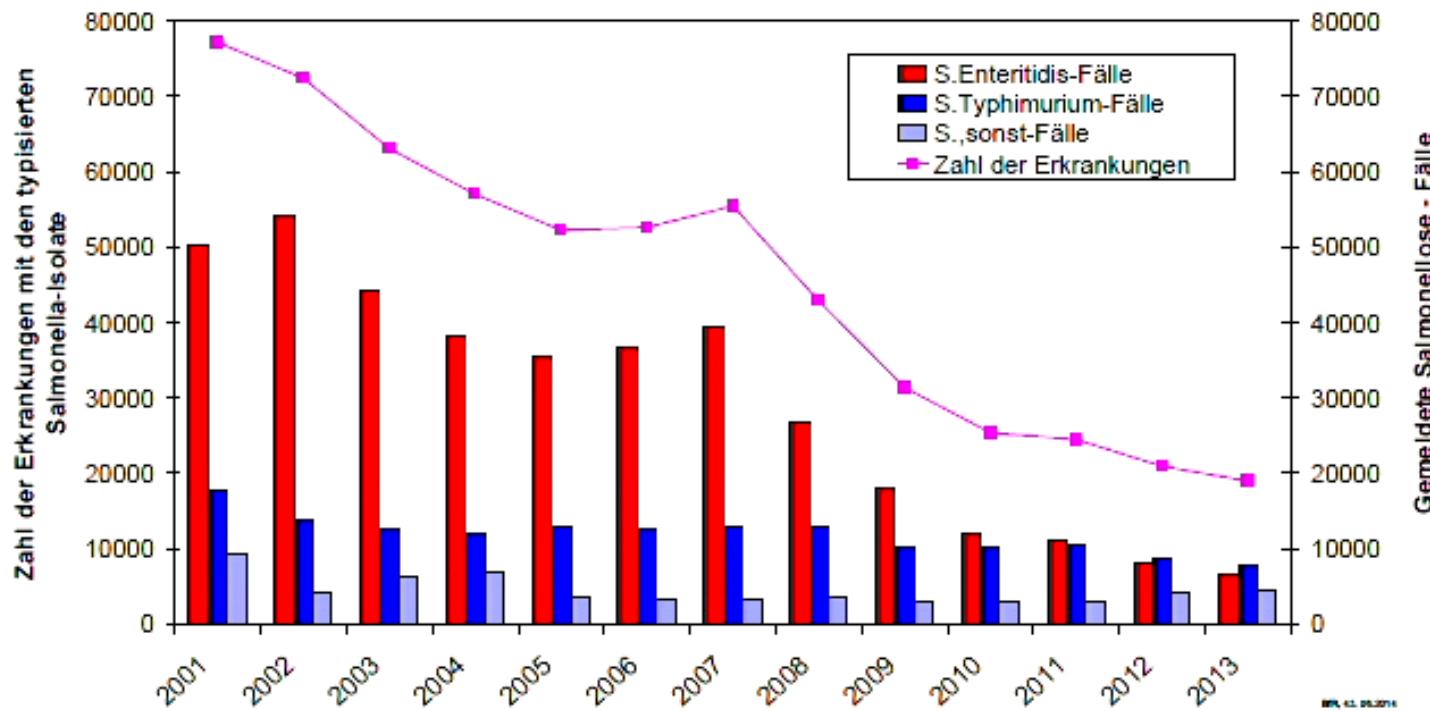


Abb. 4.2.1: Dem RKI gemeldete Fälle von Salmonellose beim Menschen 2001–2013 (n. RKI, 2013; nach IfSG, 2014)

## Total Reported Cases of Human Infections 2014

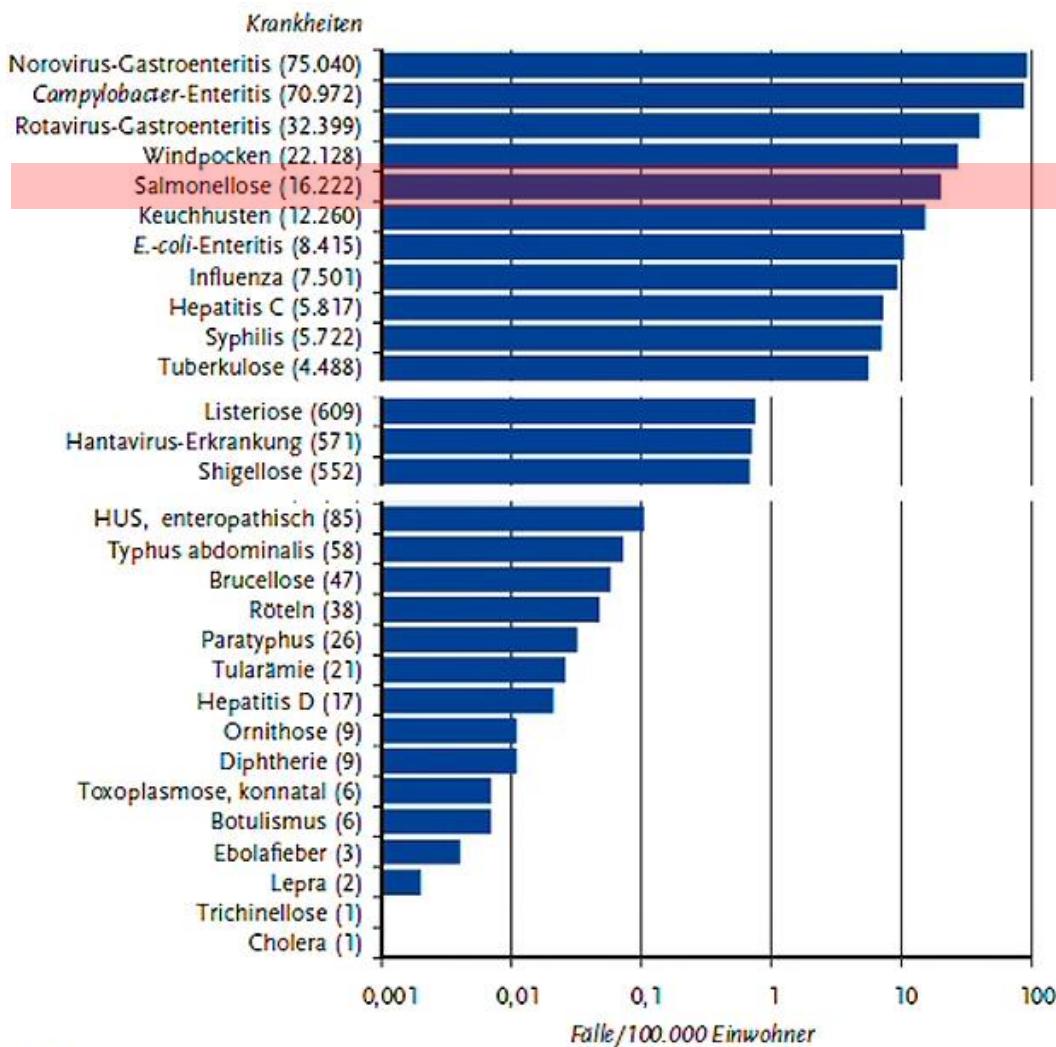
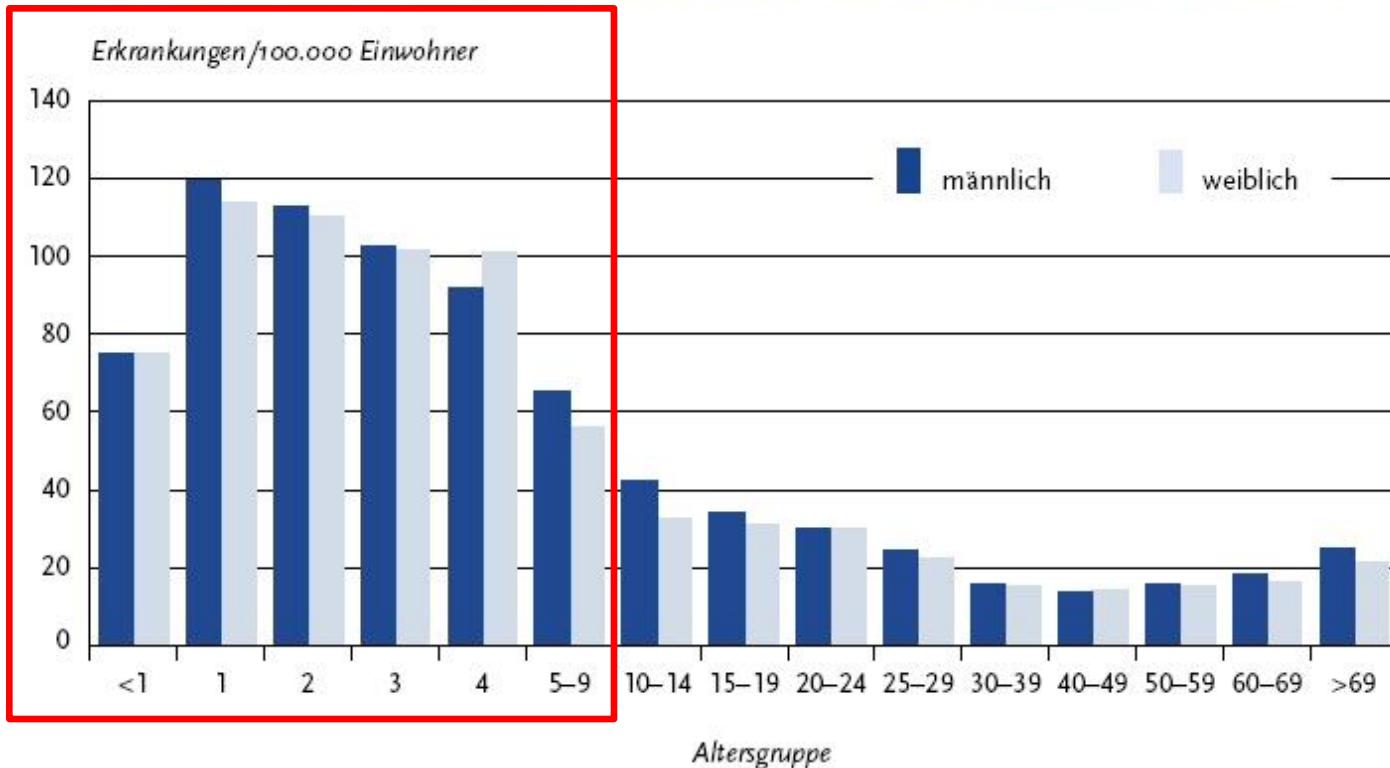


Abb. 4.1.1:  
Inzidenz (logarithmisch) und Anzahl der Fälle aller meldepflichtigen Krankheiten mit mindestens einem Fall,  
Deutschland, 2014



## Total Reported Cases of *Salmonella* Infections 2011 by Age of Patients

Übermittelte Salmonellosen pro 100.000 Einwohner nach Alter und Geschlecht, Deutschland, 2011 (n=20.809)



- Majority of reported cases among the young



## Four Main Forms of Salmonellose

*Salmonella* infections can have a number of differing consequences, often serovar-dependent:

- **Acute Enterocolitis:** short-term, intestinal inflammation, diarrhea
- **Chronic Enterocolitis:** long-term intestinal inflammation, often recurring diarrhea
- **Acute Septicemia:** involvement of the liver, spreading to the circulatory (blood) system  
may involve abortion in animals, including recurring abortion
- **Clinically Asymptomatic Shedding:**

**Active carriage** - Constant or intermittent shedding of *Salmonella* in the faeces

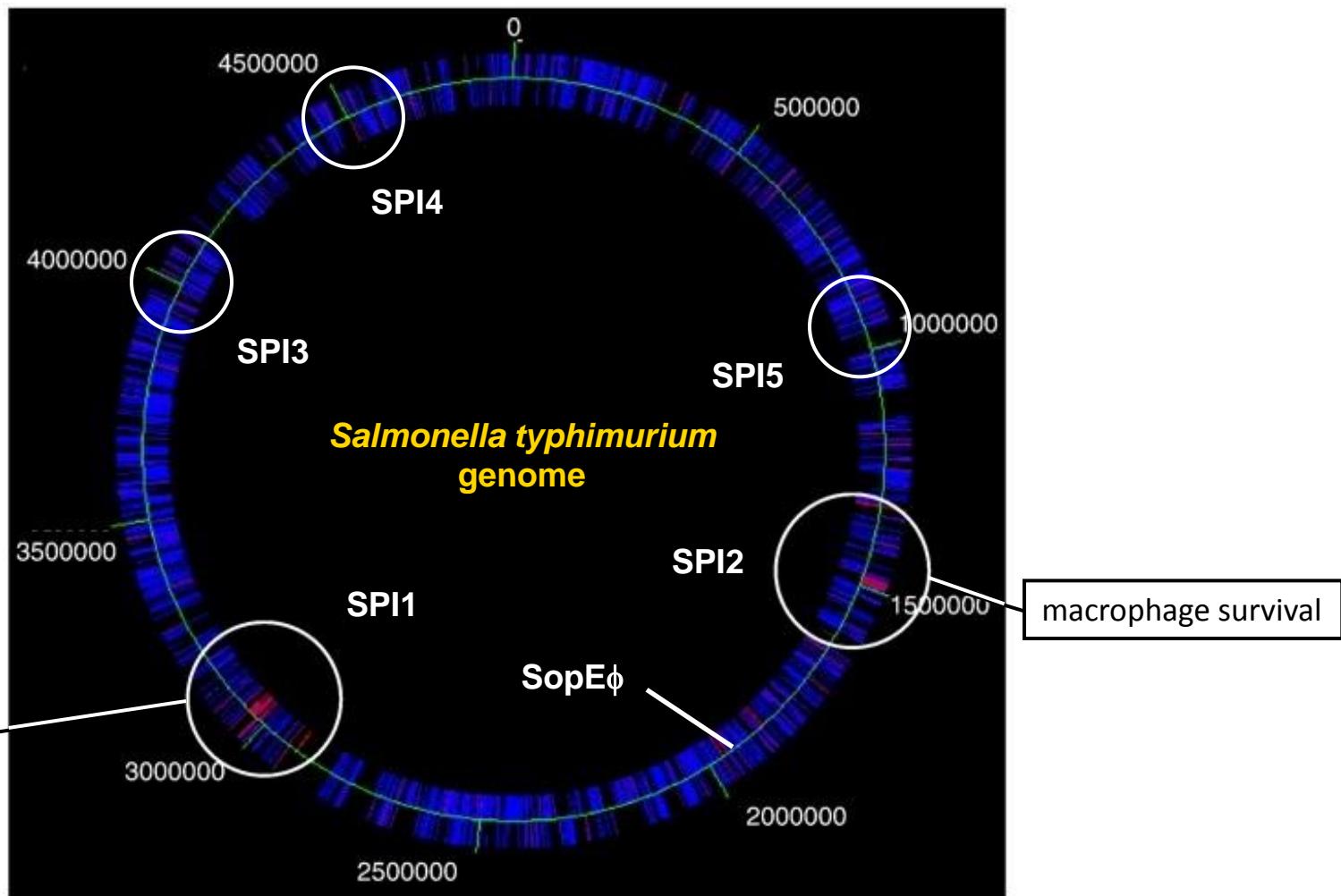
**Latent carriage** - Persistent infection of the Lymph nodes or Tonsils without shedding  
(stress can result in clinical symptoms and "active" infection)

**Passive carriage** - Continuous uptake and shedding of *Salmonella* from/into the environment  
without invasion/infection of the host. Removal of contaminated materials  
results in reduction or loss of further infections



*Salmonella* is a pathogen of the intestinal tract, but gains access to lymphatic / immune tissues for proliferation and persistence within the host...

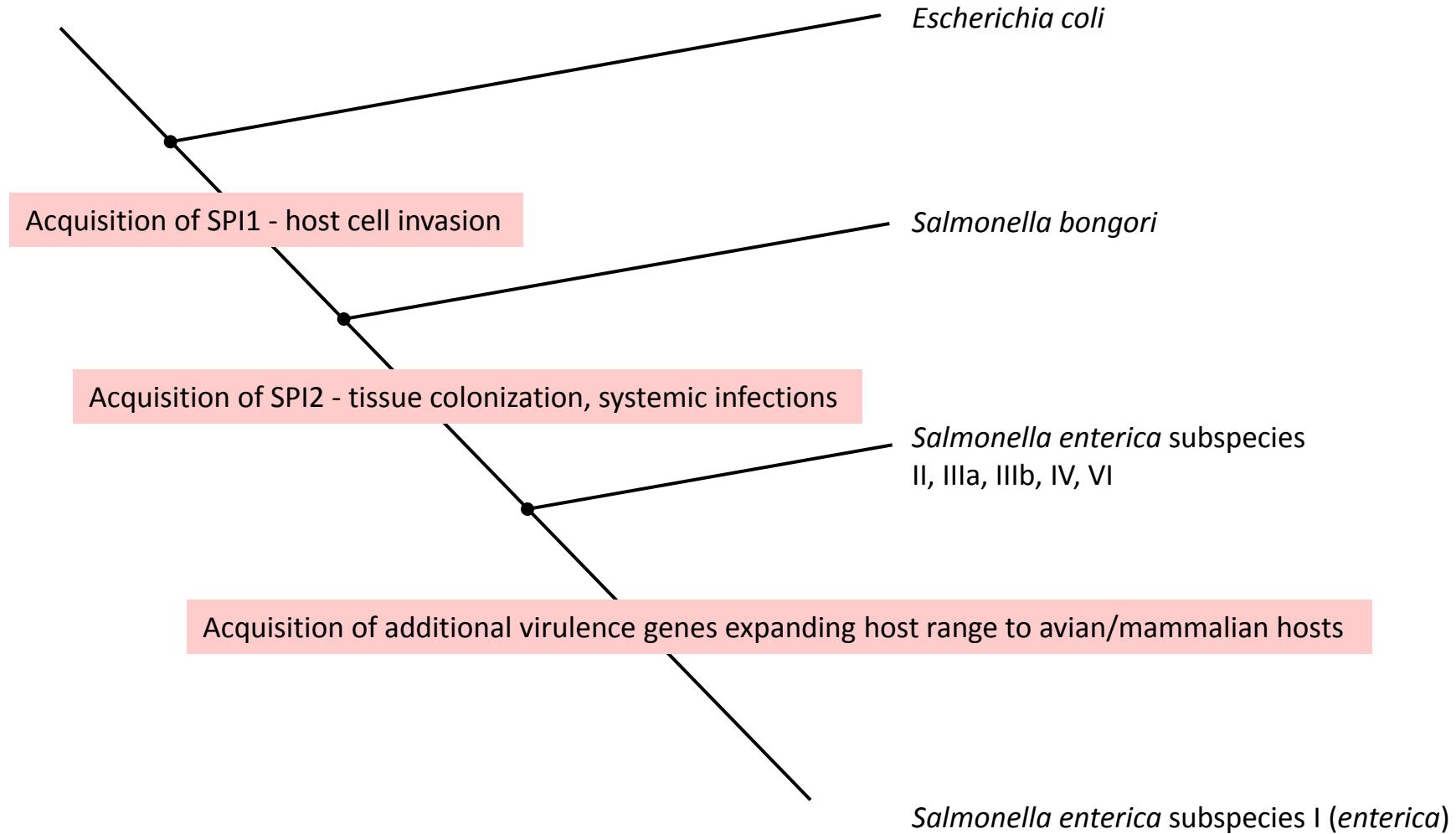
## *Salmonella* Pathogenicity Islands (SPI)\*



\*SPI1 present in all serovars, SPI2 not in *S. bongori*. Other SPI serovar/isolate-dependent

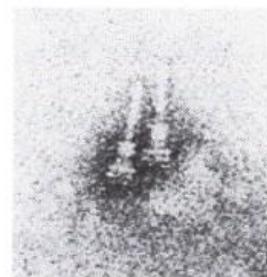
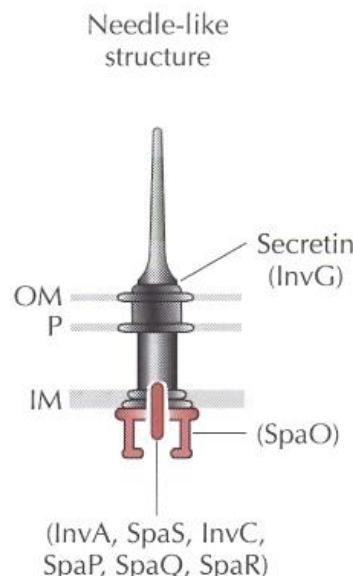
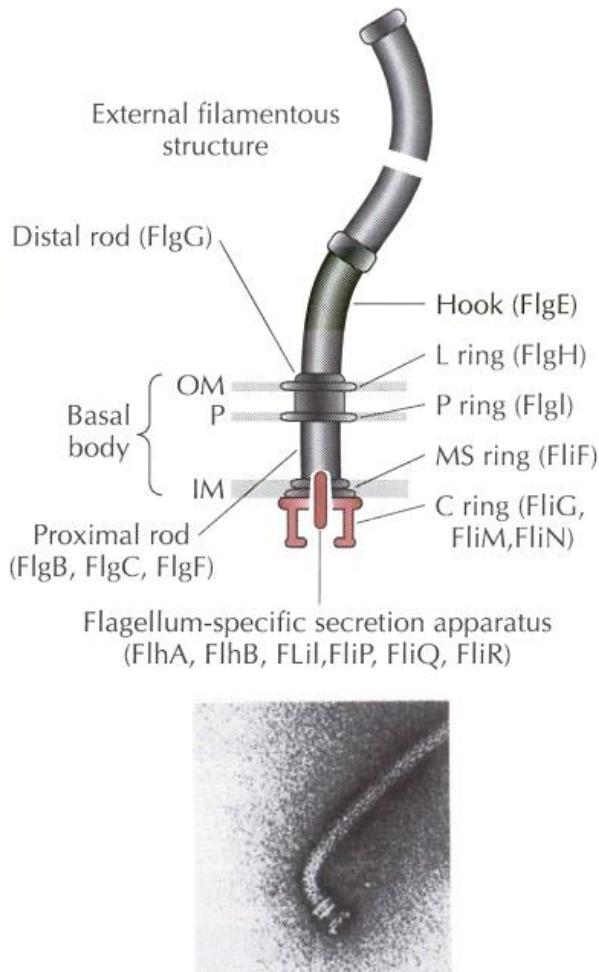


## Model for the Evolution of Virulence in the Genus *Salmonella*





## *Salmonella* Secreted Virulence/Effector Proteins

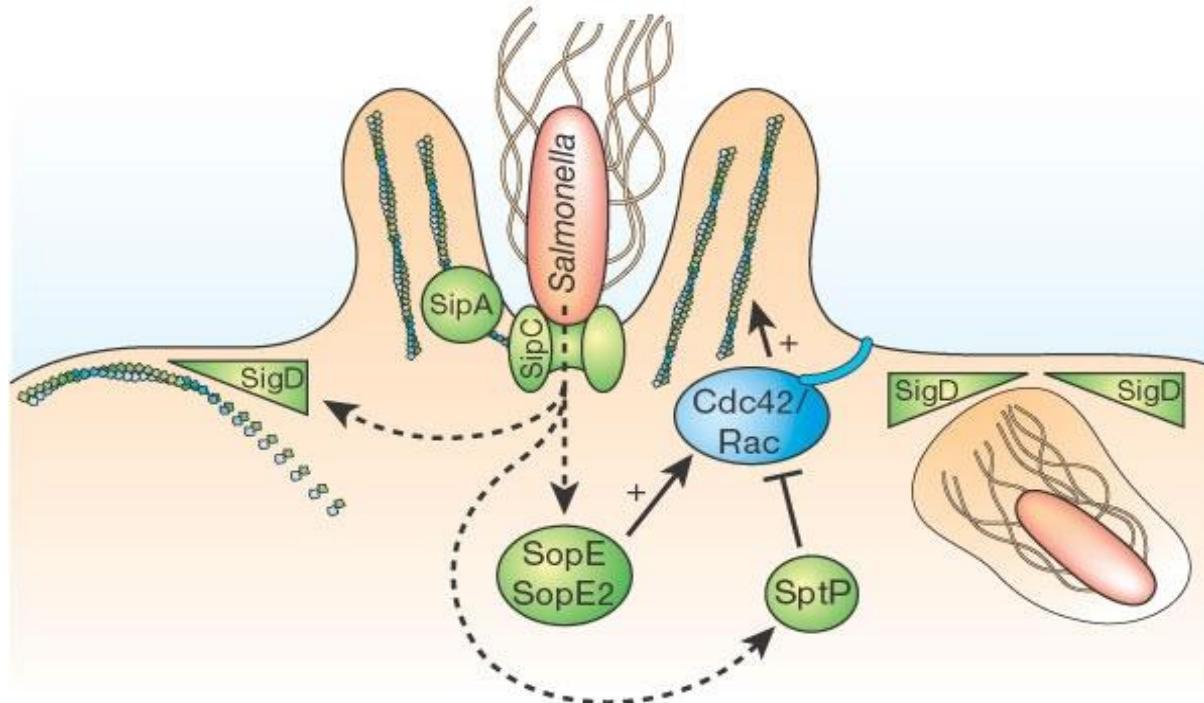


The SPI1- and SPI2-encoded **Type III Secretion Apparatus** show homology with the bacterial flagellar apparatus

Like the flagellar apparatus, the type III secretion apparatus is essentially a **transport system** for proteins

The *Salmonella* type III secretion system is designed for the **delivery (secretion)** of virulence proteins into eukaryotic host cells

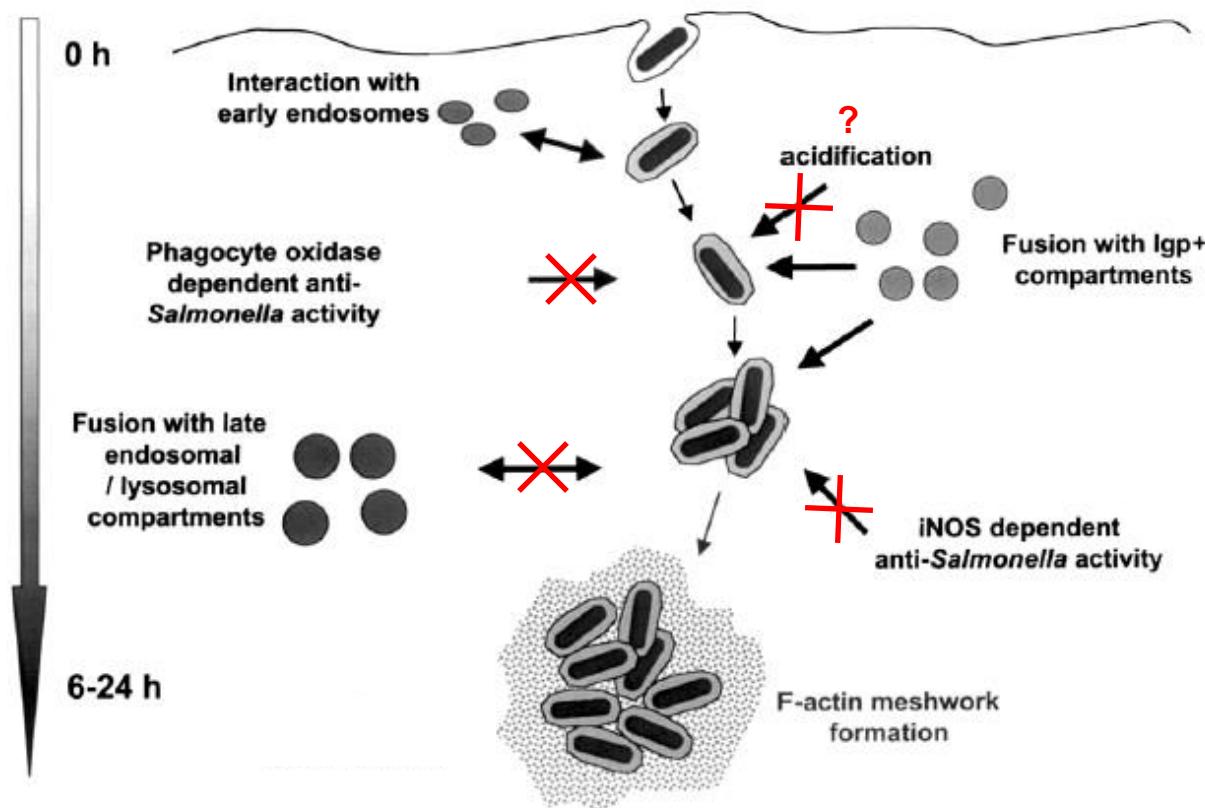
## *Salmonella* Invasion of Host (Intestinal Epithelial) Cells



- **SigD** weakens the actin cytoskeleton attachment to the membrane. **SipA** and **SipC** act as actin nucleating proteins - the stage is set for re-structuring the **actin cytoskeleton**
- **SopE** and **SopE2** activate the host Rho-family GTPases **CDC42** and **Rac1** leading to activation of Arp2/3 Complex and actin polymerization - membrane movement - **uptake**



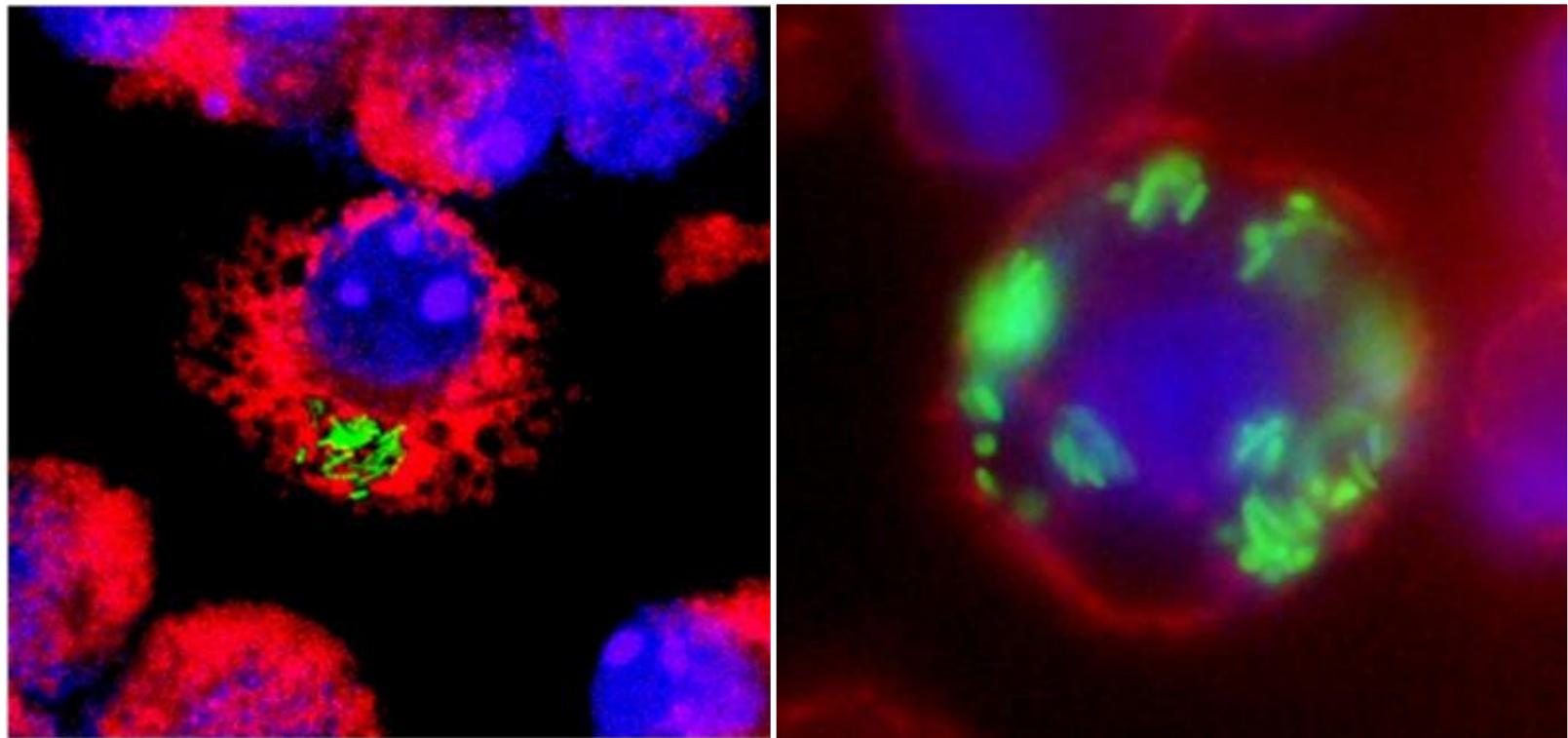
## *Salmonella* Infection of Host Cells and Establishment of the Intracellular Vacuole



✗ Processes inhibited or reduced by *Salmonella*



## *Salmonella* Survives and Proliferates within Macrophage



Murine (left) and human (right) macrophage harboring GFP-expressing *S. Typhimurium* (**green**) and counter-stained with Lysotracker (**red**) and nuclear stain DAPI (**blue**)

Photos: Dr. M. Nordhoff, Institut für Mikrobiologie und Tierseuchen, Freie Universität Berlin



## How *Salmonella* Copes With Macrophage / Phagosome Defenses

- Acquisition of the [vATPase](#) by the *Salmonella*-containing phagosome results in a reduction of the vacuolar/phagosomal pH to approx. pH 5 - 6
- *Salmonella* has a number of genes involved in an [Acid Tolerance Response](#), and is capable of surviving acidic conditions for long periods, particularly if allowed to acclimate
- The host [NADPH Oxidase](#) produces reactive oxygen species ([superoxide](#)), which is antibacterial due to strong oxidizing effects
- However, *Salmonella* encodes at least [4 Superoxide Dismutases](#), two cytoplasmic ([SodA/MnSod](#) and [SodB/FeSod](#)) and two periplasmic ([SodCI](#) and [SodCII](#)), which detoxify superoxide
- In addition, *Salmonella* also encodes a number of [Catalases](#), which can detoxify [hydrogen peroxide](#), the intermediate reaction species from superoxide detoxification



## Pathogenesis of Different *Salmonella* Serovars in Different Species

- **Broad Host-Range** Serovars - *S. Typhimurium*, *S. Enteritidis*
- **Host-Restricted** Serovars - *S. Typhi*, *S. Gallinarum*, etc.
- **Host-Adapted** Serovars - *S. Dublin*, *S. Choleraesuis*, etc.

Host \ Serovar	<i>S. Typhimurium</i> / <i>S. Enteritidis</i>	<i>S. Typhi</i>	<i>S. Dublin</i>	<i>S. Choleraesuis</i>	<i>S. Gallinarum</i>
Humans	Enterocolitis/ Diarrhea	Typhus/ Septicemia	Septicemia	Septicemia	-
Poultry	Diarrhea/ Septicemia*	-	-	-	Fowl Typhoid/ Septicemia
Cattle	Diarrhea/ Septicemia*	-	Diarrhea/ Septicemia	-	-
Swine	Diarrhea/ Asymptomatic*	-	-	Septicemia	-
Mice	Septicemia	-	Septicemia	Septicemia	-

\* age-dependent pathogenesis; - non-pathogenic. Here, "septicemia" includes bacteremia and/or sepsis



## Uneven Distribution of Systemic Forms of Salmonellosis in Humans

Serovar	Total Isolates	Blood Samples	Ratio (%)
Enteriditis	563.221	6.775	1.2
Typhimurium	183.542	2.692	1.5
Typhi	4.636	2.825	60.9
Paratyphi A	3.125	1.727	55.3
Paratyphi B*	2.463	308	12.5
Paratyphi C	37	20	54.1
Choleraesuis	147	48	32.7
Dublin	1.283	530	41.3

\* *S. Paratyphi A* and *S. Paratyphi C* are host-restricted (humans), *S. Paratyphi B* isolates are also infectious for animals

Certain serovars show a much higher rate of systemic infections in humans, including host-adapted serovars such as *S. Choleraesuis* (swine) and *S. Dublin* (cattle)

Questions:

- What bacterial or host factor(s) are responsible for these differences?
- If bacterial, can the traits be acquired by the widespread, broad host-range *Salmonella* serovars?



## Infection and Persistence of *Salmonella* in Host Cells

*Microbial Pathogenesis* 2000; **29**: 121–126  
doi:10.1006/mpat.2000.0367

# Experimental *Salmonella typhi* infection in the domestic pig, *Sus scrofa domestica*

Eleanor S. Metcalf<sup>a</sup>, Glen W. Almond<sup>b</sup>, Patricia A. Routh<sup>b</sup>,  
John R. Horton<sup>c</sup>, Richard C. Dillman<sup>c</sup> & Paul E. Orndorff<sup>c\*</sup>

- Studies with the host-restricted serovar *S. Typhi* showed carriage in porcine tonsils for up to 3 weeks with no clinical signs of infection.
- Other studies show that transport stress results in shedding of *S. Typhimurium* by market weight pigs despite no signs of infection nor faecal shedding prior to transport (e.g. Isaacson *et al.*, 1999. *Am. J. Vet. Res.* 60:1155-1158).
- Finishing pigs can show high levels of shedding of *S. Typhimurium* without developing signs of infection. *Salmonella* persist in the mesenteric lymph nodes. (Rostagno *et al.*, 2011. *Foodborne Pathog. Disease* 8:623).



# Increase in Multi-Drug Resistant *Salmonella* Serovars

JOURNAL OF CLINICAL MICROBIOLOGY, Aug. 2002, p. 2813–2822  
0095-1137/02/\$04.00+0 DOI: 10.1128/JCM.40.8.2813–2822.2002  
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Vol. 40, No. 8

## Molecular Characterization of Multidrug-Resistant *Salmonella enterica* subsp. *enterica* Serovar Typhimurium Isolates from Swine

Wondwossen Abebe Gebreyes<sup>1\*</sup> and Craig Altier<sup>2</sup>

Chiu et al. BMC Microbiology 2010, **10**:86  
<http://www.biomedcentral.com/1471-2180/10/86>



RESEARCH ARTICLE

Open Access

Characterization of 13 multi-drug resistant *Salmonella* serovars from different broiler chickens associated with those of human isolates



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f. 607 253-3943  
w. ahdc.vet.cornell.edu

Animal Health Advisory  
Multi-drug Resistant *Salmonella* Dublin in Cattle



## Systemic Forms of non-Typhoidal Salmonellosis in Humans

### Intracontinental spread of human invasive *Salmonella* Typhimurium pathovariants in sub-Saharan Africa

Chinyere K Okoro<sup>1,20</sup>, Robert A Kingsley<sup>1,20</sup>, Thomas R Connor<sup>1</sup>, Simon R Harris<sup>1</sup>, Christopher M Parry<sup>2,3</sup>, Manar N Al-Mashhadani<sup>3</sup>, Samuel Kariuki<sup>4</sup>, Chisomo L Msefula<sup>5,6</sup>, Melita A Gordon<sup>7</sup>, Elizabeth de Pinna<sup>8</sup>, John Wain<sup>8,9</sup>, Robert S Heyderman<sup>5,10</sup>, Stephen Obaro<sup>11,12</sup>, Pedro L Alonso<sup>13,14</sup>, Inacio Mandomando<sup>14,15</sup>, Calman A MacLennan<sup>16,17</sup>, Milagritos D Tapia<sup>18</sup>, Myron M Levine<sup>18,19</sup>, Sharon M Tennant<sup>19</sup>, Julian Parkhill<sup>1</sup> & Gordon Dougan<sup>1</sup>

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Recall the question posed earlier regarding serovar-specific tendencies for systemic infections:

Certain serovars show a much higher rate of systemic infections in humans, including host-adapted serovars such as *S. Choleraesuis* (swine) and *S. Dublin* (cattle)

Questions:

- What bacterial or host factor(s) are responsible for these differences?
- If bacterial, can the traits be acquired by more widespread, broad host-range *Salmonella* serovars?



## Acknowledgements



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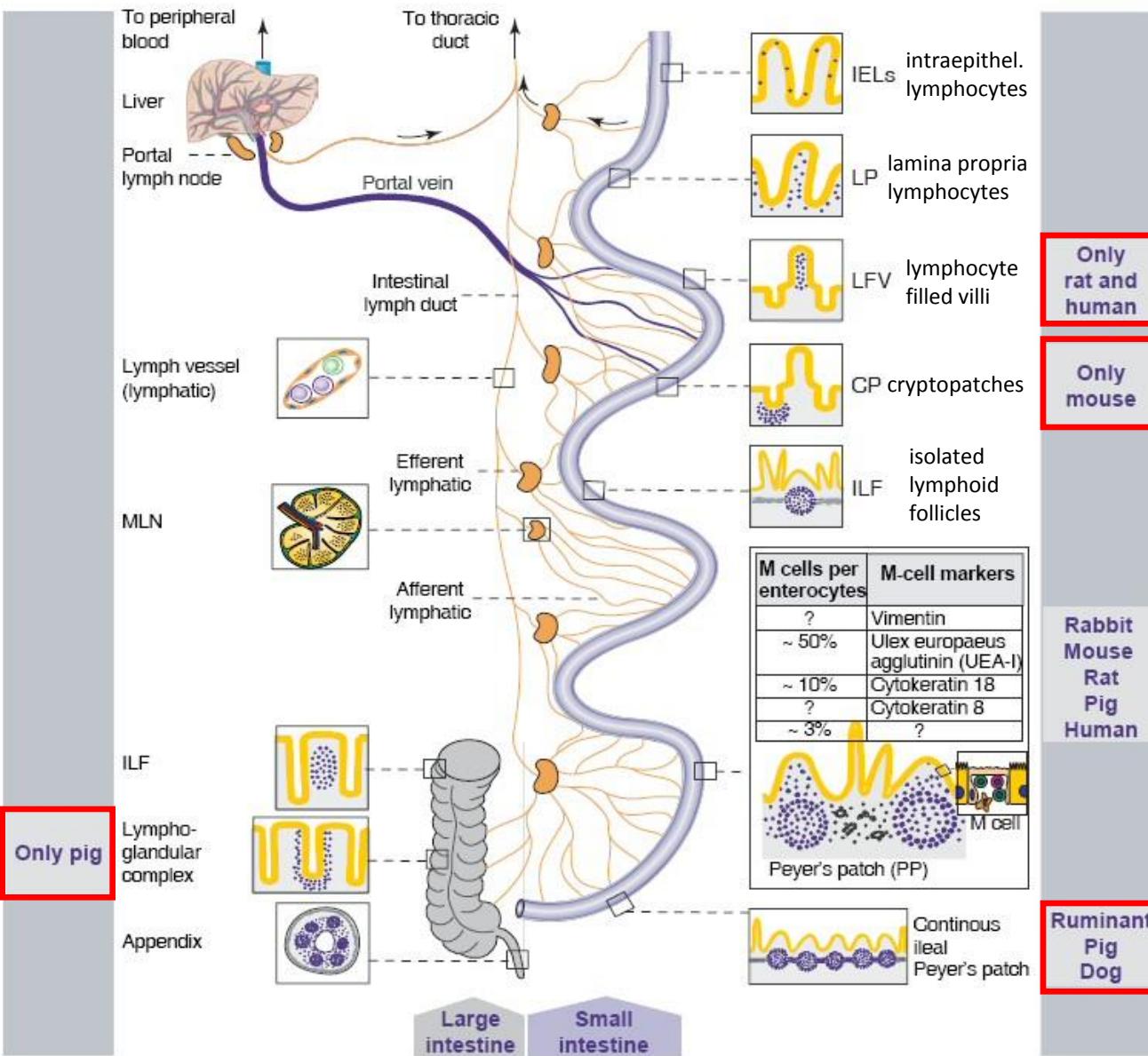


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# The Gastrointestinal Associated Lymphoid Tissue (GALT)



The GALT includes many other, additional regions of accumulated immune cells

Many of these regions are species-specific, e.g. the [cryptopatches](#) in mice or the [lympho-glandular complex](#) in swine

Other regions, such as the extended [Continuous Peyer's Patch](#), is found only in certain animal species

In addition, the [M Cells](#) show specific differences in surface and structural aspects in different animal species and humans

Suggest differences in antigen presentation or immune responses