

Emerging Microbial Hazards

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Overview

- Microbial hazard
- What is a pathogen
- Foodborne disease
- Emerging pathogens
- Factors in the appearance of emerging pathogens
- Case studies:
 - Cryptosporidium
 - E.coli O157:H7



Microbial hazard

- Term generally used to refer to the microorganisms: bacteria, fungi or viruses, that pose a risk of causing foodborne illness if they reach a sufficient number of cells on or in food contaminated with them.
- illnesses are caused by pathogenic bacteria



What is a pathogen?

PATHOGEN: "that with produces suffering"

A pathogen is a biological agent that causes disease in its host

Pathogens include:

- Bacteria (e.g. Strep throat)
- Viruses (e.g. hepatitis A)
- Fungi (e.g. athletes foot)

Foodborne pathogen: pathogen that has been acquired in food or the food chain



Foodborne disease

"a disease of an infectious or toxic nature caused by, or thought to be caused by, the consumption of food or water",

Economic burden due to costs incurred from division of resources to patient treatment

"Food and waterborne diarrhoeal diseases, for example, are leading causes of illness and death in less developed countries, killing an estimated 2.2 million people annually, most of whom are children"

WHO Global Strategy for Food Safety, 2002

Some statistics

Foodborne illness causes:

76 million illnesses

325,000 hospitalisations

5000 deaths

In the U.S., according to the CDC

The cost of five foodborne pathogens
-6.9 billion USD in 2000 USDA ERS estimates

Five foodborne outbreaks in 1996

- 300-700 million GBP in 1996 in England and Wales



Unknown pathogens

Known pathogens account for:
19% of total estimated number of cases
36% of deaths

Great majority of foodborne illness caused by unknown or emerging pathogens

Pathogens that have acquired antibiotic resistance



Emerging pathogens

- New, re-emerging or antimicrobial-resistant infections
- Incidence has increased in the last 20 years
- Incidence threatens to increase in the near future
- •13 out of 27 "top" US pathogens identified within last 25 years



Aetiology	Reservoirs	Mode of transmission	Distribution
Verocytotoxigenic Escherichia coli	Livestock and water	Meat, faeces, direct contact	Worldwide
Campylobacteriaceae - Campylobacter species - Arcobacter species - Helicobacter species	Poultry and pigs	Meat and faeces	Worldwide
Salmonella Typhimurium DT104	Livestock and wildlife	Faeces, eggs, meat and direct contact	Worldwide
Listeria monocytogenes	Livestock, wildlife, environment	Meat, milk, faeces and vegetation	Worldwide
Vibrio species - V. cholerae O1 - V. parahaemolyticus - V. vulnificus	Aquatic environment, fish and seafood	Fish, seafood, faeces	Worldwide
Listeria monocytogenes	Environment, livestock and humans	Milk and dairy products, fish, poultry, raw meat, ready-to-eat meat products and person-to-person transmission	Worldwide
Yersinia enterocolitica	Livestock, wildlife, fish and water	Meat, faeces and raw fish	Worldwide
Aeromonas species	Aquatic environments and livestock	Meat, milk, seafood and vegetables	Worldwide
Mycobacterium bovis	Cattle	Milk	Worldwide
Brucella melitensis	Sheep and goats	Milk	Worldwide
Enterobacter sakazakii	Not known	Powdered infant milk formula	Worldwide

Emerging pathogens

- -Already present in the environment
- Change in conditions give a selective advantage
 - Infective agent
 - Host
 - Environment



Why do foodborne pathogens emerge?

- Globalisation of the food supply
- Changes in agricultural practices
- Inadvertent introduction of pathogens into new geographic areas
- Travellers, refugees, and immigrants exposed to unfamiliar foodborne pathogens while abroad
- Changes in microorganisms
- Change in the human population
- Changes in lifestyle



Globalisation of the food supply

- Improvement in international travel
 - From months to hours
- Dissemination of food animals and food products worldwide
- Availability of various produce year-round



Globalisation of the food supply

- Longer food chain-negative effect on food safety
 - Increased opportunity for contamination
 - Time/temperature abuse of products
 - Rapid spread of infectious agents over large distances
 - Emergence of particular pathogens in particular areas for the first time
 - Trend towards diffuse and widespread outbreaks



Globalisation of the food supply

- Difficult to trace source of outbreak
- Outbreak caused by a wide variety of foods
 - Traditionally meat, poultry, seafood, pasteurised milk
 - Now "low risk" foods e.g. apple juice, pepperoni, fresh fruit and vegetables
- E. coli O157:H7 outbreak associated with spinach in US (Sept 06)
 - 199 people infected across 26 states
 - 3 deaths
 - Contaminated irrigation water thought to be the cause





Changes in agricultural practices

- Intensive agriculture(high input)
 - High external input e.g. use of fertilisers and pesticides
 - More food produced per acre compared to other systems
 - Reduction of biodiversity
 - Irreversible soil erosion
 - Run-off to aquatic systems



Agricultural practices

- Extensive farming(low input)
 - Reduction of external inputs
 - Antibiotics, pesticides, herbicides, synthetic fertilisers
 - Diversification of crops and animals
 - Organic farming: a type of low input farming
 - Code of principles that are regulated internationally



Intensive farming

 May have lead to emergence and increased prevalence of Salmonella, Campylobacter, E. coli O157:H7



- High stocking rates facilitates dissemination of pathogens
- Increased prevalence of pathogens in animals going to slaughter
- Spreading of animal wastes onto agricultural land
- Feeding of grains



- Emergence of antibiotic resistant strains
 - Antibiotics are widely used in high input farming systems
 - To treat animal disease
 - To promote growth



Case Studies

- Three case studies produced as part of Safe Foods project
 - Cryptosporidium
 - Escherichia coli O157:H7 in cattle

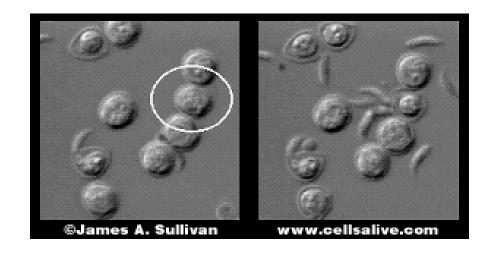


Case study: Cryptosporidium

Parasitic pathogen that causes a diarrhoeal illness called cryptosporidiosis



Cryptosporidium parvum in a fecal sample

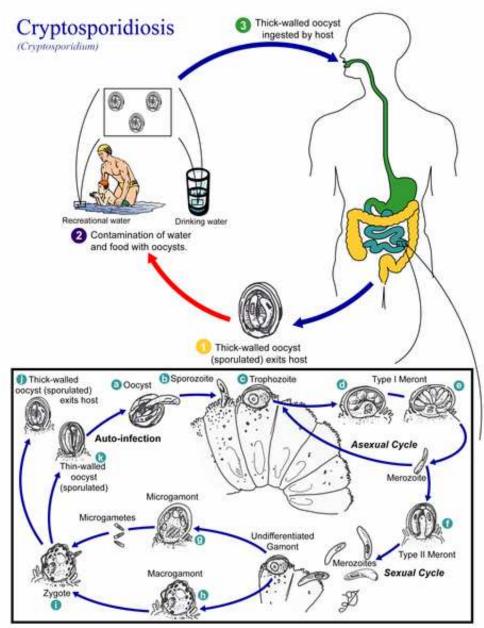


Electron micrograph of Cryptosporidium parvum



Life cycle of Cryptosporidium

- Obligate intracellular protozoan parasites
- Infective stage: highly resistant oocysts
- •Low infective dose





Cryptosporidium

- Identified as human pathogen in 1976
- 1993 Milwaukee outbreak
- Implications for
 - Immunocompromised individuals
 - Farm animals
- Consists of 13 recognised species
 - C. parvum, C. hominis, (human)
 - C. felis (cats), C. canis (dogs)



Cryptosporidium

- Most significant microbial pathogens to emerge
- Concerns because:
 - Cryptosporidium can be transmitted through water and food
 - Capable of causing a high degree of morbidity
 - No effective anti-parasitic treatment to eradicate from the GI tract in symptomatic individuals



Cryptosporidiosis

- C. hominis responsible for most outbreaks of human cryptosporidiosis in many regions
- Europe: C. parvum dominant
 - Intensive animal husbandry
 - C. meleagridis emerging as problem
 - Other species found have broad host range
- Humans major source of infections



Food borne Cryptosporidiosis

- In 1999, 10% cases via food in U.S.
- The rest contaminated water or person to person
- Raw fruits and veg, raw milk, meat and meat products, apple cider
- Hard to incriminate as Cryptosporidium hard to detect- underestimation of cases



Exposure assessment of Cryptosporidium

- Risk factors depend on
 - Quality of raw materials
 - Process steps and process environment
 - Product composition, packaging, storage conditions
 - Growth not important as Cryptosporidium cannot grow outside host, but viability reduction is important



Risk factors

- Water-raw sewage, treatment process, amount of water consumed
- Raw fruit and veg
 - Cultivation
 - Harvesting
 - Transport and storage
 - Industrial processing
 - Food preparation by consumer



Risk factors

- Meat products
 - Slaughterhouse
 - contact with contaminated water, contaminated environment, contamination of feed, cross contamination between animals, age influence of animals, influence of season
 - Storage of meat
 - Processing options



Cryptosporidium detection methods

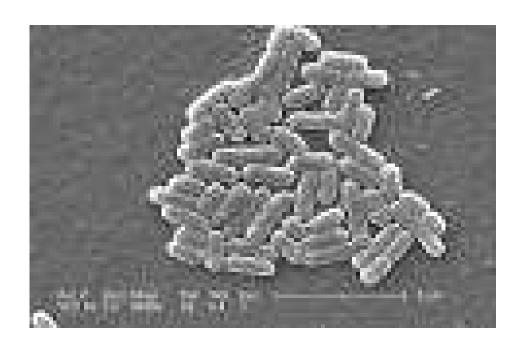
- Direct examination of stool preparations
- Epifluorescence microscopy for food, water and environmental samples
- Monoclonal antibodies conjugated to FITC
- PCR



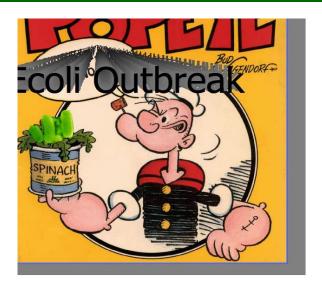
Cryptosporidium: Recommendations

- Difficult to prevent in food manufacturing
- Changes in animal husbandry and agricultural practices
- Broad host range of some strains
- Primary hazard in water supply, potential problem for fresh produce
- Not a significant problem in foods sufficiently heat treated
- More research into surveillance, epidemiology and detection methods needed
- Increased globalization of food chain increases risk
- Number of immunocompromised individuals increasing, making Cryptosporidium a greater hazard

Case study: Escherichia coli O157:H7 in cattle



Electron micrograph of E. coli



With reference to the recent outbreak of *E. coli* O157:H7in the U.S. associated with contaminated Spinach



E. coli 0157:H7

- Produce toxins:
 - Toxic to Vero cell cultures (vt1/vt2)
 - Similar to shiga toxins produced by Shigella (shiga-like toxin I or II)
- Verotoxin-producing E. coli (VTEC) or shiga-like toxin-producing E. coli (STEC)
- Survive in many environments
- Low infective dose
- Vulnerability of susceptible populations

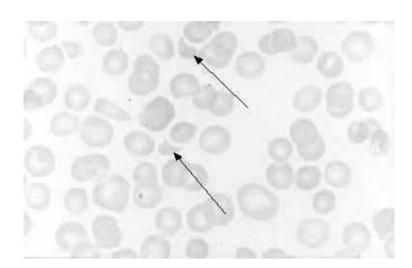
E. coli 0157:H7

- Since 1981 number of reported cases has increased
 - Improved surveillance programmes
 - Better detection methods
- Facultative anaerobic Gram-negative bacterium
- Found in GI tracts of mammals
- Somatic O157 antigen, flagellar H7 antigen



E. coli O157:H7, Complications

- Causes Haemorrhagic colitis (HC) and haemolytic uraemic syndrome (HUS) in humans
- Sometimes causes Trombotic trombocytopenic purpura (TTP)



A blood film from a patient with TTP showing relatively minor changes. Platelets are severely reduced and there are occasional red cell fragments (arrowed).



E. coli 0157:H7

- 1982, recognised as a human pathogen
- Three hypothesis for emergence
 - Recently emerged but conditions for spread always present
 - Slaughter and meat processing practices modified promoting contamination
 - Existence in meat supply, but consumer practices changed
- Changes at several levels of food chain



Virulence factors

- Most important
 - Shiga toxins I and II (encoded by stx1 and stx2); cytotoxic
 - Intimin protein (encoded by eae)
- Others include
 - EAST1 toxin (encoded by astA)
 - toxB protein
 - Associated attaching/effacing components (type III secretion system, Esp proteins, etc.)



Reservoirs/modes of transmission of *E. coli* O157: H7

- Isolated from many species
 - Sheep, goats, deer, pigs, cats, horses, gulls
- Cattle are major reservoir for *E. coli* O157:H7 and other VTEC
- Routes of transmission
 - Contaminated bovine products
 - Direct contact with infected animals
 - Consumption of contaminated foods
 - Person to person
 - Swimming/drinking contaminated water



E. coli 0157:H7

- Seasonal variation in shedding patterns
 - Increases during summer and early fall
- Young animals have higher prevalence
- E. coli O157:H7 can be isolated from healthy and ill cattle faeces
- Widespread in beef and dairy herds, variable within animal and herd
- No gender significance
- Calves on pasture may be less exposed to bacteria
- Prevalence higher in grain fed cattle; but not sure of influence on shedding of organism



E. coli 0157:H7

- Bedding material influences prevalence
- Faeces are vehicle for distribution of E. coli O157:H7
- Insects may play a role in hosts/vectors in dissemination
- Antimicrobial resistance occurs less in organic farming systems
- Example of "first world" emerging pathogen



E. coli 0157:H7 Recommendations

- Not enough information to be sure *E. coli* O157:H7 emerged
- Further research needed to elucidate mechanisms
 - Influence of bedding material, diet and starvation, milk based diets, survival in manure, seasonal variations, role of farm types, occurrence of antibiotic resistance, role of flies or other animals
- From this information
 - Modifications to control E. coli O157:H7



Summary

- Wide variety of factors involved in emergence
- Understanding of ecology of bacteria will help prediction
- To characterise emerging agents
 - Targeted surveillance systems
 - Outbreak investigations and research
 - Review pathogens that emerged in the past
 - Multidisciplinary teams

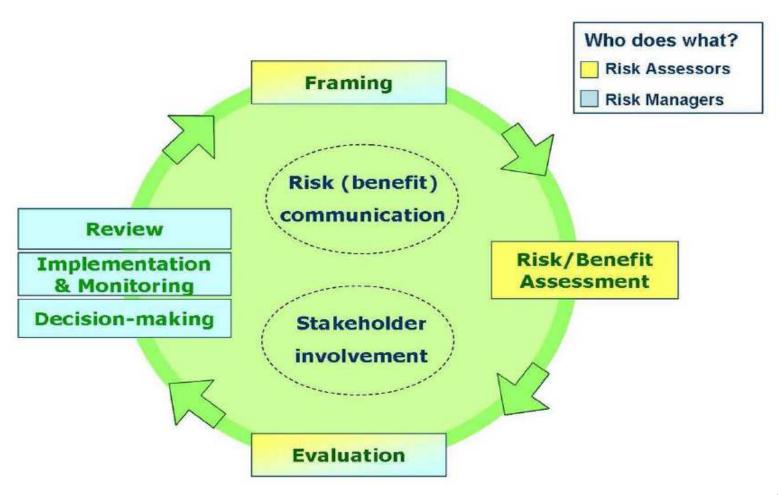


Summary

- Data used for
 - Risk analysis
 - Risk assessment
- Aid policy makers design most appropriate prevention strategies
- Safe foods overall aim
 - Assist in preparation of a new risk analysis approach for foods



Safe foods cycle: risk analysis framework





Based on the reports written for Safe Foods

To be published in Food and Chemical Toxicology

- "Early Detection of Emerging Risks Associated with Food and Feed Production", Marvin, H.J.P., Kleter, G.A, Kelly, B. G., Ossendorp, B., Vespermann, A., Beczner, J. and Prandini, A.
- "Emerging microbial hazards in food and feeds and factors that influence their emergence", Byrne, C. M., Bolton, D. J., Howlett, B., Kelly, B. G., Orlova, O., Ossendorp, B. and Vespermann, A.
- "Case Study: Cryptosporidium spp.", Orlova, O., Kelly, B. G. and Santare, D.
- "Case Study: Escherichia coli O157:H7 in cattle", Vespermann, A., Howlett, B., Bräunig, J., Käsbohrer and Bolton, D. J.
- "Case Study: Norovirus", Koopmans, M., Duizer, E., Reuter, G. and Beczner, J.



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