

WYSPY PATOGENNOSCI

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- 2.1. Wyspy genomowe a wyspy patogenności.
3. Występowanie wysp patogenności.
4. Ważniejsze bakteryjne wyspy patogenności.
- 4.1. *Escherichia coli*.
- 4.2. *Yersinia* spp.
- 4.3. *Salmonella* spp.
- 4.4. *Shigella* spp.
- 4.5. *Vibrio cholerae*.
- 4.6. *Helicobacter pylori*.
5. Podsumowanie

Pathogenicity islands

Abstract: Recent studies on bacterial virulence mechanisms revealed that some virulence determinants are located within so-called pathogenicity islands (PAIs). That term has been invented to describe large, often unstable DNA regions harboring virulence and mobility genes, usually flanked by direct repeats. PAIs allow bacteria to acquire or lose large amounts of DNA, and are regarded as elements involved in bacterial macroevolution. The PAIs of several bacterial species are reviewed herein.

1. Introduction.
2. Common characteristics of pathogenicity islands.
- 2.1. Genomic islands and pathogenicity islands.
3. Occurrence of pathogenicity islands.
4. More important bacterial pathogenicity islands.
- 4.1. *Escherichia coli*.
- 4.2. *Yersinia* spp.
- 4.3. *Salmonella* spp.
- 4.4. *Shigella* spp.
- 4.5. *Vibrio cholerae*.
- 4.6. *Helicobacter pylori*.
5. Summary

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