

Genomic Epidemiology: Whole-Genome-Sequencing-Powered Surveillance and Outbreak Investigation of Foodborne Bacterial Pathogens - DTU Orbit (08/11/2017)

Genomic Epidemiology: Whole-Genome-Sequencing-Powered Surveillance and Outbreak Investigation of Foodborne Bacterial Pathogens

As we are approaching the twentieth anniversary of PulseNet, a network of public health and regulatory laboratories that has changed the landscape of foodborne illness surveillance through molecular subtyping, public health microbiology is undergoing another transformation brought about by so-called next-generation sequencing (NGS) technologies that have made whole-genome sequencing (WGS) of foodborne bacterial pathogens a realistic and superior alternative to traditional subtyping methods. Routine, real-time, and widespread application of WGS in food safety and public health is on the horizon. Technological, operational, and policy challenges are still present and being addressed by an international and multidisciplinary community of researchers, public health practitioners, and other stakeholders.

General information

State: Published

Organisations: National Food Institute, Research Group for Genomic Epidemiology, University of Georgia, Texas Technical University

Authors: Deng, X. (Ekstern), den Bakker, H. C. (Ekstern), Hendriksen, R. S. (Intern)

Number of pages: 22

Pages: 353-374

Publication date: 2016

Main Research Area: Technical/natural sciences

Publication information

Journal: Annual Review of Food Science and Technology

Volume: 7

ISSN (Print): 1941-1413

Ratings:

Web of Science (2017): Indexed Yes

Scopus rating (2016): SJR 2.668 SNIP 2.479 CiteScore 8.29

Web of Science (2016): Indexed yes

Scopus rating (2015): SJR 2.665 SNIP 3.01 CiteScore 8.14

Scopus rating (2014): SJR 2.344 SNIP 2.292 CiteScore 6.66

Scopus rating (2013): SJR 2.45 SNIP 2.903 CiteScore 7.34

Scopus rating (2012): SJR 1.685 SNIP 2.983 CiteScore 4.81

Scopus rating (2011): SJR 1.122 SNIP 2.182

Original language: English

Food safety, Public health, Epidemiology, Subtyping, Bioinformatics, Genomics

DOIs:

10.1146/annurev-food-041715-033259

Source: FindIt

Source-ID: 277289958

Publication: Research - peer-review › Journal article – Annual report year: 2016