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Aplicaciones de las nuevas tecnologías de secuenciación en seguridad y fraude alimentario

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Who We Are

Illumina is a global leader in DNA sequencing and array-based technologies, serving customers in the research, clinical and applied markets. Our products are used for applications in the life sciences, oncology, reproductive health, agriculture and other emerging segments.



\$4.5 billion (2023)

Annual revenue



~9,100

Number of employees



Jacob Thaysen

President & CEO



San Diego, CA, USA

Headquarters



1998

Year founded

More than 20,000 Sequencing Systems Installed Around the World

Our sequencing instruments

Low-throughput



MiSeq™



MiniSeq



iSeq 100



NextSeq 500

Mid-throughput



NextSeq 550



NextSeq 1000/2000



MiSeq™Dx



NextSeq™ 550Dx

High-throughput



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OUR MISSION

Improve human health
by unlocking the power of
the genome

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Genomics applications in Food Science

Integrated in many processes and steps across the food chain

NGS and arrays are used in all the steps from gene to plate



Breeding & Selection

Production

Processing

Consumption

Research

Genomic Breeding

Parentage

Veterinary Diagnostics

Environmental Testing

Food Safety

Tracing and Surveillance

Food Authenticity

Food Traceability

Spoilage

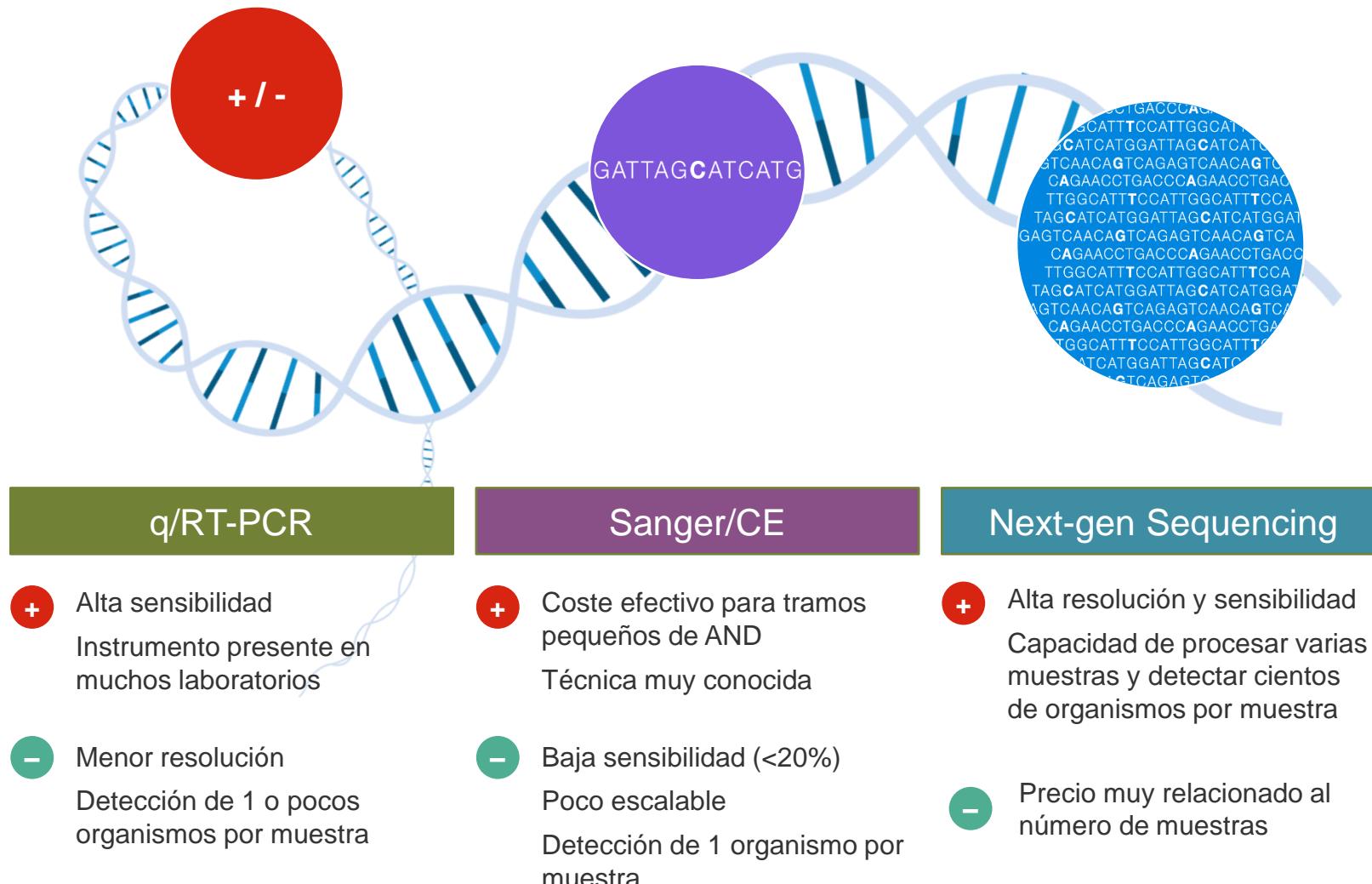


The power of NGS

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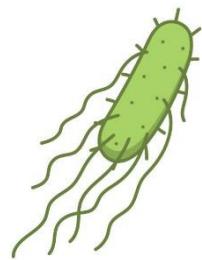
Comparación de tecnologías de biología molecular



Comparación de tecnologías de biología molecular

q/RT-PCR

Presence/ absence results



Salmonella enterica

Sanger/CE

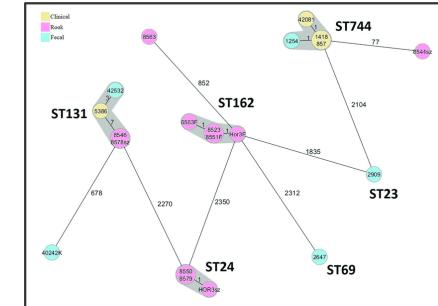
MLST of 7 loci for Typing



Salmonella enterica subsp. enterica

Next-Gen Sequencing

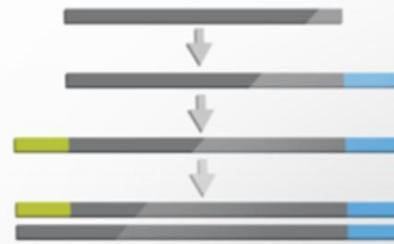
wgMLST of the Whole genome



Full ID for outbreak investigation, serotyping, AMR

The NGS approach

Integrated workflows from prepared sample to answer



Library preparation



Sequence



Analyze & interpret

Flexible workflow solutions enable markets

Simplified
library prep

Custom
content

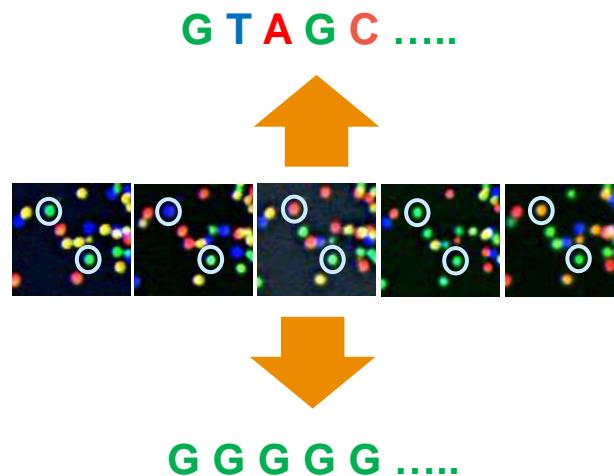
Flexible, accessible
sequencing

Integrated
analysis

Analysis

Primary analysis

- Runs in parallel to the sequencing
- Uses the images to provide base calls of the reads



Secondary analysis

- Maps the reads to the reference genome (alignment)
- Analyses the mapped reads for any differences compared to the reference genome (variant calling)

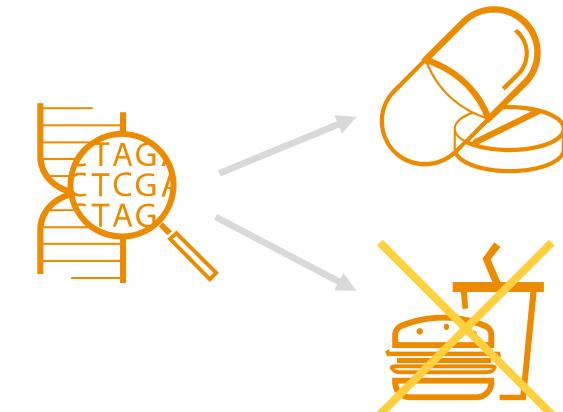
The diagram illustrates the secondary analysis process. It shows a magnifying glass focusing on a DNA double helix. The DNA sequence is aligned with a reference sequence at the bottom. The aligned sequence is: CGACGGACTCGATTA, ATACGACAGACTCGATTA, CATACTGACAGAC, CGGACTCGATTA, CATACTGACGGACTCG, CGACGGACTCGATTA, CATACTGACGGACTCGATTA. The last two lines are highlighted in orange, indicating they are the reference sequence.

CGACGGACTCGATTA
ATACGACAGACTCGATTA
CATACTGACAGAC
CGGACTCGATTA
CATACTGACGGACTCG
CGACGGACTCGATTA
CATACTGACGGACTCGATTA

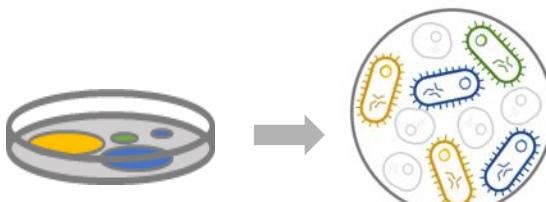
Reference Sequence

Interpretation

- Leads to knowledge and insights into basic biology
- Potential causes of genetic disease
- Possible treatment or prevention



Applications for Culture vs. Culture-Free Sequencing



Whole-Genome Sequencing (WGS)

Sequencing a **cultured isolate** for taxonomy, evolution, and function

- Food outbreaks, identify source
- Monitor for pathogens in product and facilities
- Differentiate resident, persistent, and transient pathogens
- Virulence factors, AR genes, and other genetic markers of interest
- Maintain or enhance food quality (starter cultures)



Amplicon Sequencing

Sequencing **specific genes** for taxonomy and relative abundance

- Community profiling to predict pathogen, spoilage risk, and allergen
- Detect genetic modifications (GMOs)
- Authenticate high value foods (olive oil)
- Verify speciation (kosher, halal)
- Procure correct ingredients for final product (gluten free)



Shotgun metagenomics

Sequencing "**everything**" in a sample for taxonomy, abundance, and function

Problemas relacionados con la alimentación tienen un gran impacto socio-económico

La genómica puede ayudar a prevenir o reducir ese impacto



Seguridad Alimentaria

El costo estimado de los incidentes de seguridad alimentaria para la economía de los Estados Unidos es de alrededor de **\$7 mil millones por año**, que proviene de notificar a los consumidores, retirar alimentos de los estantes y pagar daños como resultado de demandas.

Economic Impact of Food Safety Outbreaks on Food Businesses. Hussain MA, et al. Foods. 2013. [PMID: 28239140](#)



Fraude Alimentario

Las estimaciones de los expertos han encontrado que el fraude alimentario **afecta al 1% de la industria alimentaria mundial** a un costo de alrededor de 10 a 15 mil millones \$ al año, aunque algunas estimaciones de expertos más recientes sitúan el costo en \$ 40 mil millones al año.

FDA: [Economically Motivated Adulteration \(Food Fraud\)](#)



Producción Alimentaria

La producción agrícola y la seguridad alimentaria siguen siendo **una de las principales preocupaciones mundiales**. La investigación y los comentarios en este número de Nature destacan las diversas amenazas a nuestras frutas y verduras y los efectos de arrastre de las crisis alimentarias.

Genomics and our future food security. Editorial. Nat Genet. 2019. [PMID: 30705438](#)



NGS applications in Food Science

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Food safety



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Food safety applications: Using NGS to Assess Food Pathogen Outbreak

20 minutos Actualidad Nacional Internacional Deportes Cultura Opinión Más ▾ | 🔎 | ⚙

Una misma cepa de listeria procedente de la carne La Mechá ha contagiado a 144 pacientes

EFE 12.09.2019 - 15:21H

Gobierno de España | Ministerio de Ciencia e Innovación | Instituto de Salud Carlos III

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French officials confirm some E. coli cases linked to Nestlé pizza

75 pacientes, 41 con Síndrome Urémico Hemolítico.

2 niños fallecidos.

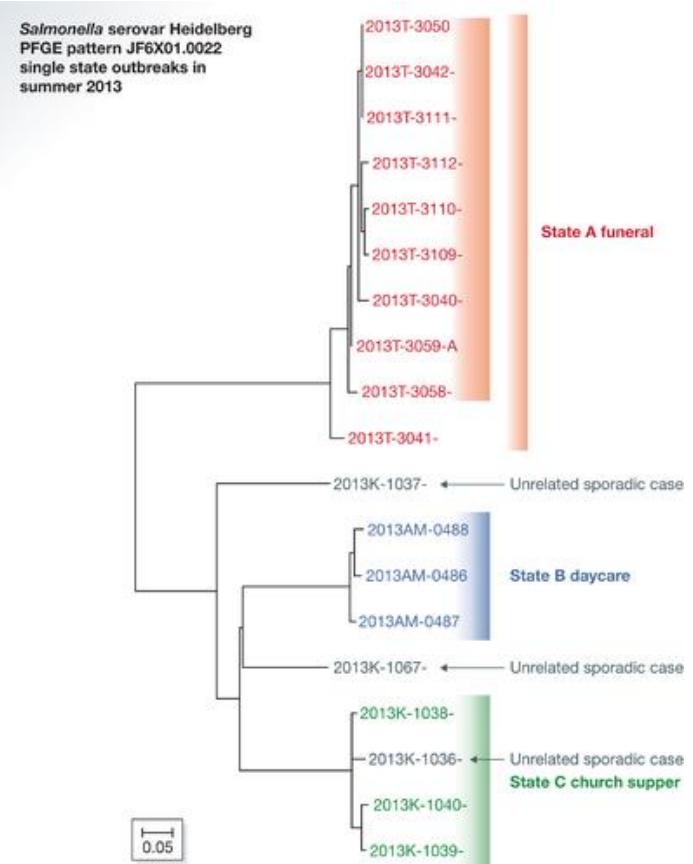
Possible origen en harina.

Detección foco en Marzo 2022.

Primeros enfermos en Enero



CDC discovers Listeria outbreak linked to enoki mushrooms

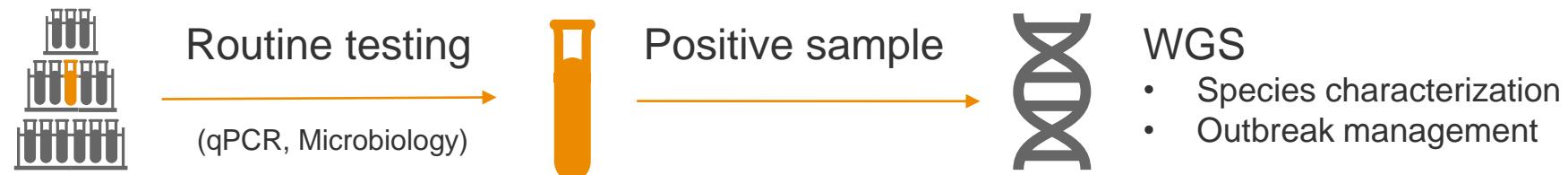


Aplicaciones en la seguridad alimentaria



La tecnología NGS puede ser una aproximación reactiva y proactiva

- Herramienta industrial para monitorizar ingredientes
- Permite evaluar la eficiencia de prevenir y de los controles sanitarios
- Se usa para determinar la persistencia de patógenos en el ambiente
- Posible indicador de resistencias antimicrobianas



Food authenticity and traceability



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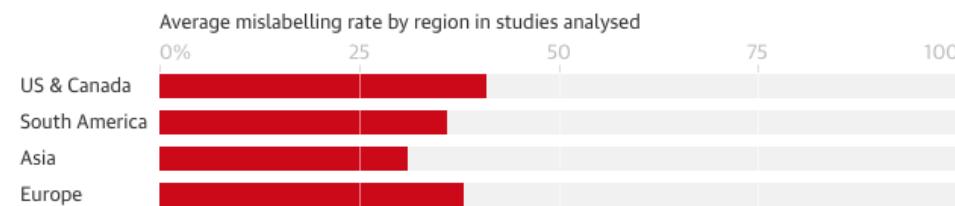
Food fraud is widespread

Adulteration, mixing, or mislabeling is extremely common globally



Revealed: seafood fraud happening on a vast global scale

Guardian analysis of 44 studies finds nearly 40% of 9,000 products from restaurants, markets and fishmongers were mislabelled



Guardian graphic. Source: Guardian review of 44 seafood studies published since 2018

A Guardian Seascape analysis of 44 recent studies of more than 9,000 seafood samples from restaurants, fishmongers and supermarkets in more than 30 countries found that 36% were mislabelled, exposing seafood fraud on a vast global scale.

The Guardian, March 15th 2021 | <https://www.theguardian.com/environment/2021/mar/15/revealed-seafood-happening-on-a-vast-global-scale>

REVIEW ARTICLE

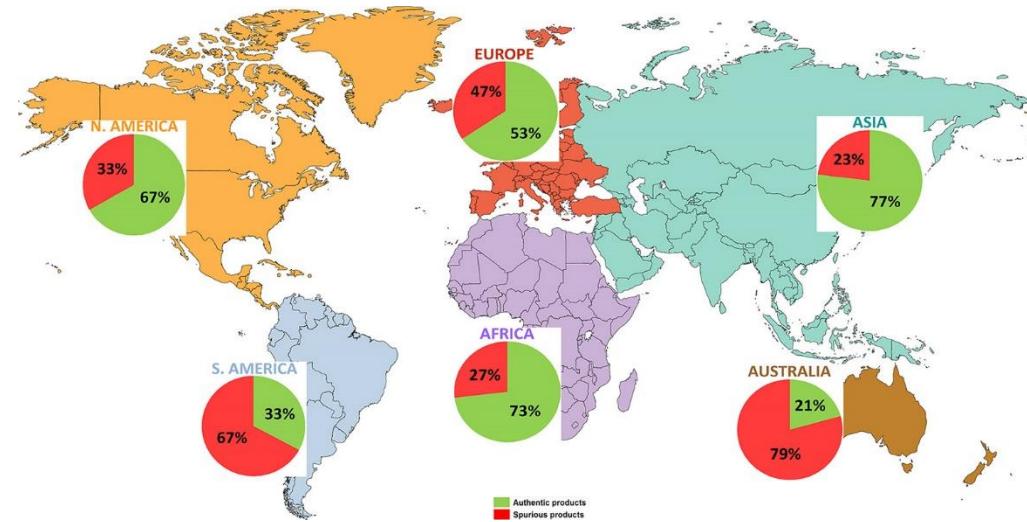
Front. Pharmacol., 24 October 2019 | <https://doi.org/10.3389/fphar.2019.01227>



The DNA-Based Authentication of Commercial Herbal Products Reveals Their Globally Widespread Adulteration

Mihail Cristin Ichim*

*"Stejarul" Research Centre for Biological Sciences, National Institute of Research and Development for Biological Sciences, Piatra Neamt, Romania



Front. Pharmacol., 24 October 2019 | <https://doi.org/10.3389/fphar.2019.01227>

Food fraud is widespread

Adulteration, mixing, or mislabeling is extremely common globally



Why NGS in Food authenticity?



Characterization of multiple species at once



Untargeted detection of thousands of organisms with no requirement for previous knowledge



Most reliable method for species identification



Allows better dealing with damaged DNA

DNA Barcoding & Metabarcoding

- Scientific foundations and databases growing
- Can provide composition + origin signatures
- Primers tailorable to specific clades
- Relationship between microbiome and food



international
BARCODE
OF LIFE

Food production

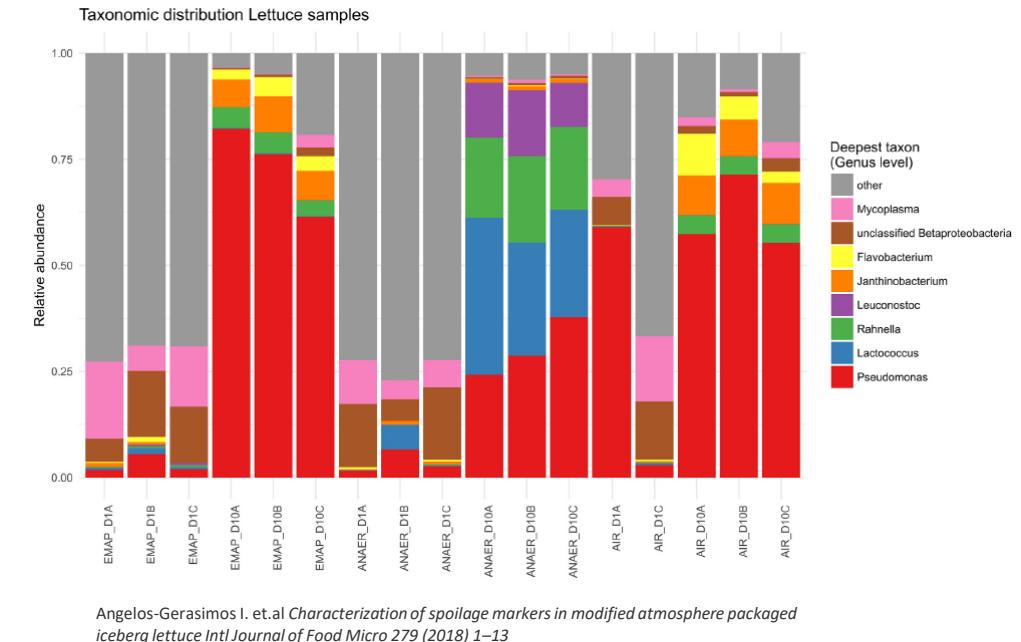


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Food production applications: Microbial Diversity in Packaging evaluation

Characterization of spoilage microbiota in packaged iceberg lettuce using culture-free amplicon sequencing

- Studied microbial and biochemical contributors to spoilage in fresh cut lettuce
- Microbial analysis of lettuce microbiome under aerobic @ 3%O₂ and perforated, and anaerobic conditions.
- Culture-free amplicon sequencing revealed *Pseudomonas* spp. as the dominating aerobic spoilers, whereas lactic acid producing bacteria (*Leuconostoc* spp. and *Lactococcus* spp.) dominated anaerobic packaging condition.
- Identification of genera with spoilage capacity to produce off odors, leading to early consumer rejection



Understanding of microbial contributors to spoilage can inform future packaging technologies and design to extend shelf life of lettuce

Food production applications:

The secret behind the beer: the genetics of industrial yeasts

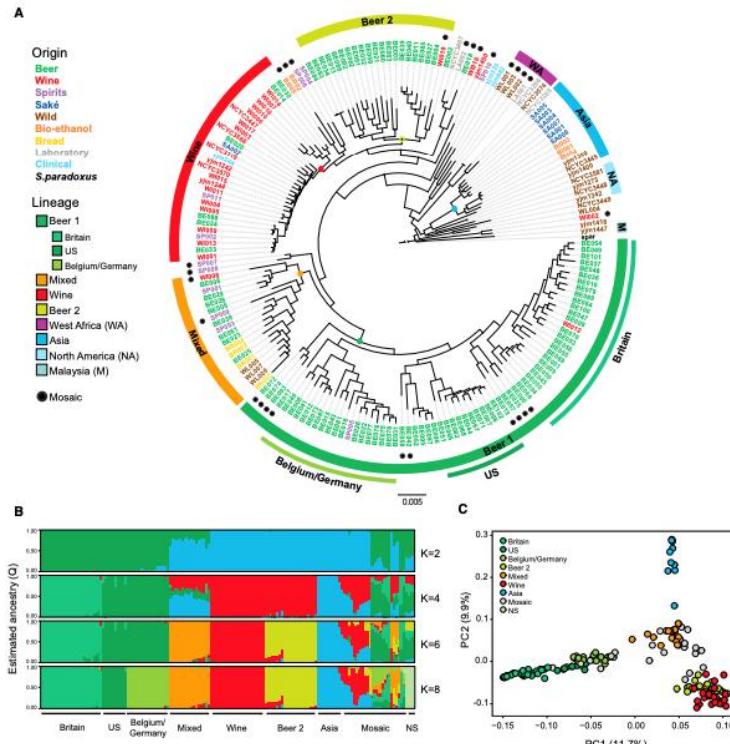


Article

Cell

Domestication and Divergence of *Saccharomyces cerevisiae* Beer Yeasts

Brigida Gallone,^{1,2,3,4,11} Jan Steensels,^{1,2,11} Troels Prahli,⁵ Leah Soriaga,⁶ Veerle Saels,^{1,2} Beatriz Herrera-Malaver,^{1,2} Adriaan Merlevede,^{1,2} Miguel Roncconi,^{1,2} Karin Voordeckers,^{1,2} Loren Miraglia,⁶ Clotilde Teiling,⁹ Brian Steffy,⁹ Maryann Taylor,¹⁰ Ariel Schwartz,⁶ Toby Richardson,⁶ Christopher White,⁵ Guy Baele,⁷ Steven Maere,^{3,4,*} and Kevin J. Verstrepen^{1,2,12,*}



KEY HIGHLIGHTS

- Analyzed the genomes and phenomes of 157 industrial *Saccharomyces cerevisiae* yeasts.
- Present-day industrial yeasts originate from only a few domesticated ancestors
- Beer yeasts show strong genetic and phenotypic hallmarks of domestication
- Domestication of industrial yeasts predates microbe discovery
- The genome sequences, phylogenetic tree, and phenotype data can be used to set up marker-assisted breeding schemes similar to those routinely used for the breeding of superior crops and livestock

The microbiota can influence cheese quality

Shotgun metagenomics to characterize the bacterial community

frontiers
in Microbiology

REVIEW
published: 23 May 2018
doi: 10.3389/fmicb.2018.01020



Sequencing of the Cheese Microbiome and Its Relevance to Industry

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³APC Microbiome Ireland, Cork, Ireland

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Sheehan JJ and Cotter PD (2018)
Sequencing of the Cheese
Microbiome and Its Relevance to
Industry. Front. Microbiol. 2018:01020.
doi: 10.3389/fmicb.2018.01020

INTRODUCTION

Cheese has a diverse microbial community, which indeed can vary within the cheese from the core to the surface that is greatly influenced by manufacturing including ripening conditions. Understanding the composition of this community (microbiota), and its impact on the quality and safety of cheese products, is of critical importance. In addition to, in the majority of cheeses, consciously added starter and adjunct bacteria (which are added as a supplement), cheese contains a heterogeneous variety of other, non-starter, microorganisms. These various microorganisms can play vital roles in the development of the organoleptic properties of cheese (Fox et al., 2008), nutrient composition, shelf-life, and safety.

Historically, culture-based microbiology techniques were used to gain an understanding of the microbial component of cheese. However, it has become increasingly clear that this approach can be limited in its ability to detect "difficult-to-culture" or sub-dominant microorganisms, thereby potentially providing misleading results. As a result, culture-independent approaches have become

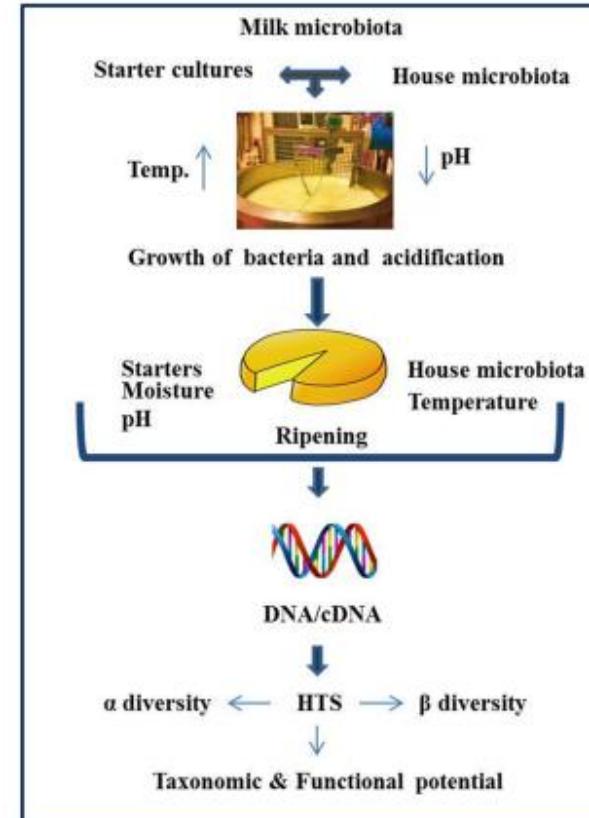


FIGURE 1 | Schematic representation of factors influencing the cheese microbiota, as revealed by HTS.

KEY HIGHLIGHTS

- Cheese has a diverse microbial community and understanding its composition, and its impact on the quality and safety of cheese products, is of critical importance.
- The Microbiota composition is influenced by several factors during cheese production
- The Microbiota play vital roles in the development of the organoleptic properties of cheese, nutrient composition, shelf-life, and safety.
- HTS is a powerful technique that can be used to provide a detailed insight into the microbiology of dairy related samples.

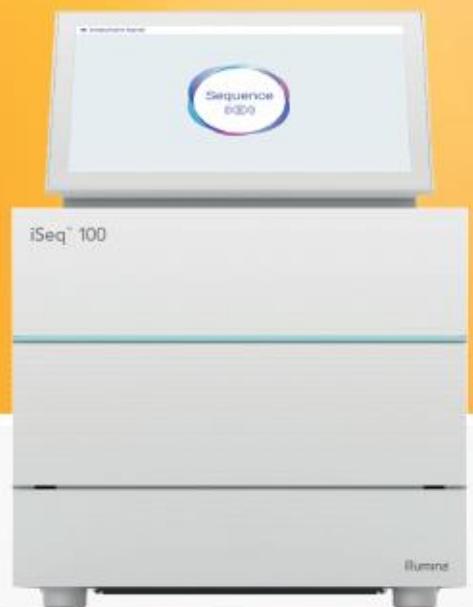
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