There's CRISPRs in my Yogurt?!

Background

Many dairy products, such as yogurt and cheese, use live bacteria in their

The bacteria used in yogurt have an adaptive immune system called the Clustered Regularly Interspaced Short Palindromic Repeat (CRISPR)-Cas System. The CRISPR System protects bacteria from repeat viral infections due to the cells' ability to store fragments of viral DNA, where it can then degrade any identical DNA.

CRISPR Technology- How it works

- Bacteriophage binds and injects its viral DNA
- 2. Cas1-Cas2 enzymes of the microbe recognize and cut out a segment of foreign DNA.
- Add protospacer DNA to genomic CRISPR region

CRISPR-associated endonuclease (Cas)



Source: Biotility: Applied Biotech Training, University of Florida



Methods

- Heirloom yogurt was made from an undefined mixture of microbes
- Commercial yogurt used was MountainHigh[™] Plain yogurt



Figure 1: MountainHigh™ and inoculated heirloom yogurt





- A polymerase chain reaction (PCR) was used to amplify the CRISPR region
- Gel electrophoresis used to analyze PCR products

Results

Gel Electrophoresis of CRISPR Samples





Figure 5: Size of CRISPR loci (amplified by PCR) from heirloom yogurt bacteria (S. Thermophilus)



Sample	Amplicon Size	
Heirloom 2	850 bp	GTTTTTG
Commercial 2	2400 bp	GTTTTTG

Table 1: Heirloom 2 bacteria from Fig. 5-4 and Commercial 2 bacteria from Fig. 6-4. Repeat sequence, number of spacers, and estimate amplicon size in CRISPR locus is shown.



• Colonies of heirloom and commercial yogurt were isolated on MRS agar plates

Figure 2: Isolation of heirloom and commercial microbes on MRS agar plates • Heirloom and commercial bacteria colonies were gram stained



Figure 4: Gram stain of heirloom yogurt bacteria (1000x oil immersion)

Figure 6: Commercial PCR gel results labeled with positive control

Repeat sequence TACTCTCAAGATTTAAGTAACTGTACAAC 6

of spacers

TACTCTCAAGATTTAAGTAACTGTACAAC 14

Commercial Yogurt Bacteria CRISPR Loci sequence

Spacer Sequences	Identity of Virus (Phage)	E-value
AAAATGGATTATCATATATTCATAAAGTGA	MAG: Bacteriophage sp. isolate 1807_32025, partial genome	2.70E+01
AACGCTTCCTAGGTTTAAGCCGTCTTTTCC	Streptococcus phage CHPC1041, complete genome 4	
TTACACAAAATTGGTTAACGTCAACGACAT	Streptococcus phage CHPC1041, complete genome 2.00E	
AAATTGTCGTTCATTGCCGTTCAACACCTC	Streptococcus phage SW23, complete genome 2.0	
ACAAAGGTTGTTATTACAGTTGGTTTATA	MAG TPA_asm: Caudoviricetes sp. isolate cthvO2, partial genome	2.40E+01
ATTTAAAGAGTTGATTAAGTGCGTTACTGT	MAG TPA_asm: Caudoviricetes sp. isolate ctJ298, partial genome	6.70E+00
AAATGGGTTATGCTGTTCAATATGCGTCCC	Streptococcus phage SW12, complete genome	2.80E-02
AAACTGAAAACAACACAGACAATTCAACAA	No significant match	
GCCCAAAATGCTAGACGTTTGAATGACGGC	Streptococcus phage CHPC663, complete genome 2.00E-0	
ATGAAGAACGTGATTCACCTACGGTATGCT	MAG TPA_asm: Caudoviricetes sp. isolate ctFZS48, partial genome	2.00E-06
TTATGTACATTTCAAAAGGTGTCATCCATA	Streptococcus phage SW12, complete genome 2.00	
AATACGATTCATTGCAAGAGTATGTTAATG	Streptococcus phage SW12, complete genome 2.00	
AATATGAAATCACAAGTAATGAGCCTAGCA	Streptococcus phage CHPC663, complete genome	7.00E-06
AAAACATTACAAGAGACTACTTTTACAAT	Streptococcus phage Javan536, complete genome	6.00E-06

identity, and its E-value.

Heirloom Yogurt Bacteria CRISPR Loci sequence

Spacer Sequences	Identity of Virus (Phage)	E-Value
AGATACTCTTGTCGCCTCTGAACAACCAG	Streptococcus phage pST	4.00E-04
TTTGATGGCTCTTGGTAGGGAACTGGATAT	Streptococcus phage pST	7.00E-06
AAAGACAAGCCCAAGGGATTGAACTAGCAA	Bacteriophage sp.	1.05E+02
CTACATTATTGATCATGTTTTTTCTCCTGT	Streptococcus phage CHPC925	2.80E-02
TAGAAGGCTCTGGAAATACAAAGCAATTCT	Caudoviricetes sp.	6.7
CGAACAGTTGGCGAGAAATCCGTCTGGCGT	Streptococcus phage CHPC929	2.0E-03

Table 3: BLAST analysis of the spacers in Heirloom 2 (See **Table 1**) for bacteriophage *identity, and its E-value.*

Our research compared the CRISPR immune systems of S. thermophilus in commercial and heirloom yogurts. We hypothesized that commercial yogurts would have superior immunity because companies want to prevent infection by viruses. However, we obtained less data from commercial samples due to less successful PCR electrophoresis, limiting our comparison.

On average, commercial samples had more spacer sequences (13.5) than heirloom samples (8.4), suggesting potentially better viral resistance. This is significant for human health as we prefer more resistant bacteria in our diet.

Our study was limited by the insufficient cultures and genetic information from commercial yogurt. Future studies could benefit from equal PCR reactions and cultures for both types.

Potential improvements could include exploring Lactobacillus CRISPR loci differences and comparing various yogurt types, such as lactose-free yogurts, or Greek yogurt. This could provide a broader understanding of the microbial differences and similarities in our food.

Barrangou, Rodolphe, and Philippe Horvath. "CRISPR: New Horizons in phage resistance and strain identification Annual Review of Food Science and Technology, vol. 3, no. 1, 10 Apr. 2012, pp. 143–162, https://doi.org/10.1146/annurev-food-022811-101134.

Zhang H, You C. 16S ribosomal RNA-depletion PCR and its application in cause analysis of yogurt package shrinkage. J Dairy Sci. 2022 Sep;105(9):7288-7297. doi: 10.3168/jds.2021-21575. Epub 2022 Aug 2. PMID: 35931476. Andrey N. Shkoporov, Colin Hill, Bacteriophages of the Human Gut: The "Known Unknown" of the Microbiome, Cell Host & Microbe, Volume 25, Issue 2, 2019, Pages 195-209, ISSN 1931-3128, https://doi.org/10.1016/j.chom.2019.01.017 CRISPR Technology- How it works. Biotility: Applied Biotech Training. University of Florida

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Table 2: BLAST analysis of the spacers in Commercial 2 (See **Table 1**) for bacteriophage

Conclusion



