

Outbreak of staphylococcal food poisoning among children and staff at a Swiss boarding school due to soft cheese made from raw milk

Sophia Johler¹, Delphine Weder², Claude Bridy², Marie-Claude Huguenin², Luce Robert², Jörg Hummerjohann³, Roger Stephan¹

¹Institute for Food Safety and Hygiene, Vetsuisse Faculty University of Zurich, Switzerland; ²Service de la Consommation et des Affaires Vétérinaires, Canton de Neuchâtel, Switzerland; ³Agroscope, Institute for Food Sciences, Bern, Switzerland

Outbreak data

Background

On October 1, 2014, children and staff members at a Swiss boarding school consumed Tomme, a soft cheese produced from raw milk. Within 7 h, all 14 persons who had consumed the cheese fell ill, among them 10 children and 4 members of the staff. Based on the short incubation time, as well as the clinical symptoms, SFP due to consumption of the raw milk cheese was considered a possible cause of the outbreak.

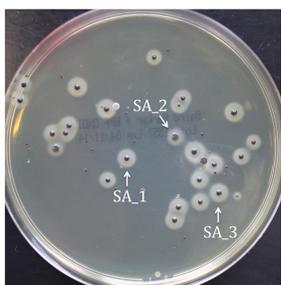
- *S. aureus* was isolated using rabbit plasma fibrinogen (RPF) agar, following EN ISO 6888-2 protocol.
- Source: Tomme raw milk cheese
- 14 patients (10 children, 4 adults)
- Symptoms: abdominal pain, emesis, (bloody) diarrhea, fever
- **Incubation time:**
 - 2.5 h (children <10y)
 - 3.5 h (children >10y)
 - 7 h (adults)

Objective

We aim to present a food poisoning outbreak and to characterize its causative agent.



Bacterial isolation & enumeration

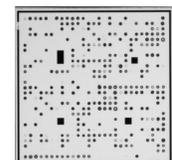


1:1,000,000 dilution

Rabbit plasma fibrinogen agar was used for enumeration of coagulase-positive staphylococci (CPS). In total, the cheese contained 10^7 CfU CPS/g. Interestingly, **three different *S. aureus* strains** (SA_1, SA_2, and SA_3) were **present at levels > 10^6 CfU/g.**

Microarray-based genotyping

All three *S. aureus* strains isolated from the Tomme cheese were characterized by DNA microarray analysis.



	major SE genes	newly described SE genes	spa type	agr type	clonal complex
SA_1	sea, sed	sej, ser	t711	I	CC8
SA_2	none	egc	t018	II	CC705
SA_3	none	egc	t458	III	CC20

We identified *S. aureus* SA_1 as the source of the outbreak.

SA_1 exhibited characteristics of a **genotype B** strain. This genotype comprises bovine *S. aureus* strains exclusively associated with very high within-herd prevalence of mastitis and has been described as a major contaminant in Swiss raw milk. It is highly likely that the raw milk used for Tomme production was heavily contaminated with *S. aureus*.

Enterotoxin detection

- SET2 mini Vidas was used to screen the cheese for SEA-SEE.
=> **SEA+** and **SED+**
- We determined by SET-RPLA that the cheese contained:
 - > **6 ng of SEA/g**
 - > **200 ng of SED/g**



Conclusions

- 1) The outbreak was caused by Tomme raw milk cheese contaminated with an *S. aureus* strain producing **SEA** and **SED**.
- 2) The outbreak was caused by a **genotype B** strain, a genotype exclusively linked to high within-herd prevalence of mastitis.
- 3) The incubation time depended on the age of the patient.

