Developing Flavours in Soft Cheeses



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Summary of the differences between lactic curd and rennet curd

Curd mainly lactic

- Acidification and only then draining
- pH 4,5
- Residual Lactose
- Deminéralisation
- Low buffer capacity
- Weak Interactions between proteins
- High moisture level
- Fragile
- Small cheeses

Curd mainly rennet

- Draining and then acidification
- pH 5,2
- No residual lactose
- High minéral concentration
- Strong buffer capacity
- Strong interactions between caseins
- Low moisture level
- Ferm and elastic
- **Big cheeses**

PLAN

1. Texture

- 2. Malt and Fruity flavours
- **3. Cabbage and Garlic flavours**
- 4. Goaty, Blue cheese to Soapy flavours
- **5. Mushroom flavours**
- 6. Plastic

Conclusions

• Context : Important changes in curd, composition, structure, aspect, texture, colour and taste due to biological activity and transport phenomena.

Lactose and lactate changes during ripening

Leclercq-Perlat et al, 2012

Microbial growth during ripening

DESACIDIFICATION

MATURATION

SURFACE

Fungi :

-Yeasts -Geotrichum -Penicillium

Fungi + ripening bacteria :

Micrococcinae : Micrococcus, Arthrobacter, Brevibacterium

Corynebacterinae : *Corynebacterium*

Staphylococci

INSIDE

pH Change at the surface and in the core of a Camembert cheese

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Buffer capacity

- Buffer capacity can be defined as the resistance to pH change, low pH decrease the buffer capacity with strong consequences on texture
 - Low pH at moulding (pH= 6.1 (traditional camembert) to 5,2 (lactic washed rind))
 - Makes easier the pH rise after the *Penicillium* growth, gives creamy underind after a while
 - Makes the cheese center tough and chalky

Evaluation of the ripening efficiency on the base of texture change

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Proposed Mechanism for δ-lactones production from triglycerides containing 5-hydroxy acids. R1=(CH2)n-CH3 ; R2 and R3=CO-(CH2)n-CH3 (Alewijn *et al.* 2007)

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Malt and Fruity flavours

They are mainly coming from amino acid breakdown

These flavour compounds are produced early in the ripening process

Yeasts are mainly responsible of these flavours

At the end of ripening, some bacteria may reinforce these flavour notes especially in washed rind cheeses

OXIDATIVE DEGRADATION OF AMINO ACIDS

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The amino acids breakdown is the main source of malt and fruity notes :

- aldehydes (malt and chocolate notes),

- alcools, esters, thioesters (fruity notes)

Starters sources of malt and fruity notes

- aldehydes (malt and chocolate notes) :

Debaryomyces hansenii, Staphylococcus xylosus

- alcools, esters, thioesters (fruity notes) :

Kluyveromyces lactis, Geotrichum candidum

Olfactive thresholds of esters

Compounds	Aromatic Notes	perception threshold
Ethyl acetate	Solvent, fruity	5 ppm ^a 22 ppm ^b
Ethyl Propionate	Pineapple	9,9 ppba
Ethyl Butyrate	Pineapple	0,13 à 45.10 ⁴ ppb ^a 0,6 ppm ^b
Ethyl Hexanoate	pineapple, banana	1 ppb ^a 0,85 ppm ^b
Isoamyle acetate	pear, banana	2 ppb ^a
2-phenylethyl acetate 2-phenylethyl propionate	floral, rose floral, fruity	18,5 ppm ^C 16,8 ppm ^C

a : in water ; b : in oil or butter ; c : in a cheese base.

Thioester formation

Acyl S Me + CoASH

Geotrichum candidum is able to produce thioesters (Berger et al, 1999)

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Degradation of amino acids also leads to :

- phenols (animal notes, role in ewes milk cheeses)
- sulfur compounds (garlic and cabbage notes- important especially in smear ripened cheeses)

Relations between sulfur amino acid metabolism and flavour production

Olfactive thresholds of sulfur compounds

Olfactive thresholds (µg/kg or ppb)

Compounds	In water	In oil
DMS	0.3*	1.2*
DMDS	18.69+	2.5*
DMTS	0.23+	2.5*
DMQS	0.06+	ND*

* Kubickova & Grosch, 1997

+ Our results

Micro-organisms able to produce sulfur compounds

- > Geotrichum candidum
- > Hafnia alvei
- > Coryneform bacteria :
 - Brevibacterium linens, B. aurantiacum
 - Micrococcus sp.
- > Staphylococcus equorum and S. lentus
- Lactobacilli

Propriétés aromatiques de Camembert fabriqués avec P. camemberti avec et sans G. candidum (Molimard et al, 1996)

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TRIGLYCERIDES

Lipases

SATURATED FATTY ACIDS

UNSATURA TED FATTY ACIDS

HYDROXY ACIDS, OXO-ACIDS, HYDROPEROXY-ACIDS

Effect of fat on the sensory properties

- Texture factor (thickener, lubricant, emulsifier,...)
- Flavour support
- Flavour precursor

Lipid degradation

Step 1 lipolysis: It can be a limiting step (Goat cheese, Blue cheese)

Step 2 : fatty acid oxidation (usually not the limiting step)

Impact of Fat

Level of lipolysis in different cheeses

B- Lipolysis in blue Cheeses compared to other types of cheeses

Table 3 Typical concentration of free fatty acids (FFA) indifferent cheese varieties

Variety	FFA (mg kg ⁻¹)	Variety	FFA (mg kg ⁻¹)
Cabrales	33200	Gruvere	1500
Danablu	32600	Brie	1300
Roquefort	32400	Cheddar	1000
Parmesan	5000	Camembert	~ 700
Provolone	2100	Mozzarella	360

Specific fatty acids are goaty

- 4-ethyl- 2octenoic acid
- 4-Ethyl octanoic acid (threshold 6ppb against
 0.9 ppm for decanoic acid)
- 4-methyl octanoic acid (threshold 20 ppb against 3.4 ppm for the decanoic acid)

Free fatty acid changes during camembert cheese ripening

In soft cheeses volatil FFA increase global flavour intensity

BUT too many fatty acids gives soapy flavours

Beta- oxidation

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Oxydation of a pentadiene motive through a Lipoxigenase

Clivage of Hydroperoxy acids through an Hydroperoxide lyase

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This pathway gives mushroom, metalic, geranium like, vegetal like flavours

Spontaneous oxydation (rare in cheese) but mainly activated by moulds (*Penicillium camemberti* mainly)

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Effect of ripening temperature on the styrene production by *P. camemberti* on a model curd

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Leclercq Perlat et al, 2004

Plastic flavour tested with a panel of 20 trained judges (camemberts made with 4 strains of *Penicillium*, with and without strains of *G. candidum*)

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CONCLUSIONS

SURFACE

INSIDE

CONCLUSIONS

- Fungi and Bacteria are able to produce a large diversity of flavour compounds which will determine the cheese flavour notes perceived
- Understanding the origin of the flavour notes
 - Understanding may help to manage flavor defects
 - May help to choose the starters
- The parameters used in the technology (management of the relative humidity, temperature, ventilation) will not only change the growth of the different species of the microbial ecosystem but also the physiology of some of the micro-organisms

