

# Effect of feeding managements on the milk concentrations of short- and medium-chain fatty acids

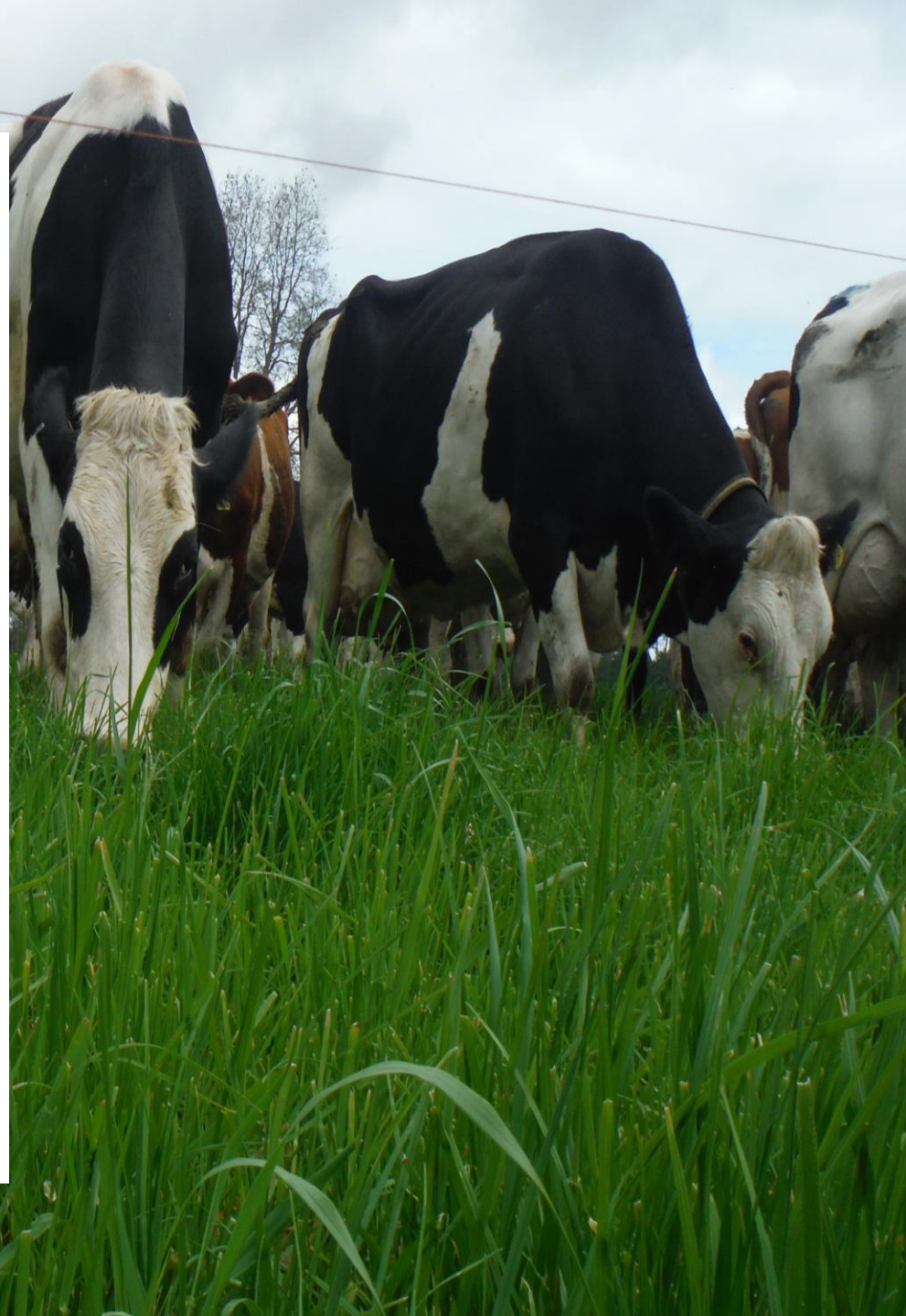
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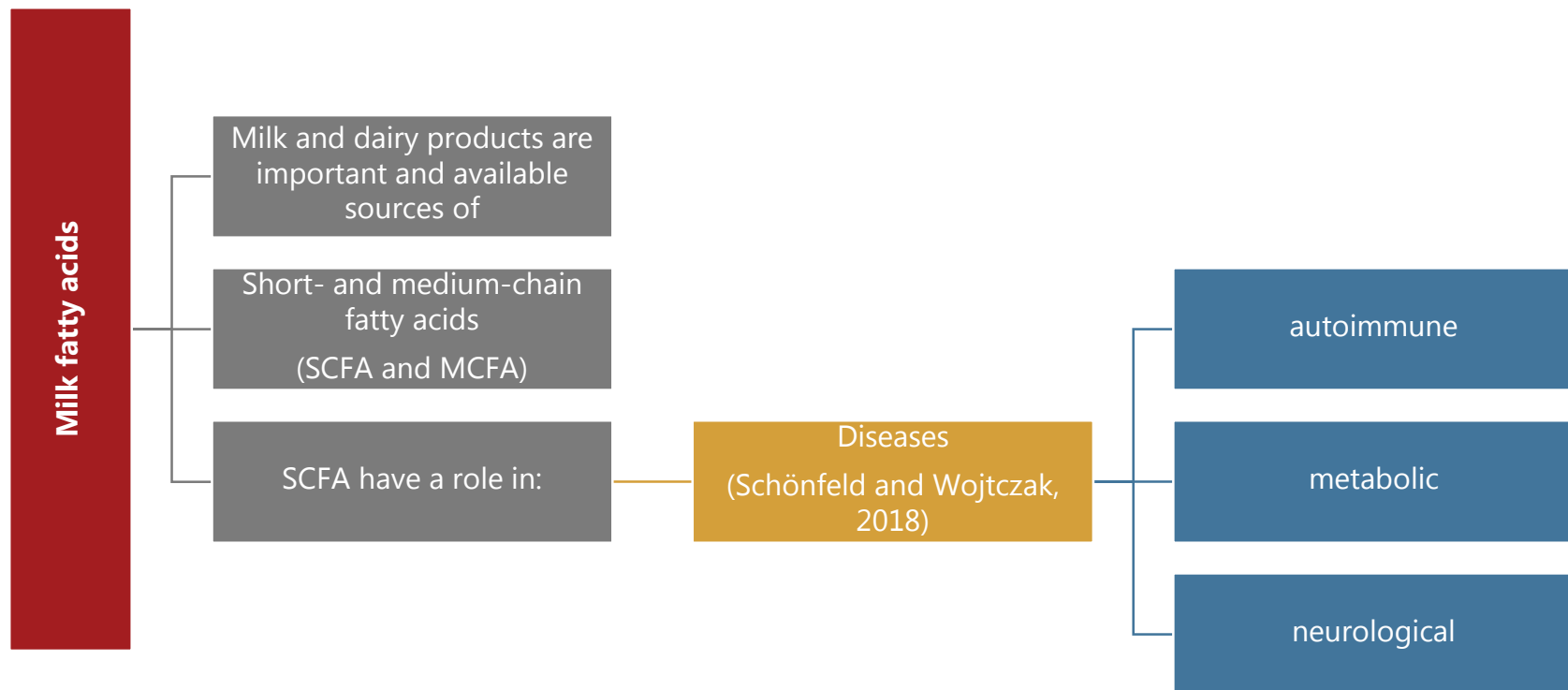
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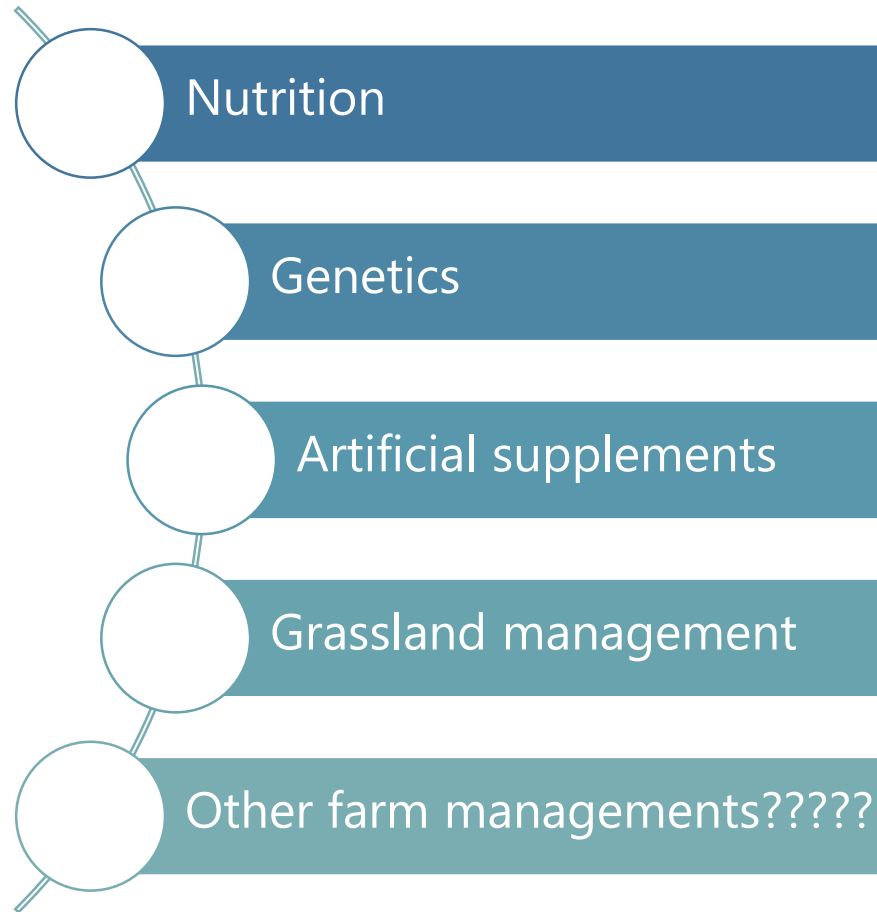
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# Background



# Background : Modulation of milk fat?



# Background



## In Denmark

- Hein et al. (2018)
- Differences in genetic correlations between:
  - Total fat and the monounsaturated FA, polyunsaturated FA, SCFA, and C16:0 contents between Danish Holstein and Jersey Holstein cows
- No correlations were made between the contents of milk SCFA and MCFA and the different **farm and feeding managements** from each herd



## Fatty acid data analysis

- Product authentication
  - Milk vs. non-dairy beverages (Vargas-Bello-Pérez et al., 2017)
  - Small-scale vs. large-scale cheese production (Vargas-Bello-Pérez et al., 2018)
- Can identify on-farm conditions that are affecting milk FA profiles (Mele et al., 2016)

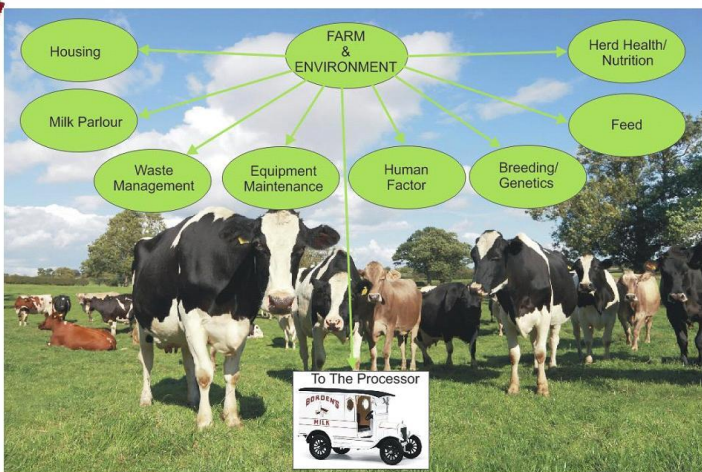
# Objective

- To identify the feeding managements (amount of dietary maize and grass silages) that are responsible for production of milk with high concentration of SCFA (4 to 6 carbons) and MCFA (from 8 to 14 carbons)

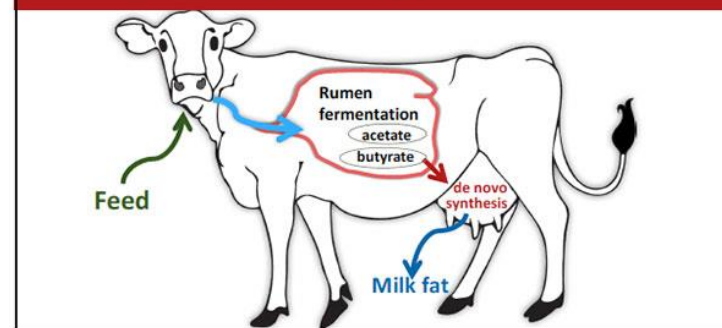


# Conceptual framework

## Testing at Farm Level



## De novo Fatty Acid Synthesis



bioactive components



**Thise Mejeri**





# Methodology

- Existing data:
  - 388 milk samples
  - 7 organic farms
  - 2 seasons
  - Milk fatty acid profiles
  - $Y = \text{Farm} + \text{season} + (\text{farm} \times \text{season}) + \text{Days in milk category} + \text{Fat percentage} + \text{Animal (random)} + \text{Error}$



Farm	Grass silage (% DM)	Maize silage (% DM)	DM intake (kg DM/d)	Somatic cell counts ( $\times 10^3/\text{ml}$ )
1	66	0	21	220
2	53	19	21	151
3	56	10	25	195
4	70	12	20	179
5	51	14	24	210
6	57	14	24	240
7	50	24	22	145

Average values from the previous 6 years

Mean, coefficient of variation (CV) of the mean, and standard deviation (SD) of the mean for the content of milk fat (g/100g), individual fatty acids, fatty acid groups and fatty acid indices (FA are given as proportions based on g/kg of FA; n = 388)

Traits	Mean	CV, %	SD	P5	P95
Milk fat	4.14	16.5	0.68	3.7	4.6
Saturated fatty acids					
C6:0	2.84	10.8	0.31	2.66	3.03
C8:0	1.58	13.4	0.21	1.43	1.73
C10:0	3.51	17.7	0.62	3.01	3.94
C12:0	3.89	18.8	0.73	3.34	4.41
C14:0	12.1	10.4	1.27	11.3	12.9
C15:0	1.14	15.4	0.18	1.01	1.25
C16:0	29.9	17.2	5.17	26.2	32.5
C17:0	0.62	13.9	0.09	0.56	0.67
C18:0	10.0	19.2	1.93	8.80	11.2
Monounsaturated fatty acids					
C14:1 cis-9	1.02	24.4	0.25	0.85	1.16
C16:1 cis-9	1.44	23.8	0.34	1.22	1.63
C18:1 trans11	1.70	36.2	0.62	1.33	1.95
C18:1 cis-9	19.6	17.1	3.38	17.3	21.5
Polyunsaturated fatty acids					
C18:2 cis-9, cis-12	2.17	27.7	0.60	1.65	2.56
C18:3 cis-9, cis-12, cis-15	0.90	28.2	0.25	0.71	1.05
C18:2 cis-9, trans-11	0.75	41.4	0.31	0.54	0.88
Fatty acid groups					
Others	6.74	12.9	0.86	6.20	7.15
$\Sigma$ Saturated fatty acids	71.2	6.51	4.63	68.0	74.5
$\Sigma$ Monounsaturated fatty acids	24.3	16.1	3.92	21.6	26.8
$\Sigma$ Polyunsaturated fatty acids	4.41	23.2	1.03	3.69	5.06
$\Sigma$ Short chain fatty acids (C6:0 - C10:0)	7.93	12.6	1.00	7.20	8.64
$\Sigma$ Medium chain fatty acids (C12:0 - C14:0)	16.1	11.2	1.80	14.7	17.3
$\Sigma$ Long chain fatty acids	36.5	15.7	5.77	32.7	40.2
$\Sigma$ Trans fatty acids	2.61	36.1	0.94	2.02	3.05

P5 = 5th percentile, P95 = 95th percentile.



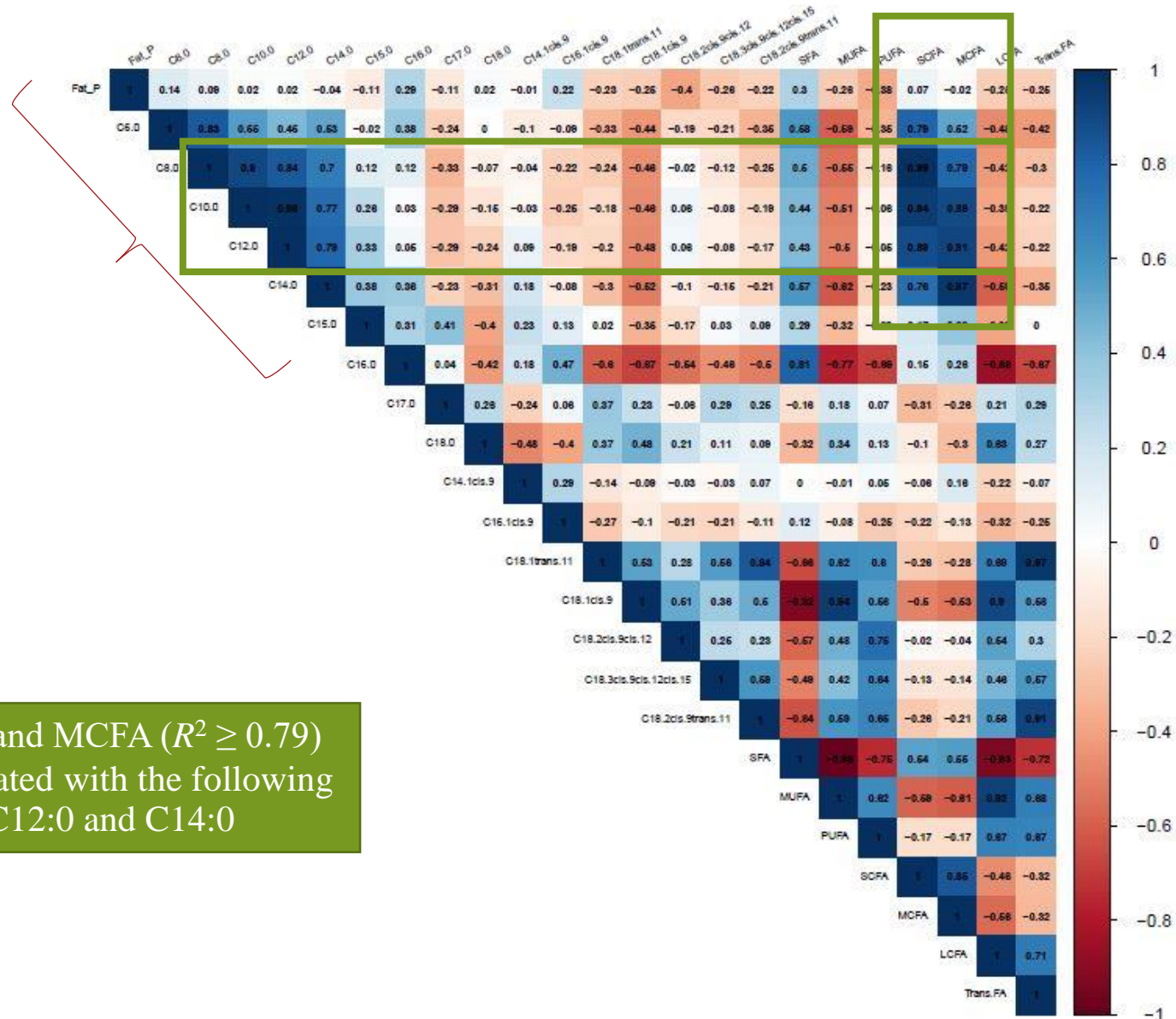
# Individual fatty acids, fatty acid groups and fatty acid indices from 7 organic farms used for milk sampling (n = 388)

Fatty acids (g/kg of FA)	Farm							SEM	Farm	Season	Fat
	1	2	3	4	5	6	7				
C6:0	2.81	2.78	2.80	2.85	2.93	2.89	2.81	0.02	NS	<0.001	<0.001
C8:0	1.58 <sup>ab</sup>	1.53 <sup>a</sup>	1.55 <sup>a</sup>	1.60 <sup>ab</sup>	1.68 <sup>b</sup>	1.59 <sup>ab</sup>	1.58 <sup>ab</sup>	0.04	<0.05	<0.05	<0.05
C10:0	3.53 <sup>ab</sup>	3.33 <sup>a</sup>	3.41 <sup>a</sup>	3.58 <sup>ab</sup>	3.84 <sup>b</sup>	3.45 <sup>ab</sup>	3.49 <sup>ab</sup>	0.11	<0.01	NS	NS
C12:0	3.89 <sup>ab</sup>	3.67 <sup>a</sup>	3.81 <sup>a</sup>	4.00 <sup>ab</sup>	4.26 <sup>b</sup>	3.79 <sup>ab</sup>	3.82 <sup>ab</sup>	0.13	<0.01	NS	NS
C14:0	11.77 <sup>a</sup>	11.74 <sup>a</sup>	12.24 <sup>ab</sup>	12.37 <sup>ab</sup>	12.71 <sup>b</sup>	12.19 <sup>ab</sup>	11.84 <sup>a</sup>	0.23	<0.01	NS	NS
C15:0	1.09 <sup>abc</sup>	1.17	1.14 <sup>abc</sup>	1.19 <sup>bc</sup>	1.20 <sup>c</sup>	1.08 <sup>ab</sup>	1.05 <sup>a</sup>	0.31	<.0001	NS	NS
C16:0	27.99	29.25	30.44	30.73	30.79	31.09	28.40	1.23	NS	NS	<0.001
C17:0	0.62 <sup>ab</sup>	0.64 <sup>b</sup>	0.62 <sup>ab</sup>	0.64 <sup>b</sup>	0.63 <sup>ab</sup>	0.61 <sup>ab</sup>	0.59 <sup>a</sup>	0.14	<0.05	NS	<0.05
C18:0	10.69	9.92	9.68	9.83	9.87	10.42	10.19	0.49	NS	NS	NS
C14:1 cis-9	1.00	1.00	1.05	1.05	0.99	1.05	0.97	0.055	NS	NS	NS
C16:1 cis-9	1.50 <sup>ab</sup>	1.45 <sup>ab</sup>	1.48 <sup>b</sup>	1.46 <sup>ab</sup>	1.38 <sup>ab</sup>	1.49 <sup>ab</sup>	1.30 <sup>a</sup>	0.059	<0.05	NS	<0.001
C18:1 trans11	1.73 <sup>ab</sup>	1.91 <sup>b</sup>	1.64 <sup>a</sup>	1.62 <sup>ab</sup>	1.52 <sup>a</sup>	1.66 <sup>ab</sup>	1.82 <sup>ab</sup>	0.12	<0.05	NS	<0.01
C18:1 cis-9	20.95 <sup>b</sup>	20.02 <sup>ab</sup>	20.13 <sup>ab</sup>	19.37 <sup>ab</sup>	18.17 <sup>a</sup>	20.22 <sup>ab</sup>	19.26 <sup>ab</sup>	0.63	<0.05	NS	<0.001
C18:2 cis-9, cis-12	2.24 <sup>ab</sup>	1.95 <sup>a</sup>	2.30 <sup>ab</sup>	2.35 <sup>b</sup>	2.17 <sup>ab</sup>	2.11 <sup>ab</sup>	2.09 <sup>ab</sup>	0.11	<0.01	<0.05	<0.001
C18:3 cis-9, cis-12, cis-15	0.91 <sup>ab</sup>	1.00 <sup>b</sup>	0.90 <sup>ab</sup>	0.87 <sup>ab</sup>	0.82 <sup>a</sup>	0.83 <sup>ab</sup>	0.93 <sup>ab</sup>	0.45	<0.05	NS	<0.001
C18:2 cis-9, trans-11	0.74 <sup>ab</sup>	0.87 <sup>b</sup>	0.76 <sup>ab</sup>	0.72 <sup>ab</sup>	0.66 <sup>a</sup>	0.70 <sup>ab</sup>	0.76 <sup>ab</sup>	0.58	<0.05	NS	<0.001
Others	6.81	6.71	6.77	6.74	6.69	6.94	6.52	0.15	NS	<0.001	NS
Saturated fatty acids	69.70 <sup>a</sup>	70.31 <sup>a</sup>	70.75 <sup>ab</sup>	71.67 <sup>ab</sup>	73.29 <sup>b</sup>	71.34 <sup>ab</sup>	71.04 <sup>ab</sup>	0.78	<0.05	NS	<0.001
Monounsaturated fatty acids	25.74 <sup>b</sup>	25.23 <sup>b</sup>	24.66 <sup>ab</sup>	23.83 <sup>ab</sup>	22.51 <sup>a</sup>	24.49 <sup>ab</sup>	24.54 <sup>ab</sup>	0.75	<0.05	NS	<0.001
Polyunsaturated fatty acids	4.55	4.45	4.58	4.50	4.20	4.16	4.42	0.19	NS	NS	<0.001
Short chain fatty acids (C6:0 - C10:0)	38.70 <sup>b</sup>	37.44 <sup>ab</sup>	36.47 <sup>ab</sup>	35.76 <sup>ab</sup>	34.40 <sup>a</sup>	36.50 <sup>ab</sup>	37.36 <sup>ab</sup>	1.12	<0.05	NS	<0.001
Medium chain fatty acids (C12:0 - C14:0)	7.92 <sup>ab</sup>	7.64 <sup>a</sup>	7.76 <sup>a</sup>	8.04 <sup>ab</sup>	8.44 <sup>b</sup>	7.93 <sup>ab</sup>	7.88 <sup>ab</sup>	0.19	<0.05	<0.05	NS
Long chain fatty acids	15.71 <sup>a</sup>	15.49 <sup>a</sup>	16.12 <sup>ab</sup>	16.44 <sup>ab</sup>	17.06 <sup>b</sup>	16.04 <sup>ab</sup>	15.72 <sup>a</sup>	0.35	<.001	NS	NS
Trans fatty acids	2.65 <sup>ab</sup>	2.98 <sup>b</sup>	2.55 <sup>ab</sup>	2.49 <sup>ab</sup>	2.33 <sup>a</sup>	2.50 <sup>ab</sup>	2.79 <sup>ab</sup>	0.18	<0.05	NS	<0.001
C14 index	7.93	8.03	7.47	7.67	7.34	8.10	7.67	0.41	NS	<0.05	NS
C16 index	4.33	4.46	4.96	4.72	4.55	4.65	4.44	0.19	NS	NS	NS
C18 index	63.86 <sup>a</sup>	65.71 <sup>ab</sup>	66.46 <sup>ab</sup>	67.10 <sup>b</sup>	66.47 <sup>ab</sup>	67.60 <sup>b</sup>	65.55 <sup>ab</sup>	0.81	<0.05	<0.001	NS
CLA index	29.60 <sup>ab</sup>	30.39 <sup>ab</sup>	29.43 <sup>a</sup>	32.78 <sup>b</sup>	30.74 <sup>ab</sup>	29.41 <sup>ab</sup>	30.61 <sup>ab</sup>	0.92	<0.05	<0.001	NS

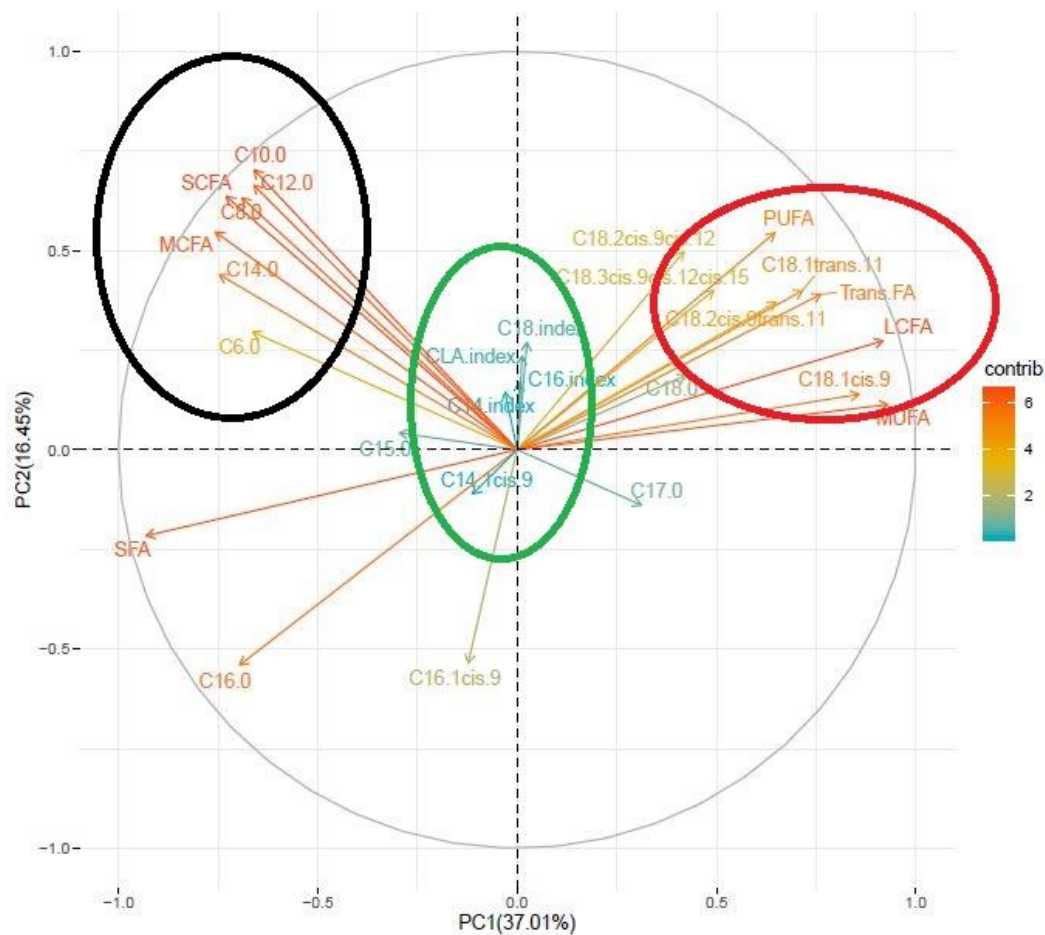
NS = Non significant

C14 index = C14:1/(C14:1 + C14:0) × 100 ; C16 index = C16:1/(C16:1 + C16:0) × 100 ; C18 index = C18:1 cis-9/(C18:1 cis-9 + C18:0) × 100 ; Conjugated linoleic acid (CLA) index = C18:2 cis-9,trans-11/(CLA cis-9,trans-11 + C18:1 trans-11) × 100.

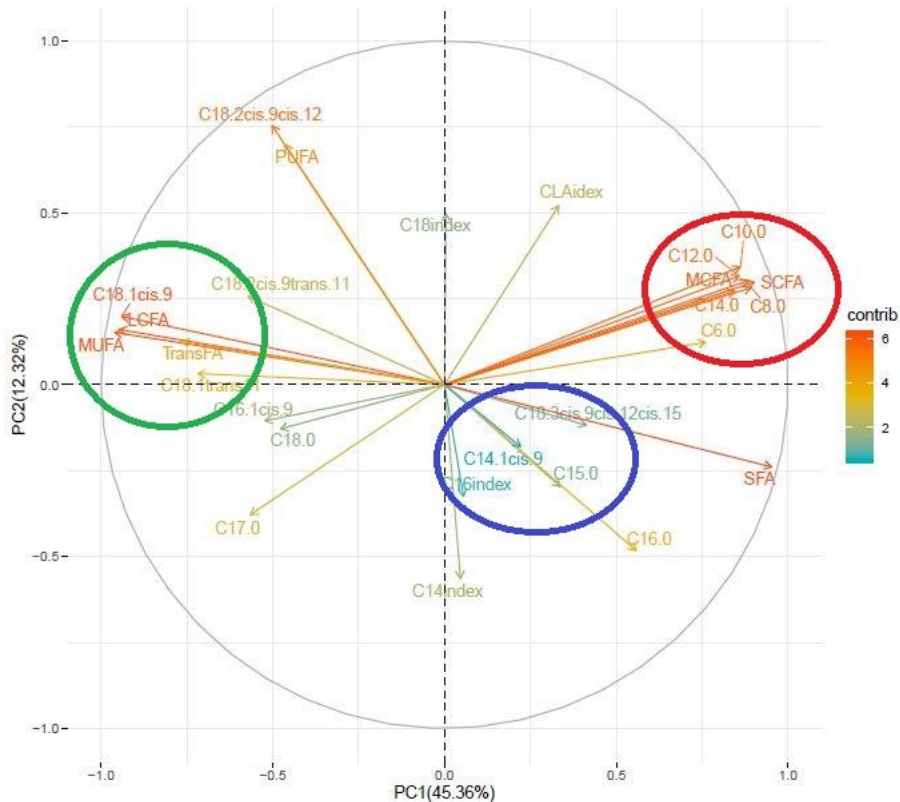
Pearson's correlation coefficient for all individual fatty acids, fatty acid groups and fatty acid indices from 7 organic farms used for milk sampling (n = 388)



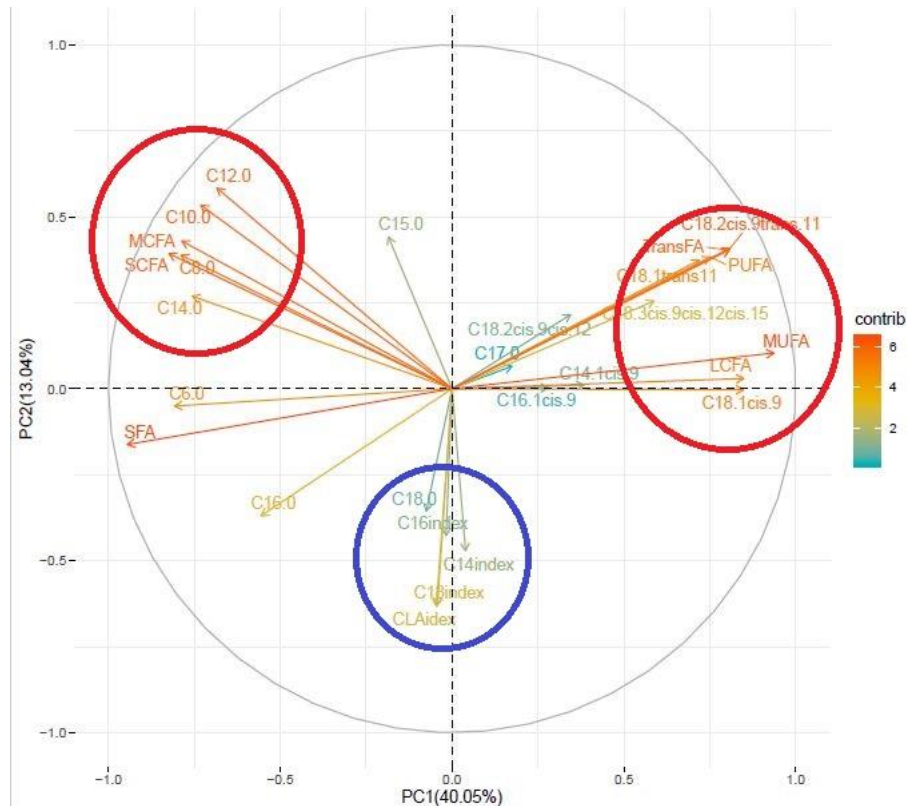
# Principal components (PC) related to the milk fatty acid profile from all farms



# Principal components (PC) related to the milk fatty acid profile from farm 1 and 5



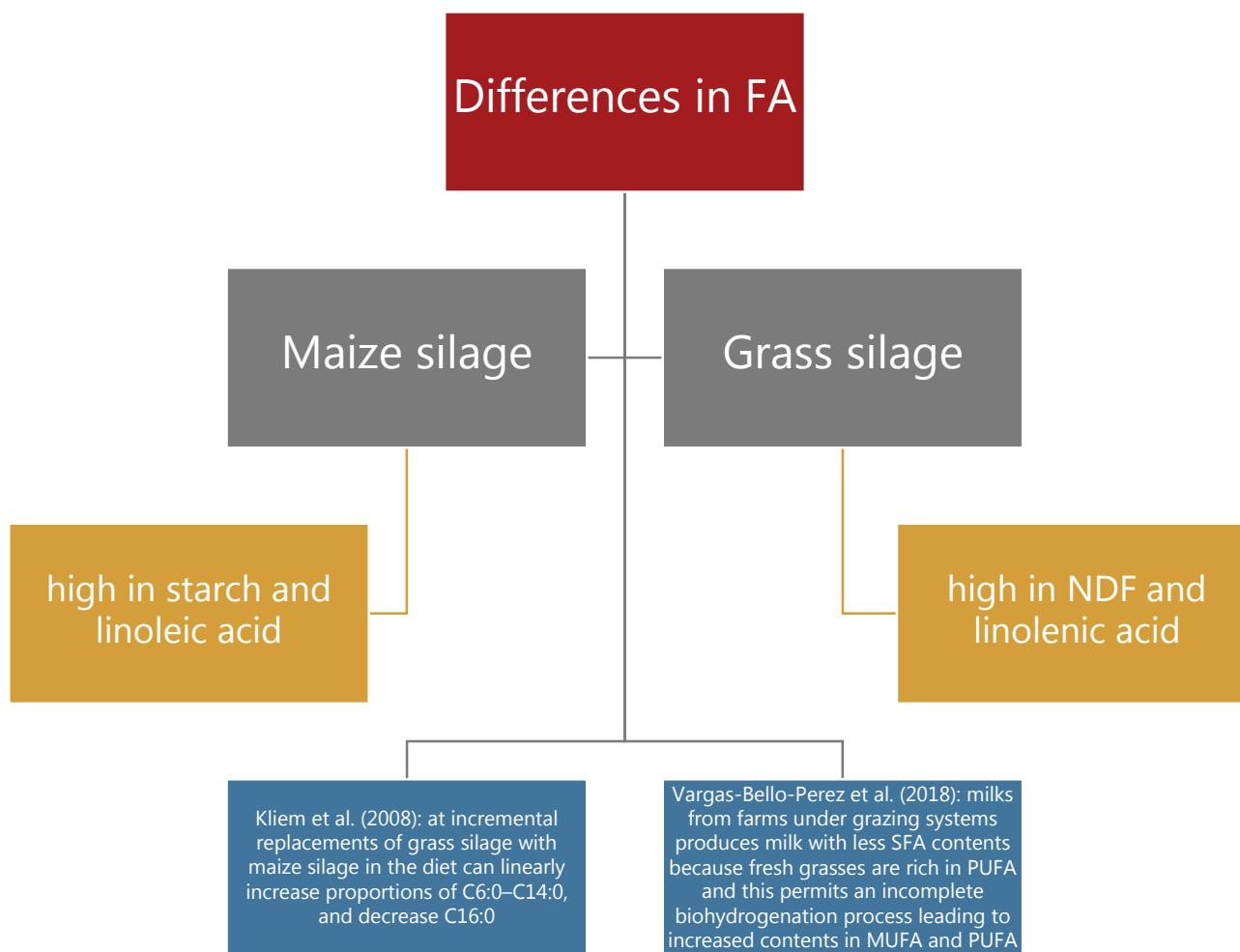
Farm 1



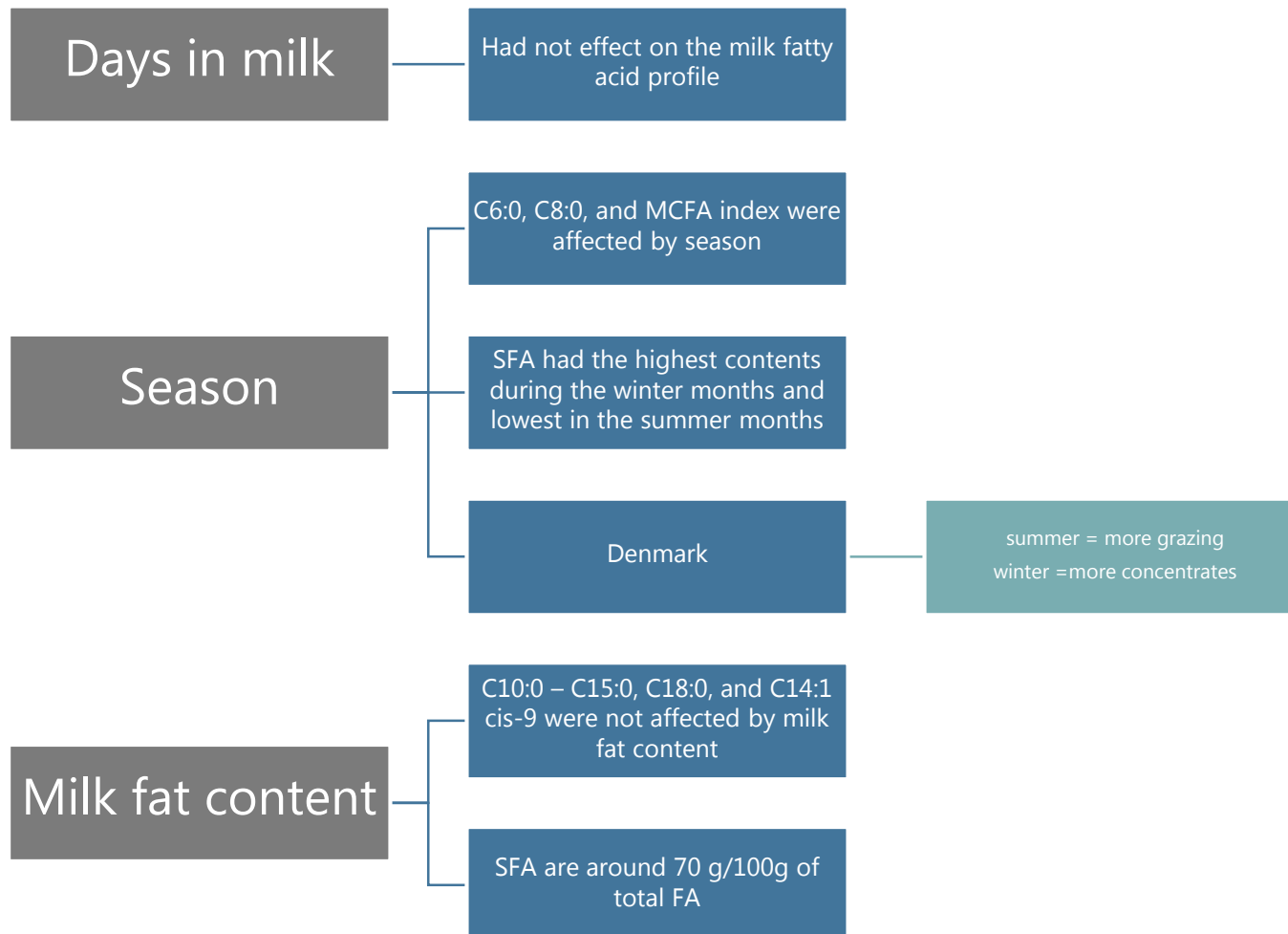
Farm 5

Farm	Grass silage (% DM)	Maize silage (% DM)
1	66	0
2	53	19
3	56	10
4	70	12
5	51	14
6	57	14
7	50	24

# Overall results



# Overall results



# General remarks

Farm	Grass silage (% DM)	Maize silage (% DM)
1	66	0
2	53	19
3	56	10
4	70	12
5	51	14
6	57	14
7	50	24



The level (% of dry matter) of dietary inclusion of grass (G) and/or maize silages (M) appeared to have an effect on SCFA (66G+no maize silage) and MCFA (51G+14M )

Increasing contents of SCFA and MCFA at the same time is very challenging due to the complexity of the cow's lipid metabolism



It will be desirable to record individual milk yields and milk compositions together with a detailed proximate analysis of diets, silages and individual feedstuffs with their corresponding FA profile



# Thank you!

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