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Original Citation:

Availability:

This version is available <http://hdl.handle.net/2318/1788940> since 2021-05-18T15:08:32Z

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Changes in goat milk fatty acids during abrupt transition from indoor to pasture diet

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ABSTRACT

Goal of this study was to evaluate the kinetics of goat milk fatty acids during abrupt transition from indoor to pasture-based diets. Twelve Valdostana goats in mid-lactation reared indoors and fed hay and concentrates for 40 days were abruptly brought outdoors on natural pasture and fed fresh grass *ad libitum*. Feed samples and individual milk samples were collected for fatty acids analysis on the last day of indoor feeding (day 0) and after 1, 2, 3, 4, 6, 9, 13, 18 and 23 days of fresh grass feeding. Milk fatty acid composition was significantly affected by sampling day. Significant changes already took place few days after transition. The most marked and consistent variations occurred at the expense of some unsaturated fatty acids. Total *trans*-octadecenoic and *trans*-octadecadienoic acids, conjugated linoleic acids (CLA) and omega-3 fatty acids constantly increased, reaching concentrations 4.0, 3.0, 3.9, and 2.2 times higher at the end of the trial than at its beginning, respectively. On the last sampling day the omega-6/omega-3 fatty

acids ratio was two times lower than its initial value. Considering individual fatty acids, the most consistent and remarkable increasing trends throughout the trial were observed for C18:1 *t*6-11, C18:1 *t*12-14+*c*6-8, C18:1 *c*14+*t*16, C18:2 *t*11*c*15, C18:2 *c*9*t*13+*t*8*c*12, CLA isomers *c*9*t*11+*t*7*c*9+*t*8*c*10 and *t*11*c*13+*c*9*c*11. Alpha-linolenic and eicosapentaenoic acids also increased significantly, but to a lesser extent. In view of the many beneficial biological effects that have been attributed to vaccenic acid (C18:1 *t*11), rumenic acid (C18:2 *c*9*t*11), and omega-3 fatty acids, results showed that, from a human health perspective, goat milk fatty acid composition consistently improved after transition from indoor to pasture feeding. Such improvements, mainly due to the high content of α -linolenic acid in pasture plants, were already significant after two or three days of fresh grass feeding. Further increases of beneficial fatty acids in milk fat were observed till about thirteen (vaccenic acid and CLA) or twenty-three (omega-3 fatty acids) days after transition. These results show that pasture can be considered a natural feeding strategy to quickly enhance the healthfulness of goat milk fat.

Keywords: goat milk, fatty acids, transition, grazing

INTRODUCTION

The positive effects of fresh grass-based diets on the fatty acid (FA) profile and the nutritional quality of dairy fat have been recognized broadly (Morand-Fehr et al., 2007). Such improvement is related to both increased milk concentrations of polyunsaturated FA (e.g., omega-3 FA and conjugated linoleic acid) which are known to exert many putative beneficial effects for human health (Barcelo-Coblijn and Murphy, 2009; Benjamin and Spener, 2009) and decreased concentrations of saturated FA (particularly lauric, myristic, and palmitic acids) able to raise risk factors for cardiovascular diseases (Ohlsson, 2010). Modifications and persistency of ruminant milk FA concentrations determined by the

transition from indoor to fresh grass feeding have been investigated in a limited number of studies in cows and sheep only. Kuzdzal-Savoie and Kuzdzal (1961) first reported a notable and fast increase in the amount of conjugated dienes and α -linolenic acid (C18:3 *c9c12c15*, ALA) in butter from dairy cows as an effect of either sudden or gradual change from a winter ration to a pasture-based diet. Similarly, Decaen and Ghadaki (1970) observed that a gradual change from a winter diet consisting of hay, silage, and concentrate to a zero-grazing system induced fast modifications in the FA composition of cow milk fat. Particularly, they observed a decrease in the proportion of short- (SCFA) and medium-chain (MCFA) fatty acids and contemporarily an increase in long-chain fatty acids (LCFA). More recently, Khanal et al. (2008) reported stearic acid (C18:0), oleic acid (C18:1 *c9*), vaccenic acid (C18:1 *t11*, VA), conjugated linoleic acid (CLA) and ALA to increase, and most of the SCFA and MCFA to decrease, during abrupt transition of dairy cows from a total mixed ration to a full-grazing diet. In their study, the majority of FA changed daily before stabilizing around 22-23 days after transition. Coppa et al. (2011) also investigated the kinetics of milk FA in dairy cows during rapid or progressive transition from hay- to alpine pasture-based diets. These authors showed that many FA (almost all saturated ones, linoleic acid – C18:2 *c9c12*, LA - and ALA) became stable after five days in both transitions, while both VA and the most abundant among CLA isomers (C18:2 *c9t11*, rumenic acid, RA) reached maximum and stable concentrations about two weeks after the maximum fresh herbage intake.

In sheep, Biondi et al. (2008) reported that the major changes in milk FA occurred during the first three days following an abrupt transition from indoor to pasture diet, being predominantly attributable to unsaturated fatty acids (UFA). Particularly, they observed significant increases of VA, RA, and ALA, as well as a notable decrease in the concentration of LA and in the omega-6/omega-3 fatty acids ratio.

To the best of our knowledge, no information is currently available on the rate of change

and persistency in goat milk FA during transition from indoor to fresh grass feeding. The objective of this work was, therefore, to examine the kinetics of responses of goat milk FA during abrupt change from a winter diet based on hay and concentrates to a full-grazing diet on natural pasture.

MATERIALS AND METHODS

Animals feeding and management

The experiment was carried out in a farm located in North-Western Italy (latitude: 45°02'51'' N; longitude: 07°19'10'' E; altitude: 643 m a.s.l.) and breeding a flock of 40 Valdostana goats. The experimental period covered a total of 24 days, from March 29 to April 21, 2011.

Twelve goats were used in the experiment (days in milk at the beginning of the trial: 126 ± 6 ; number of lactation: 2.2 ± 0.4), with an average body condition score of 3.0 ± 0.5 (Hervieu and Morand-Fehr, 1999). The goats were maintained indoor for 40 days (February 18 to March 29, pre-experimental period and first day of experimental period) when they were fed exclusively with $1.5 \text{ kg head}^{-1} \text{ day}^{-1}$ of hay (first cut) and $0.4 \text{ kg head}^{-1} \text{ day}^{-1}$ of commercial concentrate containing maize and wheat bran. Means and standard deviations of milk yield and milk fat, protein and lactose percentages at the beginning of the pre-experimental period were equal to $1.17 \pm 0.306 \text{ kg head}^{-1} \text{ day}^{-1}$, $42.2 \pm 0.52 \text{ g kg}^{-1}$, $31.3 \pm 0.22 \text{ g kg}^{-1}$ and $43.4 \pm 0.26 \text{ g kg}^{-1}$, respectively. From midday of March 30 (day 1) the goats were abruptly brought outdoor and exclusively fed with fresh grass from natural pasture (main species: *Lolium perenne* L., *Trifolium pratense* L., and *Poa* spp.) *ad libitum*. The pasture area was flat and adjacent to the farm. The goats were manually milked indoors twice a day (at 6.00 h and 18.00 h). They were allowed to graze during the milking interval while they were maintained indoors during the night. The goats had free access to water and mineralized salt blocks during both pasture and indoor housing.

Sampling procedure and laboratory analyses of milk

Individual milk yield was recorded on a daily basis; individual milk samples were collected at the afternoon milkings ten times during the trial: on the last day of indoor feeding (day 0) and on days 1, 2, 3, 4, 6, 9, 13, 18 and 23 of pasture feeding. Two aliquots of each sample were taken for laboratory analysis. One aliquot (50 mL) was immediately stored at 4°C with a preservative and subsequently transported to the laboratory for the analysis of fat, protein, and lactose (MilkoScan FT 6000, Foss Electric, Hillerød, Denmark). The second aliquot (150 mL) was frozen at -20°C and successively analyzed for FA composition as previously reported by Renna et al. (2012). Peaks were identified by injecting pure FAME standards (Sigma-Aldrich, Milano, Italy; Matreya Inc., Pleasant Gap, PA, USA and Restek Corporation, Bellefonte, PA, USA) and by comparison with the chromatogram published by Collomb and Bühler (2000). Quantification was assessed by using nonanoic acid as internal standard. The results are expressed as absolute values as g kg⁻¹ fat.

Sampling procedure and laboratory analyses of feed

The hay and concentrate fed to the goats during the indoor period were collected on day 0. For herbage sampling, the pasture area was subdivided in plots of 200 m² each. A sub-area of 200 cm² was considered for sampling at every intersection among plots. Pasture samples were collected following the same time schedule (days 1, 2, 3, 4, 6, 9, 13, 18, and 23) as for the collection of milk samples. The grazing behavior of the goats was observed at each sampling date. After closely observation, hand-plucked forage samples, simulating plants parts consumed by the goats, were collected from pasture. Two aliquots of each herbage sample were transported to the laboratory in a portable refrigerator at 4°C and then frozen at -80°C. Before the chemical analysis, the first aliquot of each herbage sample was dried at 40°C for 24 h; hay, concentrate and dried herbage samples were then ground with a cutting mill to pass a 1-mm screen sieve (Pulverisette 15 - Fritsch GmbH,

Idar-Oberstein, Germany). Samples were analyzed for dry matter (DM), crude protein (CP), ether extract (EE), ash, neutral detergent fiber (NDF), and acid detergent fiber (ADF) according to AOAC procedures (2000). The second aliquot of herbage samples was freeze-dried (Edwards MF 1000, Milano, Italy) and ground. This aliquot as well as both grounded hay and concentrate were used for the assessment of the FA composition as reported by Alves et al. (2008). FAME were separated and quantified by using the same analytical instruments and temperature program described for the analysis of milk samples. The injection volume was 0.5 µL. Peaks were identified by injecting pure FAME standards (Restek Corporation, Bellefonte, PA, USA) and by comparison with the chromatogram published by Alves et al. (2008). Quantification was assessed by using heptadecanoic acid as internal standard. The results are expressed as absolute values as mg 100g⁻¹ DM.

Statistical analysis

The goat was considered as the experimental unit. Changes in milk yield, main constituents and FA were analyzed using the PROC MIXED procedure of SAS (2006) according to the following model:

$$Y_{ijk} = \mu + D_i + G_j + \varepsilon_{ijk},$$

where Y_{ijk} = mean of response variable, μ = population mean, D_i = effect of day, G_j = random effect of goat, and ε_{ijk} = experimental error. Pairwise multiple comparisons were assessed by using the PDIFF option in SAS. Significance was declared at $P \leq 0.05$.

RESULTS

Characteristics of the feedstuffs

The chemical and FA compositions of hay, commercial concentrate, and fresh grass are presented in Table 1.

In fresh grass samples, DM and fiber contents increased while CP decreased from day 1 to day 23, following the advance of plants' age.

Hay, concentrate and fresh grass strongly differed in their FA amounts and compositions. The concentrate showed 1.8-fold and 4.0-fold higher total FA amounts when compared to fresh grass and hay, respectively. On average, the fresh grass had a total FA concentration approximately doubled than that detected in hay. Alpha-linolenic acid was the most abundant FA in fresh grass, accounting on average for the 62% of total FA. Palmitic and linoleic acids showed similar concentrations, accounting for the 16 and 14% of total FA, respectively. Alpha-linolenic, palmitic and linoleic acids were also the most representative FA of hay, together accounting for about the 83% of total FA. However, the relative contribution of ALA was lower (33% of total FA) in hay if compared to fresh grass. Linoleic acid was, instead, the prevailing FA in the concentrate, accounting for about the 57% of total FA. It was followed by oleic and palmitic acids, both representing about the 18-19% of total detected FA. ALA represented only about the 2% of total FA in the concentrate. The total FA content and both concentration and proportion of ALA decreased from the beginning till the end of the trial (data not shown).

Milk yield and gross composition

Milk yield and gross composition were significantly affected by sampling day ($P \leq 0.001$; Table 2). Milk yield maintained similar values as that observed the last day of stall feeding till day 9 (showing the absolute minimum levels on days 2 and 3 after transition), and then slightly increased reaching the absolute highest value on day 23. Fat production started to increase significantly the second day after transition and continued to increase till the end of the experiment. The fat percentage of milk significantly increased the second day after transition as well, but no significant variations were observed for this parameter in the following sampling days. Protein percentage, protein production and lactose production tended to increase as well, while no clear increasing or decreasing trend was observed for

the percentage of lactose in milk.

Milk fatty acid composition

Results of groups of FA and individual FA are reported in Tables 3 and 4, respectively.

The FA profile of milk fat was affected by days on pasture to a great extent. Only two FA (C18:1 *t*5 and C18:2 *t*10*c*12) were not significantly influenced by sampling day.

Total SCFA and MCFA started to decrease significantly the second day after transition (day 2) and they continued to decrease till day 4 (1.2 and 1.5 times lower values on day 4 relative to values on day 0, respectively). Subsequently SCFA rose again, reaching at the

end of the trial concentrations that did not significantly differ from the concentration observed the last day of stall feeding. A similar trend also occurred for MCFA, but their concentration was still significantly lower at the end of the trial than at its beginning.

Conversely, LCFA rapidly and markedly increased the second day after diet change and then significantly declined from day 6 till the end of the trial. Starting from day 9 milk LCFA concentrations were not statistically different from the value recorded on day 0.

Total saturated fatty acids (SFA) underwent a conspicuous drop from day 1 to day 4 subsequent to transition. From day 6 SFA increased again, but their levels remained generally lower if compared to day 0. Total branched-chain fatty acids (BCFA) declined as well from day 1 to day 4. From day 4 to day 9 they remained quite constant and finally slightly increased, so that in the period from day 13 to day 23 their levels did not statistically differ from the value observed on day 0.

Temporal changes in the concentration of total monounsaturated fatty acids (MUFA) showed a sharp increase from day 1 to days 2-3. MUFA levels then significantly declined till the end of the trial when they reached their absolute minimum levels. The trend

observed for total polyunsaturated fatty acids (PUFA) was much clearer, as this group of FA continuously rose until the sixth day after transition, thereafter remaining constant.

PUFA were 1.7 times higher on day 23 than on day 0.

208 On day 3 total *trans*-octadecenoic (Σ C18:1 *trans*) and *trans*-octadecadienoic (Σ C18:2
 209 *trans*) acids showed values already 1.8 and 1.6 times higher than those observed on day 0,
 210 respectively. These groups of FA continued to increase markedly till the end of the
 211 experiment. The highest values (about four and three times higher than the values
 212 recorded the last day of stall feeding, respectively) were observed the last two (Σ C18:1
 213 *trans*) or three (Σ C18:2 *trans*) sampling days.

214 Under the chromatographic conditions applied in this trial, the most abundant among
 215 *trans*-octadecenoic acids in milk fat (vaccenic acid - C18:1 *t*11) coeluted with other C18:1
 216 *trans*-isomers (C18:1 *t*6-10). This sum (C18:1 *t*6-11) as well as the values recorded for
 217 other detected *trans*-octadecenoic isomers, particularly C18:1 *t*12-14 (which coeluted with
 218 C18:1 *c*6-8 isomers) and C18:1 *t*16 (which coeluted with C18:1 *c*14), started to increase
 219 significantly three days after turning out to pasture. C18:1 *t*6-11 isomers continued to
 220 increase till day 13 and then maintained constant values (approximately four times higher
 221 than the value observed on day 0). C18:1 *t*12-14+*c*6-8 and C18:1 *c*14+*t*16 isomers reached
 222 their absolute highest concentrations on days 18 and 23, with values 2.8 and 2.1 times
 223 higher than those recorded the last day of indoor feeding, respectively.

224 Considering individual CLA isomers, in the applied chromatographic conditions C18:2
 225 *c*9*t*11, C18:2 *t*7*c*9 and C18:2 *t*8*c*10 coeluted in a single peak in the chromatogram. Their
 226 sum represented on average the 97% of total CLA. They started to increase the third day
 227 after transition and continued to increase significantly until day 13, thereafter maintaining
 228 constant concentrations. The raise was conspicuous: values observed the last three
 229 sampling days were up to 3.6 times higher than the value observed on day 0. The sum of
 230 these three CLA isomers showed high individual variability among the goats involved in
 231 the trial. In fact, it varied between 3.33 and 7.77 g kg⁻¹ fat at the beginning of the trial (day
 232 0) and between 11.79 and 22.96 g kg⁻¹ fat the last sampling day. About two-fold variation
 233 was constantly maintained among individual goats all along the experiment. However, the

ranking of individual goats for CLA content was not stable throughout the trial. The sums
 CLA $c9t11+t7c9+t8c10$ and C18:1 $t6-11$ were strongly correlated each other ($r=0.97$;
 $P\leq 0.001$).
 CLA isomers $t11c13$ and $c9c11$ coeluted in the chromatogram. The third day after
 transition from the winter diet to full grazing their sum was already about nine times
 higher than the initial value. Their sum continued to increase until day 13 when it reached
 the highest absolute concentration. The contribution of these isomers to the total CLA
 content of milk varied from 0.42% on day 0 to 2.7% on day 23. Sampling day
 significantly affected the concentration of CLA isomer $t9t11$ as well. Its lowest value was
 observed the last day of stall feeding. From day 1 to day 23 its concentration remained
 quite constant, with the exception of days 13 and 18 when the highest values were reached
 (about 3 times higher than the value observed on day 0).
 Linoleic acid significantly increased until the third day after transition to pasture feeding.
 Then it decreased, reaching concentrations up to 1.5 fold lower than those observed on
 day 0.
 Total omega-3 FA as well as ALA (which is the most abundant detected FA in this group)
 significantly and constantly increased from the last day of stall feeding to the last
 sampling day. At the end of the trial total omega-3 FA and ALA concentrations were
 about two-fold higher than values recorded on day 0. Considering other omega-3 FA, a
 clear increasing trend was also observed for C18:2 $t11c15$. A first significant raise
 (approximately doubled values with respect to concentrations detected the last day of stall
 feeding) was observed the third day after variation of the diet. Then it continued to rise
 constantly till the last sampling day, doubling again its concentrations so that on day 23 it
 was about four times higher than on day 0. A significant and positive correlation was
 found between this octadecadienoic acid and the sum of CLA isomers $t11c13$ and $c9c11$
 ($r=0.75$; $P\leq 0.001$). Among long-chain omega-3 FA, eicosapentaenoic acid (C20:5

c5c8c11c14c17, EPA) followed a similar trend as that observed for ALA and C18:2 *t11c15*. However, a significant increase was detected only after thirteen days after transition. Its highest concentrations were observed the last two sampling days (with values that almost doubled with respect to those observed at the beginning of the trial). Conversely, no clear positive trend was observed for docosapentaenoic acid (C22:5 *c7c10c13c16c19*, DPA).

The omega-6/omega-3 FA ratio started to decline the third day after switching from stall to pasture feeding. This ratio continued to decrease significantly throughout the trial up to day 13. The lowest absolute values were observed the last sampling day, when it was two times lower than its initial value.

Concerning $\Delta 9$ -desaturase activity (estimated as the ratios of C16:1 *c9* to C16:0 – **DI₁₆** – and C18:1 *c9* to C18:0 – **DI₁₈**), a significant increase was observed until day 4 after transition ($P \leq 0.001$). However, from day 6 these ratios significantly decreased again reaching the absolute lowest values at the end of the trial.

DISCUSSION

Milk yield and gross composition

The observed slight and progressive raise in milk yield and protein after transition from indoor to pasture feeding could be related to increased ingested energy as reported to occur in case of turning out, particularly when fresh grass is at an early growth stage. Similarly, an early phenological phase of pasture plants resulted in high milk fat content due to the increased intake of highly digestible fiber (Morand-Fehr et al., 2007). Besides the vegetative stage of pasture plants, the diet fed during the indoor period can significantly affect milk main constituents after turning out to pasture. If compared to maize silage, both hay or grass silage used as winter rations were found to determine an increase in milk fat content (Hoden et al., 1985). Such findings seem to be confirmed in

the current trial as milk fat significantly increased after diet change from a hay-concentrate based diet to a pasture based diet with fresh grass at early vegetative stage.

Milk fatty acid composition

The main changes in goat milk FA during lactation have been shown to occur in early lactation and have been mainly attributed to lipid mobilization as the consequence of a negative energy balance phase for the animals. A relatively stable FA pattern is instead generally observed in mid and late lactation (Ataşoğlu et al., 2009; Chilliard et al., 2003). For this reason, a confounding effect due to lactation stage on milk FA can be reasonably excluded in the current study and the observed variations are likely to be attributable to feeding aspects only.

Switching from a winter indoor diet (on average 70% hay and 30% concentrate) to a full grazing diet resulted in a higher availability of FA (particularly polyunsaturated) to be used for the synthesis of milk fat. It is known that the haymaking process notably reduces the total FA and the ALA concentrations in forages (Kalač and Samková, 2010). Consequently the availability of FA, and above all of ALA, increased switching the goats from a prevalent hay-based diet to a fresh grass-based diet. An increased supply of LCFA (especially with a high level of unsaturation) and some of their biohydrogenation intermediate products (e.g., C18:1 *trans* and C18:2 *trans* isomers) have been shown to inhibit the *de novo* synthesis of C8:0 to C16:0 FA within the bovine mammary gland, by exerting direct and/or indirect effects on the lipogenic enzymes acetyl-CoA carboxylase (ACC) and fatty acid synthase (FAS) and by reducing acetate and 3-hydroxybutyrate bioavailability for mammary lipogenesis (Chilliard and Ferlay, 2004). Nonetheless, milk fat synthesis and FA responses to dietary PUFA supplies are known to vary considerably among ruminant species (Chilliard et al., 2007). Mammary ACC and FAS mRNA abundance and/or activity have been reported to be much less affected by dietary PUFA in dairy goats than cows (Bernard et al., 2009). In the current study, the results we obtained

312 concerning SCFA and MCFA suggest a valuable inhibition of *de novo* synthesis in the
313 goat mammary gland by switching from an indoor to a grazing diet. Such inhibitory effect
314 was evident just the day after the diet change and was quite remarkable as these FA
315 decreased respectively by about 20 and 30% from day 0 to day 4. Since caproic and
316 caprylic acids are partly synthesized by metabolic pathways that are independent of ACC
317 (Chilliard and Ferlay, 2004), our results seem to confirm the existence of factors other
318 than altered mammary gene expression or lipogenic enzymes (particularly ACC) activities
319 able to exert an inhibitory effect of *de novo* synthesis of SCFA and MCFA in dairy goats.
320 The concentrations of the above-mentioned FA (particularly caproic, caprylic, and capric
321 acids) did not stabilize but significantly increased again beginning from day 6. Such
322 results suggest a possible temporal adaptation in mammary metabolism, which has already
323 been observed to occur in dairy cows even if at slower rate (Ferlay et al., 2006). In
324 addition, the observed subsequent increase in *de novo* synthesized FA could be also
325 related to changes in the physical characteristics of the sward grazed by the goats due to
326 maturation of plants with consequent notable changes in their FA concentrations and
327 proportions (particularly reductions in the total FA concentrations and in the proportion
328 and concentration of ALA) (Boufaïed et al., 2003; Cabiddu et al., 2009). Similar
329 proportions in the reduction of total FA and ALA in different grassland species with the
330 advance of the phenological phase (in a comparable period of time) have been also
331 reported by Wyss and Collomb (2010). Further investigations will be needed to better
332 elucidate both the mechanisms involved in the inhibition of *de novo* synthesis of SCFA
333 and MCFA and the persistency of such an effect in lactating goats.

334 Concerning LCFA, the trend observed for stearic acid could be ascribed to a decreasing
335 UFA content in pasture plants during the grazing period. The decrease in oleic acid seems
336 to be at least partly related to the lower availability of stearic acid. In fact, it is known that
337 more than the 50% of oleic acid is formed within the mammary gland by the activity of

338 $\Delta 9$ -destaurase on stearic acid. Moreover, the decreased $\Delta 9$ -desaturase activity (estimated
 339 by DI_{18} and probably due to a higher availability of *trans* FA (Chilliard et al., 2007)) also
 340 contributed to the decrease in the concentration of oleic acid in milk.

341 As previously reported for dairy ewes (Biondi et al., 2008), the most remarkable and
 342 consistent variations in goat milk FA concentrations after transition from a winter indoor
 343 to a pasture diet occurred at the expense of some long chain unsaturated FA and have to be
 344 mainly related to a different FA supply from the ingested feeds. The observed increasing
 345 concentrations of all *trans*-octadecenoic isomers (with the exception of C18:1 *t5*), of many
 346 non-conjugated *trans*-octadecadienoic isomers (particularly C18:2 *t11c15*, C18:2
 347 *c9t13+t8c12*, C18:2 *c9t12*, and C18:2 *ttNMID+t9t12*) and CLA isomers
 348 *c9t11+t7c9+t8c10*, *t11c13+c9c11*, and *t9t11* after the diet change are the consequence of
 349 the high content of ALA in pasture plants. In fact, ALA is well known to undergo within
 350 the rumen an intense and complex biohydrogenation process carried out by the anaerobic
 351 microbial flora. Many octadecatrienoic isomers (C18:3 *c9t11c15* and C18:3 *c9t13c15*
 352 among others) have been shown to be formed during the initial step of ruminal
 353 biohydrogenation of ALA. These trienes are subsequently hydrogenated to a multitude of
 354 non conjugated and conjugated dienes, which are in turn mainly hydrogenated to
 355 monoenoic FA. C18:2 *t11c15* is thought to be the major non conjugated octadecadienoic
 356 isomer deriving from the biohydrogenation of ALA (hydrogenation of C18:3 *c9t11c15*).
 357 Other non conjugated C18:2 *trans*-isomers are expected to be formed similarly. For
 358 example, the C18:2 *c9t13* isomer probably derives from reduction of the *c15* double bond
 359 in C18:3 *c9t13c15* (Lee and Jenkins, 2011). The $\Delta 9,12$ C18:2 isomers are thought to be
 360 mainly formed during the biohydrogenation of LA, even if a putative metabolic pathway
 361 for their formation from ALA has been also suggested (Chilliard et al., 2007). In the
 362 current trial, the observed increased concentrations of C18:2 *ttNMID+t9t12* and C18:2
 363 *c9t12* after diet change suggest their partly formation from dietary ALA. On the contrary,

364 since C18:2 *t9c12* tended to decrease after turning out to pasture, we could hypothesize
 365 that ALA biohydrogenation would not be an important metabolic way for the ruminal
 366 formation of this diene.

367 Concerning conjugated linoleic isomers, the increase in the sum of C18:2
 368 *c9t11+t7c9+t8c10* after turning out to pasture is presumably attributable to rumenic acid,
 369 the main CLA isomer in ruminant milk fat. In fact, the other two isomers (*t7c9* and *t8c10*)
 370 were previously reported to correlate positively and significantly with dietary oleic and
 371 linoleic acids, respectively, while no significant correlation was found with dietary ALA
 372 (Collomb et al., 2004). The increase in rumenic acid is due to similar increased
 373 concentrations of C18:1 *t6-11*, among which the precursor (C18:1 *t11*) for CLA *c9t11 de*
 374 *novo* synthesis within the mammary gland belongs. C18:1 *t11* is an intermediate of the
 375 biohydrogenation processes and is formed from both dietary LA and ALA. The higher
 376 supply of ALA from pasture if compared to the indoor diet suggests a higher formation of
 377 vaccenic acid in the rumen as well as a higher absorption into the bloodstream and
 378 consequently a higher availability for desaturation mediated by the $\Delta 9$ -desaturase enzyme
 379 within the mammary gland.

380 Referring to minor CLA isomers, as only very low amounts of CLA *c9c11* are usually
 381 present in dairy fat (Ferlay et al., 2008), the sum CLA *t11c13+c9c11* can be almost
 382 completely attributed to the *t11c13* isomer. Kraft et al. (2003) first hypothesized that CLA
 383 *t11c13* could be formed within the rumen at the third step of the biohydrogenation of
 384 dietary ALA, by means of an isomerization at the expense of C18:2 *t11c15*. Our results
 385 corroborate the hypothesis by Kraft et al. (2003) since both C18:2 *t11c15* and CLA *t11c13*
 386 significantly and rapidly increased after transition and they were significantly and
 387 positively correlated each other. However, the rapid increase we observed in the sum of
 388 CLA isomers *t11c13* and *c9c11* could be attributed to the latter isomer as well. In fact,
 389 more than 50% of CLA *c9c11* has also been recently reported to derive directly from the

390 biohydrogenation of dietary ALA (Lee and Jenkins, 2011). The same authors also reported
 391 that CLA *t9t11* partly derives from ALA, which could explain the observed increasing
 392 trend of this CLA isomer in goat milk fat after transition from indoor to pasture diet.
 393 As usually occurs with high-forage diets, CLA *t10c12* was detected only in traces
 394 (concentrations ≤ 0.01 g kg⁻¹ fat) and was not significantly affected by the feeding change.
 395 Besides vaccenic acid, other *trans*-octadecenoic fatty acids are known to be formed by
 396 means of various isomerizations occurring during different steps of the biohydrogenation
 397 of dietary UFA. In particular, the observed increase in the concentration of C18:1 *t12-*
 398 *14+c6-8* could be attributed to C18:1 *t13* and *t14* isomers, which were found to be formed
 399 during the biohydrogenation of both C18:2 *c9t13* and CLA $\Delta^{11,13}$ (Chilliard et al., 2007).
 400 The increased concentrations of ALA and EPA (the latter formed by means of ALA
 401 desaturation and elongation processes (Barcelo-Coblijn and Murphy, 2009)) in milk fat
 402 have to be related as well to the higher ALA supply from fresh grass. Their concentrations
 403 in milk remained however quite low since the extent of the raise was less pronounced if
 404 compared to those observed for C18:2 *t11c15*, C18:2 *t11c13*, C18:2 *ttNMID+t9t12*, C18:2
 405 *c9t13+t8c12*, C18:1 *t6-11*, and C18:2 *c9t11+t7c9+t8c10*, probably because the
 406 disappearance of ALA in the rumen is very high (usually >90% in case of high-forage
 407 diets) while only a little part of ALA is absorbed intact in the gut and secreted into milk
 408 (Chilliard and Ferlay, 2004). The observed consistent increasing concentrations of ALA
 409 despite the lower ALA supply due to the advance of the phenological phase of the grazed
 410 pasture plants suggests that the rate of disappearance of ALA probably decreased during
 411 the trial.
 412 Overall, the observed kinetics of responses of goat milk fatty acids to transition from
 413 indoor to pasture feeding were more similar to changes already observed in dairy ewes
 414 (Biondi et al., 2008) rather than dairy cows (Coppa et al., 2011; Khanal et al., 2008) under
 415 comparable feeding conditions.

CONCLUSION

A sudden transition of dairy goats from winter indoor to fresh grass feeding significantly affected the concentrations of FA in milk already two or three days after the diet change. Short- and medium-chain FA rapidly decreased by about 17% and 33% until day 4 after transition, suggesting that fresh grass feeding inhibited their *de novo* synthesis within the mammary gland. An adaptation to the new dietary conditions is hypothesized since from day 6 these FA significantly increased again. The higher availability of α -linolenic acid from pasture plants determined a notable increase in milk concentrations of its ruminal biohydrogenation intermediates, particularly conjugated (*t11c13+c9c11*, *t9t11*) and non-conjugated (*t11c15*, *c9t13+t8c12*, *c9t12*, *ttNMID+t9t12*) *trans*-octadecadienoic acids, and *trans*-octadecenoic acids (*t6-11*, *t12-14+c6-8*, *c14+t16*). The sum of CLA isomers *c9t11+t7c9+t8c10* also markedly increased (up to 261% at day 13), due to the higher absorption and availability of C18:1 *trans* isomers in the mammary gland as substrates for $\Delta 9$ -desaturase activity. Omega-3 FA (particularly α -linolenic and eicosapentaenoic acids) increased to a lesser extent (up to 93% and 85% at days 23 and 18, respectively), probably because of the high rate of disappearance of dietary ALA in the rumen. The increase in milk concentration of FA considered beneficial for human health went on till about thirteen (vaccenic and rumenic acids) or twenty-three (omega-3 fatty acids) days after transition.

ACKNOWLEDGMENTS

This research was supported by MIUR grants (ex 60%). The authors gratefully acknowledge their colleague Vanda Malfatto, Dr. Alessandro Moschietto and the farmers for their careful technical assistance.

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1 Table 1. Chemical composition (g kg⁻¹ DM, unless otherwise stated) and fatty acid profile (mg
2 100g⁻¹ DM) of the feed consumed by the goats.^a
3

	Indoor feeding		Pasture ^c
	Hay	Concentrate ^b	
Main nutrients			
DM (g kg ⁻¹)	891	886	198 ± 11.2
Ash	78	31	90 ± 9.5
CP	103	136	152 ± 25.6
EE	35	39	26 ± 2.8
NDF	568	202	458 ± 48.8
ADF	322	68	229 ± 16.9
ADL	56	17	41 ± 0.5
NE _L (MJ kg DM ⁻¹)	4.8	7.7	5.8
Fatty acids			
C12	7.9	n.d.	1.2 ± 0.48
C14	22.6	6.4	7.7 ± 1.18
C15	n.d.	n.d.	2.6 ± 0.26
C16	364.5	809.2	387.2 ± 29.03
C16:1 <i>n</i> 3	33.1	4.9	57.4 ± 12.64
C16:1 <i>c</i> 9	3.3	8.9	2.5 ± 1.18
C18	49.6	56.3	32.9 ± 2.86
C18:1 <i>c</i> 9	70.5	823.3	46.4 ± 8.81
C18:1 <i>c</i> 11	2.9	41.2	6.2 ± 1.37
C19	n.d.	n.d.	1.2 ± 0.31
C18:2 <i>c</i> 9 <i>c</i> 12 (LA)	188.4	2489.1	344.0 ± 65.49
C20	2.4	10.1	12.6 ± 1.59
C18:3 <i>c</i> 6 <i>c</i> 9 <i>c</i> 12	n.d.	n.d.	6.1 ± 1.00
C18:3 <i>c</i> 9 <i>c</i> 12 <i>c</i> 15 (ALA)	354.6	130.5	1507.3 ± 395.34
C22	n.d.	n.d.	0.4 ± 0.15
ΣSFA	446.8	881.9	445.7 ± 32.72
ΣMUFA	109.8	878.2	112.5 ± 8.56
ΣPUFA	543.1	2619.6	1857.4 ± 459.35
TFA	1099.7	4379.7	2415.6 ± 492.04

4 ^a Abbreviations: DM, dry matter; CP, crude protein; EE, ether extract; NDF, neutral detergent fiber; ADF,
5 acid detergent fiber; NE_L, net energy for lactation; LA, linoleic acid; ALA, α-linolenic acid; SFA,
6 saturated fatty acids; MUFA, monounsaturated fatty acids; PUFA, polyunsaturated fatty acids; TFA, total
7 fatty acids; n.d., not detected.

8 ^b Based on maize and wheat bran.

9 ^c Means and standard deviations of nine collected samples.

Table 2. Goat milk yield and gross composition during abrupt transition from indoor (hay and concentrate; day 0) to pasture (days 1 to 23) feeding.^a

	Days on pasture										Significance
	0	1	2	3	4	6	9	13	18	23	
Yield (kg head ⁻¹ day ⁻¹)	1.02 ^{DE}	1.05 ^{DE}	0.99 ^E	0.99 ^E	1.03 ^{DE}	1.09 ^D	1.17 ^{CD}	1.18 ^{BC}	1.25 ^{AB}	1.27 ^A	***
Fat (g kg ⁻¹)	41.8 ^B	40.2 ^B	47.3 ^A	45.4 ^A	46.6 ^A	45.8 ^A	45.2 ^A	46.4 ^A	45.3 ^A	48.4 ^A	***
Fat (g head ⁻¹ day ⁻¹)	40.7 ^F	41.7 ^F	46.8 ^{DE}	44.7 ^{EF}	46.8 ^{DE}	50.9 ^{CD}	52.9 ^{BC}	54.2 ^{BC}	57.3 ^{AB}	60.7 ^A	***
Protein (g kg ⁻¹)	30.5 ^D	31.3 ^{CD}	31.8 ^{BC}	31.7 ^{BC}	32.2 ^{BC}	33.6 ^A	32.4 ^B	34.1 ^A	32.5 ^B	34.2 ^A	***
Protein (g head ⁻¹ day ⁻¹)	31.1 ^E	32.5 ^E	31.3 ^E	31.1 ^E	32.4 ^E	36.9 ^D	37.7 ^{CD}	40.2 ^{BC}	40.7 ^B	43.2 ^A	***
Lactose (g kg ⁻¹)	40.4 ^F	40.8 ^{DEF}	40.3 ^F	40.1 ^F	41.4 ^{DE}	43.7 ^A	41.8 ^{CD}	42.6 ^{BC}	40.6 ^{EF}	43.2 ^{AB}	***
Lactose (g head ⁻¹ day ⁻¹)	41.0 ^{DE}	42.8 ^D	39.7 ^{DE}	39.3 ^E	41.7 ^{DE}	47.8 ^C	48.8 ^{BC}	50.1 ^{BC}	51.2 ^B	54.6 ^A	***

^a Total number of samples analyzed equal to 120 (12 goats × 10 sampling days).

^{A-F} Means within a row with different superscripts differ significantly. Probability: *** P≤0.001.

Table 3

Table 3. Groups of fatty acids (g kg⁻¹ fat) in goat milk during abrupt transition from indoor (hay and concentrate; day 0) to pasture (days 1 to 23) feeding.^{a,b}

	Days on pasture										Significance
	0	1	2	3	4	6	9	13	18	23	
Σ short chain ¹	125.2 ^B	123.8 ^B	108.3 ^C	97.7 ^D	103.4 ^{CD}	128.4 ^{AB}	129.3 ^{AB}	122.7 ^B	134.2 ^A	130.6 ^{AB}	***
Σ medium chain ²	367.9 ^A	376.1 ^A	307.2 ^{BC}	266.1 ^D	246.6 ^D	295.6 ^C	299.9 ^{BC}	319.2 ^{BC}	318.9 ^{BC}	320.7 ^B	***
Σ long chain ³	416.8 ^D	428.5 ^{CD}	513.3 ^A	526.6 ^A	486.8 ^{AB}	465.8 ^{BC}	422.8 ^{CD}	435.6 ^{CD}	405.6 ^D	415.5 ^D	***
Σ saturated ⁴	590.2 ^{AB}	598.2 ^A	534.2 ^{CDE}	480.9 ^F	450.9 ^G	523.4 ^{DE}	518.6 ^E	534.0 ^{CDE}	551.3 ^{CD}	554.9 ^{BC}	***
Σ branched chain ⁵	26.8 ^{ABC}	28.5 ^A	27.4 ^{AB}	24.7 ^{CDE}	22.3 ^F	22.9 ^{EF}	23.2 ^{DEF}	25.5 ^{BCD}	25.6 ^{BC}	25.8 ^{BC}	***
Σ monounsaturated ⁶	285.6 ^{CD}	294.4 ^C	353.1 ^A	362.3 ^A	337.1 ^{AB}	312.3 ^{BC}	282.7 ^{CDE}	287.3 ^C	254.3 ^E	255.3 ^{DE}	***
Σ C18:1 ⁷	273.3 ^{CDE}	281.5 ^{CD}	340.6 ^A	349.7 ^A	324.8 ^{AB}	299.4 ^{BC}	269.4 ^{DEF}	273.1 ^{CDE}	242.0 ^F	243.4 ^{EF}	***
Σ C18:1 <i>trans</i> ⁸	12.0 ^G	11.8 ^G	15.9 ^G	21.6 ^F	26.0 ^E	37.6 ^D	38.7 ^{CD}	42.7 ^{BC}	47.8 ^A	46.1 ^{AB}	***
Σ polyunsaturated ⁹	34.6 ^E	37.0 ^E	42.5 ^D	47.9 ^C	49.4 ^C	54.7 ^{AB}	51.4 ^{BC}	57.2 ^A	54.0 ^{AB}	57.7 ^A	***
Σ C18:2 ¹⁰	26.6 ^E	28.5 ^E	32.7 ^D	37.2 ^C	38.6 ^C	42.6 ^{AB}	40.8 ^{BC}	45.5 ^A	42.7 ^{AB}	43.7 ^{AB}	***
Σ C18:2 <i>trans</i> ¹¹	10.9 ^D	11.1 ^D	13.9 ^D	17.8 ^C	20.2 ^C	26.0 ^B	27.4 ^B	32.3 ^A	31.9 ^A	31.4 ^A	***
Σ <i>trans</i> without CLA ¹²	33.1 ^F	33.5 ^F	43.8 ^E	57.5 ^D	66.1 ^D	91.4 ^C	94.1 ^C	104.9 ^B	116.5 ^A	113.1 ^{AB}	***
Σ n3 FA ¹³	7.5 ^F	8.1 ^{EF}	9.4 ^E	11.1 ^D	11.2 ^D	12.8 ^{BC}	11.9 ^{CD}	13.6 ^B	13.4 ^{BC}	16.3 ^A	***
Σ n6 FA ¹⁴	23.1 ^E	25.1 ^{BCDE}	27.7 ^{AB}	28.9 ^A	27.6 ^{AB}	26.7 ^{ABC}	23.6 ^{DE}	24.3 ^{CDE}	25.2 ^{BCDE}	26.1 ^{BCD}	***
n6/n3	3.28 ^A	3.26 ^A	3.04 ^A	2.65 ^B	2.51 ^B	2.11 ^C	2.02 ^{CD}	1.79 ^{DE}	1.88 ^{CDE}	1.61 ^E	***
Σ CLA ¹⁵	5.5 ^F	5.5 ^F	7.1 ^F	9.6 ^E	12.1 ^D	17.0 ^C	18.2 ^{BC}	21.4 ^A	19.6 ^{AB}	19.4 ^{AB}	***
Σ unsaturated ¹⁶	320.3 ^{DE}	331.4 ^{DE}	395.6 ^{AB}	410.2 ^A	386.5 ^{AB}	367.0 ^{BC}	334.1 ^{CDE}	344.5 ^{CD}	308.3 ^E	313.0 ^{DE}	***
HSFA ¹⁷	568.2 ^A	582.8 ^A	445.8 ^B	369.2 ^C	345.8 ^C	459.5 ^B	449.1 ^B	477.4 ^B	478.7 ^B	481.8 ^B	***

^a Total number of samples analyzed equal to 120 (12 goats × 10 sampling days).
^b Abbreviations: CLA, conjugated linoleic acid; FA, fatty acids; HSFA, hypercholesterolemic saturated fatty acids.
^{A-G} Means within a row with different superscripts differ significantly. Probability: *** P≤0.001.

¹ C4, C5, C6, C7, C8, C10, C10:1.
² C12, C13 *iso*, C13 *aiso*, C12:1 *c*, C13, C14 *iso*, C14, C15 *iso*, C14:1 *t*, C15 *aiso*, C14:1 *c*+C15, C16 *iso*, C16, C17 *iso*, C16:1 *t*, C17 *aiso*, C16:1 *c*.
³ C17, C17:1 *t*, C18 *aiso*, C18, Σ C18:1, Σ C18:2, C20, C20:1 *t*, C18:3 *c6c9c12*, C20:1 *c9*, C20:1 *c11*, C18:3 *c9c12c15*, C18:2 *c9t11+t7c9+t8c10*, C18:2 *t10c12*, C18:2 *t11c13+c9c11*, C18:2 *t9t11*, C20:2 *c,c* n6, C22, C20:3 n6, C20:3 n3, C20:4 n6 (AA), C20:5 n3 (EPA), C22:5 n3 (DPA).
⁴ C4, C5, C6, C7, C8, C10, C12, Σ branched chain, C13, C14, C14:1 *c*+C15, C16, C17, C18, C20, C22.
⁵ C13 *iso*, C13 *aiso*, C14 *iso*, C15 *iso*, C15 *aiso*, C16 *iso*, C17 *iso*, C17 *aiso*, C18 *aiso*.
⁶ C10:1, C12:1 *c*, C14:1 *t*, C16:1 *t*, C16:1 *c*, C17:1 *t*, Σ C18:1, C20:1 *t*, C20:1 *c9*, C20:1 *c11*.
⁷ C18:1 *t5*, *t6-11*, *t12-14+c6-8*, *c9*, *c11*, *c12*, *c14+t16*.
⁸ C18:1 *t5*, *t6-11*, *t12-14+c6-8*.
⁹ Σ C18:2, C18:3 *c6c9c12*, C18:3 *c9c12c15*, C20:2 *c,c* n6, C20:3 n6, C20:4 n6 (AA), C20:5 n3 (EPA), C22:5 n3 (DPA).
¹⁰ C18:2 *t,t*-NMID+*t9t12*, *c9t13+t8c12*, *c9t12*, *c,c*-MID+*t8c13*, *t11c15*, *t9c12*, *c9c12*, *c9c15*, *c9t11+t7c9+t8c10*, *t10c12*, *t11c13+c9c11*, *t9t11*.

¹¹ C18:2 *t,t*-NMID+*t9t12*, *c9t13+t8c12*, *c9t12*, *c,c*-MID+*t8c13*, *t11c15*, *t9c12*, C18:2 *c9t11+t7c9+t8c10*, C18:2 *t10c12*, C18:2 *t11c13+c9c11*, C18:2 *t9t11*.

¹² C14:1 *t*, C16:1 *t*, C17:1 *t*, Σ C18:1 *t*, Σ C18:2 *t* (without CLA *trans*), C20:1 *t*.

¹³ C18:2 *t11c15*, C18:2 *c9c15*, C18:3 *c9c12c15*, C20:5 n3 (EPA), C22:5 n3 (DPA).

¹⁴ C18:1 *t12-14+c6-8*, C18:1 *c12*, C18:2 *t,t*-NMID+*t9t12*, C18:2 *c9t12*, C18:2 *t9c12*, C18:2 *c9c12*, C18:3 *c6c9c12*, C20:2 *c,c* n6, C20:3 n6, C20:4 n6 (AA).

¹⁵ C18:2 *c9t11+t7c9+t8c10*, *t10c12*, *t11c13+c9c11*, *t9t11*.

¹⁶ C10:1, C12:1 *c*, C14:1 *t*, C16:1 *t*, C16:1 *c*, C17:1 *t*, Σ C18:1, Σ C18:2, C20:1 *t*, C18:3 *c6c9c12*, C20:1 *c9*, C20:1 *c11*, C18:3 *c9c12c15*, C18:2 *c9t11+t7c9+t8c10*, C18:2 *t10c12*, C18:2 *t11c13+c9c11*, C18:2 *t9t11*, C20:2 *c,c* n6, C20:3 n6, C20:4 n6 (AA), C20:5 n3 (EPA), C22:5 n3 (DPA).

¹⁷ Calculated as C12+4*C14+C16.

Table 4

Table 4.
(a) Individual short- and medium-chain fatty acids (g kg⁻¹ fat) in goat milk during abrupt transition from indoor (hay and concentrate; day 0) to pasture (days 1 to 23) feeding.^{a,b}

	Days on pasture										Significance
	0	1	2	3	4	6	9	13	18	23	
C4	20.52 ^{CD}	19.09 ^E	20.26 ^{CDE}	21.13 ^{BCD}	22.96 ^A	22.42 ^{AB}	22.36 ^{AB}	19.91 ^{DE}	20.71 ^{CD}	21.36 ^{BC}	***
C5	0.07 ^{CDE}	0.11 ^{ABC}	0.12 ^{AB}	0.08 ^{BCDE}	0.07 ^{DE}	0.06 ^E	0.10 ^{BCD}	0.11 ^{ABC}	0.14 ^A	0.11 ^{ABC}	**
C6	18.71 ^{BC}	17.91 ^{CDE}	17.56 ^{DE}	16.96 ^E	18.31 ^{CD}	20.50 ^A	20.32 ^A	18.54 ^{CD}	19.72 ^{AB}	19.82 ^A	***
C7	0.17 ^{BCD}	0.20 ^{AB}	0.16 ^{CD}	0.14 ^{DE}	0.11 ^E	0.17 ^{BCD}	0.19 ^{BC}	0.18 ^{BCD}	0.24 ^A	0.21 ^{AB}	***
C8	20.57 ^{CD}	20.16 ^{DE}	18.93 ^{EF}	17.42 ^G	18.51 ^{FG}	22.61 ^A	22.12 ^{AB}	20.83 ^{BCD}	22.70 ^A	22.00 ^{ABC}	***
C10	62.36 ^B	63.41 ^{AB}	49.21 ^C	40.38 ^D	41.79 ^D	60.36 ^B	61.78 ^B	60.84 ^B	68.38 ^A	65.03 ^{AB}	***
C10:1	2.83 ^A	2.93 ^A	2.05 ^D	1.60 ^E	1.66 ^E	2.27 ^{BCD}	2.40 ^B	2.33 ^{BC}	2.28 ^{BCD}	2.08 ^{CD}	***
C12	28.83 ^A	29.99 ^A	22.47 ^D	17.36 ^E	16.68 ^E	23.69 ^{CD}	25.46 ^{BC}	25.22 ^{CD}	30.68 ^A	28.19 ^{AB}	***
C13 <i>iso</i>	0.24 ^{BC}	0.27 ^{AB}	0.17 ^{DE}	0.12 ^E	0.12 ^E	0.17 ^{DE}	0.21 ^{CD}	0.24 ^{BC}	0.25 ^{BC}	0.30 ^A	***
C13 <i>aiso</i>	0.41 ^B	0.51 ^A	0.29 ^{CD}	0.19 ^E	0.18 ^E	0.25 ^{DE}	0.30 ^{CD}	0.32 ^{CD}	0.33 ^C	0.32 ^{CD}	***
C12:1 <i>c</i>	0.69 ^{AB}	0.76 ^A	0.47 ^{EF}	0.38 ^{FG}	0.32 ^G	0.41 ^{EF}	0.45 ^{EF}	0.51 ^{DE}	0.63 ^{BC}	0.58 ^{CD}	***
C13	0.74 ^{AB}	0.80 ^A	0.51 ^D	0.31 ^E	0.31 ^E	0.56 ^{CD}	0.67 ^{ABC}	0.63 ^{BCD}	0.75 ^{AB}	0.69 ^{ABC}	***
C14 <i>iso</i>	1.23 ^A	1.31 ^A	1.02 ^{BC}	0.69 ^D	0.64 ^D	0.71 ^D	0.92 ^C	0.97 ^C	1.19 ^A	1.17 ^{AB}	***
C14	80.90 ^A	83.79 ^A	59.26 ^C	45.83 ^D	43.44 ^D	66.00 ^{BC}	62.31 ^{BC}	66.68 ^{BC}	67.41 ^B	67.91 ^B	***
C15 <i>iso</i>	2.09 ^B	2.32 ^{AB}	1.80 ^C	1.45 ^D	1.23 ^D	1.47 ^D	1.75 ^C	2.16 ^{AB}	2.29 ^{AB}	2.40 ^A	***
C14:1 <i>t</i>	0.05 ^E	0.08 ^{DE}	0.11 ^{CD}	0.17 ^{AB}	0.17 ^{AB}	0.14 ^{BC}	0.15 ^{ABC}	0.19 ^A	0.18 ^{AB}	0.15 ^{ABC}	***
C15 <i>aiso</i>	3.96 ^{BC}	4.29 ^B	3.26 ^D	2.41 ^{EF}	2.06 ^F	2.74 ^E	3.71 ^{CD}	4.16 ^{BC}	4.94 ^A	4.87 ^A	***
C14:1 <i>c</i> +C15	11.01 ^A	11.27 ^A	7.99 ^D	6.01 ^{EF}	5.13 ^F	6.15 ^E	7.81 ^D	9.01 ^C	9.92 ^B	10.41 ^{AB}	***
C16 <i>iso</i>	2.90 ^B	3.37 ^A	2.82 ^B	2.31 ^{DE}	1.93 ^F	2.07 ^{EF}	2.08 ^{EF}	2.46 ^{CD}	2.61 ^{BCD}	2.73 ^{BC}	***
C16	215.83 ^A	217.60 ^A	186.30 ^B	168.54 ^{CD}	155.33 ^D	171.84 ^{BC}	174.45 ^{BC}	185.44 ^B	178.37 ^{BC}	181.95 ^{BC}	***
C17 <i>iso</i>	3.86 ^{BCDE}	4.20 ^{ABC}	4.42 ^A	4.26 ^{AB}	3.86 ^{BCDE}	3.98 ^{BCDE}	3.83 ^{CDE}	4.09 ^{ABCD}	3.73 ^{DE}	3.63 ^E	**
C16:1 <i>t</i>	0.89 ^F	0.86 ^F	1.04 ^F	1.46 ^E	1.92 ^D	3.05 ^C	3.58 ^B	4.26 ^A	3.45 ^{BC}	3.48 ^B	***
C17 <i>aiso</i>	8.11 ^{AB}	8.28 ^{AB}	8.59 ^A	7.85 ^{ABC}	7.14 ^{CD}	6.99 ^D	7.06 ^{CD}	7.58 ^{BCD}	7.66 ^{BCD}	7.69 ^{BCD}	***
C16:1 <i>c</i>	6.24 ^A	6.36 ^A	6.66 ^A	6.73 ^A	6.14 ^A	5.36 ^B	5.13 ^B	5.34 ^B	4.47 ^C	4.22 ^C	***

Table 4.

(b) Individual long-chain fatty acids (g kg⁻¹ fat) in goat milk and desaturase indexes during abrupt transition from indoor (hay and concentrate; day 0) to pasture (days 1 to 23) feeding.^{a,b}

	Days on pasture										Significance
	0	1	2	3	4	6	9	13	18	23	
C17	6.97 ^{DE}	6.85 ^{DE}	8.59 ^{AB}	8.71 ^A	7.83 ^{BC}	7.05 ^{CD}	6.19 ^E	6.44 ^{DE}	6.33 ^{DE}	6.57 ^{DE}	***
C17:1 <i>t</i>	0.53 ^{DEF}	0.74 ^{BC}	0.99 ^A	1.14 ^A	1.05 ^A	0.80 ^B	0.67 ^{BCD}	0.59 ^{CDE}	0.44 ^F	0.50 ^{EF}	***
C18 <i>aiso</i>	4.06 ^{CD}	3.92 ^{CDE}	5.02 ^{AB}	5.46 ^A	5.16 ^A	4.49 ^{BC}	3.37 ^E	3.49 ^{DE}	2.60 ^F	2.69 ^F	***
C18	93.94 ^B	94.99 ^B	112.71 ^A	111.27 ^A	96.47 ^B	97.28 ^B	89.68 ^B	92.35 ^B	97.49 ^B	101.72 ^{AB}	**
C18:1 <i>t5</i>	0.06	0.08	0.10	0.11	0.07	0.08	0.02	0.06	0.04	0.10	ns
C18:1 <i>t6-11</i>	9.57 ^E	9.41 ^E	12.98 ^E	18.37 ^D	22.73 ^C	33.66 ^B	34.34 ^B	38.30 ^A	40.87 ^A	39.54 ^A	***
C18:1 <i>t12-14+c6-8</i>	2.33 ^E	2.29 ^E	2.86 ^{DE}	3.16 ^{CD}	3.16 ^{CD}	3.83 ^{BC}	4.30 ^B	4.38 ^B	6.87 ^A	6.51 ^A	***
C18:1 <i>c9</i>	253.68 ^D	261.88 ^{CD}	316.09 ^{AB}	318.77 ^A	290.12 ^{BC}	253.22 ^D	222.81 ^E	222.06 ^E	185.03 ^F	188.00 ^F	***
C18:1 <i>c11</i>	4.59 ^{BCD}	4.58 ^{BCD}	5.13 ^{AB}	5.49 ^A	5.30 ^A	4.89 ^{ABC}	4.29 ^{DE}	4.41 ^{CDE}	3.93 ^E	3.87 ^E	***
C18:1 <i>c12</i>	0.69 ^{ABC}	0.84 ^A	0.81 ^{AB}	0.85 ^A	0.75 ^{ABC}	0.67 ^{BC}	0.59 ^C	0.70 ^{ABC}	0.81 ^{AB}	0.81 ^{AB}	*
C18:1 <i>c14+t16</i>	2.21 ^D	2.38 ^D	2.65 ^{CD}	2.96 ^{BC}	2.64 ^{CD}	3.03 ^{BC}	3.02 ^{BC}	3.23 ^B	4.41 ^A	4.59 ^A	***
C18:2 <i>t,t</i> -NMID+ <i>t9t12</i>	0.39 ^F	0.53 ^F	0.69 ^E	0.94 ^D	0.86 ^{DE}	1.02 ^{CD}	0.97 ^D	1.25 ^{BC}	1.51 ^A	1.42 ^{AB}	***
C18:2 <i>c9t13+t8c12</i>	0.15 ^G	0.19 ^{FG}	0.22 ^{EF}	0.30 ^{EF}	0.34 ^E	0.50 ^D	0.53 ^{CD}	0.64 ^C	0.91 ^A	0.78 ^B	***
C18:2 <i>c9t12</i>	1.45 ^G	1.49 ^{FG}	1.81 ^{EF}	2.03 ^{DE}	1.95 ^{DE}	2.06 ^{DE}	2.24 ^{CD}	2.44 ^{BC}	2.89 ^A	2.75 ^{AB}	***
C18:2 <i>c,c</i> -MID+ <i>t8c13</i>	1.33 ^D	1.37 ^D	1.76 ^C	2.06 ^{BC}	2.03 ^{BC}	2.07 ^{BC}	2.13 ^B	2.36 ^{AB}	2.51 ^A	2.51 ^A	***
C18:2 <i>t11c15</i>	0.97 ^E	1.09 ^E	1.31 ^{DE}	1.91 ^{CD}	1.99 ^C	2.56 ^{BC}	2.84 ^B	3.57 ^A	3.76 ^A	3.85 ^A	***
C18:2 <i>t9c12</i>	1.04 ^{AB}	0.98 ^{AB}	1.08 ^A	0.97 ^{AB}	0.90 ^{BC}	0.76 ^{CD}	0.46 ^E	0.69 ^D	0.68 ^D	0.66 ^D	***
C18:2 <i>c9c12</i> (LA)	15.69 ^D	17.08 ^{BCD}	18.58 ^{AB}	19.19 ^A	18.18 ^{ABC}	16.50 ^{CD}	13.28 ^E	12.96 ^E	10.70 ^F	12.17 ^{EF}	***
C18:2 <i>c9c15</i>	0.04 ^D	0.27 ^A	0.15 ^{BCD}	0.19 ^{ABC}	0.19 ^{ABC}	0.10 ^{BCD}	0.10 ^{BCD}	0.20 ^{AB}	0.08 ^{CD}	0.19 ^{ABC}	***
C20	2.27 ^{AB}	2.33 ^A	1.73 ^D	1.31 ^{EF}	1.06 ^G	1.23 ^{FG}	1.19 ^{FG}	1.48 ^E	1.84 ^{CD}	2.04 ^{BC}	***
C20:1 <i>t</i>	0.21 ^{CD}	0.19 ^{CD}	0.23 ^{BC}	0.28 ^{AB}	0.31 ^A	0.29 ^{AB}	0.22 ^C	0.16 ^D	0.17 ^{CD}	0.17 ^{CD}	***
C18:3 <i>c6c9c12</i>	0.14 ^{CD}	0.10 ^{DE}	0.15 ^{BCD}	0.10 ^{CDE}	0.11 ^{CDE}	0.08 ^E	0.20 ^{AB}	0.15 ^{BC}	0.22 ^A	0.21 ^A	***
C20:1 <i>c9</i>	0.23 ^{CDE}	0.26 ^{CD}	0.32 ^{BC}	0.16 ^E	0.19 ^{DE}	0.17 ^{DE}	0.39 ^{AB}	0.32 ^{BC}	0.41 ^A	0.29 ^C	***
C20:1 <i>c11</i>	0.70 ^A	0.69 ^A	0.62 ^{AB}	0.66 ^A	0.55 ^{BC}	0.48 ^{CD}	0.32 ^E	0.43 ^D	0.34 ^E	0.39 ^{DE}	***
C18:3 <i>c9c12c15</i> (ALA)	5.16 ^{CD}	4.84 ^D	6.23 ^C	7.49 ^B	7.56 ^B	8.39 ^B	7.47 ^B	8.02 ^B	7.59 ^B	9.94 ^A	***
CLA <i>c9t11+t7c9+t8c10</i>	5.48 ^F	5.36 ^F	6.89 ^F	9.29 ^E	11.78 ^D	16.46 ^C	17.56 ^{BC}	19.76 ^A	18.95 ^{AB}	18.76 ^{AB}	***
CLA <i>t10c12</i>	<0.01	<0.01	0.01	<0.01	<0.01	<0.01	<0.01	0.01	0.01	0.01	ns
CLA <i>t11c13+c9c11</i>	0.02 ^D	0.04 ^D	0.08 ^D	0.19 ^C	0.26 ^C	0.45 ^B	0.52 ^B	0.64 ^A	0.53 ^B	0.51 ^B	***
CLA <i>t9t11</i>	0.05 ^D	0.10 ^C	0.10 ^C	0.11 ^{BC}	0.10 ^C	0.14 ^{ABC}	0.14 ^{ABC}	0.16 ^{AB}	0.16 ^A	0.13 ^{ABC}	***
C20:2 <i>c,c</i> n6	0.07 ^F	0.14 ^{EF}	0.20 ^{DE}	0.26 ^{CD}	0.30 ^{BC}	0.30 ^{BC}	0.31 ^{BC}	0.42 ^A	0.36 ^{AB}	0.39 ^{AB}	***
C22	0.97 ^{BC}	1.22 ^A	1.06 ^{ABC}	0.75 ^{DE}	0.64 ^E	0.64 ^E	0.73 ^{DE}	0.91 ^{CD}	0.98 ^{BC}	1.15 ^{AB}	***

C20:3 n6	0.13 ^{BC}	0.16 ^{AB}	0.19 ^A	0.17 ^{AB}	0.16 ^{AB}	0.13 ^{BC}	0.08 ^D	0.10 ^{CD}	0.07 ^D	0.08 ^{CD}	***
C20:4 n6	1.17 ^{CD}	1.45 ^A	1.36 ^{ABC}	1.19 ^{CD}	1.19 ^{CD}	1.38 ^{AB}	1.13 ^D	1.22 ^{BCD}	1.08 ^D	1.08 ^D	**
C20:5 n3 (EPA)	0.46 ^{DEF}	0.50 ^{DEF}	0.47 ^{DEF}	0.41 ^F	0.43 ^{EF}	0.60 ^{CD}	0.56 ^{CDE}	0.65 ^{BC}	0.85 ^A	0.75 ^{AB}	***
C22:5 n3 (DPA)	0.94 ^D	1.36 ^{AB}	1.19 ^{BC}	1.08 ^{CD}	1.06 ^{CD}	1.18 ^{BC}	0.92 ^D	1.17 ^{BC}	1.17 ^{BC}	1.53 ^A	***
DI ₁₆ ^c	0.29 ^C	0.29 ^C	0.36 ^B	0.40 ^A	0.40 ^A	0.31 ^C	0.29 ^C	0.29 ^C	0.25 ^D	0.23 ^D	***
DI ₁₈ ^d	27.4 ^{BC}	28.0 ^{BC}	28.4 ^{AB}	29.1 ^{AB}	30.4 ^A	26.0 ^{CD}	25.3 ^D	24.3 ^D	19.1 ^E	18.6 ^E	***

^a Total number of samples analyzed equal to 120 (12 goats × 10 sampling days).

^b Abbreviations: *c*, *cis*; *t*, *trans*; NMID, non methylene interrupted diene; MID, methylene interrupted diene; LA, linoleic acid; ALA, α -linolenic acid; CLA, conjugated linoleic acid; EPA, eicosapentaenoic acid; DPA, docosapentaenoic acid; DI, desaturase index.

^c Calculated as: C16:1 *c*9/C16:0.

^d Calculated as: C18:1 *c*9/C18:0.

^{A-G} Means within a row with different superscripts differ significantly. Probability: * P≤0.05; ** P≤0.01; *** P≤0.001; ns, not significant (P≥0.10).