



Changes in fatty acids in plasma and association with the inflammatory response in dairy cows abomasally infused with essential fatty acids and conjugated linoleic acid during late and early lactation

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ABSTRACT

Dairy cows are exposed to increased inflammatory processes in the transition period from late pregnancy to early lactation. Essential fatty acids (EFA) and conjugated linoleic acid (CLA) are thought to modulate the inflammatory response in dairy cows. The present study investigated the effects of a combined EFA and CLA infusion on the fatty acid (FA) status in plasma lipids, and whether changes in the FA pattern were associated with the acute phase and inflammatory response during late pregnancy and early lactation. Rumen-cannulated Holstein cows ($n = 40$) were assigned from wk 9 antepartum to wk 9 postpartum to 1 of 4 treatment groups. Cows were abomasally supplemented with coconut oil (CTRL, 76 g/d), linseed and safflower oil (EFA, 78 g/d of linseed oil and 4 g/d of safflower oil; ratio of oils = 19.5:1; n-6:n-3 FA ratio = 1:3), Lutalin (CLA, 38 g/d; isomers *cis*-9,*trans*-11 and *trans*-10,*cis*-12; each 10 g/d), or both (EFA+CLA). Blood samples were taken to measure changes in FA in blood plasma on d -63, -42, 1, 28, and 56, and in plasma lipid fractions (cholesterol esters, free fatty acids, phospholipids, and triglycerides) on d -42, 1, and 56 relative to calving, and in erythrocyte membrane (EM) on d 56 after calving. Traits related to the acute phase response and inflammation were measured in blood throughout the study. Liver samples were obtained for biopsy on d -63, -21, 1,

28, and 63 relative to calving to measure the mRNA abundance of genes related to the inflammatory response. The concentrations of α -linolenic acid and n-3 FA metabolites increased in lipid fractions (especially phospholipids) and EM due to EFA supplementation with higher α -linolenic acid but lower n-3 metabolite concentrations in EFA+CLA than in EFA treatment only. Concentration of linoleic acid decreased in plasma fat toward calving and increased during early lactation in all groups. Concentration of plasma arachidonic acid was lower in EFA- than in non-EFA-treated groups in lipid fractions and EM. The *cis*-9,*trans*-11 CLA increased in all lipid fractions and EM after both CLA treatments. Plasma haptoglobin was lowered by EFA treatment before calving. Plasma bilirubin was lower in EFA and CLA than in CTRL at calving. Plasma concentration of IL-1 β was higher in EFA than in CTRL and EFA+CLA at certain time points before and after calving. Plasma fibrinogen dropped faster in CLA than in EFA and EFA+CLA on d 14 postpartum. Plasma paraoxonase tended to be elevated by EFA treatment, and was higher in EFA+CLA than in CTRL on d 49. Hepatic mRNA abundance revealed time changes but no treatment effects with respect to the inflammatory response. Our data confirmed the enrichment of n-3 FA in EM by EFA treatment and the inhibition of n-3 FA desaturation by CLA treatment. The elevated n-3 FA status and reduced n-6:n-3 ratio by EFA treatment indicated a more distinct effect on the inflammatory response during the transition period than the single CLA treatment, and the combined EFA+CLA treatment caused minor additional changes on the anti-inflammatory response.

Key words: α -linolenic acid, conjugated linoleic acid, lipid fraction, inflammatory response

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INTRODUCTION

Essential fatty acids (**EFA**) have been shown to have beneficial effects in a wide variety of diseases in humans and animals, including dairy cows (Zamaria, 2004; Palmquist, 2010; Moallem, 2018). The immunomodulatory effects of EFA are mostly mediated through eicosanoid synthesis (e.g., prostaglandins, leukotrienes, and thromboxanes) from EFA. The main n-3 fatty acids (**FA**) are α -linolenic acid (**ALA**) with the metabolites eicosapentaenoic acid (**EPA**), docosapentaenoic acid (**DPA**), and docosahexaenoic acid (**DHA**); the main n-6 FA are linoleic acid (**LA**) and arachidonic acid (**ARA**; James et al., 2000). Generally, EFA of the n-3 series are known to be less inflammatory than n-6 EFA (James et al., 2000; Sordillo, 2016; Moallem, 2018). The changes in the n-6:n-3 FA ratio directly affect the production of mediators that are involved in the secretion of proinflammatory cytokines, such as TNF α (tumor necrosis factor- α), IL-1 β , and IL-6 (Simopoulos, 2016), and may also affect acute phase response in dairy cows around calving (Silvestre et al., 2011). The acute phase proteins (**APP**) play major roles in several aspects of the systemic reaction to inflammation and exert multiple systemic functions, including the adaptation of the energy and protein metabolism and the protection of tissue for excessive inflammatory alterations (Fleck, 1989; Ceciliani et al., 2012). Any effort to reduce the acute phase response in the transition period would be useful for optimizing the productive performance of high-yielding dairy cows (Bertoni et al., 2008; Trevisi et al., 2015). The APP response is triggered to a large extent by the production of interleukins, especially IL-1 β and IL-6 (Fleck, 1989; Bertoni and Trevisi, 2013; Trevisi et al., 2015).

The plasma lipid fractions consist of cholesterol esters (**CE**), free fatty acids (**FFA**), phospholipids (**PL**), and triglycerides (**TG**). In addition to PL in cell membranes, PL and CE in plasma are a large reservoir for EFA that supply organs and tissues and influence their functions. Incorporation of FA varies between distinct blood lipid pools, and follows fluctuation-induced changes from late pregnancy to early lactation (Christie, 1981; Malovrh et al., 2014). Essential fatty acids such as ALA and LA cannot be synthesized by mammals, including ruminants, and must be delivered by food (Palmquist, 2010). In addition, CLA isomers are produced in the rumen by EFA transformation. Therefore, rumen CLA production depends on EFA intake (Chilliard et al., 2001; Shingfield et al., 2010). The isomer-specific, health-promoting effects of CLA in humans are well known (Nagao and Yanagita, 2005; Shokryazdan et al., 2017). Some of the CLA isomers

reveal metabolic effects in dairy cows (e.g., milk fat reduction and glucose-sparing effects; Bauman et al., 2000; Hötger et al., 2013). These effects are able to improve the glucose status of dairy cows, which may have consequences for milk production and the immune response during the transition period (Baumgard et al., 2017). Therefore, especially in the transition period, the metabolic and immune function of dairy cows via EFA (especially n-3 FA) or CLA supply is of high interest.

The aim of the present study was to investigate the FA composition in plasma lipid fractions and erythrocyte membranes (**EM**) during supplementation with EFA and CLA alone and in combination. In addition, the effect of such a combined EFA and CLA treatment on the acute phase and inflammatory response of dairy cows during late pregnancy and early lactation was studied. A combined EFA and CLA supplementation refers to the supply of EFA and related rumen and tissue CLA production in dairy cows receiving fresh grass or on pasture (Kelly et al., 1998; Ferlay et al., 2006; Lahlou et al., 2014). Doses for the supplied EFA (linseed and safflower oil in a ratio of 19.5:1, providing an n-6:n-3 FA ratio of 1:3 in the supplement mixture) and CLA were recently evaluated in a companion dose-response study in mid-lactating dairy cows (Haubold et al., 2020). To avoid rumen degradation of the supplemented FA, all FA were infused into the abomasum (Vogel et al., 2020). We hypothesized that an elevated combined EFA and CLA intake would change the plasma FA pattern and affect the inflammatory response of dairy cows during the transition from late pregnancy to early lactation.

MATERIALS AND METHODS

Animals, Housing, Feeding, and FA Supplementation

The experimental procedures were carried out according to animal welfare guidelines and were approved by the relevant authorities of the state Mecklenburg-Western Pomerania, Germany (LALLF M-V/TSD/7221.3-1-038/15). For the present study, 40 German Holstein cows in the first third of their second lactation (expected milk yield of 11,000 kg/305 d) were purchased from a local farm (Agrarprodukte Dedelow GmbH, Prenzlau, Germany), kept in the experimental animal facility for cattle of the Leibniz Institute for Farm Biology (FBN, Dummerstorf, Germany), and adapted to the feeding and housing conditions. Cows were surgically equipped with a rumen cannula (10-cm center diameter cannula; Bar Diamond Inc., Parma, ID) 8 wk before the start of the experiments; access to the abomasum via the rumen for FA infusion has

recently been described (Haubold et al., 2020; Vogel et al., 2020). The presented investigations were part of a comprehensive project regarding the effects of EFA and CLA supplementation on performance and metabolism in transition dairy cows (Vogel et al., 2020). The cows were investigated from wk 9 antepartum (**ap**) until wk 9 postpartum (**pp**); animal management, performance data, and milk and carcass composition have been described recently (Vogel et al., 2020).

Feeding and housing management was the same for all cows during the experimental period. Briefly, animals were kept in a loose-housing freestall barn and were fed ad libitum with a corn silage-based TMR that was adapted for the dry period (wk 6 to 0 ap) as well as late- (wk 10 to 7 ap) and early-lactation periods (wk 1 to 8 pp). The TMR provided low amounts of fat and low n-3 FA supply, but contained a significant concentration of LA. Therefore, we obtained a high n-6:n-3 ratio in the FA profile of plasma and milk fat in cows fed with this diet (Weber et al., 2016; Revskij et al., 2019; Vogel et al., 2020). The chemical composition of the 2 different diets was determined according to the recommendations of the German Society of Nutritional Physiology (GfE, 2001, 2008; Table 1). The FA composition of the diets is shown in Table 2. The final n-6:n-3 FA ratio in the diets was increased to 7:1. Feed samples from TMR and corn silage were taken weekly and analyzed according to the Weender standard procedure (Naumann and Bassler, 2012) at the Agricultural Analysis and Research Institute (LUFÄ, Rostock, Germany). Cows had free access to water and were milked twice a day at 0630 and 1800 h during late and early lactation.

The cows were examined in 5 blocks of 8 animals each. The blocks started 1 after the other every 3 mo such that the whole study lasted ~21 mo in total. For each block, 8 cows were purchased such that expected calving date of each block was within 3 wk. In addition, milk yield of all cows had to be within 10,000 to 12,500 kg/305 d during second lactation (mean \pm SD: 11,101 \pm 1,118 kg of milk/305 d). Cows within a block did not differ in BW (mean \pm SD: 662 \pm 56 kg) and predicted calving interval (mean \pm SD: 395 \pm 39 d). Two cows per block were randomly assigned to 1 of 4 treatment groups (Vogel et al., 2020): control group (**CTRL**; coconut oil, Sanct Bernhard, Bad Ditzgenbach, Germany; n = 9); EFA (linseed oil; **DERBY**, Derby Spezialfutter GmbH, Münster, Germany and safflower oil; **GEFRO**, GC Memmingen/Allgäu, Germany; ratio of 19.5:1, providing an n-6:n-3 FA ratio of 1:3 in the supplement mixture; n = 9); CLA (Lutalin: *cis*-9,*trans*-11 and *cis*-10,*trans*-12 CLA; BASF, Ludwigshafen, Germany; n = 10); and the combination of EFA

Table 1. Ingredients and chemical composition of the diets

Item, g/kg of DM	Diet	
	Lactation	Dry period ¹
Ingredient		
Corn silage	457	421
Straw	97	223
Concentrates ² (granulated)	446	—
Dried sugar beet pulp	—	163
Extracted soybean meal	—	99
Rye grain	—	75
Minerals ³	—	10
Urea ⁴	—	9
Chemical composition		
NE _L ⁵ MJ/kg of DM	7.1	6.5
Crude fat	23	21
Crude fiber	173	219
CP	146	141
Utilizable protein ⁵	143	141
NFC	432	379
NDF	346	423
ADF	197	249
RNB ^{5,6}	0.5	0.0

¹The dry period diet was fed from wk 6 to 1 before calving.

²Ceravis AG, Malchin, Germany. Ingredients: 46.5% dried sugar beet pulp, 25.3% extracted soybean meal, 23.8% rye grain, 1.4% urea, 1.1% premix cow, 1.00% calcium, 0.37% phosphorus, 0.42% sodium, vitamins A, D₃, E, copper, ferric, zinc, manganese, cobalt, iodine, and selenium. Chemical composition: 44.4% NFC, 24.1% CP, 21.6% NDF, 12.4% ADF, 9.3% crude fiber, 8.2% crude ash, 1.8% crude fat, 7.9 MJ of NE_L/kg of DM.

³KULMIN MFV Plus (Bergophor Futtermittelfabrik Dr. Berger GmbH and Co. KG, Kulmbach, Germany): 8.5% magnesium, 7.5% phosphorus, 6.5% sodium, 3.5% HCl-insoluble ash, 1.5% calcium, additives: vitamins A, D₃, E, B₁, B₂, B₆, B₅, B₃, B₁₂, B₉, H, zinc, manganese, copper, cobalt, iodine, selenium, and *Saccharomyces cerevisiae*.

⁴Piarumin (SKW Stickstoffwerke Piesteritz GmbH, Lutherstadt Wittenberg, Germany); 99% urea, 46.5% total nitrogen.

⁵German Society of Nutrition Physiology (GfE 2001, 2008, 2009) and Deutsche Landwirtschaftliche Gesellschaft (DLG, 2013).

⁶RNB = ruminal nitrogen balance.

and CLA (**EFA+CLA**; n = 10). Two cows calved prematurely and had to be excluded from the study. Data from these 2 cows were not included in the statistical analyses. Cows were abomasally supplemented from d -63 ap until d 63 pp. The dosages of the daily abomasally infused supplements are shown in Table 3, and the FA composition of the infused lipids is presented in Supplemental Table S1 (<https://doi.org/10.3168/jds.2020-18735>). Dosages of FA applications were chosen based on a previous study (Haubold et al., 2020). Cows were supplemented with 2 equal portions of daily doses in the morning (0700 h) and afternoon (1630 h). The daily dose of infused fats was halved during the dry period. The drying period started at 6 wk before expected calving, and cows received dry period antibiotics (Benestermycin, Vetmedica GmbH, Ingelheim, Germany) and a teat concealer (Calgodip Zitzenversiegler T-Hexx dry, Calvatis GmbH, Germany).

Table 2. Fatty acid composition of the experimental diets

Fatty acid, g/kg of DM	Diet	
	Lactation	Dry period ¹
10:0	0.01	0.01
12:0	0.04	0.03
14:0	0.12	0.18
15:0	0.04	0.04
16:0	4.73	4.53
16:1 <i>cis</i> -9	0.06	0.05
17:0	0.09	0.08
17:1 <i>cis</i> -9	0.01	0.01
18:0	0.63	0.60
18:1 <i>cis</i> -9	4.82	3.84
18:1 <i>cis</i> -11	0.28	0.21
18:2 <i>cis</i> -9, <i>cis</i> -12	9.63	9.32
18:3 <i>cis</i> -9, <i>cis</i> -12, <i>cis</i> -15	1.35	1.37
18:4 <i>cis</i> -6, <i>cis</i> -9, <i>cis</i> -12, <i>cis</i> -15	0.04	0.02
20:0	0.15	0.16
20:1 <i>cis</i> -11	0.08	0.06
20:2 <i>cis</i> -11, <i>cis</i> -14	0.05	0.02
21:0	0.01	0.02
22:0	0.18	0.25
22:1 <i>cis</i> -13	0.01	—
22:2 <i>cis</i> -13, <i>cis</i> -16	0.01	0.04
23:0	0.05	0.02
24:0	0.23	0.29
SFA ²	6.27	6.21
MUFA ³	5.27	4.17
PUFA ⁴	11.08	10.77
Sum of n-3 fatty acids ⁵	1.39	1.39
Sum of n-6 fatty acids ⁶	9.69	9.38
n-6:n-3 ratio	7.00	6.76

¹The dry period diet was fed from wk 6 to 0 before calving.

²Sum of 10:0; 12:0; 14:0; 15:0; 16:0; 17:0; 18:0; 20:0; 21:0; 22:0; 23:0; and 24:0.

³Sum of 16:1 *cis*-9; 17:1 *cis*-9; 18:1 *cis*-9; 18:1 *cis*-11; 20:1 *cis*-11; and 22:1 *cis*-13.

⁴Sum of 18:2 *cis*-9,*cis*-12; 18:3 *cis*-9,*cis*-12,*cis*-15; 18:4 *cis*-6,*cis*-9,*cis*-12,*cis*-15; 20:2 *cis*-11,*cis*-14; and 22:2 *cis*-13,*cis*-16.

⁵Sum of 18:3 *cis*-9,*cis*-12,*cis*-15 and 18:4 *cis*-6,*cis*-9,*cis*-12,*cis*-15.

⁶Sum of 18:2 *cis*-9,*cis*-12; 20:2, *cis*-11,*cis*-14; and 22:2 *cis*-13,*cis*-16.

Blood Plasma Sampling, Preparation of EM, and Analyses

Blood samples were taken at 63, 42, 35, 28, 21, and 10 d before expected parturition, on d 1 after calving, and weekly thereafter up to d 56 pp from the jugular vein into evacuated tubes (Vacurette, Greiner Bio-One International AG, Kremsmünster, Austria). Blood was collected in the morning after milking and before subsequent feeding. Immediately after collection, samples were placed on ice and then centrifuged at $1,500 \times g$ at 4°C for 20 min. Extracted plasma was aliquoted and frozen at -20°C until analysis of FA and metabolites, and at -80°C until analysis of APP and interleukins. Additionally, at d 56 pp, EM were prepared according to a modified procedure of Kamata et al. (2008). After centrifugation ($2,200 \times g$ at 4°C for 15 min) and plasma collection, the buffy coat and the first layer of

the erythrocytes were discarded. The remaining erythrocytes were washed 3 times with Tris-buffered saline and centrifuged at $2,200 \times g$ at 4°C for 10 min. Thereafter, $10 \mu\text{L}$ of butylhydroxytoluol (2%) was added as an antioxidant to 2 mL of concentrated erythrocytes, and samples were stored at -80°C until analyses of EM for FA.

Bilirubin from samples containing sodium fluoride and potassium oxalate (2–4 and 1–3 mg/mL, respectively) was measured using an automatic spectrophotometer (ABX Pentra 400; Horiba ABX, Montpellier, France, #LT-BR0500). Plasma haptoglobin was determined from blood samples with lithium-heparin (12–30 IU of heparin/10 mL) used as an anticoagulant and by the method described by Skinner et al. (1991). The haptoglobin measurement was adapted to the ILAB600 condition (Calamari et al., 2016), which is based on the peroxidase activity of the methemoglobin-haptoglobin complex measured by the rate of oxidation of guaiacol (hydrogen donor) in the presence of hydrogen peroxide (oxidizing substrate). Bovine interleukins IL-1 β (cat. no. ESS0027; Thermo Scientific, Frederick, MD) and IL-6 (cat. no. ESS0029; Thermo Scientific) plasma concentrations were determined using commercial bovine-specific colorimetric sandwich ELISA kits according to Trevisi et al. (2015). The intra- and inter-assay coefficients of variation were 4.5% and 17.0%, and 3.5% and 13.4% for IL-1 β and IL-6, respectively. Plasma paraoxonase was measured by the method of Ferré et al. (2002) adapted to the ILAB 650 apparatus, as previously described by Trevisi et al. (2018). Plasma fibrinogen was determined by rapid heat precipitation according to Millar et al. (1971). Plasma concentrations of immunoglobulins IgG₁, IgG₂, and IgM were measured at d 63, 42, and 21 ap and on d 1, 14, 28, 42, and 56 pp via ELISA (no. E10–116, E10–117, E10–101, Bethyl Laboratories Inc., Montgomery, TX; Gerbert et al., 2018). The intra- and interassay coefficients of variation were 3.1% and 2.7% for IgG₁, 2.8% and 1.7% for IgG₂, and 4.2% and 2.9% for IgM. For the statistical analyses of the data, time points of sampling at which cows presented clinical symptoms of disease were retrospectively excluded from the analysis.

Plasma Lipid Extraction and FA Analyses

The FA composition in plasma was measured in blood samples containing EDTA (K₃EDTA, 1.8 g/L) at d –63, –42, 1, 28 and 56 relative to calving. Lipid extraction and separation of lipid fractions (i.e., CE, FFA, PL, and TG) from blood plasma were performed at d –42, 1, and 56 pp. The detailed sample preparation procedure was previously described by Dannenberger

Table 3. Amounts of daily abomasally infused supplements¹

Supplementation	Treatment						
	CTRL ²	EFA		CLA ²	EFA+CLA		
	Coconut oil ³	Linseed oil ⁴	Safflower oil ⁵	Lutalin ⁶	Linseed oil	Safflower oil	Lutalin
Daily infused oils, g/d							
Lactation dosage	76	78	4	38	78	4	38
Dry period dosage	38	39	2	19	39	2	19
Daily infused fatty acids (g/d) at lactation dosage ⁷							
18:3 <i>cis</i> -9, <i>cis</i> -12, <i>cis</i> -15	0.00	39.9	0.01	0.00	39.9	0.01	0.00
18:2 <i>cis</i> -9, <i>cis</i> -12	1.39	12.4	2.48	1.34	12.4	2.48	1.34
18:2 <i>cis</i> -9, <i>trans</i> -11	0.00	0.00	0.01	10.3	0.00	0.01	10.3
18:2 <i>trans</i> -10, <i>cis</i> -12	0.00	0.02	0.01	10.2	0.02	0.01	10.2

¹Cows were supplemented daily with coconut oil (CTRL), linseed and safflower oil (EFA), Lutalin (CLA, *cis*-9,*trans*-11 and *trans*-10,*cis*-12), or both (EFA+CLA).

²Addition of vitamin E (0.06 g/d), Covitol 1360 (BASF, Ludwigshafen, Germany), to compensate for vitamin E in linseed oil (0.07%) and safflower oil (0.035%).

³Sanct Bernhard, Bad Ditzgenbach, Germany.

⁴DERBY, Derby Spezialfutter GmbH, Münster, Germany.

⁵GEFRO, Memmingen/Allgäu, Germany.

⁶BASF, Ludwigshafen, Germany.

⁷The dosage for lactation was halved during the dry period.

et al. (2017). The FA analysis of the cow plasma lipids was performed using capillary gas chromatography with a CP-Sil 88 CB column (100 m × 0.25 mm, Agilent, Santa Clara, CA) that was installed in a PerkinElmer gas chromatograph CLARUS 680 with a flame ionization detector and split injection (PerkinElmer Instruments, Shelton) as described by Dannenberger et al. (2012). We used C19:0 as an internal standard for the quantification of FA. For the calibration procedure, we used the reference standard mixture Sigma FAME (Sigma-Aldrich, Deisenhofen, Germany), the methyl ester of C18:1*cis*-11, C22:5n-3, and C18:2*cis*-9,*trans*-11 (Matreya, PA), C22:4n-6 (Sigma-Aldrich, Deisenhofen, Germany), and C18:4n-3 (Larodan, Limhamn, Sweden). The 5-point calibration of a single FA ranged between 16 and 415 µg/mL and was checked after GC analysis of 5 samples.

Liver Tissue Sampling and Analyses

Liver biopsies were collected on d -63, -21, 1, 28 and 63 relative to calving. Skin was cut under local anesthesia, and liver tissue was extracted using a tailor-made biopsy needle with an outer diameter of 6 mm (Weber et al., 2013). Extracted liver tissue was immediately frozen in liquid nitrogen and stored at -80°C until analysis. After pulverization in liquid nitrogen, the RNA was extracted from homogenized liver tissue with TRIzol reagent (Life Technologies, Darmstadt, Germany) and cleaned with a RNeasy Mini Kit (Qia-

gen GmbH). The RNA quality was determined with an Agilent 2100 Bioanalyzer (Agilent Technologies). The mean RNA integrity number of all samples was 6 ± 1. The RNA integrity was checked by gel electrophoresis when the RNA integrity number was lower than 6 to make sure that RNA was not degraded. The quantity and purity of the total RNA were also measured by the optical density via a spectrophotometer (NanoPhotometer, Implen GmbH, Munich, Germany) at 260:280. Quantification of the relative mRNA abundance of selected genes was conducted as previously described (Saremi et al., 2012b). We used *LRP10* and *HPCAL1* as reference genes. Primer sequences, accession numbers, and PCR conditions for target genes are listed in Supplemental Table S2 (<https://doi.org/10.3168/jds.2020-18735>). The selected target genes related to inflammatory and immune responses were *TNFA*, *IL1A*, *IL1B*, *HP*, *SAA2*, *FGA*, *CRP*, *TLR4*, and *PON1*, and genes associated with prostaglandin synthesis were *COX1* and *COX2*. Primer products were verified by sequencing using the BigDye Terminator v1.1 Cycle Sequencing kit and an ABI 3130 Genetic Analyzer (Life Technologies, Carlsbad, CA). Relative mRNA amounts of the investigated genes were determined by real-time PCR (LightCycler 96, Roche Molecular Biochemicals, Mannheim, Germany) and SYBR green I detection. Melting curve analysis and agarose gel electrophoresis were used for verification of PCR products. Quantification cycle values and amplification efficiencies received with the use of LinRegPCR version 2017.0 (Ruijter

et al., 2013) were imported into qBASE+ version 3.1 (Biogazelle, Gent, Belgium) for all of the following calculations and quality controls. For normalization, the geometric mean of the reference gene abundance was used.

Statistical Analyses

Statistical analyses were conducted with SAS for Windows (version 9.4; SAS Institute Inc., Cary, NC). Plasma concentrations of immune and inflammatory traits, FA in plasma and plasma lipids, and hepatic mRNA abundance data were analyzed by repeated-measures ANOVA using the MIXED procedure, and a model containing EFA (levels: yes, no), CLA (levels: yes, no), time (levels: d relative to calving), block (levels: 1–5), and the respective interactions (EFA × CLA, EFA × time, CLA × time, and EFA × CLA × time) as fixed effects. The calving interval and milk yield in the second lactation were used as covariates. The most suitable covariance structure was the compound symmetry. For the analyses of plasma metabolites (haptoglobin, fibrinogen, paraoxonase, bilirubin, IL-6, and IL-1 β), the following distinct time periods were evaluated: antepartum (d 63, 42, 35, 28, 21, and 10 ap), transition (d 21, 10 ap and d 1, 7, 14, 21, and 28 pp), postpartum (d 1, 7, 21, 28, 35, 42, 49, and 56 pp), and the entire period of the study (d 63 ap to d 56 pp). The analyses of plasma FA, plasma immunoglobulins, and mRNA abundance in the liver were only calculated for the entire period due to fewer sampling time points. The least squares means (LSM) and their standard errors were computed for each fixed effect in the ANOVA model to display the results, and all group differences of the LSM were tested by the Tukey-Kramer procedure. The SLICE statement of the MIXED procedure was used to assess partitioned analyses of the LSM for interactions. Differences with $P < 0.05$ were considered significant. The CORR procedure of SAS was used to calculate Pearson correlations between data on the inflammatory response in liver and blood plasma.

RESULTS

Fatty Acids in Blood Plasma

The complete data for plasma FA concentrations and the FA composition of plasma fat are shown in Figure 1A–H and in Supplemental Tables S3 and S4 (<https://doi.org/10.3168/jds.2020-18735>). The concentrations of ALA, EPA, and DPA in plasma fat (equal to all lipids in blood plasma) increased (EFA × time: $P < 0.001$) with EFA supplementation and were higher ($P < 0.001$) in both EFA groups than in CTRL across the

entire study (Figure 1A–C). Plasma concentration of ALA was higher ($P < 0.05$) in EFA+CLA than in EFA on d –42 ap and 56 pp. Plasma EPA was lower ($P < 0.05$) on d 28 pp and plasma DPA was lower ($P < 0.05$) on d 1 pp in EFA+CLA than in EFA. Plasma DHA did not change before calving and was higher ($P < 0.05$) in EFA than in CTRL on d 1 pp, higher ($P < 0.05$) in EFA than in all other groups on d 28 pp, and higher ($P < 0.05$) in EFA+CLA than in EFA and CTRL on d 56 pp.

The plasma concentration of LA dropped toward calving, and continuously increased during early lactation (time: $P < 0.001$) in all groups (Figure 1E). On d 28 pp, plasma LA was higher in CLA than in CTRL and EFA ($P < 0.05$). The concentration of ARA in plasma fat was lower ($P < 0.05$) in both EFA groups than in CTRL and in CLA across the entire study (Figure 1F). Plasma concentrations of both CLA isomers were higher ($P < 0.05$) in CLA and EFA+CLA than in CTRL and EFA across the entire study (Figure 1G and H). Moreover, plasma *cis*-9,*trans*-11 CLA was higher ($P < 0.05$) in CLA than in EFA+CLA on d –42 ap, d 28 pp, and d 56 pp.

Lipid Fractions in Plasma

Cholesterol esters were the greatest lipid fraction in plasma fat (Figure 2A). Plasma FFA and PL proportions increased from d –42 ap to d 1 pp and then dropped again (time: $P < 0.001$; Figure 2B and C). The proportion of TG in plasma was highest in all groups during gestation and dropped toward calving ($P < 0.001$; Figure 2D). At d –42 ap, the proportion of TG in plasma was higher in CLA than in the all other groups ($P < 0.05$).

Fatty Acids in Lipid Fractions

The complete data for FA composition in the plasma lipid fractions of CE, FFA, PL, and TG are shown in Supplemental Tables S5–S8 (<https://doi.org/10.3168/jds.2020-18735>). The concentration of ALA was higher ($P < 0.05$) in EFA and EFA+CLA than in CTRL and CLA in almost all measurements of the lipid fractions (Figures 3A, 4A, 5A, and 6A). The concentration of ALA was higher ($P < 0.05$) in EFA+CLA than in EFA in CE (d –42 ap and d 56 pp) and in TG (d 56 pp). The concentration of EPA was higher ($P < 0.05$) in both EFA treatments than in CTRL and CLA in CE and PL across the entire study (Figures 3B and 5B). In the TG fraction, EPA was higher ($P < 0.05$) in EFA+CLA than in CLA and CTRL at d 56 pp (Figure 6B). The concentration of DPA in lipid fractions was only markedly increased ($P < 0.05$) in PL of the EFA

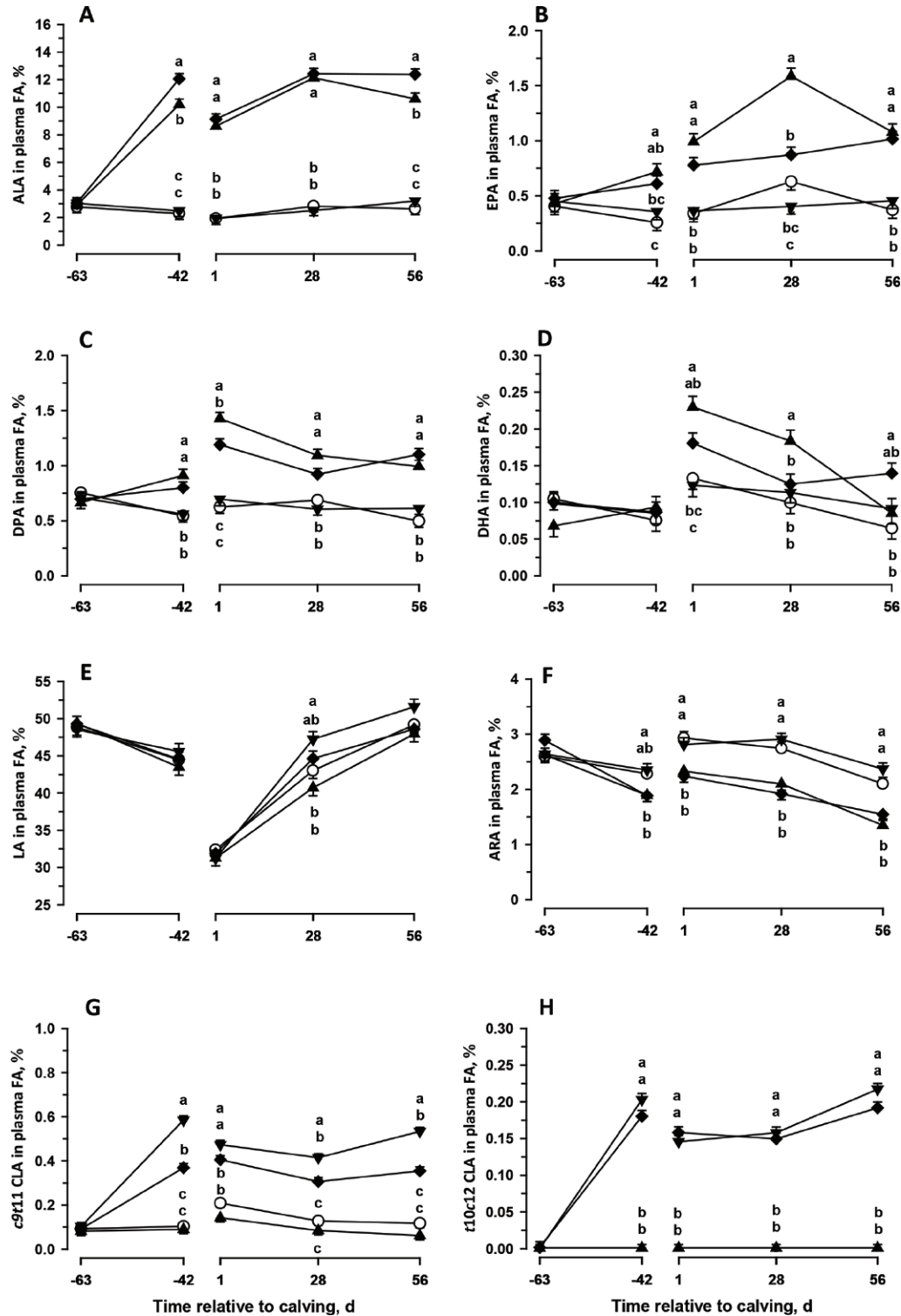


Figure 1. (A) Plasma concentrations of α -linolenic acid (ALA), (B) eicosapentaenoic acid (EPA), (C) docosapentaenoic acid (DPA), (D) docosahexaenoic acid (DHA), (E) linoleic acid (LA), (F) arachidonic acid (ARA), (G) *cis*-9,*trans*-11 CLA (*c9,t11* CLA), and (H) *trans*-10,*cis*-12 CLA (*t10,c12* CLA) from 9 wk before until 8 wk after calving in cows supplemented daily with coconut oil (\circ CTRL; $n = 9$), linseed and safflower oil (\blacktriangle EFA; $n = 9$), Lutalin (\blacktriangledown CLA *c9,t11* and *t10,c12*; $n = 10$; BASF, Ludwigshafen, Germany), or EFA+CLA (\blacklozenge ; $n = 10$) from wk 9 antepartum until wk 8 postpartum. Data are presented as the LSM \pm SE. Values with different lowercase letters (a-c) indicate significant differences ($P < 0.05$) among treatments at the respective time point. Statistically significant effects ($P < 0.05$) were observed for EFA treatment [all fatty acids (FA) except *t10,c12* CLA], CLA treatment (ALA; LA; *c9,t11* CLA; *t10,c12* CLA), EFA \times CLA interaction (*c9,t11* CLA), time (all FA), EFA \times time (all FA except LA and *t10,c12* CLA), and CLA \times time (all FA except ARA).

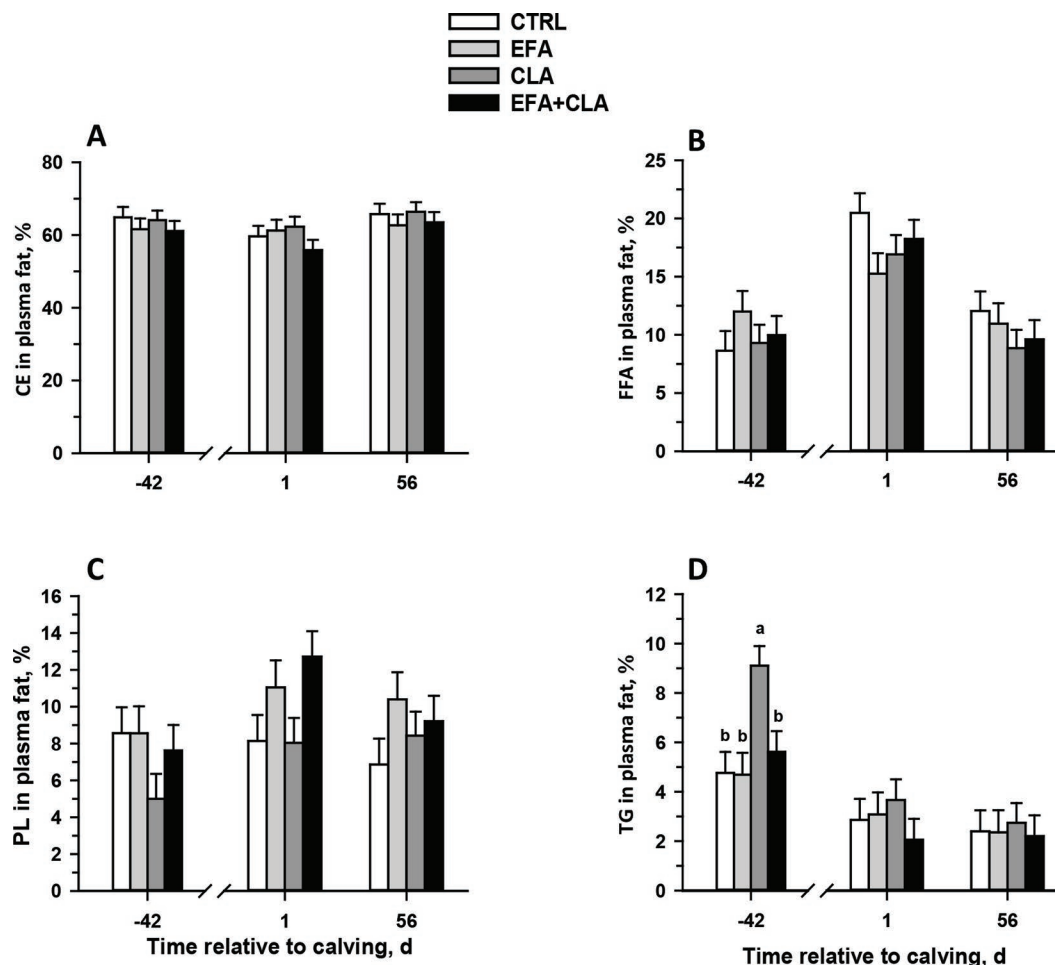


Figure 2. (A) Plasma concentrations of cholesterol esters (CE), (B) free fatty acids (FFA), (C) phospholipids (PL), and (D) triglycerides (TG) on d -42 before calving and d 1 and 56 after calving in cows supplemented with coconut oil (CTRL), linseed and safflower oil (EFA), Lutein (CLA *cis*-9,*trans*-11 and *trans*-10,*cis*-12; BASF, Ludwigshafen, Germany), or EFA+CLA. Data are presented as the LSM \pm SE. Values with different lowercase letters (a, b) indicate significant differences ($P < 0.05$) among treatments at the respective time point. Statistically significant effects ($P < 0.05$) were observed for time (all fatty acids) and for CLA \times time (PL and TG).

and EFA+CLA groups across the entire study (Figure 5C). The proportion of DHA in PL was higher ($P < 0.05$) in EFA than in CTRL and CLA on d 1 pp, and higher in EFA+CLA than in CTRL on d 56 pp (Figure 5D).

The LA proportion in EFA-treated groups was lower ($P < 0.05$) in CE on d -42 ap and 56 pp, but was higher ($P < 0.05$) in PL on d 56 pp than in non-EFA-treated groups (Figures 3E and 5E). In TG, LA was higher ($P < 0.05$) in EFA+CLA than in EFA on d 56 pp (Figure 6E). The proportion of ARA was lower ($P < 0.05$) in EFA-treated groups than non-EFA-treated groups across the entire study in CE and on d 1 and 56 pp in PL (Figures 3F and 5F). In TG, ARA was higher ($P < 0.05$) in CLA than in CTRL on d 56 pp (Figure 6F). Concerning CLA isomers in lipid fractions, *cis*-9,*trans*-11 CLA in CE was higher ($P < 0.05$) in CLA

than in CTRL across the entire study (Figure 3G). In PL, *cis*-9,*trans*-11 CLA was higher ($P < 0.05$) in both CLA-treated groups than in non-CLA groups across the entire study, and was higher ($P < 0.05$) in CLA than in EFA+CLA on d -42 ap and 56 pp (Figure 5G). In TG, *cis*-9,*trans*-11 CLA was higher ($P < 0.05$) in CLA-treated than non-CLA-treated groups on d -42 ap and d 56 pp (Figure 6G). The *trans*-10,*cis*-12 CLA was only detectable in CE and was higher ($P < 0.05$) in CLA-treated than in non-CLA-treated groups across the entire study (Figure 3H).

Fatty Acids in EM

The complete data on FA in EM are shown in Supplemental Table S9 (<https://doi.org/10.3168/jds.2020-18735>). The EFA treatment ($P < 0.001$) increased

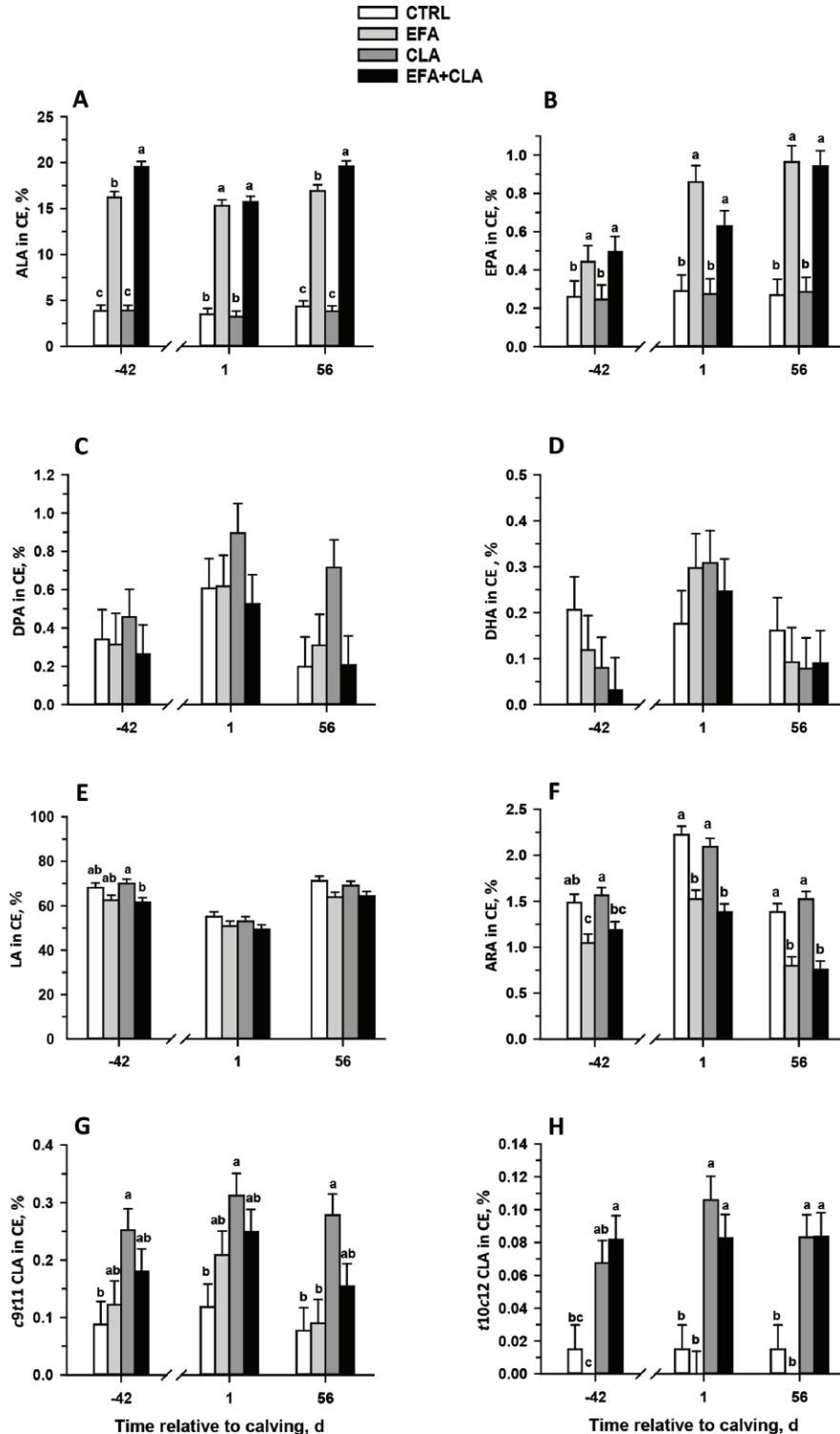


Figure 3. Fatty acid concentrations in plasma cholesterol ester (CE): (A) α -linolenic acid (ALA), (B) eicosapentaenoic acid (EPA), (C) docosapentaenoic acid (DPA), (D) docosahexaenoic acid (DHA), (E) linoleic acid (LA), (F) arachidonic acid (ARA), (G) *cis*-9,*trans*-11 CLA (*c9,t11* CLA), and (H) *trans*-10,*cis*-12 CLA (*t10,c12* CLA) on d -42 before calving and d 1 and 56 after calving in cows supplemented with coconut oil (CTRL), linseed and safflower oil (EFA), Lutalin (CLA *c9,t11* and *t10,c12*; BASF, Ludwigshafen, Germany), or EFA+CLA. Data are presented as the LSM \pm SE. Values with different lowercase letters (a–c) indicate significant differences ($P < 0.05$) among treatments at the respective time point. Statistically significant effects ($P < 0.05$) were observed for EFA treatment (ALA, EPA, LA, ARA), CLA treatment (ALA; *c9,t11* CLA; *t10,c12* CLA), EFA \times CLA interaction (ALA; DPA; *c9,t11* CLA), time (all fatty acids except *t10,c12* CLA), and EFA \times time (ALA, EPA, ARA).

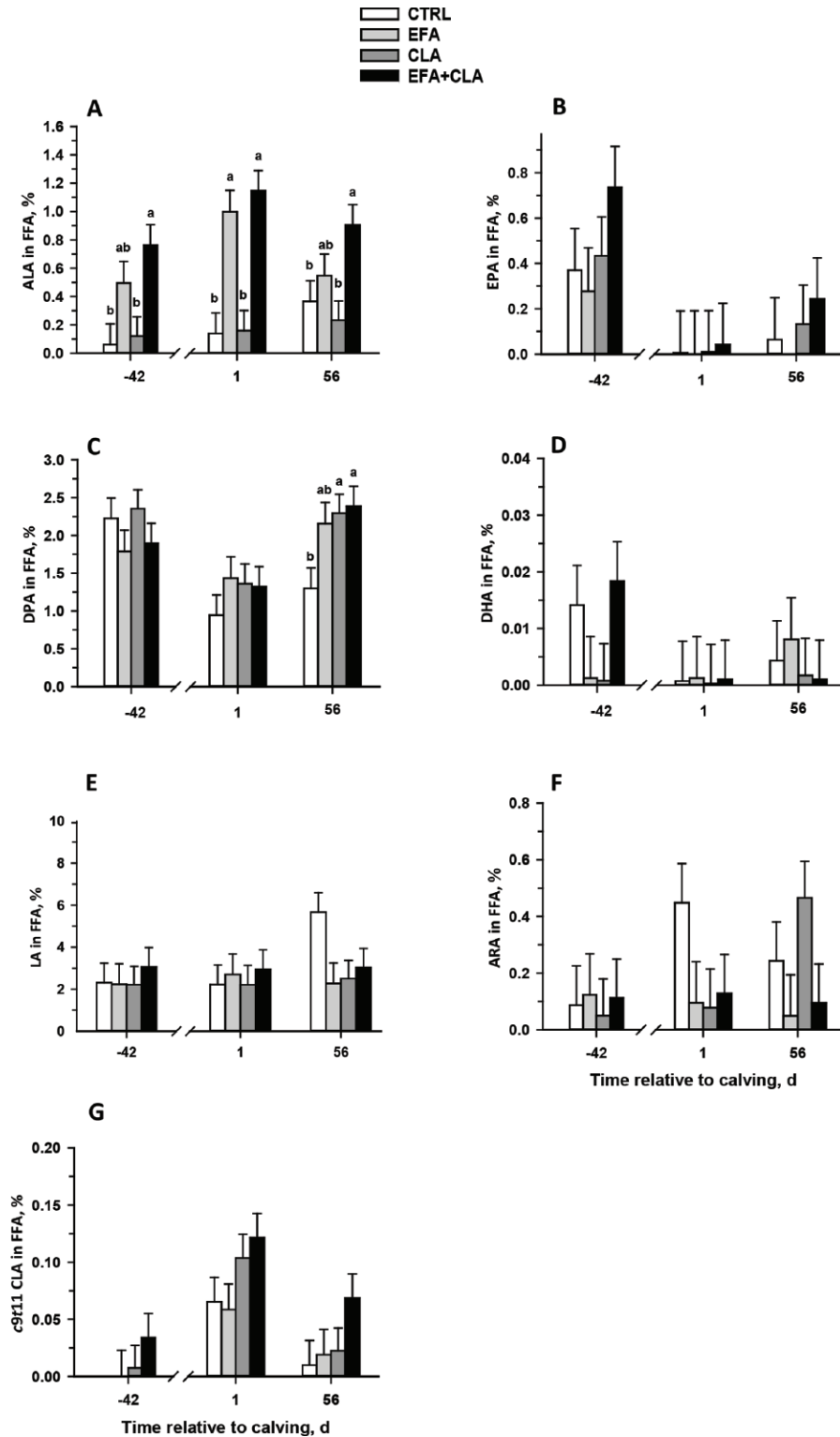


Figure 4. Fatty acid concentrations in plasma free fatty acids (FFA): (A) α -linolenic acid (ALA), (B) eicosapentaenoic acid (EPA), (C) docosapentaenoic acid (DPA), (D) docosahexaenoic acid (DHA), (E) linoleic acid (LA), (F) arachidonic acid (ARA), and (G) *cis*-9,*trans*-11 CLA (*c9,t11* CLA) on d -42 before calving and d 1 and 56 after calving in cows supplemented with coconut oil (CTRL), linseed and safflower oil (EFA), Lutalin (CLA *c9,t11* and *t10,c12*; BASF, Ludwigshafen, Germany), or EFA+CLA. Data are presented as the LSM \pm SE. Values with different lowercase letters (a, b) indicate significant differences ($P < 0.05$) among treatments at the respective time point. Statistically significant effects ($P < 0.05$) were observed for EFA treatment (ALA), CLA treatment (*c9,t11* CLA), time (ALA; EPA; DPA; *c9,t11* CLA), and EFA \times time (ALA, DPA).

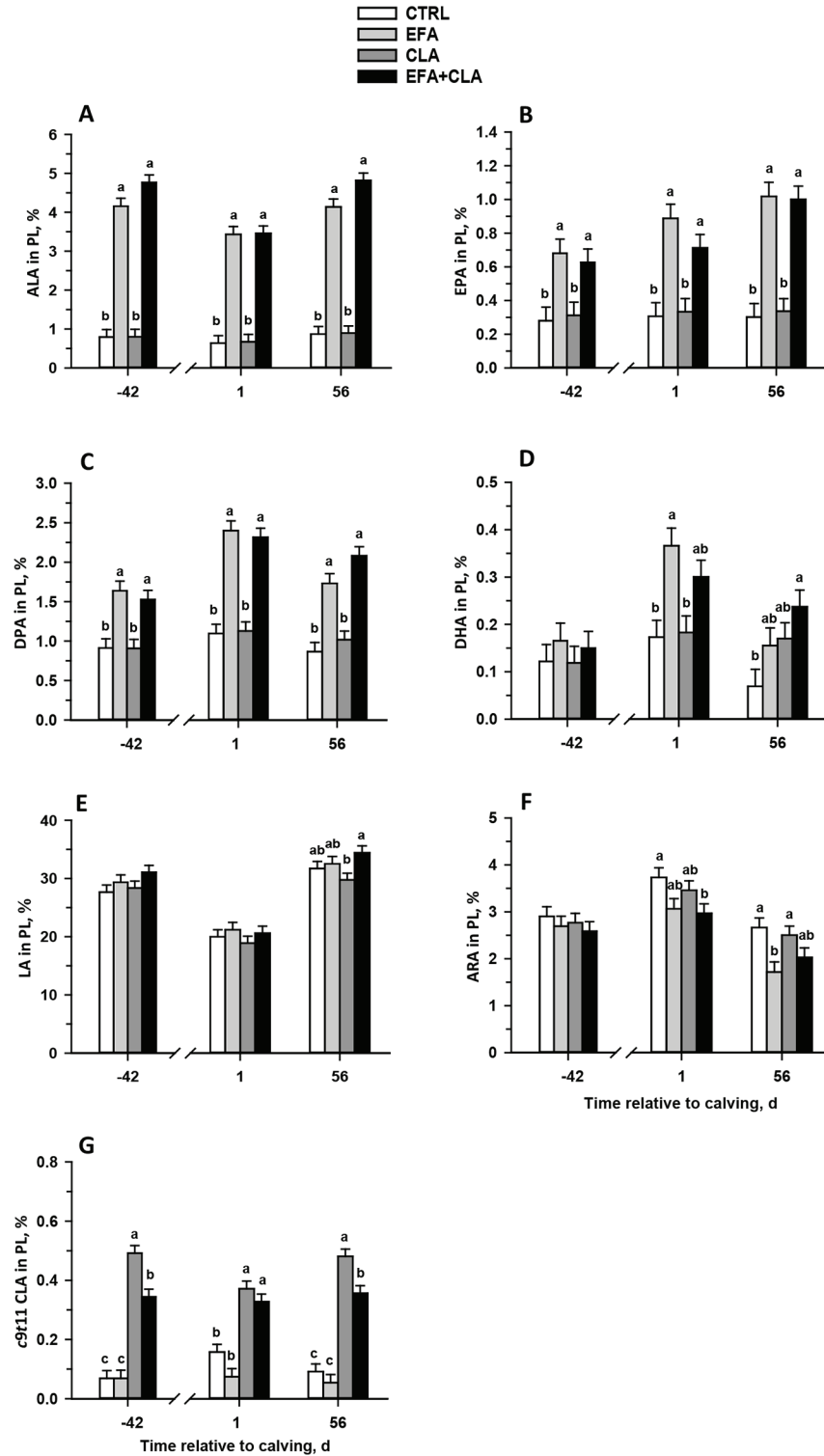


Figure 5. Fatty acid concentrations in phospholipids (PL): (A) α -linolenic acid (ALA), (B) eicosapentaenoic acid (EPA), (C) docosapentaenoic acid (DPA), (D) docosahexaenoic acid (DHA), (E) linoleic acid (LA), (F) arachidonic acid (ARA), and (G) *cis*-9,*trans*-11 CLA (*c9,t11* CLA) on d -42 before calving and d 1 and 56 after calving in cows supplemented with coconut oil (CTRL), linseed and safflower oil (EFA), Lutalin (CLA *c9,t11* and *t10,c12*; BASF, Ludwigshafen, Germany), or EFA+CLA. Data are presented as the LSM \pm SE. Values with different lowercase letters (a–c) indicate significant differences ($P < 0.05$) among treatments at the respective time point. Statistically significant effects ($P < 0.05$) were observed for EFA treatment (all fatty acids), CLA treatment (*c9,t11* CLA), time (all fatty acids except *c9,t11* CLA), EFA \times time (all n-3 fatty acids), and CLA \times time (DHA; *c9,t11* CLA).

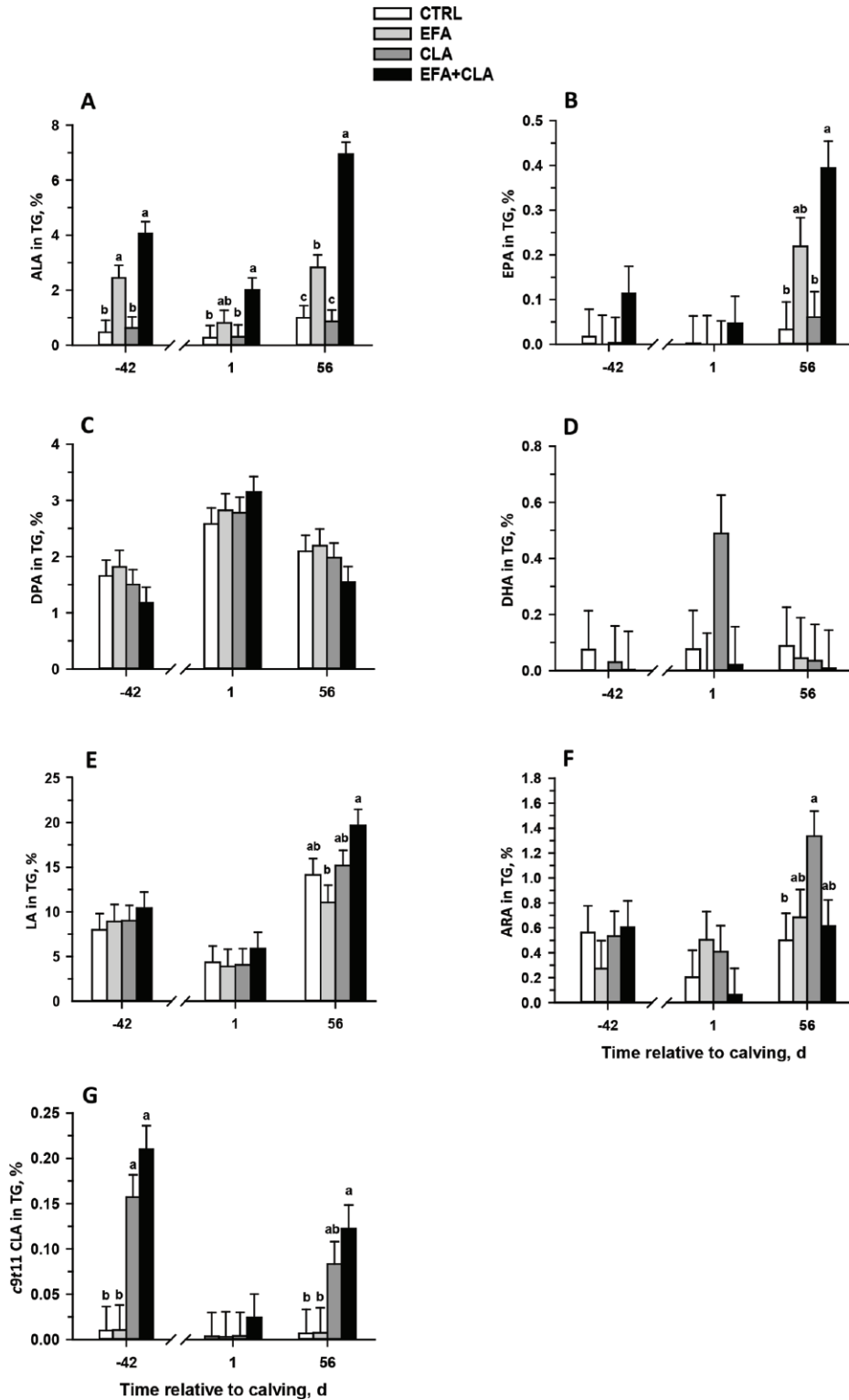


Figure 6. Fatty acid concentrations in triglycerides (TG): (A) α -linolenic acid (ALA), (B) eicosapentaenoic acid (EPA), (C) docosapentaenoic acid (DPA), (D) docosahexaenoic acid (DHA), (E) linoleic acid (LA), (F) arachidonic acid (ARA), and (G) *cis*-9,*trans*-11 CLA (*c9,t11* CLA) on d -42 before calving and d 1 and 56 after calving in cows supplemented with coconut oil (CTRL), linseed and safflower oil (EFA), Lutalin (CLA *c9,t11* and *t10,c12*; BASF, Ludwigshafen, Germany), or EFA+CLA. Data are presented as the LSM \pm SE. Values with different lowercase letters (a-c) indicate significant differences ($P < 0.05$) among treatments at the respective time point. Statistically significant effects ($P < 0.05$) were observed for EFA treatment (ALA, EPA), CLA treatment (ALA; *c9,t11* CLA), EFA \times CLA (ALA), time (all fatty acids except DHA), EFA \times time (ALA, EPA), and CLA \times time (ALA; DPA; *c9,t11* CLA).

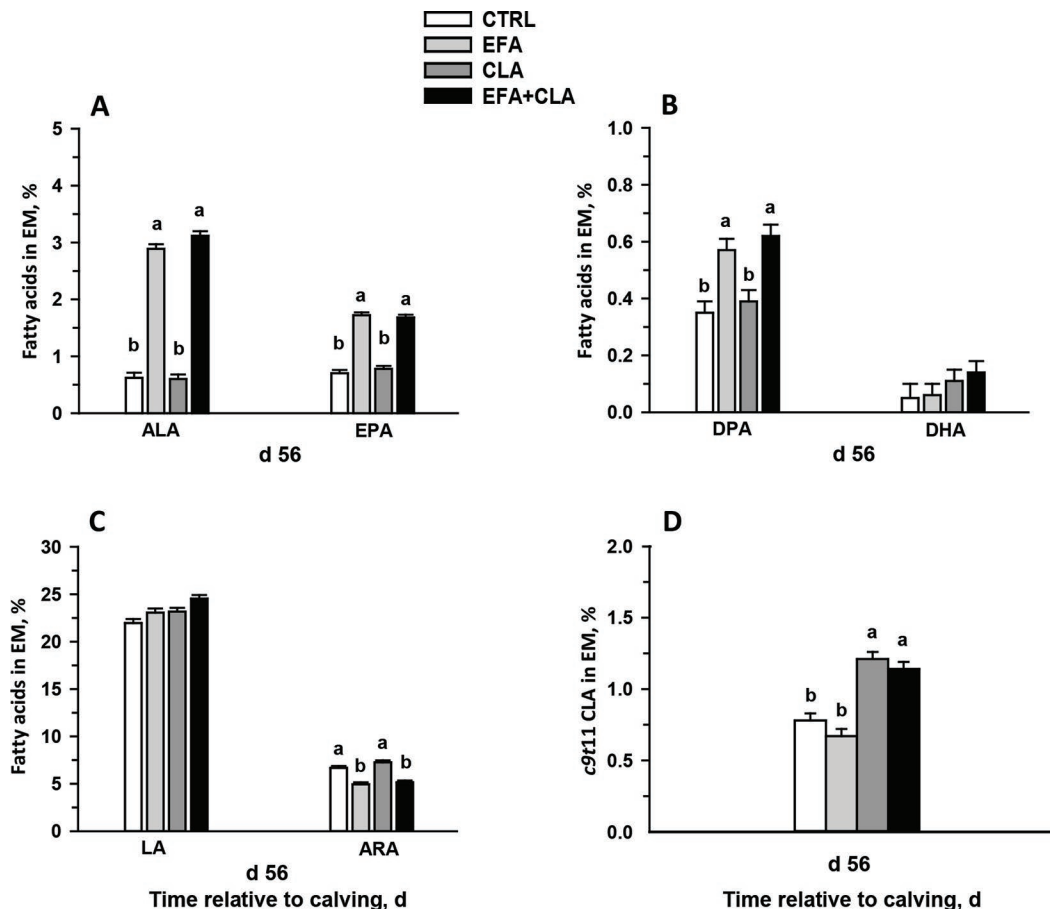


Figure 7. Fatty acid concentrations in erythrocyte membrane (EM): (A) α -linolenic acid and eicosapentaenoic acid (ALA, EPA), (B) docosapentaenoic acid and docosahexaenoic acid (DPA, DHA), (C) linoleic acid and arachidonic acid (LA, ARA), and (D) *cis*-9,*trans*-11 CLA (*c9,t11* CLA) on d 56 after calving in cows supplemented with coconut oil (CTRL), linseed and safflower oil (EFA), Lutalin (CLA *c9,t11* and *t10,c12*; BASF, Ludwigshafen, Germany), or EFA+CLA. Data are presented as the LSM \pm SE. Values with different lowercase letters (a, b) indicate significant differences ($P < 0.05$) among treatments at the respective time point. Statistically significant effects ($P < 0.05$) were observed for EFA treatment (ALA, EPA, DPA, LA, ARA), CLA treatment (ARA; *c9,t11* CLA), and EFA \times CLA (ALA).

concentrations of ALA and its metabolites (EPA and DPA) in EM (Figures 7A and 7B). Concentration of ARA was lower ($P < 0.001$) in EFA-treated than in non-EFA-treated cows ($P < 0.001$, Figure 7C). Concentration of *cis*-9,*trans*-11 CLA was higher ($P < 0.001$) in CLA-treated than non-CLA-treated cows (Figure 7D); *trans*-10,*cis*-12 CLA was not detected in EM.

Indicators of Inflammation and Hepatic Function in Blood Plasma

Plasma haptoglobin concentration peaked in all groups around calving ($P < 0.001$). Plasma haptoglobin was lowered ($P < 0.05$) by EFA treatment ap, was higher ($P < 0.05$) in CTRL than in EFA+CLA on d -21 ap, and tended to be higher ($P = 0.08$) in CTRL than in EFA on d -35 and -21 ap (Figure 8A). The plasma concentration of fibrinogen was slightly reduced

ap in both CLA-treated groups after the initiation of supplementation compared with EFA (EFA \times CLA and CLA \times time; $P < 0.05$; Figure 8B). Plasma fibrinogen was highest on d 1 pp, and then decreased ($P < 0.05$) in all groups, except in EFA. In EFA, the highest fibrinogen concentration was observed on d 14 pp. Plasma fibrinogen was lower ($P < 0.01$) in CLA than in EFA and EFA+CLA on d 14 pp. The plasma paraoxonase concentration decreased toward calving and then increased after calving in all groups (time; $P < 0.001$; Figure 8C). Plasma paraoxonase tended to be affected by EFA treatment ($P = 0.08$). On d 49 pp, the paraoxonase concentration was higher ($P < 0.05$) in EFA+CLA than in CTRL. A tendency toward higher paraoxonase concentrations in EFA+CLA compared with CTRL remained until d 56 pp ($P = 0.07$). The plasma bilirubin concentration was low before calving, markedly increased within a week after parturition, and

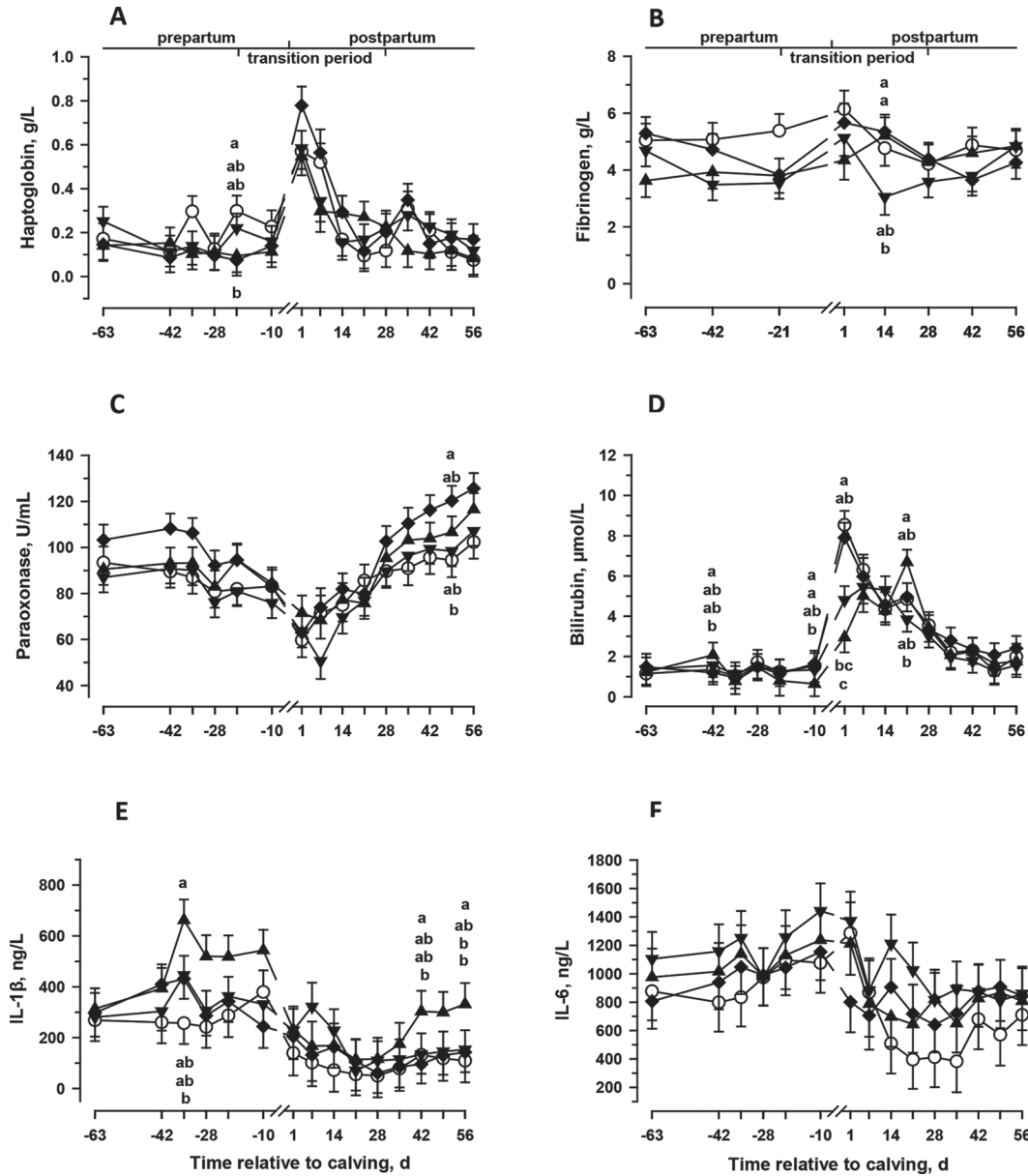


Figure 8. Plasma concentrations of (A) haptoglobin, (B) fibrinogen, (C) paraoxonase, (D) bilirubin, (E) IL-1 β , and (F) IL-6 from 9 wk before until 8 wk after calving in cows supplemented daily with coconut oil (\circ CTRL; $n = 9$), linseed and safflower oil (\blacktriangle EFA; $n = 9$), Lutalin (\blacktriangledown CLA *cis*-9,*trans*-11 and *trans*-10,*cis*-12; $n = 10$; BASF, Ludwigshafen, Germany), or EFA+CLA (\blacklozenge ; $n = 10$) from wk 9 antepartum until wk 8 postpartum. Data are presented as the LSM \pm SE. Values with different lowercase letters (a–c) indicate significant differences ($P < 0.05$) among treatments at the respective time point. Statistically significant ($P \leq 0.05$) effects for haptoglobin concentration during antepartum (EFA), transition (time), postpartum (time), and during the entire study (time). Statistically significant ($P \leq 0.05$) effects for fibrinogen concentration during antepartum (EFA \times CLA; CLA \times time), transition (EFA \times CLA; time; EFA \times time), postpartum (time), and during the entire study (time). Statistically significant ($P \leq 0.05$) effects for paraoxonase concentration during antepartum (time; EFA \times CLA \times time), transition (time), postpartum (time; EFA \times time), and during the entire study (time). Statistically significant ($P \leq 0.05$) effects for bilirubin concentration during antepartum (time; EFA \times CLA \times time), transition (EFA \times CLA; time; EFA \times CLA \times time), postpartum (EFA \times CLA; time; EFA \times CLA \times time), and during the entire study (EFA \times CLA; time; EFA \times CLA \times time). Statistically significant ($P \leq 0.05$) effects for IL-1 β concentration during antepartum (time), transition (time), postpartum (time; CLA \times time), and during the entire study (time). Statistically significant ($P \leq 0.05$) effects for IL-6 concentration during antepartum (time), transition (time), postpartum (time), and during the entire study (time).

then gradually decreased in all groups (time; $P < 0.001$; Figure 8D). Plasma bilirubin was higher ($P < 0.05$) in EFA than EFA+CLA on d -42 ap, but higher ($P <$

0.05) in EFA+CLA and CTRL compared with EFA on d -10 ap. Plasma bilirubin was lower ($P < 0.001$) in EFA and CLA than in CTRL, and was lower ($P <$

0.05) in EFA than in EFA+CLA on d 1 pp. Plasma bilirubin was higher ($P < 0.01$) on d 21 pp in EFA than in CLA. Plasma concentrations of IL-1 β and IL-6 decreased after calving ($P < 0.001$; Figures 8E and 8F). Plasma IL-1 β was higher ($P < 0.05$) in EFA than in CTRL on d -35 ap (trend remained on d -28 ap, $P = 0.09$). Another trend was observed on d -10 ap for higher concentrations in EFA than in EFA+CLA ($P = 0.05$). Plasma IL-1 β was higher ($P < 0.05$) in EFA than in EFA+CLA on d 42 and was higher ($P < 0.05$) in EFA than in EFA+CLA and CTRL on d 56 pp.

Correlations among plasma concentrations of inflammatory traits in blood plasma are presented in Supplemental Table S10 (<https://doi.org/10.3168/jds.2020-18735>). Plasma concentration of paraoxonase was positively related to the plasma concentration of fibrinogen ($r = 0.26$; $P < 0.001$) and was negatively related to the plasma concentration of haptoglobin and bilirubin ($r = -0.44$ and -0.24 ; $P < 0.001$). Plasma concentration of IL-1 β was positively related to the plasma concentration of IL-6 ($r = 0.41$; $P < 0.001$) and was negatively related to the plasma concentration of fibrinogen and bilirubin ($r = -0.32$ and -0.21 ; $P < 0.001$ and $P < 0.01$, respectively).

The plasma concentration of IgG₁ decreased ($P < 0.05$) from 6.0 ± 0.3 g/L at d -63 ap to 1.9 ± 0.3 g/L at calving, and increased again to 5.3 ± 0.3 g/L at d 56 pp in all groups. The plasma concentration of IgG₂ decreased ($P < 0.05$) from 3.6 ± 0.2 g/L at d -63 ap to 2.9 ± 0.2 g/L at d 28 pp in all groups. The plasma concentration of IgM decreased ($P < 0.05$) from 2.9 ± 0.2 g/L at calving to 2.2 ± 0.2 g/L at 14 pp in all groups. Neither EFA nor CLA, nor the combined EFA+CLA treatment affected immunoglobulin concentrations in plasma (Supplemental Figure S1, <https://doi.org/10.3168/jds.2020-18735>).

mRNA Abundance of Factors Related to Immune Response and Inflammation in Liver

Transcript abundances of all investigated genes were affected by time ($P < 0.05$) but not by EFA or CLA treatment (Table 4). The genes encoding the APP *HP*, *CRP*, *SAA*, and *FGA* showed patterns that are typical for the transition period with a rise around calving and lower abundances ap and pp. The mRNA abundance of the *HP*, *CRP*, *SAA*, *FGA*, *IL1A*, *IL1B*, and *PON1*, as well as *COX1* and *COX2* mRNA increased on d 63 pp. The abundance of *TNFA* mRNA showed neither time changes nor group differences. The abundance of *TLR-4* mRNA was greater ($P < 0.01$) in the CLA group than in the other groups on d 28 pp.

Correlations between plasma concentrations of inflammatory traits in blood plasma and liver are pre-

sented in Supplemental Table S10 (<https://doi.org/10.3168/jds.2020-18735>). Plasma concentration of paraoxonase was positively related to the mRNA abundance of *PON1* and *CRP* ($r = 0.37$ and 0.24 , respectively; $P < 0.001$ and $P < 0.01$), and was negatively related to the mRNA abundance of *SAA* and *HP* ($r = -0.17$ and -0.25 , respectively; $P < 0.05$ and $P < 0.001$). In addition, there were weak positive correlations of plasma paraoxonase concentration with mRNA abundance of *IL1A* and *IL1B* ($r = 0.16$ and 0.15 , respectively; $P < 0.05$). Plasma concentration of haptoglobin was positively related to the mRNA abundance of *HP*, *SAA*, and *FGA* ($r = 0.47$, 0.36 , and 0.25 , respectively; $P < 0.001$). Plasma fibrinogen concentration was positively related to the mRNA abundance of *PON1* and *CRP* ($r = 0.22$ and 0.23 , respectively; $P < 0.01$).

DISCUSSION

Fatty Acid Profiles in Plasma, Lipid Fractions, and EM

Changes in the FA pattern in plasma fat due to EFA and CLA infusion corresponded with changes recently published for milk fat (Moallem, 2018; Haubold et al., 2020). A larger increase in ALA induced by the combined EFA+CLA treatment compared with only EFA treatment in plasma fat supported observations in milk fat (Vogel et al., 2020), and was probably the consequence of the reduction in de novo FA synthesis by CLA supplementation (Bauman et al., 2011; Bionaz et al., 2015; Vogel et al., 2020). The LA concentration in plasma fat was only affected on d 28 pp with a higher concentration in the CLA than in the EFA and CTRL groups, highlighting a shift toward low LA in plasma fat as a result of increased middle-chain FA (CTRL) or ALA (EFA) infusion.

The concentrations of n-3 FA metabolites (EPA, DPA, and DHA) increased with EFA treatments, but the increases were lower in plasma fat of EFA+CLA than EFA-treated cows. These findings may refer to a *trans*-10,*cis*-12 CLA-related inhibition of n-3 FA desaturation in dairy cows (Harvatine and Bauman, 2011; Haubold et al., 2020), and support recent findings in milk fat (Vogel et al., 2020). The ARA concentration in plasma fat was lower in EFA and EFA+CLA than in CTRL and CLA due to the shift from n-6 to n-3 FA with the EFA treatments and proposed inhibition of n-6 desaturation (Palmquist, 2010). The increase in *cis*-9,*trans*-11 CLA was greater in CLA-treated cows than in EFA+CLA-treated cows. This shift in FA composition was not observed in milk fat (Vogel et al., 2020) and could not be explained by the relative shift of the FA pattern in plasma fat due to the additional

Table 4. Hepatic mRNA abundance (arbitrary unit) of traits related to inflammatory and immune response and prostaglandin synthesis in cows supplemented daily with coconut oil (CTRL; n = 9), linseed and safflower oil (EFA; in a ratio of 19.5:1; n = 9), CLA (n = 9), CLA (n = 10), or the combination (EFA+CLA; n = 10), from wk 9 antepartum until wk 9 postpartum¹

Gene ²	Treatment						Fixed effect, P-value					
	Time	CTRL	EFA	CLA	EFA+CLA	CLA × Time	EFA	CLA	EFA × CLA	Time	EFA × time	CLA × time
Related to the inflammatory and immune response												
<i>HP</i>	-63	4.42 ± 2.01 ^B	1.41 ± 2.00	1.06 ± 1.87	2.14 ± 1.87 ^B	0.3	0.7	0.3	0.001	1.0	1.0	
	-21	2.47 ± 1.89 ^B	1.06 ± 1.99	1.58 ± 1.99	0.66 ± 1.99 ^B							
	1	15.50 ± 2.18 ^A	8.07 ± 2.33	9.73 ± 2.56	14.03 ± 2.32 ^A							
<i>SAA2</i>	28	0.89 ± 2.00 ^B	1.20 ± 2.00	1.90 ± 1.99	1.18 ± 2.00 ^B							
	63	3.71 ± 2.16 ^B	5.36 ± 1.99	5.86 ± 1.87	2.33 ± 1.88 ^B							
	-63	1.82 ± 0.70 ^{AB}	1.30 ± 0.69	1.12 ± 0.65	1.62 ± 0.65 ^{AB}	0.5	0.8	0.6	0.001	0.9	0.9	
<i>FGA</i>	-21	1.93 ± 0.66 ^{AB}	1.01 ± 0.69	1.11 ± 0.69	0.86 ± 0.69 ^B							
	1	4.50 ± 0.76 ^A	3.20 ± 0.81	2.62 ± 0.89	4.31 ± 0.80 ^A							
	28	0.68 ± 0.69 ^B	0.92 ± 0.69	2.00 ± 0.69	0.69 ± 0.69 ^B							
<i>CRP</i>	63	1.96 ± 0.75 ^{AB}	2.54 ± 0.69	2.79 ± 0.65	1.83 ± 0.65 ^{AB}							
	-63	1.18 ± 0.23 ^{AB}	1.07 ± 0.23 ^{AB}	0.91 ± 0.21 ^B	1.12 ± 0.21 ^{AB}	0.9	0.8	0.9	0.001	0.9	0.2	
	-21	1.14 ± 0.22 ^B	1.16 ± 0.22 ^{AB}	0.86 ± 0.22 ^B	0.77 ± 0.22 ^B							
<i>PON1</i>	1	1.98 ± 0.24 ^A	1.46 ± 0.26 ^{AB}	1.29 ± 0.28 ^{AB}	1.84 ± 0.26 ^A							
	28	0.71 ± 0.23 ^B	0.81 ± 0.23 ^B	1.25 ± 0.22 ^{AB}	0.74 ± 0.22 ^B							
	63	1.25 ± 0.24 ^{AB}	1.73 ± 0.22 ^A	1.86 ± 0.21 ^A	1.50 ± 0.21 ^{AB}							
<i>IL1A</i>	-63	1.28 ± 0.40	1.39 ± 0.39 ^B	0.72 ± 0.37 ^B	1.14 ± 0.37 ^{AB}	0.6	0.4	1.0	0.001	0.5	0.6	
	-21	1.51 ± 0.37	0.88 ± 0.39 ^B	0.91 ± 0.39 ^B	0.79 ± 0.39 ^B							
	1	2.05 ± 0.43	1.51 ± 0.46 ^{AB}	1.46 ± 0.50 ^{AB}	1.76 ± 0.45 ^{AB}							
<i>IL1B</i>	28	1.24 ± 0.39	1.03 ± 0.39 ^B	2.02 ± 0.39 ^{AB}	1.08 ± 0.39 ^{AB}							
	63	2.45 ± 0.43	3.13 ± 0.39 ^A	2.63 ± 0.37 ^A	2.37 ± 0.37 ^A							
	-63	1.94 ± 0.77	1.51 ± 0.77	0.97 ± 0.72	1.48 ± 0.72 ^B	1.0	0.4	0.6	0.001	0.3	0.2	
<i>TNF</i>	-21	2.37 ± 0.73	1.14 ± 0.76	1.38 ± 0.76	1.11 ± 0.76 ^B							
	1	1.11 ± 0.83	0.81 ± 0.89	0.50 ± 0.97	2.25 ± 0.88 ^{AB}							
	28	1.28 ± 0.77	1.26 ± 0.77	3.55 ± 0.76	1.89 ± 0.76 ^{AB}							
<i>TLR4</i>	63	2.63 ± 0.83	3.42 ± 0.76	3.48 ± 0.72	4.52 ± 0.72 ^A							
	-63	1.14 ± 0.54 ^B	2.40 ± 0.53 ^{AB}	0.72 ± 0.50 ^B	0.96 ± 0.50	0.7	0.19	0.5	0.001	0.12	0.8	
	-21	1.82 ± 0.51 ^{AB}	1.10 ± 0.53 ^B	1.19 ± 0.56 ^{AB}	0.99 ± 0.53							
<i>IL1B</i>	1	2.25 ± 0.62 ^{AB}	2.00 ± 0.67 ^B	1.16 ± 0.66 ^{AB}	2.75 ± 0.84							
	28	1.57 ± 0.54 ^{AB}	1.41 ± 0.53 ^B	2.03 ± 0.53 ^{AB}	0.67 ± 0.60							
	63	2.71 ± 0.67 ^A	4.28 ± 0.53 ^A	3.23 ± 0.53 ^A	2.45 ± 0.50							
<i>TNF</i>	-63	1.43 ± 0.40	1.70 ± 0.39	0.95 ± 0.37 ^B	1.16 ± 0.37	0.9	1.0	0.2	0.05	0.7	0.4	
	-21	1.55 ± 0.38	1.30 ± 0.39	1.60 ± 0.39 ^{AB}	1.05 ± 0.39							
	1	1.40 ± 0.43	1.61 ± 0.45	1.35 ± 0.49 ^{AB}	1.65 ± 0.45							
<i>TNF</i>	63	1.49 ± 0.43	1.52 ± 0.39	1.90 ± 0.39 ^{AB}	1.26 ± 0.39							
	-63	1.39 ± 0.38	2.38 ± 0.39	2.57 ± 0.37 ^A	1.82 ± 0.37	0.2	0.3	0.5	0.5	0.7	0.9	
	-21	1.66 ± 0.36	1.47 ± 0.38	1.14 ± 0.41	1.99 ± 0.38							
<i>TNF</i>	1	2.45 ± 0.45	1.83 ± 0.50	1.25 ± 0.49	2.24 ± 0.63							
	28	1.42 ± 0.38	1.82 ± 0.38	2.02 ± 0.38	1.27 ± 0.44							
	63	1.24 ± 0.49	1.68 ± 0.38	1.22 ± 0.38	1.33 ± 0.36							
<i>TNF</i>	-63	1.84 ± 1.21	1.34 ± 1.21	1.63 ± 1.13 ^B	0.86 ± 1.13	0.07	0.6	0.8	0.4	0.5	0.1	
	-21	3.73 ± 1.22	1.39 ± 1.20	1.55 ± 1.20 ^B	1.47 ± 1.20							
	1	2.26 ± 1.32	0.71 ± 1.41	0.99 ± 1.55 ^B	2.90 ± 1.40							
<i>TNF</i>	28	1.69 ± 1.21 ^b	1.33 ± 1.21 ^b	7.45 ± 1.20 ^{aA}	2.01 ± 1.21 ^b							
	63	2.49 ± 1.30	3.03 ± 1.20	2.42 ± 1.13 ^B	1.37 ± 1.13							

Continued

Table 4 (Continued). Hepatic mRNA abundance (arbitrary unit) of traits related to inflammatory and immune response and prostaglandin synthesis in cows supplemented daily with coconut oil (CTRL; n = 9), linseed and safflower oil (EFA; in a ratio of 19.5:1; n = 9), CLA (n = 10), or the combination (EFA+CLA; n = 10) from wk 9 antepartum until wk 9 postpartum

Gene ²	Time	Treatment						Fixed effect, P-value					
		CTRL	EFA	CLA	EFA+CLA	EFA	CLA	EFA	CLA	EFA × CLA	Time	EFA × time	CLA × time
Related to prostaglandin synthesis													
COX1	-63	1.16 ± 0.36	1.21 ± 0.36 ^{AB}	0.85 ± 0.36 ^B	0.97 ± 0.34 ^B	0.5	0.7	0.8	0.001	0.4	0.4	0.4	0.4
	-21	1.42 ± 0.37	0.81 ± 0.39 ^B	1.08 ± 0.36 ^B	0.77 ± 0.36 ^B								
	1	1.38 ± 0.39	0.94 ± 0.42 ^{AB}	0.88 ± 0.46 ^B	1.26 ± 0.42 ^{AB}								
COX2	28	1.42 ± 0.36	1.39 ± 0.42 ^{AB}	2.51 ± 0.36 ^A	1.43 ± 0.36 ^{AB}								
	63	1.85 ± 0.39	2.42 ± 0.36 ^A	2.54 ± 0.34 ^A	2.50 ± 0.34 ^A	0.4	1.0	1.0	0.01	0.5	0.18	0.18	
	-63	1.23 ± 0.28	0.91 ± 0.28	0.84 ± 0.28 ^B	1.16 ± 0.26								
	-21	1.53 ± 0.28	1.15 ± 0.30	1.09 ± 0.28 ^{AB}	0.77 ± 0.28								
	1	1.47 ± 0.30	1.17 ± 0.32	1.06 ± 0.36 ^{AB}	1.36 ± 0.32								
28	1.17 ± 0.28	1.10 ± 0.28	2.07 ± 0.28 ^A	1.29 ± 0.28									
63	1.48 ± 0.30	1.98 ± 0.28	1.82 ± 0.26 ^{AB}	1.67 ± 0.26									

^{a,b}Means within a row with different lowercase superscripts differ ($P < 0.05$).

^{AB}Means within a column with different uppercase superscripts differ ($P < 0.05$).

¹Values are the LSM and SE.

EFA infusion because the absolute concentration of *cis-9,trans-11* CLA was also higher in CLA than in EFA+CLA (Supplemental Table S4, <https://doi.org/10.3168/jds.2020-18735>). Therefore, the reason for the differences in *cis-9,trans-11* CLA between CLA- and EFA+CLA-treated cows is currently unknown. In contrast, *trans-10,cis-12* CLA did not differ between EFA and EFA+CLA groups in plasma fat.

The different plasma lipid fractions exert distinct functions, and their fluctuations around parturition are connected with increased requirements of substrates for energy production during late gestation and the increased demands of FA for milk fat synthesis (Christie, 1981; Bionaz et al., 2007; Baumgard et al., 2017). In ruminants, most of the PUFA in plasma, especially LA, are localized in the CE and PL fractions, and dietary PUFA that escape rumen degradation are particularly stored in plasma CE and PL (Christie, 1981; Loor and Herbein, 2003; Palmquist, 2010). In fact, our results showed that CE contained the highest proportion of ALA and LA compared with other lipid fractions. The selective transfer of dietary LA and ALA to specific lipid fractions that do not provide significant amounts of FA to the mammary gland for milk fat production, such as CE and PL, may be a mechanism to store biologically active FA for more essential functions (Noble et al., 1982; Palmquist, 2010). In the present study, EFA and EFA+CLA treatments increased ALA in all lipid fractions because EFA was enriched for ALA. An increase in EPA concentration due to EFA and EFA+CLA treatments was predominantly observed in the PL and CE fractions. Elevated concentrations of DPA and partly of DHA after EFA and EFA+CLA treatment were mainly determined in the PL fraction. In addition, enhanced proportions of ALA, EPA, and DPA were observed in the EM of EFA- and EFA+CLA-treated cows. Cell membranes are rich in PL. Therefore, our study confirmed n-3 FA storage was preferably in the PL fraction of the blood plasma (Barceló-Coblijn and Murphy, 2009; Palmquist, 2010; Moallem, 2018).

Because all cows were fed the same diet with a high LA content, the concentration of LA in the plasma lipid fractions differed to a lesser extent among groups or even was lower in EFA- or EFA+CLA-treated cows. The concentrations of ARA in the CE and PL plasma fractions, as well as EM, were clearly reduced by EFA with or without CLA treatment. In addition to a general shift from n-6 to n-3 FA due to EFA treatment, the lower ARA proportion was probably caused by the inhibition of n-6 desaturation due to EFA infusion (Palmquist, 2010). An additional inhibitory effect of CLA treatment on $\Delta 6$ -desaturase activity, and consequently a reduced conversion of LA to ARA (as assumed in rodents), probably did not occur in the present study

because ARA proportion in lipid fractions did not differ between CLA and EFA+CLA groups. The CLA dosages used in rodent studies were much higher than in the present study, which may explain our different findings (Belury and Kempa-Steczko, 1997; Park et al., 2001). On the other hand, a higher concentration of LA in TG on d 56 in EFA+CLA than in EFA again indicated a shift toward long-chain FA due to inhibition of de novo FA synthesis caused by CLA (Bauman et al., 2011; Bionaz et al., 2015).

Only the *cis-9,trans-11* isomer was measurable in all lipid fractions of the plasma in the present study. In contrast to *trans-10,cis-12* CLA, *cis-9,trans-11* CLA is endogenously synthesized from vaccenic acid (18:1 *trans-11*) by $\Delta 9$ -desaturase (Griinari et al., 2000). However, mammalian tissue lacks the enzyme $\Delta 12$ -desaturase to reconvert *trans-10,cis-12* CLA from 18:1 *trans-10*; such that all of the *trans-10,cis-12* CLA detected in plasma seems to originate from gastrointestinal absorption (Pariza et al., 2001). The fact that the plasma FA profile is the first medium to mirror dietary changes, and that CE represents the greatest fraction of all lipids in plasma, explains why *trans-10,cis-12* CLA could only be detected in this compartment. Studies in humans (Burdge et al., 2005) and porcine EM (Malovrh et al., 2014) showed preferential uptake of the *cis-9,trans-11* CLA isomer when both *cis-9,trans-11* and *trans-10,cis-12* CLA were administered in the same doses. This may explain why we could not detect *trans-10,cis-12* CLA in the EM of the investigated cows.

Systemic and Hepatic Inflammatory Response

Our results regarding the inflammatory response in blood plasma revealed marked changes in APP and interleukins around calving, indicating a mild to moderate inflammatory reaction, which is commonly observed in the periparturient period (Bertoni et al., 2008; Trevisi et al., 2015; Jawor et al., 2016). The concentrations of haptoglobin and bilirubin decreased, while paraoxonase rose in all groups, probably because of the reduction of metabolic energy load and improvement of liver function after calving. In the present and in previous studies, higher concentrations of IL-1 β and IL-6 were collected during pregnancy, and IL-6 and IL-1 β decreased shortly after parturition (Sordillo et al., 1995; Ishikawa et al., 2004; Trevisi et al., 2015).

Our study revealed significant interactions between EFA and CLA treatments in the transit phase. The EFA treatment lowered the plasma haptoglobin concentration before calving and tended to raise paraoxonase activity throughout supplementation. Paraoxonase belongs to the negative APP and expresses a reverse time pattern during the transition period compared

with haptoglobin (Bionaz et al., 2007). Plasma bilirubin is related to the acute phase response (Trevisi et al., 2012), and bilirubinemia was lowest in the EFA group at calving. In previous studies, a milder acute phase response after LPS challenge in n-3 FA-fed cows (Trevisi et al., 2011; Greco et al., 2015) and calves (Ballou et al., 2008) has been described. The findings in the present study may point toward a better recovery after the transition phase in EFA-treated cows (Bertoni et al., 2015). Therefore, our results support the general concept of a reduced inflammatory response in the dairy cow with an elevated n-3 FA and a reduced n-6:n-3 FA status (Lessard et al., 2003; Gandra et al., 2016; Sordillo, 2016).

A higher IL-1 β plasma concentration in the EFA group throughout the dry period and after the transition was probably associated with the greater intake of n-3 FA (Lokesh et al., 1990). Given that proinflammatory cytokines stimulate the production of APP (Fleck, 1989; Bertoni and Trevisi, 2013; Trevisi et al., 2015), the fact that a low plasma haptoglobin concentration was paralleled by elevated plasma IL-1 β in the present study was not expected. The lower n-6:n-3 FA ratio in blood plasma might have impaired the stimulation of plasma haptoglobin by inhibiting the action of proinflammatory cytokines. Although IL-1 β was markedly higher in our EFA group, no stimulating response was noted for the concentration of plasma haptoglobin; however, plasma IL-1 β was negatively associated with plasma concentration of fibrinogen, which supports findings in the literature (Bode et al., 2012). The stimulatory effect of IL-1 β on haptoglobin production might differ among species and in bovine hepatocytes, IL-1 β failed to stimulate haptoglobin synthesis (Nakagawa-Tosa et al., 1995).

Concerning CLA, the *cis-9,trans-11* CLA isomer in particular has anti-inflammatory capacities (Viladomiu et al., 2016). Studies in epithelial cells in the bovine mammary gland have indicated reduced gene expression of pro- and anti-inflammatory cytokines after LPS challenge by EFA and CLA (Dipasquale et al., 2018). In the present study, the plasma fibrinogen concentration decreased in the blood plasma of the CLA-treated groups after the onset of supplementation, and plasma fibrinogen was lowest on d 14 pp in CLA-treated cows. In addition, CLA treatment alone reduced the bilirubin increase at calving. It was assumed that part of the CLA effect is based on the modulation of glucose metabolism caused by the reduction in milk fat synthesis in the mammary gland (Hötger et al., 2013). Glucose serves as an important fuel for immune cells, but at the onset of lactation most of the glucose is utilized in the mammary gland for synthesis of lactose (Baumgard et al., 2017; Gross et al., 2018). Measurement of whole glucose

turnover by [$^{13}\text{C}_6$] glucose revealed a reduction of the endogenous glucose production and a glucose-sparing effect due to CLA administration in the current study (L. Vogel, M. Gnott, and H. M. Hammon, unpublished observation), which supports previous findings (Hötger et al., 2013). However, inflammatory traits were only marginally affected by CLA treatment, which might indicate that the inflammatory status of the cows was not related to the glucose metabolism in the present study. Previous studies also showed no CLA effects on the inflammatory response during the transition period in dairy cows (Saremi et al., 2012a; Schäfers et al., 2018).

The abundance of genes involved in the acute phase response (*HP*, *SAA*, *FGA*, *CRP*, and *IL1*) increased around calving, which is consistent with our plasma results and supported by the literature (Loor et al., 2005). Correlations of mRNA abundance and plasma concentration of haptoglobin and paraoxonase clearly pointed at the importance of their production in liver. Nevertheless, no differences between groups were detected. The elevated mRNA abundance of the *APP* and *COX1* and *COX2* on d 63 pp in the liver could be initiated by the acute stress that cows experienced before slaughter (Murata and Miyamoto, 1993; Colditz, 2002).

CONCLUSIONS

Abomasal infusion of EFA with or without CLA increased the ALA concentration in all measured lipid fractions of the blood plasma. However, n-3 FA metabolites were primarily elevated in the PL fraction, indicating a different distribution of n-3 metabolites among lipid fractions in blood plasma. In addition, n-3 FA were enriched in the EM of EFA-treated cows, supporting the role of PL in retaining n-3 FA in EM. The combined EFA+CLA treatment partly reduced the enrichment of n-3 FA in blood plasma due to the inhibitory effect of CLA on EFA desaturation. The elevated n-3 FA status and the reduced n-6:n-3 ratio in blood plasma indicated a more distinct anti-inflammatory effect during the transition period in blood plasma than the CLA treatment. The combined EFA+CLA treatment barely showed additional benefits on the regulation of a presumed anti-inflammatory response; therefore, the combined CLA+EFA treatment did not affect the inflammatory response in a synergistic manner in dairy cows around calving in the present study.

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

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