# Pancreatic Lipase and Its Related Protein 2 Are Regulated by Dietary Polyunsaturated Fat during the Postnatal Development of Rats

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# ABSTRACT

The developmental gene expression of pancreatic lipase (PL) and its related proteins (PLRP1 and PLRP2) is anticoordinate. It is unknown whether dietary fat regulates the expression of these proteins in the preweanling stage. For determining the regulation of development and diet on PL, PLRP1, and PLRP2 as early as the suckling period, pregnant (Sprague-Dawley) rats consumed from day 15 (d15) of pregnancy through d9 of lactation a purified low (11% of energy) safflower oil diet [low-fat (LF)]. From d9 of lactation, dams and their respective pups were fed LF, mediumfat (MF; 40% of energy), or high-fat (HF; 67% of energy) safflower oil diets to d56. Milk fatty acid content had 15- to 100-fold less C:10 and 2.6- to 3.3-fold more C18:2 in MF and HF groups. Diet (LF < MF = HF; P < 0.002), postnatal development (d15 < d21 < d28 = d56; P < 0.001), and interaction of diet  $\times$  development significantly affected PL activity starting as early as d15. PL mRNA levels showed a parallel effect of diet (LF < HF = MF; P < 0.013) and development (P < 0.001). Both PLRP1 and PLRP2 mRNA levels were significantly affected by development (P < 0.001) and had an anticoordinate pattern compared with PL expression (d15 > d21 > d28). Reported for the first time is the significant down-regulation of PLRP2 mRNA levels by high polyunsaturated fat in suckling (d15) rats. In conclusion, PL and PLRP2 gene expression is regulated anticoordinately by the amount of dietary polyunsaturated fat starting as early as the preweanling phase of development. (*Pediatr Res* 56: 256–262, 2004)

## Abbreviations

FFA, free fatty acid
HF, high fat
LF, low fat
MF, medium fat
PL, pancreatic lipase
PLRP2, pancreatic lipase-related protein 2
PLRP1, pancreatic lipase-related protein 1
PUFA, polyunsaturated fatty acids

Dietary fat provides the major energy (~50% calories) during infancy in breast milk and formulas (1). During the past decade, special attention has been given to the type of fat necessary for normal infant development, especially to longchain polyunsaturated fatty acids (PUFA). PUFA derived from essential fatty acids (C18:2, linoleic acid, and C18:3 $\alpha$ linolenic) play a key role in brain and normal retinal development (2).

Fat digestion and absorption in the infant and premature neonate depend on the developmental pattern of lipases (3). The digestive system continues to develop after birth in infants, and especially so in premature neonates, influencing their ability to digest fat. Gastric lipase, which starts the digestion of

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dietary fat and accounts for 10-30% of fat digestion (4), pancreatic colipase-dependent lipase, and milk bile salt– stimulated lipase all have potential roles in digestion of milk fat in the neonate (3). Pancreatic lipase-related protein 2 (PLRP2) may play a role in neonatal fat digestion as well (5). Although the pancreatic lipase levels in the newborn and even more so in the premature infant are low, the digestion of milk fat depends on these four lipases with unique and only partially overlapping functions (5–7).

The exocrine pancreas synthesizes and secretes pancreatic lipase and two pancreatic lipase-related proteins; PLRP1 and PLRP2. PL and its related proteins have been identified in humans and rats. The PLRP1 is highly homologous to PL with conserved serine and histidine residues in the active site (8,9); however, this protein does not exhibit colipasedependent lipolytic activity when its full-length cDNA is expressed in COS or Sf9 cells (8,9). Although PLRP1 is secreted by the exocrine pancreas in several species (10), the function of PLRP1 remains unknown. Three reports

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(11–13) suggested that two mutations in residues 179 and 181 (alanine and proline to valine and alanine, respectively) render PLRP1 inactive. The second related protein (PLRP2) exhibits different lipolytic properties than PL. Even though PLRP2 has a limited activation by colipase in the presence of bile salts, PLRP2 has considerable lipolytic activity in the absence of colipase and has both phospholipase and galactolipase activities (14).

The postnatal developmental pattern of PL and its related proteins is anticoordinate in human and rat (9,15). The mRNA levels of PLRP1 and PLRP2 are high in the suckling only phase (days 0-14), decrease in the suckling-nibbling phase (days 14-21), and are low throughout weaning and adulthood (21 d and later). In contrast, PL is not expressed at detectable levels until postnatal day 14 (d14), and its expression increases throughout the suckling-nibbling and weanling phases to maximal levels in adulthood. It has been proposed that the anticoordinate expression of PL and its related proteins may reflect different roles in lipid digestion during postnatal development (9). Supporting their proposal, Lowe *et al.* (5) found a decreased neonatal dietary fat absorption and digestion in PLRP2 knockout mice.

PL is the main enzyme responsible for digestion of dietary triglycerides in the weaned and adult period (4). In the presence of colipase and bile salts, PL hydrolyzes dietary triglycerides in the small intestine to two-monoglycerides and free fatty acids (FFAs)-which are then absorbed. Both humans and rats adapt the synthesis of pancreatic lipase in response to dietary variations and thereby maximize the utilization of dietary fat (16). The physiologic importance of this dietary regulation has been shown. The up-regulation of PL enhances the response of cholecystokinin to increased dietary fat by increasing the rate of digestion of triglycerides and the release of FFAs in the proximal small intestine (17). In weanling animals, PL and PLRP1 adapt to increased dietary fat (41–75% of total energy) within 24 h. After 5 d, PL and PLRP1 reach higher steady-state levels of PL activity and synthesis and PL and PLRP1 mRNA (16-22).

The regulation of PL and PLRP1 by type of fat (chain length and degree of saturation) is controversial. We have previously shown that PL is regulated similarly by different types of fat above a threshold of 49% of energy as fat (21) and that highly polyunsaturated fat and fat rich in medium-chain triglycerides stimulate PL gene regulation below that threshold (20). It is unknown whether the gene expression of PL, PLRP1, and PLRP2 is modified by dietary fat in the suckling phase, when the pups consume only maternal milk, or in the sucklingnibbling phase of development, when the pups consume both maternal milk and maternal diet.

The fatty acid composition of maternal milk can be modified by altering the amount of polyunsaturated fat in the lactating dam's diet (23–25). The aim of this study was to determine whether the amount of polyunsaturated dietary fat can regulate PL, PLRP1, and PLRP2 during development including the suckling only, nibbling-suckling, weanling, and adult phases and to determine whether there is an interaction of this dietary regulation with the developmental regulation of PL and its related proteins when PUFA are introduced as early as the preweanling age.

#### METHODS

*Experimental protocol.* Fourteen-day timed pregnant female Sprague-Dawley rats (Charles River Lab, MA, U.S.A.) were housed individually in nursing cages in a temperaturecontrolled (24°C) room with a 12:12-h light-dark cycle and were fed a nonpurified commercial diet for the first 24 h. From the d2 until the d9 after delivery, dams were fed a low-fat (LF; 11% of energy as safflower oil) purified AIN-76 diet ad libitum (Table 1). For adjusting the rats to the milking procedure, pregnant rats were held routinely in milking position every day while the milking pump was on until delivery. From delivery until d9, dams and pups were disturbed only to feed and water them to reduce cannibalism. On d9, litter size was reduced to 11-13 without regard to sex of pup to standardize litter size and milk availability. On d9, lactating dams (with their respective pups) were randomly assigned to one of three diets: LF (11% of energy as safflower oil), moderate fat (MF; 40% of energy as safflower oil), or high fat (HF; 67% of energy as safflower oil; Table 1). These diets were isoenergetic and isonitrogenous but varied in the content of cellulose (to maintain the same caloric density), which has been shown not to affect the regulation of the exocrine pancreas (26). On d21 after delivery, all progeny were weaned to the respective maternal diet for the duration of the study and were housed individually in hanging cages. The dams were milked at d9 and d15 after delivery. Before milking, dams received an intraperitoneal injection of oxytocin (1 U/100 g body weight). Milk was obtained in capillary tubes attached to a vacuum pump. Suction was regulated manually. Individual milk samples were stored at  $-80^{\circ}$ C. The protocol was approved by the University of Maryland ACUC committee, and the NRC's guide for the care and use of laboratory animals was carried out.

Food consumption was measured daily. Pup body weights were measured at d9, d15, d21, d28, and d56 of the experiment when selected male pups were killed by  $CO_2$  inhalation. Sample size varied from 6 to 12 per group on d15–56 because of the variation in number of male pups per dam, which limited sampling on some days. A minimal sample size of 5 rats was estimated by power analysis to detect a 50% change in PL and PLRP1 mRNA levels with a population variance of 30% for PL

 Table 1. Dietary composition (21)

	LF	MF	HF
Fat (% kcal)	11	40	67
Carbohydrate (% kcal)	67	39	11
Component [% Diet			
(by weight)]			
Casein	20	20	20
DL-methionine	0.3	0.3	0.3
AIN-76 mineral mix	3.5	3.5	3.5
AIN-76 vitamin mix	1.0	1.0	1.0
Choline bitartrate	0.2	0.2	0.2
Cellulose	5.0	20.2	34.8
Safflower oil	5.0	17.1	28.9
Cornstarch	65.0	37.8	11.3

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and PLRP1. Because of the small pancreatic size on d9, pups were sampled for either PL mRNA or PL activity. Thus, sample size for body weight and pancreatic weight is larger (n = 20) than for either PL mRNA or PL activity. For d15 and later, pancreatic size was large enough that a portion of each pancreas was frozen immediately on dry ice and stored at  $-80^{\circ}$ C for enzyme analysis. The remainder of the pancreas was used immediately for RNA isolation as described below.

*Fatty acid analysis.* To 50 L of pooled milk samples for each treatment, 30 g of C:17 FFA, 500 L of benzene, and 3 L of acetylchloride:MeOH (1:15) were added (27). (Because of the small volumes of milk, pooling samples was necessary for sufficient volume for analysis.) The samples were incubated in a 65°C water bath for 2 h. After cooling, 10 g of C:15 fatty acid methyl ester, internal standard, 500  $\mu$ L of hexane and 1 L of H<sub>2</sub>O were added, and the samples were centrifuged to separate. The hexane layer was taken and injected into the gas chromatograph for fatty acid analysis (27).

**Pancreatic enzyme analysis.** Pancreatic fragments were homogenized in 9 vol of PBS (0.15 M of NaCl, 5 mM of PO<sub>4</sub>, pH 7.4) with a Polytron homogenizer. Homogenates were centrifuged at 14,000  $\times$  g at 4°C for 30 min. The supernatant was removed, and soybean trypsin inhibitor was added (final concentration, 0.01%). The supernatant was used for determination of enzyme activity and protein content. Lipase activity was assayed by a titrimeric method (21) with 20 mM of NaOH using a gum arabic–stabilized emulsion of neutralized triolein with excess crude colipase. Protein was determined by the method of Lowry *et al.* (28), using bovine albumin as a standard. Enzyme activities were expressed as nits (micromoles of fatty acid released per minute) per milligram of protein.

RNA extraction and hybridization studies. RNA was isolated as described by Chomczynski and Sacchi (29). This method is a single extraction with an acid guanidinium thiocyanate-phenol-chloroform mixture. Freshly isolated pancreatic fragments were immediately homogenized with a Polytron homogenizer (titanium probe) twice for 20 s, in ice-cold 4 M of guanidinium thiocyanate, 26 mM of sodium citrate, (pH 7), 0.5% sarcosyl, and 0.7% 2-mercaptoethanol. Sequentially, the RNA was extracted by adding 0.2 M of sodium acetate (pH 4) and phenol and chloroform-isoamyl alcohol mixture (49:1). RNA was precipitated with isopropanol, reprecipitated with 75% ethanol, and dissolved in sterile DEPC-treated water. RNA was quantified by UV absorption at 260 nm. The integrity of RNA was checked by 0.8% agarose gel electrophoresis for the presence of intact 18S and 28S ribosomes. Levels of PL, PLRP1, and PLRP2 mRNA and S28 RNA were quantified by dot-blot analysis. Each cDNA used yields a single band on Northern blots as reported by us and others (30,31). Dot blots are more quantitative than Northern blots and were used for this reason. Furthermore, area densitometry is highly correlated with direct Luminescence Scintillation Center (LSC) counting of blots (30). Recombinant plasmids used in these studies were the gifts of Dr. H.F. Kern [University of Marburg, Germany (PLRP1 cDNA; 0.82-kb insert in PstI site of pUC9, (32))], Dr. J. Williams [University of Michigan (PL cDNA; 1.5-kb insert in EcoRI site of pUC18, (30)] and D. Soprano [Temple University, Philadelphia (28S cDNA probe), and Dr. P. Howells (PLRP2 cDNA; 1.3-Kb insert in XdaI and KpNI site in pKSII, (33)). Previously, we reported negligible cross-hybridization of PL and PLRP1 (33). We also examined the specificity of the homologous and heterologous hybridization of random primelabeled PL, PLRP1, and PLRP2 cDNA inserts with 0.2  $\mu$ g of linearized plasmid DNA containing PLRP2 insert. Hybridization and washing conditions were identical to those described here. The PLRP2 cDNA cross-hybridization to the PL and PLRP1 plasmid averaged 0 and 0.1%, respectively, of its hybridization to the homologous PLRP2 plasmid. These results demonstrate the specificity of the hybridization of each PL probe under the highly stringent conditions used in this study. Specific mRNAs were quantified by dot-blot hybridization within the range of linear hybridization (30). Total RNA was diluted in DEPC-treated water to the appropriate concentration, verified by absorbance at 260 nm. For denaturation, the samples were mixed with 6.15 M of formaldehyde in 0.75 M of NaCl and 0.075 M of trisodium citrate [5 $\times$  SSC; 1 $\times$  SSC in 0.15 M of NaCl and 0.015 M of sodium citrate (pH 7)] and heated for 15 min at 65°C. The denatured samples were spotted onto a nitrocellulose filter using a Millipore dot-blot apparatus. The filters were cross-linked by UV radiation with optimal dosage (120 mJ/cm<sup>2</sup>) and prehybridized at 42°C for 2 h in a solution that contained 50% formamide 5× SSC, 5× Denhardt's solution, 0.1% SDS, and 0.03% tRNA. Hybridization was performed at 42°C for 16-18 h after adding the <sup>32</sup>Plabeled cDNA probe. Plasmids that contained cDNA inserts were labeled by nick translation for 28S and by random priming for PL, PLRP1, and PLRP2 (Promega labeling kits Prime-a-Gene and Nick translation). After hybridization, filters were washed under increasingly stringent conditions ( $1 \times SSC$ with 1% SDS to 0.2 M of sodium citrate with 1% SDS). The films were autoradiographed overnight at  $-80^{\circ}$ C. Autoradiographic films were quantified by an area laser densitometer (Bio-Rad) and volume integration. The data were expressed as relative absorbency units of each sample relative to the absorbency of 28S.

**Data analysis.** All data, expressed as mean  $\pm$  SE, were analyzed by two-way ANOVA (34) for the independent effects of *I*) diet, *2*) development, and *3*) the interaction effects of diet × development. Data for d9 were not included in the two-way ANOVA because only the LF group was sampled at this time. The mean for d9 LF is presented for comparison. Results were considered significantly different if P < 0.05.

# RESULTS

*Fatty acid composition of milk.* The milk fatty acid composition on d15 of lactation from rats that consumed LF (11%), MF (40%), and HF (67%) safflower oil diets is presented in Table 2. The amount of dietary safflower oil altered the milk fatty acid content with 15- to 100-fold C10:00 and 2.6- to 3.3-fold C18:2 in the milk of lactating rats that consumed MF and HF safflower oil diets, respectively, compared with lactating rats that consumed LF safflower oil diet. This change resulted in an increase in both long-chain/medium-chain ratio

 Table 2. Fatty acid composition of milk from lactating rats that

 consumed LF (11%), MF (40%), or HF (67%) safflower oil diets

 on day 15

	Diet		
Fatty acid	LF	MF	HF
10:0	24.7%	0.24%	1.6%
12:0	0.73%	5.5%	0.78%
14:0	0.55%	4.5%	0.59%
14:1	0.05%	0.0007%	Not detected
15:0	0.43%	0.003%	0.25%
15:1	0.07%	Not detected	Not detected
16:0	28.7%	11.7%	6.2%
16:1	2.19%	0.3%	0.07%
17:0	3.4%	0.16%	2.3%
17:1	0.06%	0.17%	Not detected
18:0	3.2%	2.8%	3%
18:1	13.5%	12.3%	11.7%
18:2	22%	58.9%	71.9%
18:3	0.13%	3.1%	1.3%
20:0	0.07%	4.1%	0.29%
Long/medium-chain ratio	2.8	8.6	72.5
P/S ratio	0.38	2.9	5.6

from 2.8 for LF to 72.5 for HF and the polyunsaturated/ saturated ratio (P/S) from 0.38 for LF to 5.6 for HF.

*Final body weight.* There was significant independent effect of diet (LF = MF > HF; P < 0.001) with a small (12 and 13.5%) decrease in the final body weight in weanling or adult rats fed HF safflower diets. There was also a significant effect of development (15d < 21d < 28d < 56d; P < 0.001) with increasing body weight from d15 through d56 (Fig. 1).

*Food consumption.* The average food consumption per day was measured in 28- and 56-d-old rats after they were weaned.



**Figure 1.** Final body weight of rats that consumed LF (11% as energy;  $\square$ ), MF (40% as energy;  $\blacksquare$ ), and HF (67% as energy;  $\square$ ) safflower oil diets. Results are expressed as mean  $\pm$  SE. Any bars without visible SE had SE less than the smallest unit on the graph. There was independent effect of diet (LF = MF > HF; P < 0.001; <sup>a,b</sup>diet not sharing a superscript differed significantly) and of development (d15 < d21 < d28 < d56; P < 0.001; <sup>c-f</sup>days not sharing a superscript differed significantly). There was no significant interaction of diet × development. The number of male pups per sample were 9 d (20 per group), 15 d (9–10 per group), 21 d (7–11 per group), 28 d (6–9 per group), and 56 d (8–12 per group).

 Table 3. Food consumption of rats that consumed LF, MF, or HF

 safflower oil diets

		Food intake (g/d)		
Day from birth	LF*	MF†	HF‡	
28*	9.6 ± 1.0	$9.0\pm0.4$	$7.0 \pm 1.0$	
56†	$21.0\pm0.6$	$19.0\pm1.5$	$17.0\pm1.4$	
D (	-			

Data are mean  $\pm$  SE.

There was an independent effect of diet (LF  $\leq$  MF  $\leq$  HF;  $P \leq$  0.001). Diets not sharing a symbol were significantly different.

There was an independent effect of days (d28 < d56; P < 0.001). Days not sharing a symbol were significantly different.

There was a significant independent effect of diet (LF < MF < HF; P < 0.001) and of development (28d < 56d; P < 0.001). The food consumption was significantly lower in 28- and 56-d-old rats that consumed the HF diet compared with rats that consumed the LF and MF diets (Table 3).

Pancreatic lipase activity. There was a significant independent effect of diet (LF < MF = HF; P < 0.002) and of development (15d < 21d < 28d = 56d; P < 0.001; Fig. 2) on pancreatic lipase activity expressed as U/mg pancreas. There was also a significant interaction of diet  $\times$  development (P < 0.001). In 15-d-old pups before weaning, PL activity was greater in pups that consumed LF and MF diets (55 and 65%, respectively) compared with pups that consumed the HF diet, although lipase activity was extremely low (6- to 20-fold less than d28). By d21, PL activity increased 6-fold and was greater in HF-fed rats (83 and 75%, respectively; P < 0.05) than in LF- or MF-fed rats. At 28 d, PL activity was 3-fold greater in HF- and MF-fed rats than in LF-fed rats (P < 0.05). A similar effect of diet was seen at 56 d with 2-fold greater PL activity in HF-fed rats compared with LF-fed rats and intermediate levels in MF-fed rats. As shown in Table 4, similar changes in PL activity are seen when PL activity is expressed as total units per pancreas.

**PL mRNA levels.** There were significant independent effects of diet (LF < MF = HF; P < 0.013) and of development (28d > 56d > 21d > 15d; P < 0.001) but no interaction of diet × development on PL mRNA levels (Fig. 3). PL mRNA levels were very low at 9 and 15 d but increased significantly by d21 through adulthood (2- to 4-fold, respectively). By d21, lipase mRNA levels increased 2- to 3-fold and were greater in the MF- and HF-fed rats. The maximal effect of diet was seen at d28.

**PLRP1 mRNA levels.** There was a significant independent effect of development (15d = 21d > 28d > 56d; P < 0.001). There was no independent affect of diet. The PLRP1 mRNA levels were very high during 9 and 15 d but decreased significantly by 28 d through adulthood (2- to 4-fold, respectively; Fig. 4).

**PLRP2 mRNA levels.** We report for the first time a significant independent effect of diet on the regulation of PLRP2 (P < 0.004; Fig. 5). PLRP2 mRNA levels decreased in HF-fed pups (79 and 90.3%, respectively) compared with LF- and MF-fed pups on d15. There was a significant effect of development (15d > 21d; P < 0.001). There was also a significant interaction of diet × development (P < 0.002). PLRP2 mRNA



**Figure 2.** Pancreatic lipase activity (U/mg of protein) of rats that consumed LF (11% as energy;  $\blacksquare$ ), MF (40% as energy;  $\blacksquare$ ), and HF (67% as energy;  $\square$ ) safflower oil diets. Results are expressed as mean  $\pm$  SE. Any bars without visible SE had SE less than the smallest unit on the graph. There was independent effect of diet (LF < MF = HF; *P* < 0.002; <sup>a,b</sup>diet not sharing a superscript differed significantly) and of development (d15 < d21 < d28 = d56; *P* < 0.001; <sup>c-e</sup>days not sharing a superscript differed significantly). There was significant interaction of diet × development (*P* < 0.001; <sup>f-k</sup>values not sharing a superscript differed significantly). The number of male pups per sample were 9 d (9 per group), 15 d (7–10 per group), 21 d (7–10 per group), 28 d (6–9 per group), and 56 d (6–12 per group).

 Table 4. Total pancreatic lipase of rats that consumed LF (11% of energy), MF (40% of energy), or HF (67% of energy) safflower oil diets

	Total pancreatic lipase (U/pancreas)				
Day	LF†	MF*	HF*		
9	13.3 ± 1.6	_	_		
15*	$10.3 \pm 2.0*$	$7.0 \pm 1.4*$	$5.9 \pm 1.6*$		
21†	$201.3 \pm 26.8 \dagger$	539.3 ± 101.5†‡	547.2 ± 85.4†‡		
28‡	$457.2 \pm 66.0$ †§‡	$878.5 \pm 206.0$ §¶‡	$980.4 \pm 256.8$ §¶‡		
56§	$1801.0 \pm 271.8$	$2491.8 \pm 367.2 \ $	$3649.0 \pm 512.4 \ $		

Data are mean  $\pm$  SE.

There was a significant independent effect of diet (LF < MF = HF; P < 0.02). Diets not sharing a symbol were significantly different.

There was a significantly independent effect of development (P < 0.001; d15 < d21 < d28 < d56). Days not sharing a symbol differed significantly.

There was significant interaction of diet  $\times$  development (P < 0.014). Values not sharing a symbol differed significantly.

not sharing a symbol dillered significantly.

levels decreased from d9 to d28 when the PLRP2 mRNA levels were undetectable.

## DISCUSSION

PL, PLRP1, and PLRP2 are of the same genetic family, as demonstrated by their gene homology and structural similarity (8,35,36). Both PLRP1 and PLRP2 gene expression have a different developmental pattern than PL mRNA (9). This anticoordinate regulation of PLRPs and PL suggests that during rat development, the genes for the PLRPs are under different regulatory controls than the PL gene. The relative high levels



**Figure 3.** Pancreatic lipase mRNA/S28 ratio of rats that consumed LF (11% as energy;  $\square$ ), MF (40% as energy;  $\blacksquare$ ), and HF (67% as energy;  $\square$ ) safflower oil diets. Results are expressed as mean ± SE. Any bars without visible SE had SE less than the smallest unit on the graph. There was independent effect of diet (LF < HF = MF; P < 0.013; <sup>a,b</sup>diet not sharing a superscript differed significantly) and of development (d15 < d21 < d56 < d28; P < 0.001; <sup>c-f</sup>days not sharing a superscript differed significantly). The number of male pups per sample were 9 d (12 per group), 15 d (7–10 per group), 21 d (6 per group), 28 d (6–9 per group), and 56 d (8–12 per group).

of PLRP mRNA around birth suggest that they may play an important role in the suckling and suckling-nibbling phase, when the rat is consuming maternal milk and maternal diet.



**Figure 4.** PLRP1 mRNA/S28 ratio of rats that consumed LF (11% as energy; ), MF (40% as energy; ), and HF (67% as energy; ) safflower oil diets. Results are expressed as mean  $\pm$  SE. Any bars without visible SE had SE less than the smallest unit on the graph. There was independent effect of development (d56 < d28 < d21 < d15; P < 0.001; <sup>a-c</sup>days not sharing a superscript differed significantly). There was no significant effect of diet and interaction of diet × development. The number of male pups per sample were 9 d (15 per group), 15 d (7–10 per group), 21 d (6 per group), 28 d (6–8 per group), and 56 d (6–11 per group).



**Figure 5.** PLRP2 mRNA/S28 ratio of rats that consumed LF (11% as energy; ), MF (40% as energy; ), and HF (67% as energy; ) safflower oil diets. Results are expressed as mean  $\pm$  SE. Any bars without visible SE had SE less than the smallest unit on the graph. There was independent effect of diet (LF = MF > HF; P < 0.004; <sup>a,b</sup>diet not sharing a superscript differed significantly) and of development (d21 < d15; P < 0.001; <sup>c-d</sup>days not sharing a superscript differed significantly). There was significant interaction of diet × development (P < 0.002; <sup>c-g</sup>values not sharing a superscript differed significantly). The number of male pups per sample were 9 d (10 per group), 15 d (8–10 per group), and 21 d (9 per group).

Altering the maternal dietary fat content did alter the milk fatty acid content in dams that consumed low, moderate, and high levels of safflower oil. The amount of PUFA (mainly 18:2) in milk was elevated as maternal dietary polyunsaturated fat increased. This phenomenon is in agreement with previous studies (23–25).

This significant change in the milk fatty acid composition consumed by the pups allowed us to determine whether PL and its related proteins are regulated by dietary fat in the suckling phase and the suckling-nibbling phase, when the pups consume both maternal milk and purified diet. Increased polyunsaturated fat, independent of developmental stage, significantly increased PL activity and mRNA in a parallel manner as described previously (MF = HF > LF). However, there was also a complex interactive effect between diet and developmental stage. At the suckling-nibbling phase (d15), high polyunsaturated fat in suckled milk or nibbled maternal diet decreased PL activity in pups without changing its mRNA levels. This suggests that this early dietary regulation of PL activity in the suckling-nibbling phase may be due to translational or posttranslational regulation. This is also the first report of dietary regulation of PL in the preweaned rat, as previous studies on adaptation of PL to dietary fat all have been in weaned or adult animals. The parallel changes found in PL mRNA levels and activity in the weanling and adult rat in this study are concordant with previous studies (20,22,37) and suggest that the dietary regulation of PL postweaning is pretranslational or transcriptional. PL exhibited the previously reported (9) developmental regulation, independent of diet; PL activity and mRNA levels were low at d9 and d15 and increased to maximal levels by d28.

PLRP1 mRNA levels were significantly affected by development in agreement with previous reports by Payne *et al.* (9). PLRP1 mRNA levels were high during the early postnatal period (d9 and d15) and declined sharply after weaning (d21) to low levels in adulthood. The different amount of polyunsaturated fat in the milk and the diet did not affect expression of PLRP1. The lack of influence of amount of fat in diet on PLRP1 expression in weanling rats in this work is surprising and differs from other studies (16,20,22,37), in which PLRP1 was shown to be transcriptionally regulated by dietary fat. However, the introduction of different amounts of PUFA, started earlier during the tertiary phase of pancreatic development, may alter the long-term regulation of PLRP1. Such "metabolic imprinting" has been proposed for other dietary lipids such as cholesterol but remains highly controversial.

The developmental expression of PLRP2 mRNA was high during the early postnatal period and declined sharply until the weaning phase (21 d postnatally) to undetectable levels in adulthood (d56). This agrees with the developmental pattern reported initially by Payne et al. (9). As early as, 15 d during the suckling-nibbling phase, there was a significant effect of diet (LF = MF > HF; P < 0.004) on PLRP2 mRNA levels. Pups that consumed milk with high levels of PUFA had significantly reduced PLRP2 mRNA levels compared with pups that consumed milk with low or moderate levels of PUFA. This is the first report of regulation of PLRP2 by different amounts of polyunsaturated fat introduced early during development. Given the report by Lowe et al. (5) that suckling PLRP2-deficient mice exhibit fat malabsorption and the suppression of PLRP2 by high levels of long-chain polyunsaturated fat in the early postnatal period, found in our study, it is clear that PLRP2 has an important role in digestion of fat at the early postnatal period.

In conclusion, three pancreatic lipases (PL, PLRP1, and PLRP2) are differentially regulated by diet. PL and PLRP2 are regulated anticoordinately at the suckling stage by dietary PUFA. This dietary regulation of these two enzymes at a very

early stage of development may have significance in the digestion of fat in the newborn. Diets with high levels of polyunsaturated fat and low levels of saturated fat suppress PLRP2. Whether this has an adverse impact on fat digestion and absorption needs to be determined in future studies. When considering the appropriate balance of PUFA and other fats for optimal fat digestion in early neonates, the developmental and dietary regulation of the different lipases, including PL and PLRP2, should taken into consideration.

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