

Transient changes of transforming growth factor- β expression in the small intestine of the pig in association with weaning

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It is well known that early weaning causes marked changes in intestinal structure and function, and transforming growth factor- β (TGF- β) is believed to play an important regulatory role in post-weaning adaptation of the small intestine. The present study examined the distribution and expression intensity of TGF- β in the small intestinal mucosa of pre- and post-weaning pigs using a specific immunostaining technique and Western blot analysis. The level of TGF- β in the intestinal mucosa, as estimated by Western blot analysis, did not change significantly during weaning. However, when examined by the immunostaining technique, TGF- β 1 (one of the TGF- β isoforms dominantly expressed in the tissue) at the intestinal villus epithelium, particularly at the apical membrane of the epithelium, decreased significantly 4 d after weaning, while the staining intensity increased significantly at the intestinal crypts compared with that in pre-weaning pigs. These changes were transient, with the immunostaining intensity for TGF- β 1 at the intestinal villi and the crypts returning to the pre-weaning level by 8 d post-weaning. The transient decrease in TGF- β 1 level at the intestinal villus epithelium was associated with obvious intestinal villus atrophy and marked reduction of mucosal digestive enzyme activities. Furthermore, the number of leucocytes staining positively for TGF- β 1 increased significantly in the pig intestinal lamina propria 4 d after weaning. These findings strongly suggest that TGF- β plays an important role in the post-weaning adaptation process in the intestine of the pig.

Transforming growth factor- β : Intestine: Weaning pig

It is well known that early weaning causes marked changes to the histology and biochemistry of the small intestine, such as villus atrophy and crypt hyperplasia, in both laboratory animals (Pluske *et al.* 1997) and human infants (Thompson *et al.* 1998). The change in intestinal structure often leads to reduced digestive and absorptive capacity and contributes to post-weaning diarrhoea. The aetiology of the intestinal structural alteration is complex, including enteropathogens, hypersensitivity to dietary antigens and withdrawal of milk-borne growth factors (Pluske *et al.* 1997; Xu *et al.* 2000). In most animals, the small intestine adapts to the weaning event and returns to normal functional capacity within 2 weeks after weaning (Hampson & Kidder, 1986; Pluske *et al.* 1997). The mechanisms controlling the adaptation process are not yet clear, but transforming growth factor- β (TGF- β) has been suspected to be an important modulator of postnatal intestinal adaptation (Xu *et al.* 2000).

TGF- β is a multifunctional polypeptide expressed by various cells in the intestinal mucosa, including the epithelial cells lining the intestinal villi and the lymphocytes in the lamina propria (Barnard *et al.* 1993). It is synthesized initially as an inactive precursor of 390–412 amino acids and the mature active peptide consists of 112 amino acids (Yue & Mulder, 2001). TGF- β is secreted from cells in a latent complex consisting of TGF- β in association with latent proteins and it needs to be activated to exert biological function (Roberts, 1995). TGF- β regulates enterocyte proliferation and differentiation and helps to maintain

intestinal integrity of the epithelial surface along the villi (Dignass & Sturm, 2001). During injury or disease, TGF- β stimulates epithelial cell migration (Ciacci *et al.* 1993) and extracellular matrix production (O’Kane & Ferguson, 1997), thereby promoting wound healing. TGF- β is also known as a potent immunoregulator and plays a critical role in maintaining mucosal immune homeostasis (Letterio & Roberts, 1998), and local production of TGF- β in the intestinal mucosa increases in response to mucosal inflammation (Babyatsky *et al.* 1996). These findings indicate that TGF- β may be involved in relieving weaning-associated villus damage and mucosa inflammation. The aim of the present study was to examine the endogenous TGF- β expression and its distribution in the small intestine of the pig during the weaning period.

Materials and methods

Tissue sample collection

Twelve Large White \times Landrace pigs from two litters were used in the present study. The animals were allowed access to creep feed from 14 d of age. At 21 d of age, all animals were removed from the sows, four of the pigs designated pre-weaning pigs were killed for tissue sample collection within 2 h after removal and the remaining eight pigs were weaned to a soyabean- and maize-based weaning diet containing 22.2% crude protein. Four of the weaned pigs were killed for tissue sample collection 4 d

after weaning and the remaining four weaned pigs were killed 8 d after weaning. Prior to tissue sample collection, all experimental animals were first anaesthetized by peritoneal injection of pentobarbital sodium (Alfasan, Wberden, Holland) and then euthanased by intracardiac injection of an overdose of the drug. Immediately after death, the small intestine was removed and placed in chilled saline. The small intestine was then freed from the mesentery and divided into the duodenum, jejunum and ileum as described previously (Xu *et al.* 1992).

A block of tissue (about 1 cm in length) was taken from the middle region of each intestinal segment for histological studies. Intestinal mucosa samples were collected from each segment of the small intestine by scraping with a glass slide. The tissue samples were immediately frozen in liquid N and then stored at -70°C until used for further analyses of enzyme activities, protein and DNA contents and Western blot analysis.

Immunohistochemical examination

Tissue blocks collected for immunohistochemical study were immediately fixed in Bouin's fluid for 24 h. The tissue blocks were then dehydrated through graded alcohol and embedded in paraffin wax. Cross-tissue sections of $5\ \mu\text{m}$ thickness were dewaxed in xylene, re-hydrated in alcohol solution of increased dilution, and then immersed in PBS. Endogenous peroxidase activity was eliminated by incubation with 1% H_2O_2 for 30 min, and non-specific binding was blocked by incubation for 30 min with 2% normal goat serum (DAKO A/S, Glostrup, Denmark). The tissue sections were then incubated overnight at 4°C with rabbit polyclonal antibodies specifically against TGF- β 1, β 2 or β 3. The antibodies were obtained from a commercial source (Santa Cruz Biotech, Santa Cruz, CA, USA) and were raised against a carboxy-terminus peptide of 15–26 amino acids of human TGF- β 1, β 2 or β 3. According to the information provided by the supplier, the antibodies detect both precursor and mature TGF- β 1, β 2 or β 3 from various species including the pig. After three washes in PBS, sections were further incubated for 2 h at room temperature with biotin-conjugated secondary antibodies, followed by three washes in PBS, and incubation for 2 h at room temperature with streptavidin–biotinylated horseradish peroxidase (Amersham Pharmacia Biotech, San Francisco, CA, USA). Subsequently, the slides were developed in 3,3-diaminobenzidine solution (0.5 mg/ml) in the presence of 0.1% H_2O_2 for 10 min. After washing in distilled water, the slides were counterstained with haematoxylin (Sigma, St Louis, MO, USA). The negative control sections were stained with the same procedures except that the primary antibodies were replaced with non-specific rabbit IgG (DAKO A/S). The tissue sections were then examined under light microscope (Carl Zeiss Inc., Werk Göttingen, Germany), and positive staining was visualized as dark brown spots.

The immunostaining intensity was semi-quantified following the procedures described by Ruifrok *et al.* (1997). The method has been validated in our laboratory recently (Mei *et al.* 2004). Briefly, TGF- β immunoreactivity was quantified using a computerized image-analysis system (Quantimet 500; Leica Cambridge Ltd, Cambridge, UK). This system consisted of an IBM-compatible computer with image-analysis software, a Leica microscope (Leica Cambridge Ltd.), a colour video camera and a 'live image' monitor. The image was converted to an analogue electronic signal by the video camera and digitized by the computerized imaging board. The mean specific optical density (OD), reflecting the

immunostaining intensity, was calculated from the formula: $\text{OD} = \log_{10}[255/(255 - \text{grayscalevalue})]$, as described by Ruifrok *et al.* (1997). The integrated OD, which is the product of the area stained and the mean specific OD, was taken as a semi-quantitative measure of the total amount of TGF- β protein present.

Morphological analyses

Paraffin tissue sections ($5\ \mu\text{m}$) were stained with haematoxylin and eosin. Following previous descriptions (Xu *et al.* 1992), morphological parameters, including the wall thickness, villus height, villus width and crypt depth, were analysed under a light microscope (Carl Zeiss Inc.) connected to an image processing and analysis system (Quantimet 500; Leica Cambridge Ltd).

Measurements of digestive enzyme activities and protein and DNA contents

Samples of small intestinal mucosa were homogenized with chilled saline using a polytron homogenizer (Kinematica AG, Littau/Lucerne, Switzerland). The activities of the sucrase (E.C. 3.2.1.48) and lactase (E.C. 3.2.1.23) in the small intestinal mucosa were determined by the method described by Dahlqvist (1964). Alkaline phosphatase (orthophosphoric monoester phosphohydrolase, EC 3.1.3.1) was determined by the method of Forstner *et al.* (1968). Protein concentration was determined with the Lowry method using bovine serum albumin as standard (Lowry *et al.* 1951). DNA was extracted from the tissue homogenates following the procedures described by Johnson & Chandler (1973) and the DNA content was determined using the diphenylamine method (Giles & Myers, 1964). Calf thymus DNA (Sigma) was used as standard.

Western blot analysis

Small intestine mucosa samples were defrosted and homogenized at 4°C in 50 mM-Tris-HCl buffer, pH 6.8, containing 5 mM-EDTA, 10 $\mu\text{g}/\text{ml}$ phenylmethylsulphonyl fluoride and 0.2% NP-40, using a polytron homogenizer (Kinematica AG). Tissue debris was removed by centrifugation at 4500 rpm for 30 min. Protein concentrations of the tissue homogenates were measured with the Lowry method as described previously (Xu *et al.* 1992).

Molecular characteristics of immunoreactive TGF- β peptides in the tissue homogenates were evaluated by SDS-PAGE and Western blot analysis as described by Rao *et al.* (2000). Briefly, tissue homogenates were diluted in 50 mM-Tris-HCl buffer, pH 6.8, containing 2% SDS, 10% glycerol, 5% 2-mercaptoethanol and 0.05% bromophenol blue, and boiled for 5 min. The prepared samples of equal amounts of protein (50 μg) were then loaded onto 12% SDS-PAGE gel and separated in an electrophoresis unit (Mini-Protein II Cell; Bio-Rad, Hercules, CA, USA) at 120 V (constant voltage) for 1–2 h. Separated proteins were then transferred onto nitrocellulose membranes (Amersham Pharmacia Biotech) at 100 V/250 mA for 120 min at 4°C . The nitrocellulose membranes were subsequently blocked with Tris-buffered saline (100 mM-Tris-HCl, pH 7.5, 0.9% NaCl) containing 0.1% Tween 20 and 5% skimmed milk powder for 2 h at room temperature followed by overnight incubation at 4°C with polyclonal antibodies against TGF- β 1, 2 or 3, respectively (Santa Cruz Biotech). The membranes were then washed with Tris-buffered saline, followed by further 2 h incubation at room temperature with biotin-conjugated goat anti-

rabbit IgG (Santa Cruz Biotech). After removing non-bound antibody, the membranes were then incubated for 2 h in a solution of streptavidin–biotinylated horseradish peroxidase complex. Subsequently, the membranes were developed in 3,3-diaminobenzidine solution (0.2 mg/ml) in the presence of 0.1 % H_2O_2 for 30 min. The membranes were then air-dried and stored in light-protected boxes at room temperature until further examination. The molecular weights of the immunoreactive bands were determined using a molecular weight standard (Invitrogen, Carlsbad, CA, USA).

For quantitative analysis, the stained membranes were scanned and the intensity of the positive bands was measured using Bio-Rad's image-analysis system. The measured intensity of each sample was calibrated and converted to mg TGF- β /g protein and mg TGF- β /mg DNA by comparison with the intensity of known amounts of TGF- β proteins (Santa Cruz Biotech).

Statistics

The results are expressed as mean and standard error of the mean. One-way ANOVA was used to compare variances among various groups. Statistical differences between two groups were assessed by the least significant difference test. All analyses were conducted using SPSS for Windows software, version 11.0 (SPSS Inc., Chicago, IL, USA). A difference with P value < 0.05 was considered significant.

Results

Changes of intestinal morphology and digestive enzyme activities in association with weaning

Weaning caused significant changes in the intestinal morphology and digestive enzyme activities, as shown in Figs. 1 and 2 and Tables 1 and 2. The changes were as expected and observations were consistent with reports in the literature.

Changes of transforming growth factor- β expression and distribution in the small intestine

In pre-weaning pigs, immunostaining with specific antibodies for TGF- β 1 was seen on both apical and basal membranes of the villus epithelium in the duodenal region of the small intestine (Fig. 1). The same staining pattern was seen in the jejunal region of the small intestine (photo not shown). However, in the ileal region of the small intestine, immunostaining with TGF- β 1-specific antibodies was observed predominantly on the basal side of the villus epithelium (Fig. 1). There were numerous leucocytes that were positively stained with TGF- β 1 antibodies and scattered in the lamina propria of the small intestine (Fig. 2). The immunostaining patterns with antibodies specific for TGF- β 2 and TGF- β 3 at the villus epithelium were similar to those of staining with specific antibodies for TGF- β 1. The epithelial cells at the intestinal crypts along the small intestine were positively stained with antibodies against the three TGF- β isoforms. No staining was observed in duodenal Brunner's glands.

Four days after weaning, the immunostaining for TGF- β 1 at the apical membrane of the villus epithelium became weaker in the duodenal and jejunal region of the small intestine compared with that in pre-weaning pigs (Fig. 1), while the staining for TGF- β 1 at the intestinal crypts became much stronger (Fig. 2). No obvious distribution change was noted in the ileal villus epithelium following weaning (Fig. 1). Similar to that in the duodenum and the jejunum, the epithelial cells at the ileal crypts stained strongly for TGF- β 1 in pigs 4 and 8 d after weaning. In the lamina propria of the small intestine, the number of leucocytes positively stained with TGF- β 1 antibodies increased significantly 4 d after weaning (Fig. 3).

Quantification of immunohistochemical staining by image analysis showed a transient decline in the positive staining area and integrated staining intensity (OD) for TGF- β 1 at the villus epithelium in the duodenal and jejunal region of the pig small intestine following weaning (Fig. 4), while the integrated OD

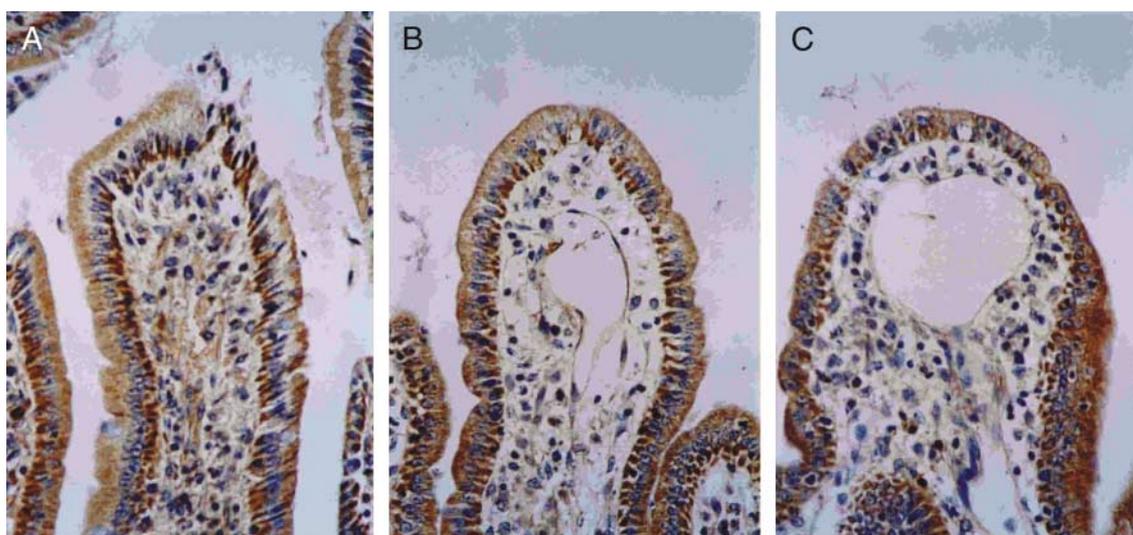


Fig. 1. Representative photomicrographs of duodenal (upper panels) and ileal (lower panels) villi from pre-weaning (A), 4 d post-weaning (B) and 8 d post-weaning (C) pigs (magnification $350\times$). Tissue sections were incubated with polyclonal antibodies against transforming growth factor- β 1 (TGF- β 1), and colour was developed using the avidin–peroxidase system. In the duodenum, immunostaining of TGF- β 1 was seen on both apical and basal membrane of the villus epithelium in pre-weaning pigs. The immunostaining intensity at the apical membrane declined markedly in 4 d post-weaning pigs and then recovered to the pre-weaning value in 8 d post-weaning pigs. In the ileum, TGF- β 1 was located predominantly on the basal side of the villus epithelial membranes in pre-weaning pigs. The immunostaining declined gradually following weaning.

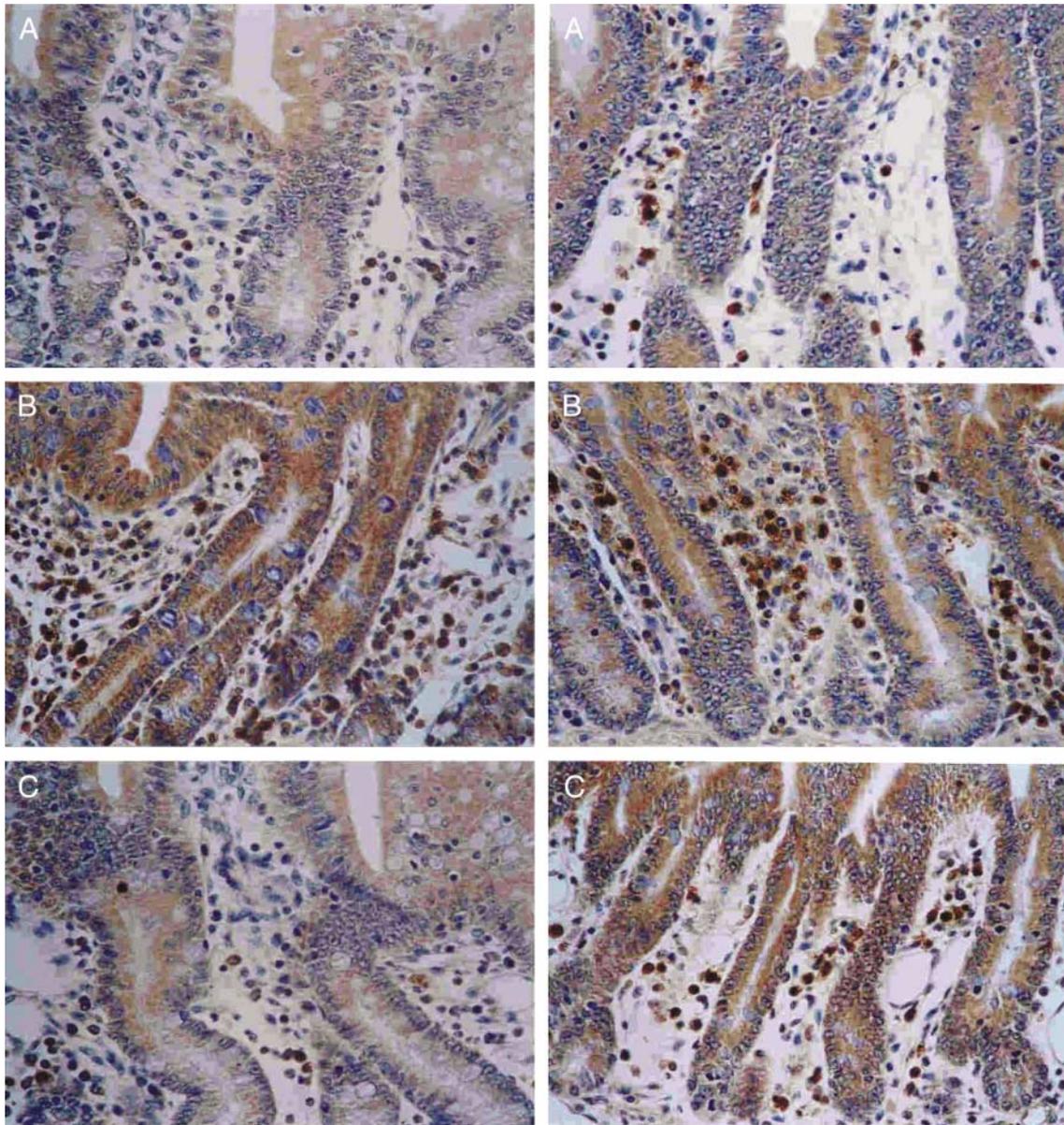


Fig. 2. Representative photomicrographs of duodenal (left panels) and ileal (right panels) crypts from pre-weaning (A), 4 d post-weaning (B) and 8 d post-weaning (C) pigs (magnification $350\times$). Tissue sections were incubated with polyclonal antibodies against transforming growth factor- β 1 (TGF- β 1), and colour was developed using the avidin–peroxidase system. In the duodenum, the immunostaining at the crypt epithelium became much stronger in 4 d post-weaning pigs. In the ileum, there was an increase in staining for TGF- β 1 following weaning. The number of cells stained positively for TGF- β 1 in the lamina propria increased following weaning.

for TGF- β 1 at the crypts increased significantly 4 d after weaning. The integrated OD for TGF- β 1 at intestinal villi and crypts returned to the pre-weaning value by the eighth day after weaning (Fig. 4). In the ileal region of the small intestine, the OD for TGF- β 1 at villus epithelium decreased progressively until the eighth day after weaning. In contrast, the staining intensity for TGF- β 1 at the ileal crypts increased progressively after weaning (Fig. 4). The immunostaining intensities for TGF- β 2 and β 3 were relatively constant in most segments of the small intestine and did not change significantly following weaning (Fig. 4).

Western blot analysis revealed a positively stained band in all homogenized mucosal tissue samples and the band corresponded to the standard proteins of TGF- β 1, β 2 and β 3. The expression levels of TGF- β 1, β 2 and β 3 in homogenized small intestinal

mucosa were also measured by Western blot analysis, and showed no significant changes in association with weaning (Fig. 5).

Discussion

Weaning is a physiologically important event that influences intestinal structure and function in the pig. Reports on the small intestine changes in morphology and brush-border hydrolase activities that occur in the weaning period are numerous. Studies in both laboratory animals (Pluske *et al.* 1997; van Beer-Schreurs *et al.* 1998) and human infants (Thompson *et al.* 1998) have shown that weaning is associated with villus atrophy, crypt hyperplasia and decrease in hydrolase-specific activities. The most severe reduction in villus height was observed in pigs 2–5 d post-weaning (Kelly *et al.*

Table 1. The effects of weaning on intestinal morphology in the pig (Mean values and standard error of the mean for four animals in each group)

Region	Pre-weaning pigs		4 d post-weaning pigs		8 d post-weaning pigs	
	Mean	SEM	Mean	SEM	Mean	SEM
Duodenum						
Villus height (μm)	221.1 ^a	9.4	122.4 ^b	5.1	183.6 ^c	8.8
Villus width (μm)	50.8 ^a	1.6	63.2 ^b	2.1	58.4 ^{ab}	1.0
Crypt depth (μm)	85.5 ^a	3.4	113.4 ^b	4.6	87.6 ^a	2.7
Villus: crypt ratio	2.6 ^a	0.1	1.1 ^b	0.0	2.2 ^a	0.2
Jejunum						
Villus height (μm)	226.6 ^a	9.4	126.9 ^b	3.5	142.7 ^c	1.9
Villus width (μm)	52.9	1.3	58.3	1.0	56.9	1.2
Crypt depth (μm)	78.2 ^{ab}	4.5	83.6 ^a	1.8	76.4 ^b	2.3
Villus: crypt ratio	3.1 ^a	0.3	1.5 ^b	0.0	1.9 ^c	0.1
Ileum						
Villus height (μm)	155.5 ^a	3.2	117.6 ^b	3.5	120.6 ^b	6.3
Villus width (μm)	55.1 ^a	1.2	55.1 ^a	0.9	65.5 ^b	1.7
Crypt depth (μm)	66.3 ^a	1.5	79.5 ^b	1.0	72.8 ^c	2.7
Villus: crypt ratio	2.4 ^a	0.1	1.5 ^b	0.1	1.7 ^b	0.1

^{a,b,c} Mean values in the same row with unlike superscript letters were significantly different among animal groups: $P < 0.05$.

1991; van Beers-Schreurs *et al.* 1998). In agreement with earlier reports, the present study showed that weaning caused significant villus shortening and crypt hyperplasia (Table 1) as well as significant declines in intestinal digestive enzyme activities (Table 2).

The causes of weaning-associated intestinal dysfunction have been a focal point of intensive research and have been related to reduced food intake, enteropathogens, hypersensitivity to dietary antigens and withdrawal of milk-borne growth factors (Pluske *et al.* 1996, 1997; McCracken *et al.* 1999). One of the milk-borne growth factors, TGF- β , is of particular interest as it has known effects on enterocyte proliferation and differentiation and it modulates intestinal mucosal immune reactions (Massague, 1990). It has been speculated that TGF- β may play an important role in post-weaning adaptation of the intestine (Xu *et al.* 2000). An earlier study in the rat showed that during the period of weaning the level of TGF- β in maternal milk decreased markedly while the endogenous production of TGF- β in the small intestine of the pup increased significantly (Penttila *et al.* 1998). In the present study, the expression levels of TGF- β in the intestinal mucosa, when

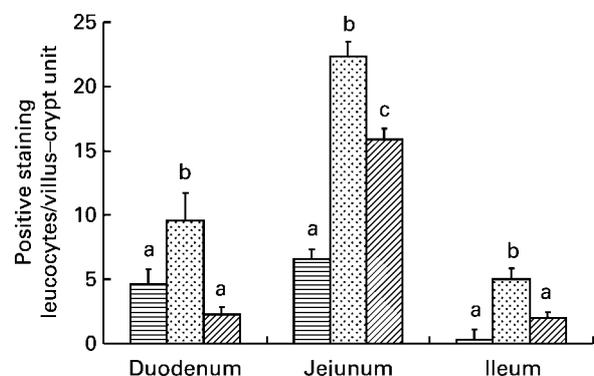
estimated by Western blot analysis, did change significantly during weaning in pigs (Fig. 5). However, when examined by an immunostaining technique, the expression intensity for TGF- β 1 at the intestinal villus epithelium, particularly at the apical membrane of the epithelium, decreased significantly in pigs 4 d after weaning while the staining intensity at the intestinal crypts increased significantly in the same animals (Fig. 4). This finding indicates a topical change of TGF- β expression in the small intestine of pigs during weaning and the change may relate to the withdrawal of milk-borne TGF- β after weaning.

Our observation of topical changes in TGF- β 1 expression in the small intestine in association with weaning, with a transient increase in TGF- β expression in the intestinal crypts and a decrease in the intestinal villi, is in contrast with the earlier report in rats that TGF- β 1 expression was most abundant in terminally differentiated villus tip cells and least abundant in the less differentiated crypt cells (Barnard *et al.* 1989). It has also been reported in the pig that the expression of inflammatory cytokines such as IL-1, IL-6 and TNF- α increases in the intestine during weaning (McCracken *et al.* 1995; Pie *et al.* 2004). The apparent discrepancy between the present observations and the earlier

Table 2. The specific activities ($\mu\text{mol/g}$ protein per min) of lactase, sucrase and alkaline phosphatase in intestinal mucosa of pre- and post-weaning pigs (Mean values and standard error of the mean for four animals in each group)

Enzyme	Pre-weaning pigs		4 d post-weaning pigs		8 d post-weaning pigs	
	Mean	SEM	Mean	SEM	Mean	SEM
Lactase						
Duodenum	249.8 ^a	5.8	101.9 ^b	9.2	49.8 ^c	1.2
Jejunum	267.8 ^a	10.2	100.4 ^b	8.5	105.9 ^b	16.7
Ileum	208.1 ^a	26.6	34.6 ^b	3.6	9.5 ^c	1.9
Sucrase						
Duodenum	48.7 ^a	3.1	14.0 ^b	1.5	9.2 ^b	1.2
Jejunum	84.0 ^a	2.2	35.8 ^b	4.5	46.9 ^b	5.9
Ileum	89.8 ^a	7.9	35.7 ^b	1.8	14.8 ^c	1.2
Alkaline phosphatase						
Duodenum	3.7 ^a	0.1	1.3 ^b	0.1	1.6 ^b	0.1
Jejunum	3.5 ^a	0.3	1.3 ^b	0.1	1.1 ^b	0.1
Ileum	4.6 ^a	0.7	1.7 ^b	0.2	0.7 ^b	0.1

^{a,b,c} Mean values in the same row with unlike superscript letters were significantly different among animal groups: $P < 0.05$.

**Fig. 3.** The number of leucocytes staining positively for transforming growth factor- β 1 per villus-crypt unit in the intestinal lamina propria in pre-weaning (□), 4 d post-weaning (▨) and 8 d post-weaning (▩) pigs. Values are means with the standard error of the mean shown by vertical bars for four animals in each group. ^{a,b,c} Mean values with unlike superscript letters were significantly different among animal groups: $P < 0.05$.

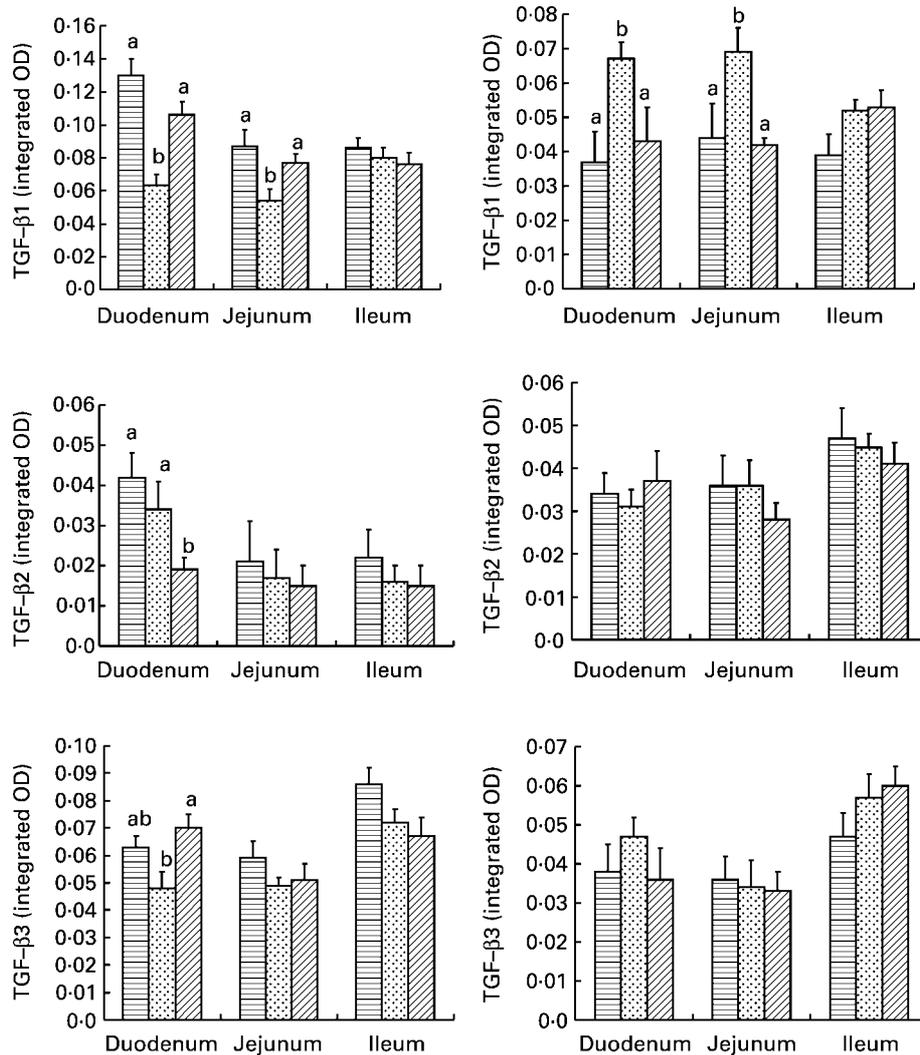


Fig. 4. The levels of transforming growth factor (TGF)-β1, TGF-β2 and TGF-β3 in the villi (left panels) and crypts (right panels) of pre-weaning (□), 4 d post-weaning (▨) and 8 d post-weaning (▩) pigs estimated by immunostaining image analysis (for details see p. 38). Values are means, with the standard error of the mean shown by vertical bars for four animals in each group, of the integrated optical density (OD), which is the product of the area stained and the mean specific OD, and represents immunostaining intensity. ^{a,b,c}Mean values with unlike superscript letters were significantly different among animal groups: *P* < 0.05.

reports may be related to species and cytokine specificities. It is well known that the expression and the effect of different cytokines are tissue-specific. Further examination of the present results (Fig. 4) also showed that the expression level of TGF-β1 was much greater in intestinal villi than in intestinal crypts in pre-weaning pigs, which is inconsistent with the earlier report in rats (Barnard *et al.* 1989).

The relationship between the decline in TGF-β level of the intestinal villus epithelium and the reduction in intestinal villus height and mucosal digestive enzyme activities in post-weaning pigs is unclear. It has been reported in the literature that TGF-β plays an important role in maintaining intestinal mucosa integrity and that it stimulates the mucosal wound healing process. TGF-β stimulates the migration of epithelial cells from the wound margin and enhances rapid intestinal epithelial restitution (Ciacci *et al.* 1993). It has also been shown that TGF-β enhances wound closure by stimulating collagen synthesis (Perr *et al.* 1996) and increasing the deposition of extracellular matrix (O’Kane & Ferguson, 1997). Oral supplementation with a bovine milk fraction enriched for TGF-β2 significantly reduced intestinal villus atrophy in rats

exposed to the chemotherapy drug methotrexate (van’t Land *et al.* 2002). From the earlier discussion, it may be speculated that the transient decline in TGF-β1 level of the intestinal villus epithelium following weaning contributed to the post-weaning intestinal villus atrophy and to the reduction of digestive enzyme activities.

It was also observed in the present study that the number of TGF-β1-positive leucocytes in pig intestinal lamina propria increased significantly 4 d after weaning (Fig. 3). The increment is likely due to an increased mucosal immune reaction to food antigens following weaning. It has been reported that the gut of early-weaned pigs reacts to food antigens in a hypersensitive manner (Li *et al.* 1990) and the local production of TGF-β in gut-associated lymphoid tissue increases significantly in response to mucosal inflammation in patients with ulcerative colitis or Crohn’s disease (Babyatsky *et al.* 1996; Hart *et al.* 2004). In mice TGF-β production in gut-associated lymphoid tissue increased markedly following oral antigen challenge (Gonnella *et al.* 1998).

The transient increase of TGF-β1-positive leucocytes in the intestinal mucosa in post-weaning pigs may play an important

TGF- β expression in weaning pig intestine

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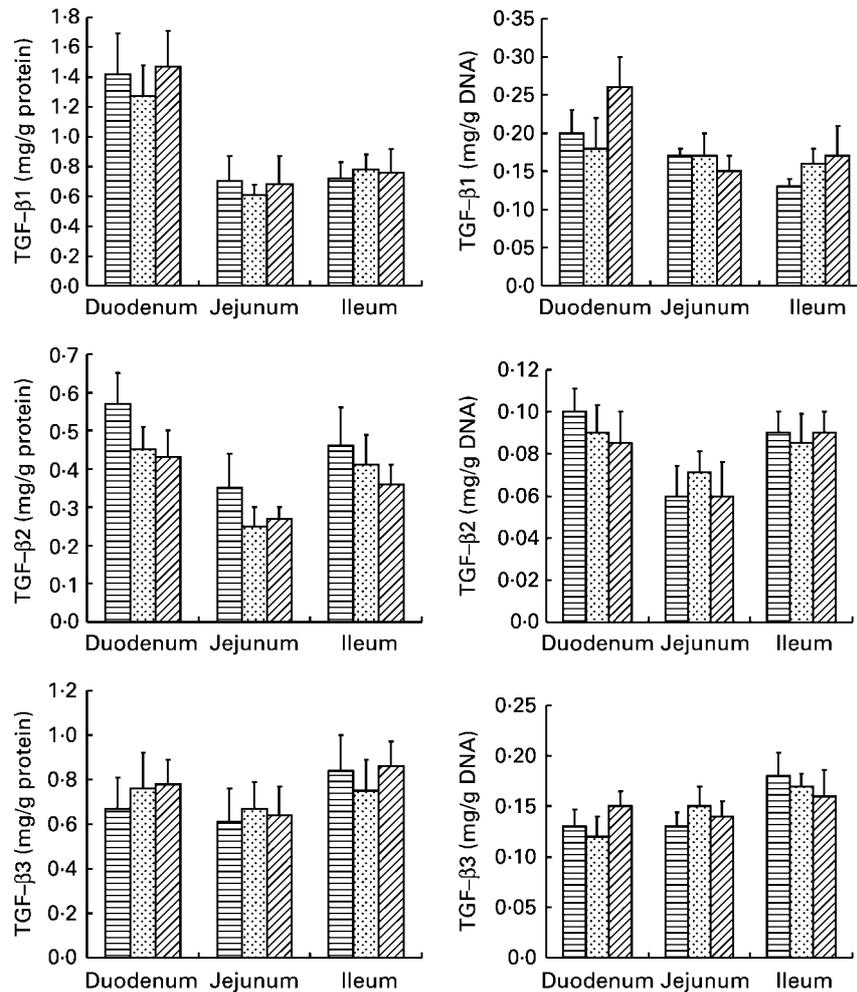


Fig. 5. The levels of transforming growth factor (TGF)- β 1, TGF- β 2 and TGF- β 3 in intestinal mucosa homogenates of pre-weaning (■), 4 d post-weaning (▨) and 8 d post-weaning (▩) pigs estimated by Western blot analysis. Values are means, with the standard error of the mean shown by vertical bars for four animals in each group, of mg TGF- β /g protein (left panels) and mg TGF- β /mg DNA (right panels). There were no significant differences in TGF- β levels among animal groups.

role in the process of post-weaning intestinal adaptation. TGF- β is an important cytokine controlling mucosal inflammation and immune response. It has been shown that TGF- β inhibits leucocyte proliferation and activation (Letterio & Roberts, 1998). Abrogation of TGF- β signalling in T cells induces T-cell-mediated inflammatory lesions in various organs including the intestine (Gorelik & Flavell, 2000). *In vitro* studies have shown that TGF- β inhibits macrophages and glioma cells from expressing MHC class II antigens (Czarniecki *et al.* 1988). TGF- β 1 and TGF- β 2 decrease secretion of pro-inflammatory cytokines (IL-6, IL-8) induced by both TNF- α and IL-1 β in a human intestinal epithelial cell line (Claud *et al.* 2003; Walia *et al.* 2003). Mice with TGF- β 1 deficiency die from cardiac, pulmonary and gastric inflammation, suggesting that TGF- β has a vital role in suppressing the activation and proliferation of inflammatory cells (Shull *et al.* 1992). TGF- β also plays an important role in oral tolerance. Oral administration of recombinant TGF- β or milk whey extract containing a high concentration of TGF- β to neonatal mice strongly inhibits the immune response to an oral antigen challenge (Penttila *et al.* 2001).

The present study did not find significant changes in TGF- β 2 and β 3 expression in the small intestine of the pig during the weaning period. The significance of the different expression

patterns of the three TGF- β isoforms is unknown. Although highly homologous in structure and similar in biological activities in many bioassay systems, the TGF- β isoforms are encoded by different genes and have different distributions and biological actions within the body (Blobe *et al.* 2000). TGF- β 1 is the most abundant isoform in all tissues and in human platelets. More than 85% of the TGF- β in the wound fluid in adult animals is TGF- β 1. TGF- β 2 is the most abundant isoform in body fluids, including the aqueous and vitreous fluid of the eye, saliva, amniotic fluid and breast milk, while TGF- β 3 is usually the least abundant isoform in both tissues and body fluids (Massague, 1990). The different physiological functions of TGF- β isoforms have been demonstrated in gene knockout mice. TGF- β 1 null mice can survive in the perinatal period when nursed by their dams but die of excessive inflammatory reactions after weaning (Shull *et al.* 1992). In contrast, TGF- β 2 null mice display a wide range of developmental heart, lung, craniofacial, limb, spinal column, eye, inner ear and urogenital defects (Sanford *et al.* 1997), while TGF- β 3 null mice die during the early postnatal period owing to defects in pulmonary development and palatogenesis (Proetzel *et al.* 1995). Neither TGF- β 2 nor TGF- β 3 can overcome the deficit of TGF- β 1 (McCartney-Francis & Wahl, 1994). It has

been shown that TGF- β 1 and TGF- β 3 are significantly more effective in inducing migration of epithelial cells across the wound edge and enhancing epithelial restitution in wounded IEC-6 monolayers than TGF- β 2 (McKaig *et al.* 2002). The different expression patterns of the three TGF- β isoforms during the weaning period may reflect a tissue-specific and spatio-temporal mechanism of TGF- β in regulating intestinal structure and function.

It was observed in the present study that the level of TGF- β in the intestinal mucosal homogenate estimated by Western blot analysis did not change significantly in the pig during the period of weaning (Fig. 5), which is in contrast with the results from image analysis of the immunostaining (Fig. 4). The possible explanation for this discrepancy is the different approaches with the two analytical methods. Western blot analysis estimated the TGF- β level in the homogenated mucosal tissue while image analysis of the immunostaining estimated the TGF- β level in different tissue locations in the small intestinal mucosa. The decrease in TGF- β level in the villus epithelium in association with weaning was accompanied by increased TGF- β concentration in the crypt region as observed by the immunostaining, thus the net effect of these two opposing responses would be no change in the TGF- β concentration in the homogenized mucosal tissue.

In summary, the present study demonstrated that TGF- β distribution and expression intensity in the small intestine change markedly in pigs in association with weaning. There was a transient decline in TGF- β 1 level in the intestinal villus epithelium but a significant increase in TGF- β 1 expression in the intestinal crypts in pigs following weaning. In addition, the number of TGF- β 1-positive leucocytes in the intestinal mucosa increased significantly in pigs 4d after weaning. These findings, together with the earlier reports of TGF- β effects on intestinal epithelial integrity and mucosal immune reaction, strongly suggest a regulatory role of TGF- β in the post-weaning adaptation process of the intestine. This knowledge may facilitate our understanding of the aetiology of post-weaning intestinal dysfunction in the pig.

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