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Feeding colostrum or a 1:1 colostrum:milk mixture for 3 days postnatal increases small intestinal development and minimally influences plasma glucagon-like peptide-2 and serum insulin-like growth factor-1 concentrations in Holstein bull calves

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ABSTRACT

This study evaluated how feeding colostrum- or a colostrum-milk mixture for 3 d postnatal affects plasma glucagon-like peptide-2 (GLP-2), serum insulin-like growth factor-1 (IGF-1), and small intestinal histomorphology in calves. Holstein bulls (n = 24) were fed colostrum at 2 h postnatal and randomly assigned to receive either colostrum (COL), whole milk (WM), or a 1:1 COL:WM mixture (MIX) every 12 h from 12 to 72 h. A jugular venous catheter was placed at 1 h postnatal to sample blood frequently for the duration of the experiment. Samples were collected at 1, 2, 3, 6, 9, 11, and 12 h. Following the 12-h meal, blood was collected at half-hour intervals until 16 h and then at 1-h intervals from 16 to 24 h. A 27-h sample was taken, then blood was sampled every 6 h from 30 to 60 h. Again, blood was taken at half-intervals from 60 to 64 h, then at 65 and 66 h, following which, a 2-h sampling interval was used until 72 h. Plasma GLP-2 (all time points) and serum IGF-1 (at time points: 1, 6, 12, 18, 24, 36, 48, and 72 h) were both analyzed. Duodenal, jejunal, and ileal tissues were collected at 75 h of age to assess histomorphology and cellular proliferation. Feeding COL, rather than WM, increased plasma GLP-2 by 60% for 2 h and tended to increase GLP-2 by 49.4% for 4 h after the 60-h meal. Insulinlike growth factor-1 area under the curve (from 12 to 72 h) tended to be 27% greater for COL than WM calves but was otherwise unaffected by treatment. Ileal crypts tended to proliferate more with MIX than WM, whereas ileal crypt proliferation did not differ for COL compared with MIX or WM and was not different between treatments in the proximal jejunum. Villi

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height was increased 1.8 and $1.5 \times$ (COL and MIX vs. WM) in the proximal and distal jejunum, respectively, whereas MIX duodenal and ileal villi height tended to be 1.5 and $1.4 \times$ that of WM. Crypt depth did not differ in any region. Surface area of the gastrointestinal tract was reduced for WM by 60 and 58% (proximal) jejunum) and 38 and 52% (ileum) relative to COL and MIX and was 54% less than MIX in the distal jejunum. Overall, extended COL feeding minimally increased plasma GLP-2 and serum IGF-1 compared with WM feeding. As COL and MIX similarly promoted small intestinal maturation, feeding calves transition milk to promote intestinal development could be a strategy for producers.

words: colostrum, glucagon-like peptide-2, Kev insulin-like growth factor-1, histomorphology

INTRODUCTION

The gastrointestinal tract (GIT) of neonatal ruminants rapidly matures after birth (Blum and Hammon, 2000; Guilloteau et al., 2009) as it adjusts to absorbing the orally provided nutrients necessary to meet the high energy and protein requirements of the neonatal phase (Davis et al., 1996). Most producers offer colostrum for the first day postnatal (Shivley et al., 2018; Winder et al., 2018), after which only milk or milk replacer is fed. The calf is deprived of the nonnutritive, bioactive fraction of transition milk (milkings 2 to 6; Blum and Hammon, 2000) that could further intestinal maturation while simultaneously being provided a diet that contains less energy and protein (Blum and Hammon, 2000). As such, the neonatal calf may receive inadequate nutrients and have suboptimal intestinal development. Feeding colostrum for an extended period of time does increase preprandial calf plasma glucose and total protein (Steinhoff-Wagner et al., 2011), an effect that is partially attributed to improved GIT

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development (Bühler et al., 1998; Steinhoff-Wagner et al., 2014; Yang et al., 2015). Similarly, transition milk or colostrum-milk mixes are more beneficial to GIT development than milk or whole milk feeding (Bühler et al., 1998; Yang et al., 2015). Although prolonged colostrum feeding may be more beneficial than transition milk feeding (Steinhoff-Wagner et al., 2014; Yang et al., 2015), it is important to note that prior research has not characterized the neonatal gut response to the natural transition from colostrum to whole milk by providing a primary meal of colostrum in advance of transition or whole milk.

Numerous nonnutritive bioactive peptides and hormones are present in colostrum and transition milk (Blum and Hammon, 2000; McGrath et al., 2016) to stimulate postnatal development of the neonatal calf gastrointestinal tract (Blum and Hammon, 2000). Notably, IGF-1 is one of the most abundant bioactive factors present in colostrum (Blum and Hammon, 2000; McGrath et al., 2016) and is known to promote small intestinal maturation (Burrin et al., 1996; Georgiev et al., 2003). Colostrum consumption may potentiate a positive feedback cycle by which local endocrine cascades are stimulated and promote GIT maturation. For instance, glucagon-like peptide (GLP)-2 can enhance the mitogenic action of IGF-1 to synergistically benefit GIT development, as demonstrated in rodent species (Dubé et al., 2006; Rowland et al., 2011; Leen et al., 2011). Glucagon-like peptide-2 has not been identified in colostrum (Petersen et al., 2003). However, it is co-secreted intra-luminally with GLP-1 after feeding (Burrin et al., 2003a,b), and as colostrum consumption stimulates GLP-1 secretion in neonatal calves (Inabu et al., 2018, 2019), it is likely that GLP-2 is similarly released and may cooperate with IGF-1 to promote small intestinal development. To the authors' knowledge, the relationship between small intestinal development, IGF-1, and GLP-2 has not been characterized in neonatal calves. However, Inabu et al. (2019) reported that calves fed colostrum for 3 d rather than milk had increased plasma GLP-1, but only on d 3 postnatal when they were compared with counterparts that consumed a colostrum-milk mixture (Inabu et al., 2019). Because GLP-2 is co-secreted with GLP-1 (Burrin et al., 2003a,b), the work of Inabu et al. (2019) suggests that that extended colostrum feeding may stimulate GLP-2 release, thereby promoting intestinal development.

The objective of the study was to determine if feeding colostrum, a mixture of colostrum and whole milk, or only whole milk after an initial colostrum meal influences GIT histomorphology and plasma GLP-2 and serum IGF-1 concentrations in neonatal Holstein calves. It was hypothesized that feeding colostrum for 3 d would benefit GIT development and elevate serum IGF-1 and GLP-2 concentrations more than feeding a mixture of colostrum and whole milk or only whole milk.

MATERIALS AND METHODS

Experiment design and procedures used in this study were approved by the Animal Care and Use Committee (University of Alberta, AUP00001595) before the start of the experiment. The current study was performed in conjunction with Inabu et al. (2019), in which experimental design have been previously described. As such, they will be briefly detailed here.

Animal Husbandry and Feeding

This experiment was conducted in the Dairy Research and Technology Centre at the University of Alberta. Holstein bull calves (n = 24; birth BW: 35 to 55 kg) were removed from their dam immediately following parturition and transferred into clean, plastic-paneled, calf pens $(1.2 \times 1.8 \text{ m})$ bedded with fresh shavings and straw. The standard birth care protocol was to stimulate calves for 20 min postnatal by drying them with clean towels and to sanitize their navel using a 7% iodine solution. Pooled and pasteurized colostrum (62 g of IgG/L) was supplied by Saskatoon Colostrum Company Ltd. (Saskatoon, SK, Canada) and fed (7.5% BW) to all calves at 2 h postnatal by bottle. The colostrum was thawed and warmed to precisely 39°C in a water bath before feeding. The same meal preparation protocol was used for the remaining meals. Before the experiment it was determined that 8 calves per treatment were required to detect a 20% difference in gastrointestinal measurements with an 80% statistical power (Berndtson, 1991). Following the first colostrum feeding, calves were randomly assigned to 1 of 3 treatments: colostrum (COL; n = 8, BW = 41.5 \pm 3.3 kg), a 1:1 mixture of colostrum and whole milk (MIX; n = 8, $BW = 39.8 \pm 3.3$ kg), or whole milk (**WM**; n = 8, BW $= 43.3 \pm 3.3$ kg). The colostrum fed in the COL and MIX treatments was the same supply of colostrum that was fed at 2 h. The whole milk fed in the MIX and WM treatments was pooled from cows at the Dairy Research and Technology Center (University of Alberta, Edmonton, AB, Canada) before the start of the experiment. For both colostrum and whole milk, a single batch was prepared and stored in bags frozen at -20° C until use to ensure that nutrient compositions and provisions were consistent within treatments. The batches were analyzed before the start of the study by the Central Milk Testing laboratory (Dairy Herd Improvement, Edmonton, AB, Canada) for fat, protein, lactose, and

urea-N. Colostrum and whole milk were freeze-dried and DM (%) was calculated from the weight of the dried sample as a proportion of the wet sample weight. Colostrum IGF-1 was analyzed using a commercially available immunoassay kit (DRG IGF-1 600 ELISA, DRG International Inc., Springfield Township, NJ) by the Saskatoon Colostrum Company Ltd. (Saskatoon, SK, Canada). The DM, macronutrient, and IGF-1 composition of COL, MIX, and WM treatments is shown in Table 1. Meals were offered by bottle (5% BW, 39°C) at 12-h intervals from 12 until 72 h after birth. If they were not consumed within 30 min, the refusal was fed using an esophageal tube. Calves that consecutively refused their full meal twice were removed from the study, and as a result, 4 calves were removed from the study. Two had been assigned to COL and the other 2 had been assigned to MIX and WM. The study continued until the necessary number of replicates (n = 8/treatment) were attained. All calves achieved adequate passive transfer (maximum concentration: COL = 30.4 ± 1.3 g IgG/L; MIX = 27.2 ± 1.3 g IgG/L; WM = 23.9 \pm 1.3 g IgG/L).

Jugular Catheterization and Blood Sampling

To facilitate ease of blood collection and reduce the stress response associated with repeated jugular venipuncture, calves underwent jugular catheterization (Fischer et al., 2018) at 1 h after birth using a 2 inch (5.08 cm), 16-gauge i.v. catheter (Terumo Medical Corporation, Somerset, NJ) fitted with a catheter extension line. Baseline blood samples were taken at birth, in conjunction with catheterization, and then at 2, 3, 6, and 9 h postnatal. An intensive postprandial sampling period (12 h in length), in which blood was sampled every 30 min for 4 h, was initiated at 12 h (first treatment meal). After 16 h, blood was collected at 60-min intervals until the sampling period was completed. One sample was then taken at 27 h, whereas the following samples were collected at 6-h intervals from 30 to 60 h. The second intensive postprandial sampling period began at 60 h (fifth treatment meal; 30-min sampling intervals for 4 h). After the 65 and 66 h samplings, the collection interval was lengthened to 120 min until the sampling was completed at 72 h. All collections that coincided with feedings were performed before the meals were fed. The total volume of blood collected (BW > 43 kg = 10 mL/collection, BW < 43 kg = 8mL/collection) per calf in one day did not exceed 1% of BW as per Animal Care regulations at the University of Alberta. The catheter and extension lines were flushed with 6 mL of saline and 1.5 mL of heparinized saline (20 IU/mL heparin, Fresenius Kabi, Toronto, ON, Canada) after each collection to prevent the formation of blood clots within the catheter. Blood was collected in 10-mL syringes and transferred into vials that were either coated with an anticoagulant (158 IU of heparin, Becton Dickinson, Franklin Lakes, NJ) or were not (Becton Dickinson) for separation of plasma and serum, respectively. A competitive serine protease inhibitor (Aprotinin, Sigma-A1153, Millipore-Sigma, Oakville, ON, Canada) was added to blood at a rate of $5 \,\mu \text{g/mL}$ immediately after the blood was transferred to heparin-coated vials. Subsequently, these vials were centrifuged at 3,000 \times q at 4°C for 20 min, whereas those intended for serum were allowed to clot for 3 h before centrifugation $(3,000 \times g \text{ at } 4^{\circ}\text{C} \text{ for } 20 \text{ min}).$ Supernatants were aliquoted (1.5 mL per aliquot) into 3 microcentrifuge tubes, then stored at -20° C until analysis of plasma GLP-2 and serum IGF-1.

Table	• 1. Diet nutrient	composition ;	and intake	from 1	12 to	72 h for	calves fe	d poole	d and	pasteurized	colostrum
(COL), a 1:1 mixture o	of colostrum:w	vhole milk	(MIX)	, or	whole m	ilk (WM)) after a	n init	ial meal of	colostrum

T,	COT	MIX	337N.C	CEM	D 1
Item	COL	MIA	VV IVI	SEM	<i>P</i> -value
DM, %	23.0	18.1	13.2		
Nutrient composition, % DM					
Fat	20.0	29.7	39.4		
Protein	57.0	39.8	22.7		
Lactose	12.2	21.2	30.3		
Urea-N, mg/dL	34.4	24.3	14.2		
Gross energy, Mcal/kg as fed	1.25	1.03	0.81		
IGF-1, $\mu g/L$	360.9	180.5			
Intake, g/meal					
DM	$485.2^{\rm a}$	373.3^{b}	294.9°	16.6	< 0.001
Fat	97.0^{b}	101.1^{ab}	$116.2^{\rm a}$	4.4	0.013
Protein	276.3^{a}	166.0^{b}	67.0°	8.0	< 0.001
Lactose	59.1°	70.1^{b}	89.4^{a}	3.0	< 0.001
Urea-N, mg/meal	695.7^{a}	484.2^{b}	308.9°	22.0	< 0.001
Gross energy intake, Mcal/meal	2.65^{a}	2.12^{b}	1.80^{b}	0.1	< 0.001
IGF-1, $\mu g/meal$	729.9	359.7		24.0	< 0.001

^{a-c}Means within a row with different superscripts differ (P < 0.05).

Slaughter and Intestinal Tissue Collection

Calves were killed by exsanguination 75 h postnatal after reaching a surgical plane of anesthesia by administration of euthanyl (30 mg of pentobarbital/kg of 75 h BW; Euthanyl, Vetoquinol, Lavaltrie, QC, Canada) via jugular catheter. The dissection was performed once death was confirmed by the absence of a heartbeat.

All equipment used for the dissection was sterilized before use by autoclaving. The abdomen was opened from the sternum to the anus to access the digestive tract. Cable ties were used to occlude the esophagus and rectum from the body and excising incisions were performed cranially and caudally of these ties, respectively. The GIT was removed, placed on a sterile tray, and specific, predetermined sections of the small intestine were located for sample collection. Duodenal and proximal jejunal tissue were respectively collected 7.5 and 100 cm distal of the abomasal pyloric sphincter. Distal jejunal tissue was collected 30 cm proximal to the cranial mesenteric artery. Ileal tissue was taken 30 cm in advance of the ileo-cecal junction (Malmuthuge et al., 2015). Tissues were preserved in 15 mL of formalin and embedded within 48 h (described below) for histomorphometric analysis by brightfield microscopy.

Brightfield Microscopy and Measurements

Histomorphometric measurements of the duodenal, proximal and distal jejunal, and ileal villi and crypts were performed to detect differences in intestinal growth and developmental state (Wongdee et al., 2016). Small intestinal tissue was dehydrated overnight by a series of ethanol solution incubations. Once dehydrated, the samples were embedded and solidified in paraffin wax, sectioned (5 μ m slice per section, 2 sections per sample) using a rotary microtome (Leica, Wetzlar, Germany), and stained with hematoxylin and eosin. Slides were mounted using an upright brightfield microscope (Microscope Axio Scope.A1, Carl Zeiss Canada Ltd., Toronto, ON, Canada), whereas ImageJ software (ImageJ, National Institute of Health, Bethesda, MD) was used to quantify villi height and width and crypt depth and width (μm) in the duodenum, proximal and distal jejunum, and ileum. An auto-generated grid was overlaid on each image for visual accuracy and consistency. Villi were measured from the fully attached epithelial cell to the villus-crypt junction. Crypts were measured from the villus-crypt junction to the tip of the crypt (Wongdee et al., 2016). Two measurements from 5 different images per slide (10 measurements/small intestinal region) were taken under $100 \times$ magnification. Widths were measured perpendicularly from the mid-section of 2 fully attached villi and crypts/image, resulting in 10 measurements/calf of villi and crypt width for duodenal, proximal and distal jejunal, and ileal regions. All measurements were averaged by calf and analyzed as one observation per dependent variable per experimental unit.

Villi height and crypt depth were used to determine the villi:crypt ratio and the mucosal surface area index was estimated from the mucosal-to-serosal amplification ratio (equation 1), originally presented in Kisielinski et al. (2002). The formula to estimate mucosal surface area index by the mucosal-to-serosal amplification ratio, using histomorphometric measures of villi and crypts is as follows:

$$M = \frac{\left[\left(a \ \times \ b\right) + \left(\frac{a}{2} + \frac{c}{2}\right)^2 - \left(\frac{a}{2}\right)^2 \right]}{\left(\frac{b}{2} + \frac{c}{2}\right)^2}$$

where M = mucosal surface area index, a = villi width (μm), b = villi height (μm), and c = crypt width (μm).

Proliferating Cell Nuclear Antigen Quantification

Proliferating cell nuclear antigen (**PCNA**) immunohistochemistry was used to quantify the proportion of proliferative cells (Hall et al., 1990) in the proximal jejunum and ileum. Samples were prepared from the same paraffin-embedded blocks of proximal jejunum and ileum as previously described. Blocks were cut in 5-µm sections, which were rehydrated on a Leica CV 5030 autostainer program 2 (Leica). Antigen retrieval was performed with citrate for 16 min at 80%, followed by cooling for 30 min. The sections were then washed twice in PBS, followed by quenching for 6 min in a solution of 10 mL of 50% H_2O_2 in 40 mL of 100% methanol. Sections were rinsed with distilled water for 5 min followed by 1-min rinse in PBS. Nonspecific adherence of antibodies was blocked with 20% normal goat serum (NGS) for 60 min. Normal goat serum was aspirated and the rabbit anti-PCNA antiserum was added for 60 min. The anti-PCNA antisera (Ab18197-Rabbit anti PCNA; Abcam Inc., Toronto, ON, Canada) was diluted 1:2,000 with 20% goat serum (NGS, Cedarlane, Burlington, ON, Canada). A biotinylated goat anti-rabbit antibody (Abcam, Cambridge, MA) was diluted 1:200 with NGS and applied as a secondary antibody for 40 min, then washed with PBS 3 times. Avidin-biotin complex (Cedarlane) was added for 40 min, followed by PBS washing 3 times before 3,3'-diaminobenzidine (Cedarlane) solution was applied. When adequate color intensity was reached, sections were washed with distilled water and counterstained with hematoxylin. Proliferating cell nuclear antigen samples were quantified using an upright brightfield microscope (Microscope Axio-Scope A.1, Carl Zeiss Canada Ltd.), whereby color intensity thresholds were manually set for individual sections with the ImageJ program (ImageJ, National Institutes of Health, Bethesda, MD). Five images of each slide were captured. The cells that were opaque and fully stained were considered to be PCNA⁺, proliferative cells. The proportion of PCNA⁺ cells/total cells from 2 crypts were measured per image under $400 \times$ magnification (10 measurements per GIT region). Proliferative index measurements were averaged within GIT region by calf to represent one observation per dependent variable per experimental unit.

Plasma GLP-2 and Serum IGF-1

Plasma GLP-2 was measured using a solid-phase competition assay (Sugino et al., 2004; Inabu et al., 2018). Human GLP-2 labeled with Europium (Peptide Institute Inc., Osaka, Japan), polyclonal anti-rat GLP-2 serum, and goat-anti-rabbit- γ -globulin coated polystyrene microtiter strips (Yanaihara Institute Inc., Shizuoka, Japan) were used to target the N-terminal of GLP-2, as described by Inabu et al. (2017) and validated by Elsabagh et al. (2017). The intra- and interassay coefficients of variation were 6.5 and 10.6%, respectively. Serum was frozen at -20° C and shipped overnight on dry ice to the Endocrine Service Lab (Western College of Veterinary Medicine, University of Saskatchewan, Saskatoon, SK, Canada) for analysis of IGF-1 by an automated solid-phase chemiluminescent immunoassay using an Immulite 1000 Immunoassay System (Siemens AG, Munich, Germany). The Immulite/Immulite 1000 system was calibrated following the instructions provided by the manufacturer. The intra- and interassay coefficients of variation for the IGF-1 analysis were 5.0 and 4.1%, respectively.

Area Under the Curve Calculation

Area under the curve (AUC) was calculated for GLP-2 (at 6-h intervals) and IGF-1 (by measurement interval) using an equation based on the trapezoidal rule and cumulated to represent AUC_{0-72} and AUC_{12-72} . Following the first (12 h) and fifth (60 h) treatment meals, 2-, 4-, and 12-h AUC intervals were calculated for GLP-2.

Statistical Analysis

All statistical analyses were performed using SAS 9.4 (SAS Institute Inc., Cary, NC). PROC UNIVARIATE

was used to analyze normality. Shapiro-Wilk P > 0.05 was used to identify that residuals were normally distributed, and data were confirmed to be homoscedastic by visual appraisal of distribution plots. All data were found to be normally distributed.

Small intestinal histomorphometric measurements and PCNA analysis were assessed as a one-way ANOVA using PROC MIXED defining treatment as a fixed effect within the model (shown below). Tukey's post-hoc test was used to distinguish means that differed.

$$Y_{ij} = \mu + \text{treatment}_i + \varepsilon_{ij},$$

 Y_{ij} = the *j*th observation in the *i*th treatment, μ = the overall mean, treatment_i = the fixed effect of the *i*th treatment, and ε_{ij} = the random error associated with the *j*th observation in the *i*th treatment.

Measurements conducted over time (i.e., GLP-2 and IGF-1) additionally included the fixed effects of time and the treatment by time interaction:

$$\mathbf{Y}_{ijkl} = \boldsymbol{\mu} + \text{treatment}_i + \text{time}_j + (\text{Trt} \times \mathbf{T})_{ij} + \text{calf}_k + \varepsilon_{ijkl},$$

where Y_{ijkl} = the *l*th observation of the *k*th calf in the *i*th treatment at the *j*th time, μ = the overall mean, treatment_i = the fixed effect of the *i*th treatment, time_j = the fixed effect of the *j*th time, calf_k = the random effect of the *k*th calf, (Trt × T)_{ij} = the fixed effect of the interaction of the *i*th treatment at the *j*th time, ε_{ijkl} = the random error associated with the *l*th observation of *k*th calf in the *i*th treatment at the *j*th time.

Degrees of freedom were defined using the Kenward-Roger approximation to account for unbalanced observations at specific time points. Covariance was modeled to clarify the auto-correlation between observations from one experimental unit repeated over time. Multiple covariances structures were tested and their appropriateness of fit to the model was determined (by the lowest Bayesian and Akaike information criterions) individually for dependent variables. Variance components, compound symmetry, unstructured, ante-dependence, and heterogeneous compound symmetry were tested as covariance structures with data that were unequally spaced, whereas equally spaced data were additionally tested with autoregressive, heterogeneous autoregressive, Toeplitz, and heterogeneous Toeplitz covariance structures. Outlier detection was performed after variance had been properly partitioned within the abovedescribed statistical model. Outliers were detected in both the GLP-2 and IGF-1 repeated measures by using PROC REG to regress the observation by the predicted response and generate the studentized residual and studentized-deleted residual. Only observations whose

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studentized-deleted residuals were >3 standards deviations from the predicted residual were removed from the data set. The IGF-1 measurement of one MIX calf at 12 h after birth was determined to be an outlier and was removed from the IGF-1 data set. All repeated measurements were corrected using the Bonferroni test. Means are considered different by P < 0.05 and trends are discussed when 0.05 < P < 0.10.

RESULTS

Dry Matter, Nutrient, Gross Energy, and IGF-1 Intake

When calves were fed treatment meals from 12 to 72 h, COL-fed calves consumed 111.9 and 190.6 \pm 16.6 g of DM/meal more (P < 0.001) than MIX- and WM-fed calves. Milk-fed calves consumed 78.4 ± 23.5 g DM/d less than MIX-fed calves. Fat intake per meal was 19.8% greater (P = 0.013; Table 1) when calves were fed WM rather than COL, though neither COL nor WM fat intake differed from MIX. Colostrum-fed calves had increased protein $(1.64 \times MIX, 4.12 \times WM;$ P < 0.001), urea (1.44 × MIX, 2.25 × WM; P < 0.001), and gross energy (GE) intake (1.25 \times MIX, 1.47 \times WM; P < 0.001) but decreased lactose intake (0.84) \times MIX, 0.66 \times WM; P < 0.001) per meal than their counterparts. Protein and urea intake were respectively greater (P < 0.001) by 99.0 \pm 11.4 g/meal and 175.3 \pm 31.2 mg/meal when calves were fed MIX instead of WM, whereas lactose intake was 19.3 ± 3.0 g/meal less (P < 0.001) for MIX- than WM-fed calves. Gross energy intake per meal did not differ between MIX (2.12) \pm 0.13 Mcal/meal) and WM (1.80 \pm 0.13 Mcal/meal).

When the contribution of the 2-h colostrum meal to total daily intake was considered, COL calves continued to consume 191.8 and 302.7 ± 37.8 g of DM/d more (P < 0.001; Table 2) than MIX and WM calves, respectively; however, MIX and WM calves did not differ in their daily DMI. Fat intake per day was increased (P= 0.034) by 16.6% for WM relative to COL, and MIX calves consumed similar quantities of fat as both WM and MIX. Colostrum protein intake remained elevated $(1.45 \times \text{MIX}, 2.32 \times \text{WM}; P < 0.001)$, whereas lactose consumption was only less (P < 0.001) than WM-fed calves by 52.2 ± 6.3 g/d. Feeding MIX rather than WM similarly resulted in calves consuming $1.60 \times \text{more}$ (P < 0.001) protein and $0.19 \times \text{less}$ (P < 0.001) lactose per day. Again, daily urea consumption was greatest (P < 0.001) when calves were fed COL as intake was 624.0 \pm 51.2 mg/d lower for WM calves, with MIX calves consuming intermediary quantities of urea that were $360.2 \pm 51.2 \text{ mg/d}$ less than COL but 263.8 ± 51.2 mg/d more than WM. Daily GE intake was highest (P < 0.001) for COL (5.73 \pm 0.21 Mcal/d) compared with MIX (4.83 \pm 0.21 Mcal/d) and WM calves (4.40 \pm 0.21 Mcal/d). Between MIX and WM, GE intake per day was not different. By purposely lowering meal provision from 7.5 to 5% BW at 12 h, all calves had consistently reduced (P < 0.001) DM (-31.4%), fat (-16.9%), protein (-42.4%), lactose (-10.3%), urea (-36.1%), and GE intake (-29.5%) during d 2 and 3 compared with d 1. Dry matter, fat, protein, lactose, urea, and GE intakes were not different between d 2 and 3. Numerous treatment × time interactions were present (P < 0.001) in DM, nutrient, and GE intake between treatments among days and are presented in Supplemental Figure S1 (https://doi.org/10.3168/jds .2019-17219) and described in Supplemental File S1 (https://doi.org/10.3168/jds.2019-17219).

Insulin-like growth factor-1 intake per meal for COL was twice that (P < 0.001) of the MIX calves and nonmeasurable for the WM calves. As such, WM IGF-1 intake could not be compared with that of COL and MIX. For the COL and MIX calves, IGF-1 intake was $1.5 \times$ greater (P < 0.001) on d 1 than both d 2 and 3. A treatment by time interaction was detected (P < 0.001) for IGF-1 intake between COL and MIX calves and is presented in Supplemental Figure S2 (https://doi.org/ 10.3168/jds.2019-17219).

Small Intestine Histomorphology and Proliferation Index

Proximal jejunum histomorphology is visually (Figure 1; image A: COL; image B: MIX; image C: WM) distinguishable among treatments with the most development of villi and mucosa being discernable in COL and MIX images. Villi height was greater in the proximal (P < 0.001; Table 3) and distal (P = 0.008)jejunum by 79.7 and 52.1%, respectively, when calves were fed COL compared with milk. Similarly, proximal and distal jejunum villi height were 84.5 and 53.1%increased and tended to be 49.1 and 35.1% greater in the duodenum (P = 0.067) and ileum (P = 0.098) for MIX relative to WM calves. These metrics did not differ between calves fed COL and MIX. Regardless of GIT region, crypt depth was not affected (P > 0.26) by feeding COL, MIX, or WM to calves. The villi height: crypt depth ratio in the proximal jejunum was lessened (P = 0.003) for WM calves to 0.52 and 0.57 times than that of the COL and MIX calves, whereas the MIX duodenal villi:crypt ratio tended (P = 0.072) to be 53.3% greater than the WM. The ratio of villi height:crypt depth did not differ $(P \ge 0.12)$ by treatment in the distal jejunum or ileum. Colostrum and MIX calves had a greater mucosal surface area than WM calves in the proximal jejunum (P = 0.001) and ileum (P = 0.010); however, they did not differ from each other. Calves fed

Item COL Intake, g/d	MIX				Day				P-value	
Intake, g/d		WМ	SE	1	2	ŝ	SE	Trt^{1}	Time	$Trt \times time^2$
DMI 1,051.2 ⁻	$859.4^{ m b}$	748.5^{b}	37.8	$1,121.2^{\mathrm{a}}$	$768.9^{ m b}$	768.9^{b}	22.9	< 0.001	< 0.001	< 0.001
Fat $210.2^{\rm b}$	215.9^{ab}	245.0^{a}	9.3	252.1^{a}	209.5^{b}	$205.9^{ m b}$	5.4	0.034	< 0.001	0.008
Protein 598.7 ^a	411.8^{b}	258.1°	18.9	589.4^{a}	339.6^{b}	339.6^{b}	11.0	< 0.001	< 0.001	< 0.001
Lactose 128.0 ^b	145.8^{b}	180.2^{a}	6.3	162.5^{a}	145.7^{b}	$145.7^{ m b}$	2.6	< 0.001	< 0.001	< 0.001
Urea-N, mg/d 1,507.4 ^a 1	$1,147.2^{\rm b}$	883.4°	25.5	$1,552.7^{\mathrm{a}}$	992.6^{b}	$992.6^{ m b}$	29.7	< 0.001	< 0.001	< 0.001
Gross energy intake, Mcal/d 5.73 ^a	$4.83^{\rm b}$	$4.40^{ m b}$	0.21	6.21^{a}	4.38^{b}	4.38^{b}	0.12	< 0.001	< 0.001	< 0.001
IGF-1 intake, $\mu g/d$ 1,581.4	956.3		55.7	$1,627.5^{a}$	$1,089.6^{ m b}$	$1,089.6^{ m b}$	39.6	< 0.001	< 0.001	< 0.001

Table 2. Nutrient, urea-N, and gross energy intake on d 1, 2, and 3 postnatal for calves fed pooled and pasteurized colostrum (COL), a 1:1 mixture of colostrum: whole milk (MIX)

Trt = treatment.

²Treatment by time interactions are shown in Supplemental Figure S1 (https://doi.org/10.3168/jds.2019-17219)

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MIX had $1.54 \times \text{greater}$ (P = 0.001) surface area than WM in the distal jejunum, whereas COL surface area within this region was similar to both MIX and WM. Duodenal surface area index was unaffected (P = 0.27)by feeding COL, MIX, or WM from 12 to 72 h.

Although treatment only tended (P = 0.054) to cause the proportion of PCNA⁺ cells in the ileal crypts to increase by 5.2% units for MIX compared with WM calves, Tukey's post-hoc test indicated that these means were separate (P = 0.044, MIX vs. WM). Calves consuming COL did not have a different $(P \ge 0.36)$ ileal crypt cell proliferative index than MIX or WM. The percentage of $PCNA^+$ crypt cells did not differ (P = 0.34) by treatment in the proximal jejunum.

Plasma GLP-2 and Serum IGF-1

Plasma GLP-2 concentration did not differ ($P \ge 0.85$; Table 4) between COL, MIX, and WM calves from 0 to 72 h or 12 to 72 h. Similarly, the treatment by time interaction did not affect $(P \ge 0.69)$ GLP-2 concentrations during these time intervals. However, plasma GLP-2 concentration between 0 to 72 h and 12 to 72 h fluctuated (P < 0.001; Figure 2) over time. Regardless of considering either the 60 or 72 h timeline, GLP-2 was lower during d 2 at 27, 30, 36, and 42 h compared with the periods immediately following 12 and 60 h, although by 48 h postnatal, plasma GLP-2 began to recover and was similar to concentrations during the high-frequency measurements (12-h periods) at 12 and 60 h. At 27, 30, 36, and 42 h, GLP-2 was equivalent to the low values $(0.49 \pm 0.1 \text{ ng/mL})$ observed immediately before the first meal after birth at 2 h.

An interaction of treatment and time (P = 0.028)was present between 12 to 14 h. Calves that were fed WM had GLP-2 concentrations that were $1.32 \times$ greater at 12.5 h than their own concentrations at 14 h, but did not differ from COL or MIX at any time point in the 2 h postprandial. Otherwise, treatment did not affect (P = 0.58) GLP-2 concentrations from 12 to 14 h, and similarly, treatment and treatment by time had no influence $(P \ge 0.57)$ on GLP-2 from 12 to 16 h or 12 to 24 h. Time affected (P = 0.022) GLP-2 concentrations between 12 to 14 h, but means were not separate after post-hoc adjustment. Plasma GLP-2 concentrations changed (P < 0.001) between 12 and 24 h, but changes were marginal and reflected increased GLP-2 concentrations at 12.5 h compared with 24 h.

For 2 h postprandial of 60 h, COL calves GLP-2 was 59.4% greater (P = 0.012) than WM. During this timeframe, MIX-fed calves GLP-2 concentration was equivalent to both COL and WM. Glucagon-like peptide-2 tended (P = 0.052) to remain elevated (46.4% greater) from 60 to 64 h for COL calves compared with WM



Figure 1. Brightfield microscopy images $(200 \times \text{magnification})$ of proximal jejunal tissue from calves that consumed either pooled and pasteurized colostrum (A), a 1:1 colostrum:whole milk mixture (B), or whole milk (C) from 12 to 72 h after an initial meal of colostrum at 2 h postnatal. Jejunal tissue is stained with hematoxylin and eosin for contrast.

calves. From 60 to 72 h, treatment differences were no longer present (P = 0.17). Additionally, no treatment by time interactions ($P \ge 0.30$) were detected from 60 to 62 h, 60 to 64 h, or 60 to 72 h. Time-dependent response patterns in GLP-2 were detected from 60 to 62 h (P = 0.034), 60 to 64 h (P = 0.041), and 60 to 72 h (P = 0.011), but means did not separate after Bonferroni adjustment.

Glucagon-like peptide-2 AUC did not change throughout the entire experimental period ($P \ge 0.41$; Table 5) or in the 2-, 4-, and 12-h intervals at 12 and 60 h ($P \ge 0.22$; Supplemental Table S1; https://doi.org/ 10.3168/jds.2019-17219) by the effect of treatment, time, or the interaction of treatment and time. The treatment × time effect ($P \le 0.048$) on GLP-2_{12h} AUC after 12 and 60 h was not different once corrected for multiple comparisons.

Neither treatment nor the interaction of treatment and time affected $(P \ge 0.14; \text{ Table 4})$ serum IGF-1 from 1 to 72 h or 12 to 72 h. Serum IGF-1 concentration was elevated (P < 0.001; Figure 3) for the first 24 h postnatal, peaking at 18 h ($83.8 \pm 3.4 \text{ ng/mL}$) and thereafter declining by 9.9, 74.9, 98.7, and 75.1% relative to 24, 36, 48, and 72 h postnatal. Concentration of IGF-1 at 72 h was 28.4 ng/mL less than concentrations at 24 h yet remained unchanged relative to 36 and 48 h. The same time-dependent changes (P < 0.001; data not shown) in IGF-1 concentration were observed when evaluating only the interval that calves were fed their respective treatments (12 to 72 h). Yet, IGF-1 AUC from 12 to 72 h tended to be greater (P = 0.088; Table 5) for COL than WM calves (3,771.04 vs. 2,973.17 \pm 248.7 ng/mL \times h), whereas MIX did not differ from either. This difference was not present in the 1 to 72 h serum IGF-1 AUC.

DISCUSSION

The current study aimed to compare the maturation of the neonatal calf GIT and relevant endocrine signals to GIT development (i.e., GLP-2 and IGF-1) when the natural transition in postnatal macronutrient and bioactive compound ingestion was mimicked. Though the composition of transition milk gradually shifts to that of mature milk by 3 to 4 d postpartum (Blum and Hammon, 2000), in the current study transition milk was simulated in the MIX treatment by combining and feeding equal parts COL and WM. Providing a fixed nutrient intake for the MIX treatment was necessary to remove the confounding influence of inconsistent daily nutrient or bioactive factor intake on measurements performed over time (i.e., plasma GLP-2 and serum IGF-1 concentrations). In addition, the MIX macronutrient composition (29.7% crude fat, 39.8% CP, 21.2% lactose, % DM) is similar to the reported transition milk compositions (milkings 2 to 6; Blum and Hammon, 2000) that correspond to the 72-h experiment duration. Thus, the MIX treatment adequately simulated transition milk and the experimental model was appropriate to study neonatal physiological responses to colostrum, transition milk, or whole milk provision.

Histomorphology has been used as a proxy to indicate development and functional capacity of the small intestine and evidence very clearly shows that consumption of milk or milk replacer instead of colostrum impedes the capacity of the neonatal GIT to achieve its developmental potential (Bühler et al., 1998; Steinhoff-Wagner et al., 2014; Yang et al., 2015). It is not unexpected that reductions in villi height, villi height:crypt depth ratio in the duodenum and proximal jejunum, and mucosal surface area in the proximal and distal jejunum and il-

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Table 3. Histomorphometric measurements of villi and crypts within the duodenum, proximal and distal jejunum, and ileum and the proportion of proliferative cell nuclear antigen positive (PCNA⁺) crypt cells in the proximal jejunum and ileum of neonatal Holstein bull calves fed either pooled and pasteurized colostrum (COL), a 1:1 mixture of colostrum:whole milk (MIX), or whole milk (WM) from 12 to 72 h after an initial meal of colostrum at 2 h postnatal

		Treatment		D 1 1		
Variable	COL	MIX	WM	- Pooled SEM	<i>P</i> -value	
Villi height, ¹ µm						
Duodenum	$395.0^{\operatorname{AB}}$	440.8^{B}	295.6^{A}	41.5	0.067	
Proximal jejunum	$468.3^{\rm a}$	480.7^{a}	260.6^{b}	31.3	< 0.001	
Distal jejunum	$463.3^{\rm a}$	$466.4^{\rm a}$	304.6^{b}	37.4	0.008	
Ileum	428.8^{AB}	453.7^{B}	335.8^{A}	39.4	0.098	
Crypt depth, ² µm						
Duodenum	189.1	201.0	211.2	18.3	0.71	
Proximal jejunum	195.5	211.4	202.7	14.1	0.73	
Distal jejunum	194.3	203.5	177.5	11.0	0.26	
Ileum	195.9	178.6	191.3	13.5	0.65	
Villi height:crypt depth ³						
Duodenum	2.1^{AB}	2.3^{B}	1.5^{A}	0.2	0.072	
Proximal jejunum	2.5^{a}	2.3^{a}	1.3^{b}	0.2	0.003	
Distal jejunum	2.5	2.4	1.8	0.3	0.14	
Ileum	2.2	2.7	1.8	0.3	0.12	
Mucosal surface area index ^{4,5}						
Duodenum	6.5	6.9	5.0	0.8	0.27	
Proximal jejunum	8.0^{a}	7.9^{a}	$5.0^{ m b}$	0.5	0.001	
Distal jejunum	7.5^{ab}	9.1^{a}	$5.9^{ m b}$	0.5	0.001	
Ileum	8.3^{a}	9.1^{a}	$6.0^{ m b}$	0.7	0.010	
PCNA ⁺ crypt cells/total cells						
Proximal jejunum	0.35	0.36	0.33	0.01	0.34	
Ileum ⁶	$0.41^{\rm ab}$	0.43^{a}	$0.38^{ m b}$	0.01	0.054	

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

^{A,B}Means within a row with different upper case superscripts tend to differ (0.05 $\leq P < 0.10$).

¹Villi height (n) by gastrointestinal tract (GIT) region and treatment (COL, MIX, and WM, respectively) as follows: duodenum, n = 8, 8, and 7; proximal jejunum, n = 8, 8, and 8; distal jejunum, n = 7, 8, and 8; and ileum, n = 7, 8, and 8.

²Crypt depth (n) by GIT region and treatment (COL, MIX, and WM, respectively) as follows: duodenum, n = 8, 8, and 6; proximal jejunum, n = 8, 8, and 8; distal jejunum, n = 8, 8, and 8; and ileum, n = 7, 8, and 8. ³Villi height:crypt depth ratio (n) by GIT region and treatment (COL, MIX, and WM, respectively) as follows: duodenum, n = 8, 8, and 6; proximal jejunum, n = 8, 8, and 8; distal jejunum, n = 7, 8, and 8; and ileum, n = 7, 8, and 8.

⁴Surface area (n) by GIT region and treatment (COL, MIX, and WM, respectively) as follows: duodenum, n = 7, 8, and 8; proximal jejunum, n = 8, 7, and 8; distal jejunum, n = 7, 8, and 8; and ileum, n = 7, 8, and 8. ⁵Surface area calculated according to the equation described by Kisielinski et al. (2002).

⁶Tukey's post-hoc test separated means despite the overall *P*-value being nonsignificant (P > 0.05).

eum were observed in WM calves when compared COL and MIX calves. However, that WM feeding tended to lessen the population of PCNA⁺ crypt ileal cells is surprising since it was only relative to MIX, and not COL calves. Blättler et al. (2001) similarly found proliferative indices were greater in calves consuming transition milk (milkings 1 to 6; d 1 to 3) instead of a milk-based formula of similar macronutrient composition (d 1 to 3) before all calves being fed milk replacer from d 4 to 7. Their results differ from those of the current study, as proliferation of duodenal epithelia was also lower when calves consumed only first-milking colostrum (as opposed to transition milk; Blättler et al., 2001) for the first 3 d postnatal, indicating that extensive colostrum consumption could adversely affect small intestinal development relative to transition milk. Unfortunately, small intestinal proliferation of colostrum- relative to formula-fed calves was not reported by Blättler et al. (2001) and their results cannot be further compared with our own. However, Steinhoff-Wagner et al. (2014) found that calves that consumed colostrum as opposed to a colostrum-based formula had lessened crypt depth but greater villi circumference, area, and height than their counterparts (Steinhoff-Wagner et al., 2014), aptly coinciding with the results of the current study. Both Blättler et al. (2001) and Steinhoff-Wagner et al. (2014) proposed that the effect of extended colostrum feeding on intestinal crypt formation was positive and demonstrated furthered maturation, or potentially resulting from greater intestinal mass (Blättler et al., 2001). Increased migration of crypt epithelial cells to villi, reductions in apoptosis, and enhanced rates of crypt fission have been suggested as potential mechanisms for the decreased crypt proliferative rates and reduced depths (as reviewed by Blum, 2006) but it is still unclear which mechanism predominates.

It was hypothesized that GIT maturation would be promoted foremost by COL compared with MIX (intermediary) and WM. Colostrum and MIX feeding did indeed further GIT development to a greater extent than WM. Yet, extensive colostrum feeding did not provide an advantage compared with MIX feeding, as evidenced by the equivalent villi height and increased mucosal surface area between COL and MIX calves. Thus, the hypothesis was partially rejected as COL feeding did not provide additional benefit to small intestine maturation compared with MIX. These results are in contrast to previous research. Overall improved gut development has been reported in 8-d-old calves that had consumed colostrum as opposed to pooled transition milk on d 1 and 2 postnatal (Yang et al., 2015), whereas feeding first-milk colostrum until d 3 postnatal (rather than colostral milkings 1 to 6) only increased duodenal development in neonatal calves despite the colostrum-calves consuming a serially diluted colostrum from d 4 to 7 (Blättler et al., 2001). The latter study is more pertinent to the present than that of Yang et al. (2015) by the fact that calves were provided a primary colostrum feeding. Bühler et al. (1998) similarly fed an initial meal of colostrum to calves that then received milk replacer until euthanasia at d 8 postnatal and contrasted these results to calves that were fed 6 colostrum meals before receiving milk replacer until d 8 postnatal. A lack of transition milk or a transition milk-mimic provision after the initial colostrum meal stunted intestinal development compared with calves that had received colostrum for 3 d (Bühler et al., 1998), coinciding with observations from the current study whereby COL small intestinal development was enhanced compared with WM regardless of WM calves consuming a primary meal of colostrum.

Based on the preceding discussions, it is possible that one timely feeding of colostrum may compensate to preserve and promote the functionality of the neonatal gut once consumption of transition milk begins. This is likely mediated in part by colostral bioactive compounds, yet the greater nutrient density of COL and MIX must be considered. Increased nutrient intake, and thus, availability of anabolic substrates for both COL and MIX calves on d 1 postnatal, likely supported the enhanced GIT development. This notion highlights the importance and potency of colostral bioactive compounds when they are consumed early in life, because MIX small intestine growth was preserved despite MIX calves having decreased nutrient and GE intake on d 2 and 3 postnatal compared with COL calves. However, the suggestion that a single colostrum feeding is sufficient to benefit calf development is currently limited solely to the GIT and likely only relevant when transition milk, not milk or milk replacer, is fed afterward. Still, this proposition has important practical implications for the dairy industry as preweaning morbidity (34%; Urie et al., 2018) and mortality (6.4%;Winder et al., 2018) is primarily caused by gastroin-

Table 4. Plasma glucagon-like peptide-2 (GLP-2) and serum IGF-1 concentrations in Holstein bull calves that were fed either pooled and pasteurized colostrum (COL), a 1:1 colostrum:whole milk mixture (MIX), or whole milk (WM) from 12 to 72 h after an initial meal of colostrum at 2 h postnatal

		Treatment				<i>P</i> -value	
Variable	COL	MIX	WM	SEM	Trt^1	Time^2	$\mathrm{Trt}\times\mathrm{time}$
Plasma GLP-2, ng/mL							
Entire period _{0-72b}	1.28	1.35	1.24	0.1	0.89	< 0.001	0.70
Post-treatment meal _{12–72h}	1.36	1.43	1.33	0.1	0.85	< 0.001	0.69
Plasma GLP-2: 12 h, postprandial,							
ng/mL							
12 to 14 h	1.43	1.59	1.85	0.3	0.58	0.002	0.028
12 to 16 h	1.34	1.48	1.75	0.3	0.60	< 0.001	0.57
12 to 24 h	1.22	1.46	1.68	0.3	0.59	< 0.001	0.71
Plasma GLP-2: 60 h, postprandial,							
ng/mL							
60 to 62 h	1.66^{a}	1.51^{ab}	1.04^{b}	0.1	0.012	0.034	0.30
60 to 64 h	1.64	1.51	1.12	0.1	0.052	0.041	0.93
60 to 72 h	1.68	1.57	1.23	0.1	0.17	0.011	0.93
Serum IGF-1, ng/mL							
Entire period _{1-72h}	72.9	67.5	61.2	4.0	0.14	< 0.001	0.15
Post-treatment meal _{12-72h}	67.6	66.3	59.4	3.8	0.28	< 0.001	0.41

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

 1 Trt = treatment.

²The effect of time is expressed as hours postnatal. GLP-2 time effects are shown in Figure 2; and IGF-1 time effects are shown in Figure 3.

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Table 5. Total area under the curve (AUC) of glucagon-like peptide-2 (GLP-2) and IGF-1 over the entire experiment period (0 to 72 h) and after the first treatment meal (12 to 72 h) for Holstein bull calves fed pooled and pasteurized colostrum (COL), a 1:1 colostrum:whole milk mixture (MIX), or whole milk (WM) from 12 h onward after an initial meal of colostrum at 2 h postnatal

		Treatment			<i>P</i> -value
Item	COL	MIX	WM	SEM	Treatment
GLP-2 AUC, 1 ng/mL \times h					
AUC _{0-72h}	88.2	87.0	80.8	9.3	0.83
AUC _{12-72h}	77.4	73.7	62.7	7.9	0.41
IGF-1 AUC , ¹ ng/mL × h					
AUC _{1-72h}	4,658.7	4,444.4	3,860.4	294.6	0.16
$\mathrm{AUC}_{12-72\mathrm{h}}$	$3,771.0^{A}$	$3,\!613.5^{\operatorname{AB}}$	$2,973.2^{B}$	248.7	0.088

^{A,B}Means within a row with different superscripts tend to differ $(0.05 \le P < 0.10)$.

¹AUC is calculated by the formula $[(a + b) \times c]/2$; a = GLP-2 or IGF-1 concentration, b = GLP-2 or IGF-1 concentration, c = duration of time between measurements.



Figure 2. Plasma glucagon-like peptide-2 (GLP-2) concentrations of neonatal Holstein bull calves from 0 to 72 h and 12 to 72 h postnatal. Calves were offered treatments of pooled and pasteurized colostrum, a 1:1 mixture of colostrum:whole milk, or whole milk from 12 to 72 h after birth after all calves received an initial colostrum meal at 2 h postnatal. Black arrows represent when meals were fed, and shaded timeframes (12 to 14 h, 12 to 16 h, and 12 to 24 h; 60 to 62 h, 60 to 64 h, and 60 to 72 h) demonstrate periods of frequent measurements that underwent additional statistical analysis relative to 12- and 60-h meals. Plasma GLP-2 from 0 to 72 h and 12 to 72 h varied (P < 0.001) over time, such that GLP-2 concentrations were depressed from 27 to 42 h compared with their post-12 h concentrations, but began to recover by 42 h and were fully recovered by 48 h postnatal. Treatment and the treatment × time interaction did not affect ($P \ge 0.69$) plasma GLP-2 from either 0 to 72 h or 12 to 72 h after birth. Error bars represent SE.



Figure 3. The change in serum IGF-1 over time for Holstein bull calves from 1 to 72 h postnatal. Calves were offered treatments of pooled and pasteurized colostrum, a 1:1 mixture of colostrum:whole milk, or whole milk from 12 to 72 h postnatal after all calves received an initial colostrum meal at 2 h postnatal. Time affected (P < 0.001) serum IGF-1 concentration as it was elevated from 1 to 18 h, declined afterward, and was lowest at 48 and 72 h postnatal. Neither treatment (P = 0.14) nor the interaction of treatment and time (P = 0.15) influenced serum IGF-1 concentrations. Means with different letters differ (P < 0.05). Error bars represent SE.

testinal disorders (56 and 32%, respectively; Urie et al., 2018). Promoting gastrointestinal development by feeding transition milk following colostrum provision could be a viable strategy for producers to decrease the incidence of preweaning digestive-associated morbidity. Further research is needed to determine how transition milk feeding, as compared with colostrum, whole milk, or milk replacer feeding, affects the incidence of digestive morbidity and preweaning mortality.

It is indisputable that colostrum is meant to further neonatal GIT ontogenesis (Guilloteau et al., 1997; Blum and Baumrucker, 2002; Zabielski et al., 2008; Ontsouka et al., 2016), as evidenced by its extensive nonnutritive bioactive factor content (reviewed by Blum and Hammon, 2000; McGrath et al., 2016) and capacity to stimulate the release of endogenous endocrine factors by nutrient delivery to the small intestine (Hadorn et al., 1997; Hammon et al., 2000; Inabu et al., 2018). Of particular interest to the current study is the oral delivery of IGF-1 in colostrum (Pakkanen and Aalto, 1997; Blum and Hammon, 2000) to the neonatal calf by its small intestine-targeted mitogenic capacity to increase epithelial cell proliferation (MacDonald, 1999; Shen et al., 2004). Because the mitogenic action of IGF-1 can be enhanced by GLP-2 (Dubé et al., 2006; Rowland et al., 2011; Leen et al., 2011), these 2 hormones were promising targets to determine their relationship to neonatal intestinal development and postnatal nutrient consumption.

Yet, study findings did not support the hypothesis that elevated plasma GLP-2 and serum IGF-1 concentrations would reflect and coincide with enhanced intestinal development by COL compared with MIX or WM feeding. These results are surprising because plasma GLP-2 is strongly correlated with nutrient intake and intestinal development in piglets (Burrin et al., 2000; Stoll et al., 2000) and mature humans (Holst et al., 1976; Xiao et al., 1999) and serum IGF-1 is positively influenced by adequate colostrum consumption (Hammon and Blum, 1997; Egli and Blum, 1998; Hammon et al., 2000). The previously discussed lack of difference in COL and MIX intestinal histomorphology could be related to the similar GLP-2 and IGF-1 concentrations for both groups. However, IGF-1 was also similar in WM-fed calves, despite these calves having hindered villi and surface area growth relative to their counterparts. The tendency for greater IGF-1 AUC from COL compared with WM consumption insufficiently relates to improved GIT development, particularly as calves consuming colostrum have been consistently demonstrated to have elevated serum IGF-1 and better small intestinal maturation than their counterparts (Hadorn et al., 1997; Hammon and Blum, 1997; Rauprich et al., 2000; Schäff et al., 2014). The tendency for increased IGF-1 AUC more likely relates to greater total nutrient and GE intake (Hammon et al., 2002; Inabu et al., 2019) when COL (compared with WM) was fed for an extended period of time (discussed below), particularly since negligible quantities of ingested IGF-1 cross the neonatal GIT in calves (Baumrucker and Blum, 1994; Vacher et al., 1995; Hammon and Blum, 1997) or pigs (Burrin, 1997; Donovan et al., 1997). The comparatively greater post-60 h plasma GLP-2 concentrations for COL rather than WM calves can also be considered minimal given that increases were only observed for up to 4 h after feeding and were not different between MIX and WM calves. To counterpoint, Burrin et al. (2003b) suggested circulating physiological GLP-2 concentrations may not accurately represent luminal GLP-2 concentrations and activity. This concept can be applied to our own results and may be one explanation for the apparent lack of plasma GLP-2 response to treatment provision or intestinal development. Yet, we cannot prove that GLP-2 was differentially secreted intra-luminally. Ultimately, plasma GLP-2 and serum IGF-1 did not correlate to improved small intestinal development within this study.

Calf IGF-1 concentrations are comparably lesser at birth than in later life (Greenwood et al., 2002; Quigley et al., 2006) but still decrease in the neonatal period when nutrient intake is restricted (Hadorn et al., 1997; Hammon and Blum, 1997; Hammon et al., 2000). This is a response that is consistent across species (Thissen et al., 1994; Greenwood et al., 2002) regardless of maturity (Ronge and Blum, 1988; Smith et al., 2002; Brown et al., 2005). In the current study, the maximum observed IGF-1 concentration (83.8 \pm 3.4 ng/mL, 18 h postnatal) was far lower than prior reported ranges of approximately 140 to 200 ng/mL during the first 24 h postnatal (Hadorn et al., 1997; Hammon and Blum, 1997). In fact, observed circulating IGF-1 concentrations in the current study are more similar to concentrations observed in calves that are purposefully nutrientrestricted (Hadorn et al., 1997; Hammon and Blum, 1997). Total nutrient and GE provision were greatest during the first 24 h postnatal by the combined feeding of colostrum and the first treatment meal. Afterward, total nutrient delivery (regardless of treatment) was reduced during d 2 and 3 compared with 1. The initial decline in serum IGF-1 was observed at 24 h postnatal, corresponding to the second treatment meal provision and comparably (to meal 1) less nutrient and GE consumption. Nutrient intake is the initial regulator of the somatotropic growth axis (Breier and Gluckman, 1991; McGuire et al., 1995) and insufficient nutrient intake depresses circulating IGF-1 (Thissen et al., 1994; Greenwood et al., 2002). As calves were not provided nutrients throughout the study at quantities equivalent to their allocation during d 1, it is not surprising that serum IGF-1 concentrations did not recover during the remainder of the study. It is possible that calves within the current study did not consume adequate nutrients to meet their requirements.

Glucagon-like peptide-2 is similarly reduced when nutrient intake is inadequate. A minimal enteral energy consumption of 40 to 60% is necessary to stimulate a GLP-2 response in 16-d-old piglets when compared with those fed 100% of their daily energy aliquot parenterally (Burrin et al., 2000) and, in 10- to 13-d old calves, reducing the milk replacer feeding rate to 1.31 (75% of control), 0.88 (50% of control), and 0.44 (25%)of control) from 1.75% BW as DM (control) caused corresponding decreases in circulating GLP-2 of 13, 28, and 51% compared with the control (Castro et al., 2016). Calf GLP-2 concentration in the present study was depressed during d 2 by 44.3 and 46.5% relative to d 1 (2 h post-colostrum feeding) and 3, respectively. By the above argument, these results indicate that calf nutrient consumption during d 2 did not satisfy requirements relative to d 1 when the DM feeding rate was lessened on average from 1.32 to 0.91% BW. The mean reduction in DM feeding rate was 31.9% as COL, MIX, and WM feeding rates were reduced by 20.0, 31.2, and 44.7% from d 1 to d 2 and 3. Conversely, the recovery in GLP-2 concentration during d 3 is peculiar as the DM feeding rate remained static relative to d 2. To the authors' knowledge, the influence of time on plasma GLP-2 during a prolonged, imposed nutrient reduction, has not been investigated, making our findings novel but challenging to interpret.

However, it is possible that the appearance of GLP-2 in blood signals the degree to which energetic demands of the GIT are satisfied (Burrin et al., 2000; Castro et al., 2016), rather than satiation of whole-body energy requirements. On this basis, we theorize that the pattern in calf plasma GLP-2 between d 1, 2, and 3 signal the balance between energetic demand and satisfaction of the intestinal tract. The recovery in plasma GLP-2 on d 3 would then suggest retrogressive adaptation (Dierksen et al., 1985; Zhang et al., 2013; Pederzolli et al., 2018) of the GIT, most likely to conserve splanchnic energy expenditure (Reynolds, 2002). Retrogressive intestinal adaptation in response to reduced feed intake has been previously demonstrated in neonatal calves (Le Huerou et al., 1992; Guilloteau et al., 2009; Steinhoff-Wagner et al., 2015). Total small intestine and small intestine mucosa mass are noted to be 31.8 and 19.2% less, respectively, on d 2 than at birth (Le Huerou et al., 1992; Guilloteau et al., 2009). Similarly, mucosal villi size (but not crypt depth) decreased when calves were fed colostrum (milkings 1, 3, and 5; 5%

BW) for 4 d postnatal compared with term-born calves fed 1 meal of colostrum before slaughter at 26 h (Steinhoff-Wagner et al., 2015). Because calves in the study had decreased total energy consumption (-1.83 Mcal)and reduced meal volume (19.8% of average d 1 meal volume) during d 2, it is plausible that energy consumption was inadequate relative to intestinal requirement. As gastrointestinal mass is affected by ME intake and feed-induced workload relative to requirements (McLeod and Baldwin, 2000; Reynolds, 2002; Hersom et al., 2004), the abrupt, combined exposure to these 2 factors and the phenotypic elasticity of neonates could cause rapid intestinal regression in calves within the current study. Consequently, GIT retrogressive adaption by d 3 would lessen the magnitude of imbalance between calf feed consumption and splanchnic energy expenditure (Reynolds, 2002), causing an apparent recovery of plasma GLP-2. However, the theory that plasma GLP-2 could signal splanchnic energy balance is currently speculation and the authors are unaware of any studies having addressed this notion.

CONCLUSIONS

Provision of COL for 3 d postnatal was anticipated to benefit small intestinal maturation to a greater extent than 3 d of MIX (intermediate) or WM feeding. Findings from this study indicate that no additional benefit is gained from COL than MIX feeding, though both had improved small intestine development compared with WM. Further, changes in serum IGF-1 and plasma GLP-2 were hypothesized to reflect intestinal development within treatments but are likely more related to total energy and nutrient consumption. The temporal decreases in IGF-1 and GLP-2 (the latter followed by recovery) are concerning as they suggest neonatal calves may consume insufficient quantities of nutrients by current industry practices. It is necessary to clarify neonatal calf metabolizable nutrient requirements relative to provision. For the present, study observations offer dairy producers the viable strategy of feeding neonatal calves transition milk or diluting colostrum with milk or milk replacer (after an initial colostrum meal) to promote intestinal development and function.

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