Fertility of lactating Holstein cows submitted to a Double-Ovsynch protocol and timed artificial insemination versus artificial insemination after synchronization of estrus at a similar day in milk range

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ABSTRACT

Our objective was to compare the AI submission rate and pregnancies per artificial insemination (P/AI) at first service of lactating Holstein cows submitted to a Double-Ovsynch protocol and timed artificial insemination (TAI) versus artificial insemination (AI) to a detected estrus after synchronization of estrus at a similar day in milk range. Lactating Holstein cows were randomly assigned to receive their first TAI after a Double-Ovsynch protocol (DO; n = 294) or to receive their first AI after a synchronized estrus (EST; n = 284). Pregnancy status was determined 33 ± 3 d after insemination and was reconfirmed 63 ± 3 d after insemination. Data were analyzed by ANOVA and logistic regression using the MIXED and GLIMMIX procedures of SAS (SAS Institute Inc., Cary, NC). By design, days in milk at first insemination did not differ between treatments (76.9 ± 0.2 vs. 76.7 ± 0.3 for DO vs. EST cows, respectively), but more DO cows were inseminated within 7 d after the end of the voluntary waiting period than EST cows (100.0 vs. 77.5%). Overall, DO cows had more P/AI than EST cows at both 33 d (49.0 vs. 38.6%) and 63 d (44.6 vs. 36.4%) after insemination, but pregnancy loss from 33 to 63 d after insemination did not differ between treatments. Primiparous cows had more P/AI than multiparous cows 33 and 63 d after insemination, but the treatment by parity interaction was not significant. Synchronization rate to the hormonal protocols was 85.3%, which did not differ between treatments; however, synchronized DO cows had more P/AI than EST cows because of an intrinsic increase in fertility after submission to a fertility program.

Key words: first artificial insemination, timed artificial insemination, estrus, fertility

INTRODUCTION

Hormonal synchronization protocols that allow for timed artificial insemination (TAI) have been incorporated widely into reproductive management programs by dairy farms (Caraviello et al., 2006; Norman et al., 2009); however, AI based on detection of estrus continues to be an important part of the overall reproductive management program on most dairy farms (Caraviello et al., 2006; Miller et al., 2007). Results from the first field trial evaluating the Ovsynch protocol on a commercial dairy farm reported that Ovsynch and TAI yielded similar pregnancies per artificial insemination (P/AI) to that of cows receiving AI after a detected estrus (39 vs. 37%; Pursley et al., 1997) but that median days to first AI (54 vs. 83) and days open (99 vs. 118) were decreased for cows submitted for first insemination using an Ovsynch protocol than for cows submitted for AI after a detected estrus, respectively. Thus, the initial effect of TAI protocols on 21-d pregnancy rates in US dairy herds was to increase the AI service rate, with little to no effect on P/AI (Norman et al., 2009).

Since the development of the Ovsynch protocol by Pursley et al. (1995), several modifications to the original Ovsynch protocol have been tested in an attempt to increase P/AI to TAI. These modifications include increasing ovulatory response to the first GnRH treat-
ment (Carvalho et al., 2015b), presynchronization using 2 PGF$_{2\alpha}$ treatments (i.e., Presynch-Ovsynch; Moreira et al., 2001; Navanukraw et al., 2004; Ribeiro et al., 2011), presynchronization using a combination of GnRH and PGF$_{2\alpha}$ (i.e., G-6-G and Double-Ovsynch; Bello et al., 2006; Souza et al., 2008; Carvalho et al., 2014a), and addition of a second PGF$_{2\alpha}$ treatment 24 h after the first within the Ovsynch protocol to induce complete luteal regression (Carvalho et al., 2015a; Wiltbank et al., 2015; Santos et al., 2016). Taken together, these modifications have yielded P/AI at first service that exceed 50% in high-producing Holstein cows (Carvalho et al., 2014a, 2015b). Thus, the latest iterations of hormonal synchronization protocols for submitting lactating dairy cows for first service can be best described as fertility programs for high-producing dairy cows because all cows can be inseminated within 7 d of the end of the voluntary waiting period and yielding more P/AI at first service.

The idea that fertility programs and TAI can yield greater fertility than AI to estrus at first insemination in high-producing dairy cows has not been definitively tested. Several experiments compared P/AI of cows inseminated after TAI with cows inseminated after a detected estrus at first AI, with some studies reporting no differences in P/AI (Pursley et al., 1997; Chebel and Santos, 2010; Dolecheck et al., 2016), whereas others reported more P/AI for cows receiving TAI (Gumen et al., 2012; Fricke et al., 2014; Stevenson et al., 2014). In all of these studies, however, DIM at first service differed between cows submitted to TAI and cows submitted to AI after a detected estrus. Further, DIM is a clear confounder because cows with more DIM at first service have more time for uterine involution and for resumption of cyclicity, both of which affect P/AI (Buch et al., 1955; Chebel and Santos, 2010).

Our objective was to compare the AI submission rate and P/AI at first service of lactating Holstein cows submitted to a Double-Ovsynch protocol and TAI versus AI to a detected estrus after synchronization of estrus at a similar DIM range. Our hypothesis was that the AI submission rate and P/AI at first service would be greater for cows submitted to first TAI after a Double-Ovsynch protocol than for cows receiving AI to a detected estrus after submission to a hormonal protocol for synchronization of estrus.

**MATERIALS AND METHODS**

All animal handling and experimental procedures were approved by the Animal Care and Use committee of the College of Agriculture and Life Sciences at the University of Wisconsin–Madison.

**Cows, Housing, and Feeding**

This study was conducted from October 2015 to June 2016 on 1 commercial dairy farm in Portugal. Lactating Holstein cows (n = 578) were milked twice daily at approximately 12-h intervals. Cows were fed twice daily a TMR consisting of corn silage and alfalfa hay as forage with corn and soybean meal-based concentrate formulated to meet or exceed the minimum nutritional requirements for high-producing dairy cows (NRC, 2001). Cows were housed in freestall barns bedded with mattress and had ad libitum access to feed and water. Primiparous and multiparous cows were housed in separate pens, and barns were equipped with fans and sprinklers that were automatically activated when the temperature inside the barns exceeded 28 and 32°C, respectively. The rolling herd average and daily milk production were 10,719 kg and 35.6 kg/cow per day with 4.9% fat and 3.3% protein during the experiment.

**Experimental Treatments**

Each week, cows at 50 ± 3 DIM (d 0) were stratified by parity (primiparous vs. multiparous) and were randomly assigned to 2 treatments to receive first insemination (Figure 1). Cows (n = 294) in the first treatment were submitted for first TAI after a modified Double-Ovsynch (DO) protocol that included a second PGF$_{2\alpha}$ treatment 24 h after the first in the Breeding-Ovsynch portion of the protocol. Briefly, on d 0 cows received the first GnRH treatment of the Pre-Ovsynch portion of the Double-Ovsynch protocol, followed by treatment with PGF$_{2\alpha}$ 7 d later and GnRH 72 h after PGF$_{2\alpha}$. Seven days later, cows received a GnRH treatment followed by 2 PGF$_{2\alpha}$ treatments administered 7 and 8 d later, with the last GnRH treatment administered 56 h after the first PGF$_{2\alpha}$ treatment followed by TAI 16 to 20 h later. Cows (n = 284) in the second treatment were submitted to a hormonal synchronization protocol for synchronization of estrus and were inseminated if detected in estrus (EST). Briefly, 3 d after d 0 (Figure 1), cows were treated with GnRH followed by treatment with PGF$_{2\alpha}$ 7 d later. Fourteen days later, cows received 2 PGF$_{2\alpha}$ treatments administered 24 h apart. Cows detected in estrus from 2 d before until 7 d after the first of the 2 PGF$_{2\alpha}$ treatments at the end of the protocol were inseminated within 12 h of detection of estrus based on visual detection of estrus, which was aided using pedometers attached to the rear legs of the cows (Westfalia Separator, GEA, Lisbon, Portugal).

The GnRH (100 μg/dose of gonadorelin diacetate tetrahydrate, Ovarelin) and the PGF$_{2\alpha}$ (25 mg/dose of dinoprost tromethamine, Enzapro-T) used in this
experiment were from Ceva Santé Animale (Libourne, France). Four high fertility sires, with a sire conception rate >0 and more than 6,000 observations each, were used for insemination, and sires were equally balanced between treatments. Cows were inseminated by 2 farm personnel, and inseminations were equally balanced between inseminators and treatments. Sires 1, 2, 3, and 4 were used in 24.1, 23.8, 24.5, and 27.6% of the inseminations for cows in the DO treatment and in 27.7, 24.5, 23.6, and 24.1% of the inseminations for cows in the EST treatment.

**Blood Sampling and Progesterone Assay**

Blood samples were collected from all cows via puncture of the median caudal blood vessels into 8-mL evacuated serum collection tubes (Vacuette, Greiner Bio-One North America Inc., Monroe, NC). Blood samples were collected on the day of enrollment (d 0) immediately before administration of any hormonal treatments, before treatment with PGF$_{2\alpha}$ (d 24), at the last GnRH treatment for DO cows or on the day of AI for EST cows, and 7 d after the last GnRH treatment for DO cows or 7 d after AI for EST cows (Figure 1). After collection, blood samples were refrigerated until centrifuged (20 min at 1,600 $\times$ g; 4°C). Serum was harvested and stored at −20°C until assayed for progesterone (P4) concentrations using a chemiluminescent assay (Immulite, Siemens, Wales, UK). Average sensitivity for the 8 assays was <0.2 ng/mL. Average intraassay and interassay coefficients of variation were 6.3 and 4.9%, respectively, based on a quality control sample (2.50 ng/mL of P4) that was replicated within each assay.

**Pregnancy Diagnosis**

Pregnancy diagnosis was performed 33 ± 3 d after AI using a portable scanner (Easi-Scan, BCF Technology Ltd., Livingston, UK) equipped with a 7.5-MHz linear-array transducer. A positive pregnancy diagnosis was based on visualization of a corpus luteum (CL) on the ovary ipsilateral to the uterine horn containing an embryo with a heartbeat. Pregnancy status for cows diagnosed pregnant at the first examination was reconfirmed 63 ± 3 d after AI using the same ultrasound machine and transducer. Cows diagnosed pregnant and subsequently diagnosed not pregnant at the pregnancy reconfirmation were classified as having undergone pregnancy loss.

**BCS Evaluation and Milk Yield**

Body condition score was evaluated for all cows on the day of the last PGF$_{2\alpha}$ treatment of each protocol using a 5-point scale with 0.25 increments where 1 = thin and 5 = fat (Ferguson et al., 1994). One person performed all BCS evaluations throughout the experiment. Based on a previous study (Carvalho et al., 2014b), BCS were categorized as either low (≤2.50) or high (≥2.75) for statistical analysis.

Milk weights for the 7 d before the last PGF$_{2\alpha}$ treatment of each protocol were recorded daily and stored in the on-farm computer software program (BoviSync, Dairy LLC, Eden, WI). For statistical analysis, average milk weights of the 7 d before insemination were used, and cows were categorized by parity (primiparous vs. multiparous) according to milk production as being greater than or less than mean milk production.

![Figure 1](https://example.com/figure1.png)

*Figure 1.* Schematic diagram of treatments and experimental endpoints. Each week, lactating Holstein cows at 50 ± 3 DIM (d 0) were stratified by parity (primiparous vs. multiparous) and randomly assigned to receive their first insemination as a timed artificial insemination (TAI) after a Double-Ovsynch (DO) protocol or as an AI to a detected estrus after a protocol for synchronization of estrus (EST).
**Statistical Analyses**

The experimental design was a completely randomized design balanced by parity (primiparous vs. multiparous) as the stratifying factor. All statistical analyses were performed using SAS computational software, version 9.4 for Microsoft Windows (SAS Institute Inc., Cary, NC).

Analysis of binary response data (P/AI, pregnancy loss, pregnancy rate, percentage of cows with low (≤2.5) BCS near AI, % of cows inseminated, synchronization rate) was performed by logistic regression using the GLIMMIX procedure of SAS. Fixed effects included in the initial models were treatment, parity, level of milk production, and BCS category except for analysis of % of cows with low (≤2.5) BCS near AI. Selection of the fixed effects model that best fit the data for each variable of interest was performed by finding the model with the lowest value for the Akaike information criterion using a backward elimination procedure that removed from the model all variables with a P-value exceeding 0.10. Effects of treatment and parity were forced to remain in the statistical model; parity was forced to remain the final model because it was used as a stratifying factor when randomizing cows to treatments. The final models included the fixed effects of treatment and parity. For P/AI and pregnancy loss, BCS category was also included in the final model. The effect of P4 concentration at d 0 on P/AI was analyzed using the GLIMMIX procedure of SAS using a model that contained only the fixed effect of P4 category. The distribution of DIM at insemination was performed using the LOGISTIC procedure of SAS with a model containing the fixed effects of treatment and parity.

Differences in milk production, BCS, average DIM, and P4 concentrations were determined by ANOVA using the MIXED procedure of SAS. The model contained the fixed effects of treatment, parity, and the treatment × parity interaction. Cows were distributed into 9 categories using P4 concentrations on d 0 and 24 (from 0.0 to ≥7.0 ng/mL), and into 5 categories using P4 concentrations near insemination (from 0.0 to ≥0.5 ng/mL) and 7 d after insemination (from 0.0 to ≥4.0 ng/mL). Differences in the proportion of cows within each P4 category were analyzed by logistic regression using the LOGISTIC procedure of SAS.

A significant difference between the levels of a classification variable was considered when \( P \leq 0.05 \), whereas differences between \( P > 0.05 \) and \( P \leq 0.10 \) were considered a statistical tendency.

**RESULTS AND DISCUSSION**

**Milk Production and BCS**

To test whether cows were appropriately randomized to treatments, we compared milk production and BCS of cows between treatments (Table 1). Average milk production of cows during the 7 d preceding insemination did not differ (\( P = 0.72 \)) between treatments; however, primiparous cows produced less (\( P < 0.01 \)) milk than multiparous cows (35.3 vs. 44.9 kg/d, respectively). Average BCS did not differ (\( P = 0.85 \)) between treatments; however, BCS was greater (\( P < 0.01 \)) for primiparous than multiparous cows (2.98 vs. 2.78, respectively). The proportion of cows with low (≤2.50) BCS did not differ (\( P = 0.54 \)) between treatments, and fewer (\( P < 0.01 \)) primiparous cows had low (≤2.50) BCS than multiparous cows [7.0% (15/215) vs. 30.9% (112/363), respectively].

**Effect of Treatment on DIM at Insemination and Submission Rate**

Several experiments have compared P/AI at first AI for cows inseminated after synchronization of ovulation and TAI with cows receiving AI after a detected estrus, with some studies reporting no differences in P/AI (Pursley et al., 1997; Chebel and Santos, 2010; Dolecheck et al., 2016), whereas others reported more P/AI for cows receiving TAI (Gumen et al., 2012; Fricke et al., 2014; Stevenson et al., 2014). In all of

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<th>Table 1. Mean (±SEM) milk production and BCS of lactating Holstein cows enrolled in the experiment</th>
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<td><strong>Treatment</strong></td>
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<td>Cows with BCS ≤2.50 at PGF2α, % (no./no.)</td>
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1Each week, lactating Holstein cows at 50 ± 3 DIM (d 0) were stratified by parity (primiparous vs. multiparous) and were randomized to receive first insemination as a timed AI after a Double-Ovsynch (DO) protocol or AI to a detected estrus after a hormonal protocol that synchronized estrus (EST).
these studies, however, DIM at which insemination occurred differed for cows submitted to TAI versus cows submitted to AI after a detected estrus. For example, in the study by Pursley et al. (1997), cows receiving TAI were inseminated with fewer DIM than cows receiving AI after estrus detection (54 vs. 83 DIM, respectively). Similarly, in the study by Dolecheck et al. (2016) cows receiving TAI were inseminated earlier after the voluntary waiting period than cows inseminated after detection of estrus (6 vs. 17 d, respectively). By contrast, in the studies by Chebel and Santos (2010), Fricke et al. (2014), and Stevenson et al. (2014), cows submitted to TAI had more DIM than cows submitted to AI after detection of estrus (74 vs. 65 DIM in the study by Chebel and Santos, 2010; 75 vs. 59 DIM in the study by Fricke et al., 2014; and 75 vs. 65 DIM in the study by Stevenson et al., 2014).

The experimental design used in the present experiment allowed us to compare the AI submission rate of cows inseminated at a similar DIM range, which was a primary objective of the experiment. By design, average DIM at first insemination did not differ \( (P = 0.37) \) between treatments (76.9 vs. 76.7 d for DO and EST cows, respectively). In addition, distribution of DIM at which cows were inseminated did not differ between treatments (Figure 2; lower panel). The AI submission rate was greater \( (P < 0.01) \) for DO than for EST cows \( [100\% \ (294/294) \ vs. \ 77.5\% \ (220/284); \ Figure \ 2; \ upper \ panel] \), but the proportion of cows inseminated did not differ \( (P = 0.64) \) between parities \( [91.2\% \ (196/215) \ vs. \ 87.6\% \ (318/363) \ for \ primiparous \ vs. \ multiparous \ cows, \ respectively] \).

Because the rate of anovulation assessed in 5,818 lactating Holstein cows from 13 herds averaged 23.3% (Bamber et al., 2009), we attempted to decrease the proportion of anovular EST cows by initiating the synchronization of estrus protocol with a GnRH treatment (Figure 1), which induced ovulation in 88% of anovular cows in another study (Gümen et al., 2003). In our study, estrus detection was performed by visual observation of cows standing to be mounted by herd-mates and was aided using a pedometry system that resulted in a 77.5% AI submission rate. By comparison, AI submission rates of lactating Holstein cows based on detection of increased activity using an activity monitoring system were 69 to 70% after 2 PGF\(_{2\alpha}\) treatments administered 14 d apart (Fricke et al., 2014), and 71% for cows treated with GnRH followed by PGF\(_{2\alpha}\) 7 d later (Valenza et al., 2012). Thus, although 70 to 80% AI submission rates can be achieved using visual detection of estrus, pedometry, or activity monitoring systems, they do not approach the 100% AI submission rate achieved for cows submitted to a fertility program and TAI for first insemination.

**Effect of Treatment on Pregnancies per AI and Pregnancy Loss**

In support of our hypothesis, DO cows had 27% more \( (P = 0.02) \) pregnancies than EST cows 33 d after insemination (Table 2). In addition, primiparous cows had more \( (P < 0.01) \) P/AI than multiparous cows \( [54.6\% \ (107/196) \ vs. \ 38.4\% \ (122/318)] \), but the treatment by parity interaction on P/AI 33 d after insemination was not significant \( [57.3\% \ (63/110) \ and \ 44.0\% \ (81/184) \ for \ primiparous \ vs. \ multiparous \ DO \ cows; \ 51.2\% \ (44/86) \ and \ 30.6\% \ (41/134) \ for \ primiparous \ vs. \ multiparous \ EST \ cows, \ respectively] \). The effect of treatment on P/AI was maintained at the pregnancy reconfirmation with DO cows having 23% more \( (P = 0.05) \) pregnancies than EST cows 63 d after insemination (Table 2). In addition, primiparous cows had more \( (P < 0.01) \) P/AI than multiparous cows \( [51.0\% \ (100/196) \ vs. \ 34.9\% \ (111/318)] \), but no treatment by parity interaction
was observed on P/AI 63 d after insemination [52.7% (58/110) and 39.7% (73/184) for primiparous and multiparous DO cows; 48.8% (42/86) and 28.4% (38/134) for primiparous and multiparous EST cows]. Submission of cows for first insemination using a fertility program such as the Double-Ovsynch protocol used in the present study yielded not only a greater AI submission rate, but 23% more P/AI than submission of cows to AI after a detected estrus at a similar DIM range. Taken together, the increased submission rate and P/AI for DO cows in the present study yielded 64% more pregnant cows at 33 d after insemination and 58% more pregnant cows at 63 d after insemination than EST cows (Table 2). A recent meta-analysis of 3 controlled studies that included 1,689 cows reported that incorporation of insemination of cows to estrus after the second PGF2α treatment of a Presynch-Ovsynch protocol decreased the odds of pregnancy by 35% compared with 100% TAI after a Presynch-Ovsynch protocol (Borchardt et al., 2016). Taken together, results from Borchardt et al. (2016) and from the present study support that insemination of high-producing Holstein cows after a detected estrus results in lower fertility than when ovarian function and timing of AI is manipulated during a fertility program such as the Double-Ovsynch protocol used in the present study followed by TAI.

In agreement with another study (Carvalho et al., 2014b), cows in the present experiment with BCS ≤2.5 had fewer (P < 0.01) P/AI than cows with BCS ≥2.5 [29.1% (30/103) vs. 48.4% (199/411)]. Cows with BCS ≤2.5 had fewer (P = 0.02) P/AI than cows with BCS ≥2.5 [27.2% (28/103) vs. 44.5% (183/411)]. Pregnancy loss from 33 to 63 d after AI did not differ (P = 0.43) between treatments [9.0% (13/144) vs. 5.9% (5/85) for DO vs. EST cows, respectively] or between parities [P = 0.47; 6.5% (7/107) vs. 9.0% (11/122) for primiparous vs. multiparous cows, respectively]. In addition, pregnancy loss did not differ (P = 0.67) between BCS categories [6.7% (2/30) vs. 8.0% (16/199), for BCS ≤2.5 vs. BCS ≥2.75, respectively]. A larger sample size would be required to perform a valid comparison of the effects of treatment or parity on pregnancy loss in the present study.

**Progesterone Profiles**

Blood samples were collected at d 0 to determine P4 status at the onset of the synchronization protocols and at d 24 of the protocols (Figure 1), at insemination, and 7 d later to assess timing of AI and to determine the proportion of cows synchronized during the protocols. **P4 at d 0.** Mean P4 concentration at d 0 was 2.8 ± 0.1 ng/mL and did not differ (P = 0.71) between treatments (2.8 vs. 2.8 ng/mL, for DO vs. EST cows, respectively), and primiparous cows tended (P = 0.07) to have lower P4 concentrations than multiparous cows (2.5 vs. 2.9 ng/mL, respectively). As expected, the distribution of cows based on P4 concentrations on the day of enrollment did not differ between treatments (Figure 3, upper left panel). Overall, the proportion of cows with P4 <0.5 ng/mL on d 0 was 27.1% (155/572) and did not differ (P = 0.88) between treatments (Table 1; Figure 3, upper left panel). Cows with <0.5 ng/mL of P4 at the onset of the protocols should consist of anovular cows in addition to a few cycling cows at a stage of the estrous cycle when P4 would normally be low. In support of this idea, more (P = 0.03) primiparous cows had P4 <0.5 ng/mL than multiparous cows [31.9% (68/213) vs. 24.2% (87/359), respectively], which agrees with data from other studies in which primiparous cows had a greater incidence of anovulation than multiparous cows (Gümen et al., 2003; Santos et al., 2004; Silva et al., 2007).

**P4 at d 0 for EST Cows Not Detected in Estrus.** Estrus cows not detected in estrus at the end of the protocol were further analyzed (Figure 4). Mean
P4 concentrations at d 0 for EST cows not detected in estrus after the final PGF₂α treatment of the protocol for synchronization of estrus was 2.7 ng/mL. Overall, only 38% of EST cows not detected in estrus at the end of the protocol had P4 concentrations <0.5 ng/mL. Thus, anovulatory status at the onset of the protocol did not fully account for failure of these cows to be detected in estrus at the end of the protocol. This is similar to findings of other studies that speculated that issues other than cyclicity status affected efficiency and accuracy of detection of estrus in lactating dairy cows submitted to a synchronization protocol (Chebel and Santos, 2010; Fricke et al., 2014). For cows with P4 <1.0 ng/mL on d 0, P/AI did not differ \( (P = 0.67) \) between treatments [40.0% (40/100) vs. 36.7% (25/68), for DO vs. EST cows, respectively]; however, for cows with P4 ≥1.0 ng/mL on d 0, DO cows had more \( (P < 0.01) \) P/AI than EST cows [53.4% (102/191) vs. 38.3% (57/149), respectively].

**P4 at d 24.** Overall, DO cows had greater P4 concentrations on d 24 than EST cows (Figure 3, upper right panel). On d 24, mean P4 concentration was 4.1 ± 0.1 ng/mL, and mean P4 concentration was greater \( (P < 0.01) \) for DO cows than for EST cows (4.6 vs. 3.5 ng/mL, respectively). Mean P4 concentrations did not differ \( (P = 0.61) \) between primiparous and multiparous cows (4.0 vs. 4.1 ng/mL, respectively). When cows were divided into 9 categories based on P4 concentrations at d 24 (Figure 3, upper right panel), more \( (P < 0.01) \) EST cows had P4 <0.5 ng/mL than DO cows [9.6% (26/271) vs. 2.8% (8/288)], and more \( (P < 0.01) \) cows not detected in estrus after a hormonal protocol for synchronization of estrus.
DO cows had P4 ≥ 4.0 ng/mL than EST cows [65.9% (190/288) vs. 37.3% (101/271), respectively].

In the present study, DO cows should have greater P4 concentrations at d 24 than EST cows due to a treatment effect on CL number. A Double-Ovsynch protocol consists of 2 successive Ovsynch protocols; the first is the Pre-Ovsynch portion of the protocol, which is followed by the Breeding-Ovsynch portion of the protocol (Souza et al., 2008; Giordano et al., 2013). Although not assessed in our study, TAI cows ovulating after the last GnRH treatment of the Pre-Ovsynch portion of the protocol and at the first GnRH treatment of the Breeding-Ovsynch part of the protocol should have 2 CL at the PGF$_{2α}$ treatment of the Breeding-Ovsynch portion of the protocol as reported in another study using the same Double-Ovsynch protocol (Fricke et al., 2016). By contrast, EST cows should have only 1 CL at this time during the protocol for synchronization of estrus. Progesterone concentrations before spontaneous or induced luteal regression are associated with P/AI in both cows receiving TAI and cows receiving AI after a detected estrus. For example, Nebel et al. (1987) reported that P/AI increased as P4 concentrations increased during the diestrus immediately preceding first AI for cows inseminated after a detected estrus. Similarly, in an analysis by Fricke et al. (2015) that evaluated the association between P4 concentrations at the time of the PGF$_{2α}$ treatment in 3,383 TAI services, cows with low P4 concentrations at a similar point during a Double-Ovsynch protocol had 20% P/AI compared with 41% P/AI for cows with high P4 concentrations at this time.

**P4 at the Time of AI.** Accuracy of detection of estrus has a profound effect on conception rates in dairy cows (Heersche and Nebel, 1994; Sturman et al., 2000); thus, accuracy of detection of estrus must be accounted for when assessing P/AI. Blood samples were collected at the time of AI to evaluate accuracy of detection of estrus for EST cows. Although the correct timing of AI cannot definitively be established for cows with low P4 at the time of AI, these cows were inseminated in the absence of a functional CL, a stage of the cycle when estrus and ovulation should occur. By contrast, cows with high P4 near the time of AI had a functional CL and were inseminated at an incorrect stage of the cycle. Similarly, P/AI for cows submitted to a Double-Ovsynch protocol are profoundly affected by lack of complete luteal regression, and low P4 at TAI for DO cows is an indicator of luteal regression at this point during a Double-Ovsynch protocol (Brusveen et al., 2009; Fricke et al., 2015).

At the time of insemination, mean P4 concentration was 0.25 ± 0.0 ng/mL. Overall, DO cows had lower (P = 0.05) P4 concentrations than EST cows (0.22 vs. 0.29 ng/mL, respectively), but mean P4 concentration was 2.1 ± 0.1 ng/mL. Mean P4 concentration was greater (P < 0.01) for EST than for DO cows (2.3 vs. 1.9 ng/mL, respectively), but mean P4 concentration 7 d after insemination did not differ (P = 0.20) between primiparous and multiparous cows (2.2 vs. 2.1, respectively).

When cows were divided into 5 categories based on P4 concentrations 7 d after insemination, more (P < 0.01) DO cows had P4 concentrations <2.0 ng/mL, whereas more (P < 0.01) EST cows had P4 ≥2.0 ng/mL (Figure 3, lower right panel). The lower P4 distribution 7 d after insemination for DO cows is possibly due to smaller ovulatory follicles because preovulatory follicular size is positively correlated with P4 concentrations after ovulation (Vasconcelos et al., 2001). High-producing Holstein cows have increased hepatic metabolism of steroid hormones (Vasconcelos et al., 2003; Wiltbank et al., 2006). For cows ovulating spontaneously after luteolysis, the dominant follicle requires a longer period of growth for estradiol concentrations to exceed the threshold required to induce a GnRH surge from the hypothalamus, which subsequently induces an LH surge from the anterior pituitary, thereby resulting in ovulation of larger/older dominant follicles (Wiltbank et al., 2006). By contrast, cows submitted to a Double-Ovsynch protocol ovulate smaller follicles because the LH surge and ovulation is induced by the final exogenous GnRH treatment at the end of the protocol, which occurs earlier during development of a synchronized follicular wave (Cerri et al., 2009; Wiltbank and Pursley, 2014).

**Effect of Treatment and Synchrony to the Protocols on P/AI**

The effect of treatment and synchrony to the hormonal protocols on P/AI was assessed for a subset of cows (n = 490) in which we had a complete set of P4 concentrations at insemination did not differ (P = 0.93) between primiparous versus multiparous cows (0.26 vs. 0.25 ng/mL, respectively). When cows were divided into 5 categories based on P4 concentrations at insemination (Figure 3, lower left panel), more (P = 0.01) EST cows had P4 <0.2 ng/mL, whereas more DO cows had P4 concentrations between 0.2 and 0.39 ng/mL. Nonetheless, the proportion of cows with P4 >0.4 ng/mL did not differ (P = 0.91) between treatments or parities [P = 0.27; 2.6% (2/194) vs. 4.5% (14/311) for primiparous vs. multiparous cows, respectively].
samples collected at d 24 (i.e., at the time of PGF$_{2\alpha}$), at the time of insemination, and 7 d after insemination (Table 3). Synchrony to the hormonal synchronization protocols was defined as cows with high P4 (i.e., >0.5 ng/mL) on d 24 (Figure 1), low P4 (<0.4 ng/mL) at insemination, and high P4 (≥1.0 ng/mL) 7 d after insemination. Using a similar methodology to other experiments (Giordano et al., 2012; Lopes et al., 2013; Carvalho et al., 2014a), P/AI of synchronized versus nonsynchronized cows was compared, and nonsynchronized cows were removed in a stepwise manner before making the same comparison at the next time point during the protocols. The percentage of cows with P4 <0.5 ng/mL on d 24 was 5.3% (26/490) and did not differ between treatments (Table 3). Overall, P/AI for cows with P4 <0.5 ng/mL was 7.7% (2/26) and did not differ between treatments; however, for cows with P4 ≥0.5 ng/mL, DO cows had more ($P = 0.03$) P/AI than EST cows (50.1 vs. 40.5%). After removal of cows with P4 <0.5 ng/mL at d 24, the percentage of cows with P4 ≥0.4 ng/mL at the time of insemination was 3.0% (14/464) and did not differ between treatments. For cows with P4 ≥0.4 ng/mL near the time of insemination, P/AI was 21.4% (3/14), and did not differ between treatments. For cows with P4 <0.4 ng/mL at the time of insemination, DO cows had more ($P = 0.03$) P/AI than EST cows (30.0 vs. 0.0%). Cows with P4 >0.5 ng/mL on d 24 and P4 <0.4 ng/mL at the time of insemination were further analyzed for P4 7 d after insemination. The percentage of cows with P4 <1.0 ng/mL 7 d after insemination was 7.1% (32/450) and did not differ between treatments. For cows with P4 <1.0 ng/mL, P/AI was 54.7 vs. 44.5%). The analysis in Table 3 resulted in an overall synchronization rate of 85.3%, which did not differ between treatments. Overall P/AI for cows considered not synchronized was 11.1% (8/72); however, for cows that were considered synchronized, 23% more ($P = 0.04$) DO cows were pregnant 33 d after insemination than EST cows (54.7 vs. 44.5%).

Synchronization rate measured by response to each of the sequential hormonal treatments within a TAI protocol has a profound effect on P/AI, and differences in P/AI to TAI between protocols have been attributed to differences in the proportion of cows considered synchronized using a similar methodological approach to that used in the present study (Giordano et al., 2012; Lopes et al., 2013; Carvalho et al., 2014a). The results of the synchronization analysis in Table 3 support that the effect of treatment on P/AI was not due to a difference in synchronization rate, but rather resulted from a 23% increase in P/AI 33 d after insemination for DO than for EST cows. Thus, because the proportion of synchronized cows did not differ between treatments, DO cows had more P/AI than EST cows because of an intrinsic increase in fertility suggesting that inseminating high-producing Holstein cows after a detected estrus results in lower fertility than when cows receive TAI after submission to a fertility program.

**Table 3.** Effect of treatment on percentage (no./no.) of cows synchronized and effect of synchronization status on pregnancies per AI (P/AI) of lactating Holstein cows

<table>
<thead>
<tr>
<th>Item</th>
<th>Treatment</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>P4 $^1$ on d 24 (i.e., at PGF$_{2\alpha}$)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cows with P4 &lt;0.5 ng/mL</td>
<td>4.5 (13/287)</td>
<td>6.4 (13/203)</td>
</tr>
<tr>
<td>P/AI for cows with P4 &lt;0.5 ng/mL</td>
<td>7.7 (1/13)</td>
<td>7.7 (1/13)</td>
</tr>
<tr>
<td>P/AI for cows with P4 ≥0.5 ng/mL</td>
<td>50.1 (140/274)</td>
<td>40.5 (77/190)</td>
</tr>
<tr>
<td>P4 at insemination</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cows with P4 ≥0.4 ng/mL</td>
<td>3.7 (10/274)</td>
<td>2.1 (4/190)</td>
</tr>
<tr>
<td>P/AI for cows with P4 ≥0.4 ng/mL</td>
<td>30.0 (3/10)</td>
<td>0.0 (0/4)</td>
</tr>
<tr>
<td>P/AI for cows with P4 &lt;0.4 ng/mL</td>
<td>51.9 (137/264)</td>
<td>41.4 (77/186)</td>
</tr>
<tr>
<td>P4 7 d after AI</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Cows with P4 &lt;1.0 ng/mL</td>
<td>7.2 (19/264)</td>
<td>7.0 (13/186)</td>
</tr>
<tr>
<td>P/AI for cows with P4 &lt;1.0 ng/mL</td>
<td>15.8 (3/19)</td>
<td>0.0 (0/13)</td>
</tr>
<tr>
<td>P/AI for cows with P4 ≥1.0 ng/mL</td>
<td>54.7 (134/245)</td>
<td>44.5 (77/173)</td>
</tr>
<tr>
<td>Estrous cycle synchronization</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Synchronized cows</td>
<td>85.4 (245/287)</td>
<td>85.2 (173/203)</td>
</tr>
<tr>
<td>P/AI for cows not synchronized</td>
<td>16.7 (7/42)</td>
<td>3.3 (1/30)</td>
</tr>
<tr>
<td>P/AI for synchronized cows</td>
<td>54.7 (134/245)</td>
<td>44.5 (77/173)</td>
</tr>
</tbody>
</table>

$^1$Each week, lactating Holstein cows at 50 ± 3 DIM (d 0) were stratified by parity (primiparous vs. multiparous) and were randomized to receive first insemination as a timed AI after a Double-Ovsynch (DO) protocol or AI to a detected estrus after a hormonal protocol that synchronized estrus (EST).

$^2$P4 = progesterone.
The specific physiological mechanisms by which Double-Ovsynch and TAI increases fertility in high-producing Holstein cows was not assessed in the present study. Cerri et al. (2009) evaluated embryos flushed from cows inseminated after an induced estrus and cows submitted to TAI after a Double-Ovsynch protocol at a similar DIM. Treatment did not affect fertilization rate (89 vs. 88%, for cows inseminated after an induced estrus versus cows receiving TAI after a Double-Ovsynch protocol); however, the percentage of embryos classified as degenerated was greater for cows inseminated after an induced estrus than for cows submitted to TAI after a Double-Ovsynch protocol (24 vs. 5%). Therefore, it is possible that increased P4 concentrations during growth of the dominant follicle coupled with ovulation of smaller dominant follicles when cows are submitted to TAI after a Double-Ovsynch protocol increased embryo quality resulting in more P/AI than when cows were inseminated after a detected estrus (Wiltbank et al., 2011a,b). In addition, timing of insemination relative to ovulation is more precise when cows are submitted to a synchronization protocol than when they are inseminated after a detected estrus (Valenza et al., 2012).

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