

Use of repopulation for optimizing sow reproductive performance and piglet loss

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Abstract

The objective of this study was to analyse sow reproductive performance and piglet loss from birth to weaning before and after repopulation of a selected farm. The observation was carried out on a productive sow farm; 160 sows were included in the experiment. Before repopulation, 80 sows from the 1st to the 5th litter were evaluated. The original population of sows was removed. Repopulated (newly stocked) group consisted of 80 gilts at the 1st litter. Newly delivered gilts with a status of minimum disease were placed into a decontaminated stable with a strict batch, black and white breeding system with stringent hygienic provisions. The piglets were weaned at the mean age of 28 ± 3 days. A very highly significant difference ($P \leq 0.001$) in favour of the repopulated group of sows was found in the evaluation of the total number of piglets, the number of live-born piglets and the number of stillborn piglets in %; a significant difference ($P \leq 0.05$) in favour of the repopulated group of sows was found in the number of stillborn piglets. In reared piglets and losses of piglets, both in numbers and percentage, a highly significant difference ($P \leq 0.001$) was found in favour of the repopulated group. This study brings important information on the benefits of repopulation pig breeding.

Diseases, health of pigs, rearing, reproduction traits, status with minimum disease

An essential condition for effective breeding of sows is to ensure good health and high performance of sows, which determine the number of reared piglets per sow (Boudný and Špička 2012; Horký et al. 2013). Rodríguez et al. (2012) consider the number of reared piglets a major economic effect of breeding sows. Optimal reproductive management is beside various endogenous and exogenous factors influenced by the health condition, which is then reflected in rearing and fattening of pigs and therefore in whole profitability of a farm. Poor health situation in herds negatively influences the farm economy (Lambert et al. 2012).

According to Opriessnig et al. (2011) great economic losses are caused worldwide by infectious or bacterial diseases of the respiratory tract of piglets. Smola and Masaříková (2007) report that these diseases occur in various combinations, causing the porcine respiratory disease complex (PRDC).

Bad health situation on farms can be solved by the system of radical repopulation method (Guedes et al. 2002). According to O'Donoghue and Ballantyne (1965) the principle of the repopulation method lies in the disruption or restriction of the contact of born piglets with their mothers and isolated rearing of the young. The major advantage of this method is the possibility of herd recovery from all diseases which are not transmissible *in utero* while maintaining the genetic potential. The method consists of extracting piglets shortly before birth either by Caesarean section or by extraction of the whole uterus (hysterectomy) or by aseptic capture of piglets (Young 1960). According to Nevrkla et al. (2013), the disease's life cycle can be interrupted this way as there is no contact between piglets and sow. This method is known as status with minimum disease (MD). The method is economically more demanding than conventional rearing of piglets by sows. Therefore, it is recommended

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for production of pigs on breeding or reproductive farms. On production farms sows give birth naturally (Jorsal and Thomsen 1988). It is necessary to follow the rules of biosecurity to prevent reinfection in a repopulated farm (Laanen et al. 2013). Jorsal and Thomsen (1988) state that the method of repopulation and creating MD herds should have a positive effect on reproductive performance of sows, piglet loss reduction, improvement of productive properties of pigs, improvement of the herd health situation and therefore reduction of treatment costs.

The objective of the work was an analysis of sow reproductive performance and piglet loss from birth to weaning after repopulation of a selected farm.

Materials and Methods

The observation was carried out on a productive sow farm, where 160 sows were included in the experiment. The experiment was approved by the Ethics Committee of Mendel University Brno (accreditation no. 57890/2012-MZE-17214).

Before repopulation, 80 sows from the 1st to the 5th litter were evaluated.

In the group of sows before repopulation, virological and serological tests (Table 1) were performed in the diagnostic laboratory of the State Veterinary Institute in Olomouc and Jihlava for the presence of porcine respiratory and reproductive syndrome (PRRS), porcine circovirus (PCV-2), porcine parvovirus (PPV), Aujeszky's disease (AD), brucellosis (BA) and classical swine fever (CSF). ELISA test was used for the diagnostics of PRRS, PCV-2, AD and CSF. Haemagglutination-inhibition test was used for the presence of PPV. BA was diagnosed by the complement fixation (CF) and Rose Bengal test (RBT).

Table 1. Diagnostics of the presence of disease in blood samples.

Disease	Samples tested	Positive samples	Negative samples
PRRS (E.s.)	5	5	0
PCV-2*	5	3	2
PPV	5	5	0
AD	2	0	2
BA	2	0	2
CSF	2	0	2

PRRS – porcine respiratory and reproductive syndrome, E.s. – European strain, PCV-2 – porcine circovirus, PPV – porcine parvovirus, AD – Aujeszky's disease, BA – brucellosis, CSF – classical swine fever, * Data relating to IgG antibodies confirming PCV-2 infection in the past. IgM antibodies indicative of acute infection were not detected in any sample.

The original population of sows was removed. Repopulated (newly stocked) group consisted of 80 gilts at the 1st litter. Animals of both groups were of the same hybrid combination.

Newly delivered MD gilts were placed into a decontaminated stable with a strict batch, black and white breeding system with stringent hygienic provisions. Stable entry was via one main entrance with mandatory showering, clothing and footwear exchange for all nursing staff and visitors. Each building entrance was equipped with a disinfection mat for disinfection and cleaning of footwear. A strict control of the movement of persons and visitors in the area of the farm was applied and the entrance of those who came into contact with other pigs during the last 3 days or who breed pigs at home was prohibited. Gilts were brought from approved source farm applying the same strict measures as the observed farm. The group of newly brought breeding gilts were first acclimated and then stabled in quarantine. Vehicles were properly cleaned and disinfected before entering the farm, drivers were not allowed to move either in the area of the farm or in the stables. Vectors such as insect and rodents which are considered viral infection carriers had to be regularly eliminated by means of disinfection and deratization.

Movement of piglets among litters was disabled, except for the first 24 h after birth when necessary. Injection needles and other supplies were used only for one litter. Windows of stables for both served and pregnant sows were equipped with nets against birds and insect. Thorough cleansing and disinfection of stables was performed after batch emptying of each section.

The piglets were weaned at the mean age of 28 ± 3 days. In both groups of sows (before and after repopulation) the phenotypic level of selected reproductive properties was observed: namely, the total number of born piglets,

the number of live-born piglets, the number of stillborn piglets, the number of reared piglets, and piglet losses from birth to weaning.

The obtained reproductive indicators and piglet losses before repopulation were compared to the indicators obtained after repopulation, and elementary statistical characteristics for differences in the evaluated indicators between the groups of gilts were analysed; namely, mean, standard deviation (mean \pm S.D.), and significance based on *t*-test. The symbol *** stands for $P \leq 0.001$, ** stands for $P \leq 0.01$, * stands for $P \leq 0.05$ and NS stands for $P \geq 0.05$. The statistical evaluation was done using the STATISTICA version 10.0 software and Microsoft Excel 2010.

Results

The repopulation of sows led to an increase of reproductive performance. Table 2 presents the phenotypic level of sow performance in the total number of piglets, the number of live-born and stillborn piglets per litter before and after repopulation of the farm. A highly significant difference ($P \leq 0.001$) in favour of the repopulated group of sows was found in the total number of piglets, the number of live-born piglets, and the number of stillborn piglets in % ; a significant difference ($P \leq 0.05$) in favour of the repopulated group of sows was found in the number of stillborn piglets. Application of farm repopulation brought an increase of the total number of piglets by 2.72 per litter in repopulated sows. The number of live-born piglets increased by 3.25 piglets per litter and the number of stillborn piglets decreased by 0.54 (5.36% difference).

Table 2. Basic statistical characteristics of the total number of piglets, the number of live-born and stillborn piglets.

Indicator	Criterion	N. of piglets	Mean \pm S.D.	Significance
Total number of piglets / litter	I	1 057	13.21 \pm 2.51	***
	II	1 274	15.93 \pm 2.20	
Number of live-born piglets / litter	I	910	11.38 \pm 2.20	***
	II	1 170	14.63 \pm 2.09	
Number of stillborn piglets / litter	I	147	1.84 \pm 1.57	*
	II	104	1.30 \pm 1.36	
Number of stillborn piglets (% / litter)	I	147	13.20 \pm 10.44	***
	II	104	7.84 \pm 7.94	

*** $P \leq 0.001$; * $P \leq 0.05$; I – before repopulation; II – after repopulation

Table 3. Basic statistical characteristics of the number of reared piglets and losses of piglets from birth to weaning.

Indicator	Criterion	N. of piglets	Mean \pm S.D.	Significance
Number of live-born piglets / litter	I	910	11.38 \pm 2.20	***
	II	1 170	14.63 \pm 2.09	
Number of reared piglets / litter	I	752	9.40 \pm 1.80	***
	II	1 075	13.44 \pm 1.78	
Loss of piglets / litter	I	161	2.01 \pm 1.45	***
	II	95	1.19 \pm 1.47	
Loss of piglets (% / litter)	I	161	16.10 \pm 10.68	***
	II	95	7.51 \pm 9.02	

*** $P \leq 0.001$; I – before repopulation; II – after repopulation

Table 3 shows the numbers of reared piglets and piglet losses from birth to weaning per one litter. Between both weaned piglets and their losses, in numbers and in percentage, a highly significant difference ($P \leq 0.001$) was proved in favour of the repopulated group.

The repopulation of sows increased the number of reared piglets from one sow per litter by 4.04 piglets. An important criterion for evaluation of the application of farm repopulation is piglet loss. The difference between the groups was 0.82 piglets in favour of the repopulated sows. In percentage, the difference between the groups of piglets was 8.59%.

Discussion

Damgaard et al. (2003) point out that litter size affects the survival of piglets after birth. Olanratmanee et al. (2010) report 12.1 born piglets per litter for sows in good hygienic conditions vs. 11.7 born piglets for sows in poor hygienic conditions which indicates the need for a good health status of breeding sows; they further found 10.3 live-born piglets per litter for sows with health problems compared to 11.1 live-born piglets for sows without health problems. Lewis et al. (2009) state that PRRS virus influences reproductive performance of sows and gilts. Their results show that healthy gilts at the first litter had over 9 live-born piglets compared to only 7 live-born piglets of gilts with health issues; furthermore they found 3.0 stillborn piglets per litter for diseased gilts and 0.6 stillborn piglets per litter for healthy sows. Their observation highlighted higher incidence of stillborn piglets in gilts. According to Cozler et al. (1998) the number of reared piglets is used for expressing sow performance. These authors note that sow productivity depends mainly on the genetics and farm management, which includes also appropriate health programs. Lewis et al. (2009) found 7.5 reared piglets per litter for sows with health problems compared to 9.25 reared piglets per litter for healthy sows.

O'Donoghue and Ballantyne (1965) report that specific pathogen free (SPF) sows are characterized by lower losses of piglets before weaning, but they emphasize that repopulation itself is not sufficient and that it is necessary to ensure strict hygiene in the herd. Munsterhjelm et al. (2006), Andersen et al. (2009), and Oliviero et al. (2010) state that appropriate health programs in sow herds minimize piglet loss after birth. According to Rootwelt et al. (2012) the loss of piglets from the live-born to the weaned in problematic herds reaches up to 16.20%. Jung et al. (2008) state that viral infections present in some herds can increase mortality of newborn piglets, adding that rotavirus infections are the cause of acute diarrhoea in suckling piglets and outbreaks of these infections are associated with an impaired immune system response. Vaillancourt et al. (1992) report that intensive sow production is accompanied by certain critical phases. Loss of piglets from birth to weaning is considered an important one, either as a result of infectious diseases or nonpathogenic causes; therefore, monitoring of piglets allows its optimization. They also point out that in problematic herds the losses can be very high. For example, piglet loss before weaning in the worst herds reached 12–30% in England, 17.6% in Croatia, and 22.2% in Slovenia.

Values of the observed reproductive indicators and piglet loss can be considered very competitive from the current perspective after repopulation, therefore the recovery of farms by means of repopulation and creating MD farms can be recommended. Strict rules of biosecurity are a necessity on repopulated farms for maintaining a good health status of sows and piglets, which can have an impact on better sow performance of sows and piglet survival.

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