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Juarez, JD.; Marco-Jiménez, F.; Vicente Antón, JS. (2021). Evaluation of foetal growth, litter size and reproductive performance in rabbit after 18 generations of selection for growth rate using cryopreserved embryos. Livestock Science. 253:1-7. https://doi.org/10.1016/j.livsci.2021.104702



The final publication is available at

https://doi.org/10.1016/j.livsci.2021.104702

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Additional Information

- Evaluation of foetal growth, litter size and reproductive performance in rabbit after 18
 generations of selection for growth rate using cryopreserved embryos
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8 Abstract

In livestock, adverse effects on reproductive performance, health traits and robustness have been demonstrated in animals selected for high production and efficiency. Using embryo cryopreservation and rederivation, we compared phenotypic traits between rabbit populations separated for 18 generations under growth rate selection pressure (R18 vs R36). To do so, embryos from the ancestral population (R18) and the most recent population (R36) were vitrified in 2000 and 2015, respectively, and rederived and grown together in a randomized controlled environment in 2015. To eliminate confounding maternal and embryo handling effects, traits were measured in the second generation after rederivation (R20 and R38 generations). Our study suggests that selection for growth rate has no adverse effect on litter size components. Thus, in the R38 generation we observed a significant increase in embryo implantation (7.2±0.71 vs 5.1±0.79) and litter size (7.1±0.29 vs 6.5±0.32). Besides, the foetal sac area at day 12 of gestation (2.44±0.070 vs 2.07±0.071 mm², for R38 vs R20, respectively), and foetal placenta area (136.7±6.14 vs 116.0±6.31 mm², for R38 vs R20, respectively) and crown-rump length of the foetus (38.0±0.68 vs 35.8±0.68 mm, for R38 vs R20, respectively) at day 19 of gestation were higher in the R38 generation. Altogether, these results show that selection for growth rate does not adversely affect components of litter size,

foetal growth and reproductive performance. However, the extent to which foundation criteria play a role in the high prenatal and perinatal mortality rate remains unclear in paternal lines.

Keyword: foetal growth, reproductive performance, paternal line, rederived population, rabbit.

1. Introduction

Rabbit lines specialized for growth rate or feed efficiency and litter size are a common selection objective, as in other animal farms, due to their economic interest (Blasco et al., 2018). Nevertheless, the relationship between growth and maternal traits and their correlated responses, unclear or not, is always positive. Although maternal effects have long been acknowledged as potentially important factors in artificial selection, their magnitude of response remains unclear. In mammals, the role of maternal effects is especially complex due to the fact that progeny experience two distinct maternal environments (prenatal uterine and postnatal nursing) influenced by numerous factors, such as the number of foetuses or litter size, parity, age, breed, heat stress and nutrition (Vuguin, 2007; Wolf et al., 2011).

The success of selection for high prolificacy in polytocous species is related with negative consequences in survival, foetal growth and birth weight in rabbit (Vicente et al., 1995; Argente et al., 2003 and 2008) and pigs (Damgaard et al., 2003; Foxcroft et al., 2006; Wolf et al., 2008). Argente et al. (2003) found a reduction in placental and foetal development in rabbits with each additional implanted embryo at 25 days of gestation. Moreover, higher intrauterine crowding has been correlated with higher foetal mortality at 18 days of gestation (Argente et al., 2008). In pigs, it has been shown that selection for increased litter size entails the possibility of various degrees of intrauterine growth retardation associated with impaired foetal and placental growth, which can result in lower birth weights (Town et al., 2004). Thus,

postnatal variation in growth performance variation may be pre-programmed during foetal development in the uterus. Furthermore, it is likely that these pre-programmed limitations in growth performance will only finally express themselves in the late grower and not at the early finisher stage of production (Du et al., 2010).

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Some studies suggest that long-term selection for growth rate results in physiological and reproductive changes (Rauw et al., 1998). For example, mice selected for increased early body weight gains showed a decreased response to superovulation and oestrous synchronization, and when they were used as recipients, they produced pups that were significantly larger with respect to body weight and tail length compared with litters gestated in females non-selected or selected divergently (Ernst et al., 2000). These authors suggested that retarded pre-pubescent reproductive development results in reproductive uterine horns more efficient for sustaining pregnancy, foetal development and growth. Thus, cattle breeds with postnatal different growth impetus and muscularity show differences in foetal development, especially in muscle tissue deposition and development (Mao et al., 2008). In rabbit paternal line, the reproductive performance traits are not a selection factor, as reproductive traits like parity and litter size had a negligible effect on growth rate (Piles and Blasco, 2003), but is unclear whether a selection programme for growth rate affects reproductive traits in rabbits. Low or null correlations between litter size at birth and postnatal growth rate have been observed in rabbit (García and Baselga, 2002; Mocé and Santacreu, 2010; Drouilhet et al., 2013; Mínguez et al., 2016; Peiró et al., 2019) and an uncertain pattern was found in pigs (Damgaard et al., 2003 Wolf et al., 2008; Zhang et al., 2016). Meanwhile, a positive genetic correlation was estimated between postnatal growth rate and ovulation rate in pig and rabbit (Bidanel et al., 1996; Ruiz-Flores and Johnson, 2001; Drouilhet et al., 2013; Peiró et al., 2019). In rabbit, reproductive differences have been described between maternal

and paternal lines. Females from the paternal line showed widespread failures, such as altered LH and steroidogenic patterns, low response to ovulation frequency and high losses in implantation during foetal development and birth (Vicente et al.; 2012, Naturil-Alfonso et al., 2015 and 2016). Paternal line males did not present normal sexual behaviour, observing low libido, lower sperm production (Pascual et al., 2004; Rosell and De La Fuente, 2009) and changes in seminal and sperm proteome (Lavara et al., 2011; Casares-Crespo et al.; 2018, Juárez et al., 2020).

The aim of this study was to evaluate whether the selection programme for daily gain in fattening period has changed foetal growth and prenatal survival, using two rederived populations separated for 18 generations. To this end, embryos from the population R18 and the most recent population (R36) were vitrified, rederived and grown together in a randomized controlled environment. To rule out confounding maternal and embryo handling effects, prenatal growth traits and litter size components were measured in the second generation after rederivation (R20 and R38).

2. Materials and methods

The animal study protocol was reviewed and approved (code number 2015/VSC/PEA/00061) by Ethical Committee of the Universitat Politècnica de València before initiating the study. All experiments were performed following guidelines and regulations outlined in Directive 2010/63/EU EEC. Animal experiments were conducted at an accredited animal care facility (code: ES462500001091).

2.1. Animals

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A rabbit paternal line (R) selected at the Universitat Politècnica de Valencia was used. This line was founded in 1989 from two closed paternal lines selected according to individual weigh gain from weaning to end of fattening period (77 days old) during 12 and 9 generations (Estany et al., 1992). Since then, the line has been selected for individual weight daily gain from 28 days (weaning) to 63 days of age (end of fattening). For the present study, two populations (R19V and R37V) separated for 18 generations of selection were used. R19V population was rederived from 256 embryos of 25 donors belonging to 10 different sire families of 18th generation and vitrified in 2000. R37V population was rederived from 301 embryos from 28 donors belonging to 15 different sire families of 36th generation, which were vitrified in 2015. Both populations were rederived at the same time in 2015 (see details in Marco-Jiménez et al., 2018). Offspring were bred in non-overlapping generations; 101 females from generations 20 and 38 were used in this experiment (named R20 and R38, respectively). Environmental conditions were maintained using a control system for light (16:8 light/dark photoperiod), ventilation and temperature (18-25 °C) and relative humidity: 60 to 75% by a forced ventilation system. Rabbits does were fed ad libitum throughout the gestation and lactation period with a commercial pelleted diets (2900 kcal of digestible energy / kg, 3.5% crude fat, 15.5% crude fibre and 17% crude protein dry matter). Non-pregnant rabbit does were fed with 140 g/animal/day until a positive pregnancy diagnosis.

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2.2. Reproduction management

One hundred and forty-two females were inseminated by males from the corresponding generation (60 from R20 and 82 from R38). To control inbreeding, males and females sharing a grandparent were avoided. Receptivity of does was improved with 12-15 UI

of eCG via intramuscular 48 hours before insemination. First insemination was performed at 20 weeks of age and then 10-12 days after parturition. Fourteen days post-insemination, pregnancy was diagnosed by abdominal palpation and, if they were non-pregnant, females were inseminated again 7 days later. Young rabbits were weaned at 28 days of age.

Insemination was performed after evaluation of ejaculate with 0.5 ml and 20 to 40 million sperm per seminal dose. Only ejaculates with more than 70% of motility rate and less than 20% of abnormal sperm were used. Ejaculates were diluted with tris-citric-glucose diluent to adjust the concentration (Viudes-de-Castro and Vicente, 1997). Immediately after insemination, ovulation was induced by an intramuscular injection of 1µg of Buserelin Acetate (Suprefact, Hoechst Marion Roussel, S.A., Madrid, Spain). Reproductive status of does at insemination time (nulliparous, primiparous lactating, multiparous lactating and non-lactating does), total litter size, liveborn and litter size at weaning were recorded for each female.

2.3. Laparoscopy and evaluated litter size components

A total of 85 does were used and 103 laparoscopies were carried out on females from fourth and fifth parity (38 does from R20 and 47 does from R38). In brief, the females were sedated with intramuscular injection of 5 mg xylazine/kg (Rompun, Bayer AG, Leverkusen, Germany) and 3 mg/kg morphine chloride (). Five minutes later, 35 mg/kg Ketamine hydrochloride (Imalgene®, Merial, S.A., Lyon, France) was administered intravenously. After laparoscopy, does were treated with antibiotics (200,000 IU procaine penicillin and 250 mg streptomycin, Duphapen® Strep, Pfizer, S.L.), 0.03 mg/kg of buprenorphine hydrochloride every 12 hours and 0.2 mg/kg of meloxicam every 24-h for 3 days. The number of corpora lutea, the number of implanted embryos at 12 days (IE) and litter size at birth (LS) were recorded per female. The following variables were calculated using the above data. Ovulation

rate (OR), defined as the number of corpora lutea, was recorded to determine ovulation rate (OR), embryonic loss rate (ELR), estimated as (OR-IE)/OR, and foetal loss rate (FLR), estimated as (LS-EI)/LS.

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2.4. Foetal growth. Ultrasound measurement

Thirty-one pregnant does from laparoscopised females (15 from R20 and 16 from R38) were examined on day 12, 19 and 26 of gestation using a portable colour Doppler ultrasound device (Esaote, Spain) with a 7.5 MHz linear probe (4-12 MHz range). Does were sedated according to the procedure described above and placed in a polystyrene cage where they were prevented from moving. The ultrasound examination was performed from right to left with the probe in sagittal orientation and, after localization of different foetal sacs, 4-6 whole foetal sac examinations per doe were performed. The identifiable structures (foetal sac, foetus and foetal and maternal placenta) were measured from frozen frame pictures on the monitor, using the Esaote 16 ultrasound software. Measurements on different days of gestation are illustrated in Fig. 1. Briefly, foetal sac (FS, A, C and E) measurements were taken when the largest surface area appeared on the screen. For whole foetus measurements, crown-rump length (CRL) was determined as the maximum distance from crown to tail base, with the foetus on a sagittal plane (Fig. B, D and F). Placental size was difficult to assess, but placental measurements were determined when the maximal placental surface with the twolobed foetal (L1FP and L2FP) and maternal placenta (MP) were visible on screen (Fig. 1A, 2A, 3A).

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2.5. Statistical analyses

Reproductive performance was analysed with a linear general model including as fixed effects rederived generation group (R20 and R38), reproductive status of does (nulliparous, primiparous lactating, multiparous lactating and non-lactating does), month-year in which insemination was done (18 levels) and the interaction between generation group and reproductive status of the mothers. Delivery rates were analysed using a probit link with binomial error distribution, included in the generalized model described above.

Litter size components (ovulation rate, implanted embryos, litter size, liveborn and rates of embryo and foetal losses) were analysed by a generalized linear model including as fixed effects rederived generation group (R20 and R38) and lactating or non-lactating status and their interactions.

To analysis foetal sac area, crown-rump length of foetus, foetal and maternal placenta areas at days 12, 19 and 26 of gestation and, weight of liveborn kits, a mixed linear mode was used:

$$Y_{ijkl} = \mu + P_i + R_j + PR_{ij} + CO_k + Cov X_l + e_{ijkl}$$

, where Yijkl was the trait to analyse, μ was the general media, P_i was the fixed effect of the rederived generation group (R20 and R38), R_j was the fixed effect of reproductive status of the doe used to analysis of weaning weight (lactating and non-lactating doe); PR_{ij} was the effect of interaction between rederived population and reproductive status of the mothers used to analysis of weaning weight, CO_k was the random effect of common litter, $Cov X_l$ was the covariate of number of implanted embryos and eijklm was the error term of the model.

Values were considered statistically different at P < 0.05. Results were reported as least square means with standard error of the mean (SEM). All analyses were performed with SPSS 26.0 software package (SPSS Inc., Chicago, Illinois, USA, 2012).

3. Results

3.1. Reproductive performance between rederived populations

Of the different reproductive parameters evaluated, only litter size at parturition was significantly different between the two rederived populations, with the litter size being larger in the most recent generation R38 (6.5 0.32 vs 7.1 0.29, Table 1). However, the high mortality in both groups around delivery and during the lactation period (more than 20% and 40%, respectively, Table 1) make this difference irrelevant. In addition, the delivery rate was also similar in both groups (Table 1).

Considering the reproductive status of the doe at the time of insemination, nulliparous females had the highest delivery rates, the lowest litter size and the highest mortality rate during lactation (Table 1). Furthermore, it is necessary to highlight that the females with 1 or more deliveries that were not pregnant presented problems to gestate again, obtaining the lowest delivery rate (Table 1).

Non-significant effects of year-month and interactions between generational group and week of reproductive status were observed for all evaluated traits.

3.2. Litter size components between rederived groups

Litter size components evaluated such as ovulation rate, foetal and embryo losses were not different between the rederived populations, despite the 18 generations of selection that separate them (Table 2). However, a significant increase in the number of implanted embryos was observed in R38 vs R20 population if the number of implanted embryos included those females that did not implant any. Both rederived populations showed high rates of loss at implantation and from implantation to parturition, but not significantly different (Table 2). Total implantation failure was 38.6% (17) in R20, while this occurred at 23.7% (14) in the R38

population. Non-significant interactions between generational group and lactation status were observed for evaluated traits.

3.3. Foetal growth during gestation

Early differences in foetal sac growth, foetal placenta and Crown-rump length of foetuses were observed at day 12 and 19 of gestation, respectively, between both populations (Table 3). At the onset of gestation, the lactation status affected the foetal sac size at 12 and 19 days and the foetal placenta size at 19 days. On day 26 of gestation, neither the generation nor the lactation status affected any trait. The bodyweight of the liveborn kits was significantly different between generations, but was not influenced by the lactating status. The highest birth weight was obtained in the offspring of R38 population (57.2±3.47 vs 69.3±3.89 for R20 and R38 from 48 and 42 liveborn kits, respectively. Data not shown in tables).

4. Discussion

To the best of our knowledge, this is the first in-depth study to evaluate the effect of selection for growth on reproductive parameters or traits using populations rederived (reestablished) from cryopreserved embryos. Although rarely used in selection experiments, cryopreserved populations offer the advantages of optimizing the experimental facilities and reducing genetic drift (García and Baselga, 2002; Piles and Blasco, 2003), and it is a successful strategy to re-establish a population to continue the breeding programme (Marco-Jiménez et al., 2018). Our study suggests that selection for growth rate has no adverse effect on litter size, foetal growth and reproductive performance, and we even observed a slight improvement in the embryo implantation and litter size and bodyweight at birth after 18 generations of selection.

Livestock animals selected for high production efficiency (litter size, growth, feed efficiency or carcass composition and meat quality) have impaired reproductive performance (Rauw et al., 1998; Bunger et al., 2005) or health traits and robustness (Rauw et al., 1998; Prunier et al., 2010; Phocas et al., 2014; Rauw and Gomez-Raya, 2015). These adverse effects of selection are often difficult to reveal as a consequence of not being registered or because the feeding or environmental conditions are being modified. For example, in swine a longterm selection for a combined breeding goal (growth, feed efficiency, body composition and litter size) has been accompanied by an improvement in litter size and weight, but unfavourable effects of selection for several traits such as an increase in stillbirths and in postnatal mortality, reduced longevity and productive life, a reduced milk production and robustness (Silalahi et al., 2016 and 2017). It is known that rabbit selection for growth traits has negative genetic correlations on ejaculate traits such as mass motility, volume, abnormal sperm rate or head sperm morphometry (Brun et al., 2006: Lavara et al., 2012 and 2013) and the female contribution to fertility has been found (Piles and Tussel, 2012). Moreover, several studies showed an impaired reproductive performance of paternal line R when it was compared with maternal lines, with high embryonic, foetal, perinatal losses and during the lactation period (Vicente et al., 2012 and 2013; Naturil-Alfonso et al., 2016). Nevertheless, the present study showed that selection for growth rate does not adversely affect litter size and reproductive performance. It is worth mentioning that, after 18 generations of the selection process, females increased in implanted embryos, ending in improved litter size. Moreover, similar prenatal losses were observed between both generations, in line with our previous results in which implantation and gestational losses were around 20-30% and 50%, respectively (Vicente et al., 2012). In maternal or crossed rabbit lines, embryonic and foetal loss rates are 10 and 20%, respectively, and up to 15% for perinatal losses (Santacreu et al.,

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1992; Fortun et al., 1993; García and Baselga, 2002; Santacreu et al., 2005; Mocé et al., 2005; Vicente et al., 2012; Ragab et al., 2014). Some of the causes of the high implantation failures and foetal losses of this paternal line could be linked to high levels of IGF-1 and leptin, lower oestrogen and progesterone levels and lower mRNA expression levels of insulin-like growth factor II receptor (IGF-IIR) at endometrial tissue found in the females (Vicente et al., 2012; Llobat et al., 2012; Naturil-Alfonso et al., 2016).

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Additionally, perinatal and lactation mortality rates were similar and high between both generations. Perinatal and lactation losses found in both can be associated with impaired maternal behaviour and were already reported for this line by Lavara et al. (2002). Moreover, this could be related to the abnormal levels of oestradiol and progesterone during gestation observed in this line (Vicente et al., 2013) and with a low litter size. This endocrine disruption might trigger a cascade of events that would affect the construction of the nest, the pheromonal cues, nursing behaviour and, finally, milk production. González-Mariscal et al. (2016) reviewed maternal behaviour and sibling interactions in rabbits, describing the role of changing concentrations of oestradiol, progesterone and prolactin throughout gestation to prime the mother's brain to respond to the newborn and as regulators of nest-building, and how the duration and periodicity of nursing will depend on the stimulation of the teats by the kits (suckling young). This should be evaluated in a subsequent study in order to improve reproductive efficiency, among other things, carry out adoptions at birth to prevent the number of young rabbits from being less than 6 so that an adequate nursing behaviour develops. Regarding negative outcomes of non-lactating does, it is probable that in spite of feed control, these females tended to accumulate fat and had difficulty mobilizing during gestation or lactation. R line does seem to prioritize maintaining their heavier body size rather than litter development, a difference from other lines (Arnau-Bonachera et al., 2018 a and b),

which may be further aggravated if does do not become pregnant and continue to gain weight (Rommers et al. 1999) and, consequently, negatively affects their long-term reproductive function (Castellini et al., 2006).

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This study also enabled us to explore whether selection for growth rate affects placentae and foetal growth. Our findings revealed that the foetal sac and foetal placenta area at day 12 of gestation and foetal placenta area at day 19 of gestation was higher in the R38 generation. Nevertheless, no differences between generations were found at day 26 of gestation, indicating that the possible effects of both selection and the gestation-lactation overlap were compensated at the end of gestation but not in the weight of liveborn kits. This could be because during the last week of gestation the fastest increase in bodyweight takes place (Vicente et al., 1995 and 2013; Argente et al., 2003). So, we have observed evidence that growth selection influenced foetal structures and final weight of foetuses. The changes in foetal structures take place at a critical period between day 12 and 19 of gestation, in which organogenesis is defined, which could be important in the final growth of gestation and during postnatal life (Vuguin, 2007; Sartori et al., 2020). During foetal development, extrauterine signalling provides a link between environmental changes and physiology of the foetus as the impetus to prepare the organism for the postnatal environment, guided mainly by epigenetic changes (Gluckman et al., 2005; Sarkies, 2020). If these adaptive responses are directed to a nutritionally deficient postnatal environment, they could potentially affect muscle, adipose and reproductive tissue development (Ford et al., 2007; Ford and Long, 2012). Skeletal muscle or reproductive tissue have a lower priority in nutrient partitioning compared to the brain and heart in response to challenges to the foetus during development, and are particularly vulnerable to nutrient deficiency (Caton et al., 2019; Crouse et al., 2019). Females from the paternal rabbit line used in this study, regardless of the repercussions due to the selection

process, shows some negative phenotypic characteristics at endocrine level and a different nutritional partition that could be triggered during prenatal development and in the first steps of postnatal development (suckling). So, a more in-depth study of these stages is necessary.

In conclusion, selection for growth rate does not adversely affect components of litter size, foetal growth and reproductive performance in rabbit does. Nevertheless, this study reinforces some significant reproductive problems, such as high prenatal and perinatal mortality in this paternal line, that were already present in generation 18. Therefore, further studies must elucidate how the founders and not the selection process could play a fundamental role in the adverse reproduction outcomes.

5. Acknowledgements

This research was supported by AGL2017-85162-C2-1-R research project funded by the Ministry of Economy, Industry and Competitiveness (MICINN, Spain). English text version revised by N. Macowan English Language Service.

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Graphical abstract.

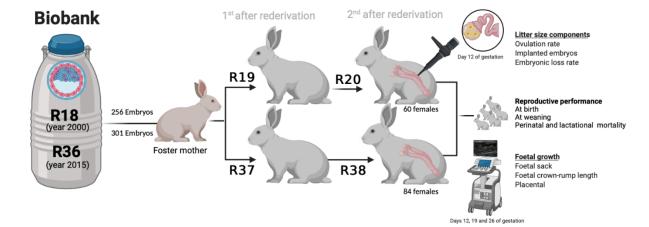


Table 1. Effect of selection for growth rate on total litter size, liveborn, litter size at weaning, perinatal and lactation mortality rates and delivery rate from two rederived population separated by 18 generations.

Туре		n	Litter size			Mortality rate (%)		Delivery rate (%)
			Total	Liveborn	At weaning	Perinatal	Lactation	Delivery rate (70)
Generation	R20	179	6.5±0.32 ^b	5.1±0.36	3.1±0.34	23.2±3.33	46.1±4.11	0.57±0.029
Generation	R38	278	7.1±0.29 ^a	5.8±0.33	3.7±0.31	21.9±3.03	42.5±3.68	0.59±0.024
	Nulliparous	142	5.9±0.37 ^b	4.1±0.35 ^b	2.3±0.39 ^b	29.8±3.86	56.7±4.80 ^b	0.76±0.032 ^a
Lactation status	Primiparous Lactating	75	7.1±0.46 ^a	5.7±0.52 ^a	3.2±0.49 ^{ab}	23.3±4.76	46.4±5.91 ^{ab}	0.59±0.044 ^b
Lactation status	Multiparous lactating	143	7.1±0.36 ^a	6.1±0.41 ^a	4.2±0.38 ^a	17.3±3.72	33.8±4.60 ^a	0.58±0.032 ^b
	Non-lactating	97	7.1±0.37 ^a	5.9±0.42 ^a	3.9±0.39ª	19.9±3.86	40.3±4.72 ^a	0.37±0.031 ^{ca}

n: Number of inseminations. Data are expressed as least squared mean \pm standard error of means. ^{a,b} Values with different superscripts in column differ significantly (p < 0.05).

Table 2. Effect of selection and lactation status on litter size components from two rederived population separated by 18 generations.

543

544			Implanted embryo		Loss rate (%)			
545	Туре		Ovulation rate ¹	Implanted	² Implanted			Litter size
546				embryo	embryos	¹ Embryonic	Foetal	
547		R20	12.1±0.45	5.1±0.79 ^b	8.1±0.73	37.0±5.19	42.8±7.31	5.0±0.80
548		(n)	(44)	(44)	(27)	(27)	(27)	(26)
549	Generation	R38	12.7±0.39	7.2±0.71 ^a	9.4±0.60	28.5±4.26	33.4±6.00	6.7±0.65
550 551		(n)	(59)	(59)	(45)	(45)	(45)	(44)
552		Non-	12.8±0.47	6.5±0.86	8.3±0.71	36.7±5.11	40.0±7.19	5.7±0.79
553		lactating						
54	Lactation status	(n)	(36)	(36)	(28)	(28)	(28)	(27)
555		Lactating	12.0±0.37	5.8±0.64	9.1±0.61	28.9±4.36	36.2±6.14	6.1±0.66
56		(n)	(67)	(67)	(44)	(44)	(44)	(43)

Data are expressed as least squared mean \pm standard error of means. n: number of laparoscopies.

561

560

557

¹It was determined as the number of corpora lutea.

² Implanted embryos in pregnant does. 559

^{a,b} Values in the same column and factor with different superscripts are statistically different (P < 0.05).

Table 3. Effect of selection and lactation status on foetal sac and placentae area and foetal size at 12, 19 and 26 days of gestation from two rederived population separated by 18 generations.

		Day of	n	Foetal sac	Maternal	Foetal	Crown-rump
	Group	gestation		area (cm²)	placenta area	placenta area	length of
		gestation		area (em)	(mm²)	(mm²)	foetus (mm)
	R21	12	82	2.07±0.071 ^b	44.9±2.46	42.0±2.29	11.2±0.35
	R39	12	95	2.44±0.070 ^a	50.7±2.44	46.7±2.28	11.5±0.34
Generation	R21	19	77	5.81±0.178	111.8±6.30	116.0±6.31 ^b	35.8±0.68 ^b
Generation	R39	19	94	6.27±0.176	118.2±6.13	136.7±6.14ª	38.0±0.68ª
	R21	26	66	10.01±0.530	197.8±12.21	247.6±15.67	73.1±1.85
	R39	20	62	10.24±0.560	193.1±13.57	244.5±17.59	72.0±1.92
	Non-lactating	12	70	2.36±0.072 ^a	48.4±2.49	46.8± 2.32	11.3±0.35
	Lactating	12	107	2.15±0.058 ^b	47.2±2.02	42.0±1.88	11.4±0.29
Lactation status	Non-lactating	19	69	6.49±0.181ª	116.2±6.23	139.0±6.27ª	36.9±0.71
Lactation status	Lactating	19	104	5.59±0.147 ^b	113.8±5.13 ^b	113.7±5.14 ^b	36.9±0.57
	Non-lactating	26	54	10.15±0.695	211.7±17.01	233.2±22.04	71.6±2.42
	Lactating	20	74	10.10±0.602	179.1±13.90	258.9±17.91	73.6±2.08

a,b Values in the same column and factor with different superscripts are statistically different (P < 0.05).

Figure. 1. Ultrasonography measurements of the foetal sac (FS), crown-rump length (CRL) of foetus and the placental measurements of the two-lobed foetal (L1FP and L2FP) and maternal (MP) at 12, 19 and 26 day of gestation.

