



Production of Pigs From Porcine Embryos Generated *in vitro*

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Generating porcine embryos *in vitro* is a critical process for creating genetically modified pigs as agricultural and biomedical models; however, these embryo technologies have been scarcely applied by the swine industry. Currently, the primary issue with *in vitro*-produced porcine embryos is low pregnancy rate after transfer and small litter size, which may be exasperated by micromanipulation procedures. Thus, in this review, we discuss improvements that have been made to the *in vitro* porcine embryo production system to increase the number of live piglets per pregnancy as well as abnormalities in the embryos and piglets that may arise from *in vitro* culture and manipulation techniques. Furthermore, we examine areas related to embryo production and transfer where improvements are warranted that will have direct applications for increasing pregnancy rate after transfer and the number of live born piglets per litter.

Keywords: pig, porcine embryo, *in vitro*, pregnancy rate, micromanipulation

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INTRODUCTION

In vitro-produced (IVP) embryos are essential for employing assisted reproductive technologies to establish porcine models. Recent development of genome editing techniques has fueled considerable interest in expanding porcine models for biomedical and agricultural purposes. As pigs have similar physiology and more relevant body and organ sizes to humans compared to other laboratory species, porcine models can be used to understand disease progression and to test relevant doses of therapeutics or medical devices. To date, several porcine biomedical models have been generated, including those for cystic fibrosis (Rogers et al., 2008), cardiovascular disease (Turk et al., 2005), cancer (Schook et al., 2015; Hendricks-Wenger et al., 2021), phenylketonuria (Koppes et al., 2020), immunodeficiency (Suzuki et al., 2012; Lee et al., 2014), viral infection (Lei et al., 2016), and xenotransplantation (Lai et al., 2002). Moreover, gene editing has been used to improve carcass traits (Lai et al., 2006) of pigs and to confer resistance to viruses that plague the swine industry (Whitworth et al., 2016, 2019).

The majority of porcine embryo production systems are comprised of three main steps: oocyte maturation, *in vitro* fertilization, and embryo culture. Initially, oocytes are aspirated from ovaries (gilts or sows) and go through the *in vitro* maturation process. The oocytes that reach the metaphase II (MII) stage are selected for *in vitro* fertilization (IVF). Afterwards, presumptive zygotes are cultured *in vitro* prior to embryo transfer. Intensive studies have been conducted to develop porcine embryo culture media (Lee et al., 2013; Spate et al., 2015; Redel et al., 2016b; Chen et al., 2018), and the development of different culture systems has dramatically impacted when embryo transfers are able to be performed. Moreover, the embryo production system is essential for designing

porcine models. To create genetically modified pigs, manipulation of oocytes and embryos is required. The most common methods for introducing mutations into target genes or transgenes are microinjection of the CRISPR/Cas9 system (mRNA or ribonucleoprotein complex) into the porcine zygote or transfection of fetal fibroblasts to generate donor cells for somatic cell nuclear transfer (SCNT) (Figure 1). Both methods are routinely used to produce genetically modified pigs; however, SCNT is technically demanding which can lower the efficiency of obtaining live piglets. After the manipulation procedures, embryos are cultured and subsequently transferred into a surrogate gilt. Gene editing of embryos by microinjection results in pigs with heterogeneous mutations, while SCNT results in pigs with the same mutation as the donor cell (Figure 1).

Although IVP of embryos has proven to be beneficial and has been adopted for species, such as cattle, small ruminants, and horses, the swine industry has been reluctant to adopt these technologies. Similar to other species, *in vitro*-produced porcine embryos exhibit reduced quality and ability to establish a pregnancy after transfer compared to their *in vivo*-derived counterparts (Macháty et al., 1998; Bauer et al., 2010). Thus, a considerable amount of effort has been devoted to improving each step of the IVP process, which is the main focus of this review. Furthermore, developmental and epigenetic aberrations have been noted after *in vitro* culture (IVC) of porcine embryos, possibly resulting in abnormalities during gestation and after birth. Other issues, including polyspermy, embryo transfer procedures, and cryopreservation, are discussed as barriers to widespread use of IVP porcine embryos.

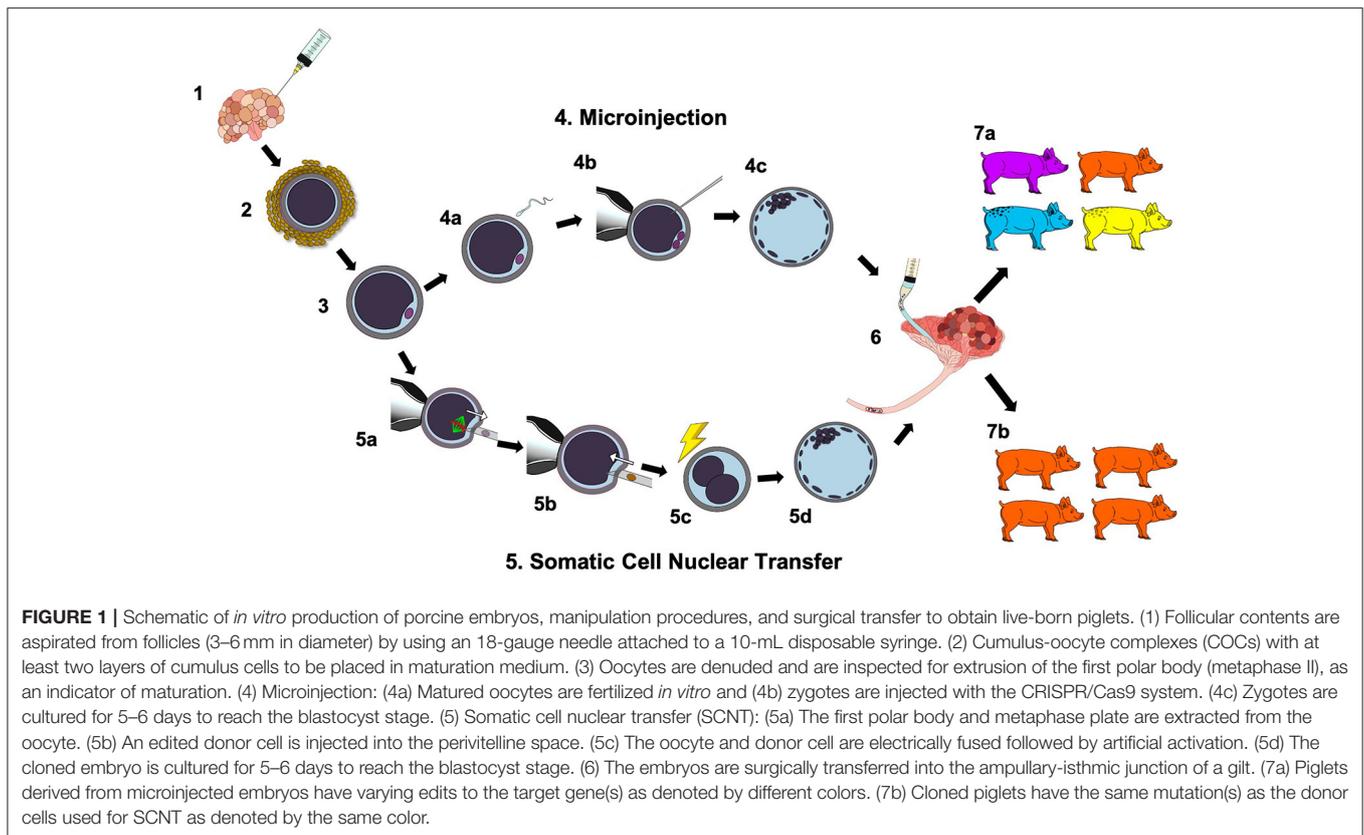
ADVANCES IN *IN VITRO* MATURATION

Oocyte maturation is a complex process that involves the coordination of events that together allow the oocyte to resume meiosis, undergo successful fertilization, and support normal embryo development and subsequent development to term (Gilchrist and Thompson, 2007). Detailed underlying mechanisms and signaling events that promote oocyte development *in vitro* are not fully understood. *In vivo*, the immature oocyte and surrounding cumulus cells (CCs) maintain an intimate relationship which allow the oocyte to gradually acquire meiotic and developmental competence (Gilchrist et al., 2008). The bi-directional communication between the oocyte and its surrounding CCs is critical in activating the necessary signaling pathways needed for competence to be attained. Starting the *in vitro* embryo production (IVP) process with the most competent oocytes is critical as the oocyte's intrinsic quality sets the stage for proper embryo development (Loneragan et al., 2003).

Oocytes remain under meiotic arrest because of high levels of oocyte secreted cyclic adenosine monophosphate (cAMP) (Mehlmann et al., 2004). The act of removing a cumulus oocyte complex (COC) from its follicular environment and placing it into culture will lead to spontaneous resumption of meiosis. This is likely due to the physical disruption of gap junctions, which leads to a reduced influx of cyclic guanosine

monophosphate (cGMP) from the CCs. Cyclic GMP plays an important role as an inhibitor of cAMP hydrolysis (Norris et al., 2009; Vaccari et al., 2009), and the reduction of cGMP in oocytes prematurely removed from follicles may cause oocytes to resume meiosis and complete nuclear maturation. However, the premature progression of meiosis often results in defective cytoplasmic maturation, therefore compromising development (Sela-Abramovich et al., 2006). Phosphodiesterases are a group of enzymes that hydrolyze cyclic nucleotides which can lead to a decrease in cGMP and cAMP levels and results in the resumption of meiosis. Treatment of COCs removed from follicles with phosphodiesterase inhibitors, such as milrinone, has shown promising results in improving oocyte nuclear and cytoplasmic competence (Roy et al., 2021). Furthermore, cAMP negatively regulates maturation promoting factor (MPF), and decreases in cAMP levels with concomitant increases in MPF is critical for successful meiosis (Yu et al., 2008). Upon fertilization, intracellular calcium oscillations in the oocyte cytoplasm promote degradation of MPF and allow meiosis to be completed (Madgwick et al., 2004).

Another issue often encountered during *in vitro* maturation (IVM) is improper activation of the mitogen activated protein kinase-3 and -1 (MAPK3/1) signaling pathway, which is a direct result of a decreased response to luteinizing hormone (LH). Oocytes derived from small or medium sized antral follicles often respond poorly to LH as the CCs surrounding oocytes from immature follicles do not possess adequate LH receptors (Eppig et al., 1997). *In vivo*, acquisition of an appropriate numbers of LH receptors in CCs followed by the LH surge activates a signaling cascade that promotes oocyte maturation, which incorporates actions of epidermal growth factor (EGF)-like factors and the downstream MAPK3/1 pathway (Shimada et al., 2003, 2006; Hsieh et al., 2007). Activation of MAPK3/1 plays an integral role in successful oocyte maturation and is needed for optimal cumulus cell expansion (Su et al., 2003; Fan et al., 2009; Yuan et al., 2017). To improve the response of COCs to LH *in vitro*, follicle stimulating hormone (FSH) is added to the medium to increase in LH receptor numbers, and EGF is supplemented to promote CC expansion and maturation (Procházka et al., 2003; Kawashima et al., 2008). Recently, addition of three cytokines, fibroblast growth factor 2 (FGF2), leukemia inhibitory factor (LIF), and insulin like growth factor 1 (IGF1) together (termed "FLI") during maturation improved oocyte competency by influencing the induction of MAPK3/1 activation in the cumulus cells (Yuan et al., 2017). Supplementation of FLI into porcine oocyte maturation medium dramatically increased the number of porcine oocytes that completed nuclear maturation, doubled the number of embryos that reached the blastocyst stage, and quadrupled the number of piglets born compared to the traditional form of oocyte maturation (Yuan et al., 2017). The FLI matured COCs showed a dramatic increase in cumulus cell expansion, decreased the number of transzonal projections as time in culture increased, and demonstrated a difference in the timing of MAPK3/1 activation in cumulus cells. Effectiveness of this novel maturation system has been validated for pigs (Procházka et al., 2021) and has been incorporated into



maturation systems for cattle and sheep (Stoecklein et al., 2021; Tian et al., 2021).

A recent study also suggests that FLI provides necessary signals to promote oocyte maturation without the presence of gonadotropins, specifically LH and FSH. Porcine oocytes placed into a maturation medium supplemented with FLI and without LH and FSH were able to complete nuclear maturation at an equivalent frequency but with only negligible CC expansion (Redel et al., 2021). The findings go against a conventional dogma that CC expansion is necessary or a marker for successful oocyte maturation. Results in the study suggest that FLI can drive oocyte competence in the absence of gonadotropins and has a downstream role in enhancing CC expansion when gonadotropins are present (Redel et al., 2021). More research is needed to completely understand the mechanisms by which FLI is promoting oocyte maturation and if other players can be supplemented to continue to improve our IVM system.

Outcomes of IVM are highly influenced by the oocyte source, specifically depending on whether the oocytes are derived from either prepubertal gilts or sexually mature sows. Bagg et al. (2007) observed that ovaries from gilts have increased numbers of follicles that are 3 mm in diameter, whereas ovaries from sows have more follicles ranging from 4 to 8 mm. Oocytes derived from small follicles (3 mm) gave rise to blastocyst-stage embryos with decreased cell numbers compared to larger follicles of either source, indicating lower developmental quality (Bagg et al., 2007). Although both oocyte sources have been used to

generate embryos that result in live piglets, oocytes from gilts have also been shown to be less responsive to FSH during maturation, in turn decreasing progression to MII, and exhibit decreased development to the blastocyst stage compared to sow-derived oocytes (Marchal et al., 2001). However, improvements in porcine maturation systems, such as the addition of FLI, have dramatically increased oocyte quality and subsequent developmental competence of gilt-derived oocytes (Yuan et al., 2017). Understanding these key pathways will assist us in designing an optimal IVM system that enhances availability of *in vitro*-matured porcine oocytes.

HISTORY OF PORCINE EMBRYO CULTURE

The process of embryo transfer in pigs necessitates embryo culture, if even for a brief period. The ability to maintain viable gametes or embryos *in vitro* is one of the enabling technologies that facilitates genetic manipulation of pigs and is central to advancements in our understanding of reproductive biology. Prior to the 1990s, zygotes, 2-cell stage embryos, compact morula, and blastocyst-stage embryos could be recovered from a donor sow and transferred into a surrogate to produce offspring. Culture from the zygote to the blastocyst stage followed by production of offspring was a rare occurrence (Davis and Day, 1978). While there were a few reports of development through the 4-cell stage (e.g., Menino and Wright, 1982), there was a so-called *in vitro* “block” to development that occurred during

the 4-cell stage. A similar “block” to development was observed in most mammalian species that corresponded to the stage at which significant amounts of RNA synthesis first began after fertilization. Embryos could be cultured before and after this critical stage while retaining viability but could not be cultured through it and maintain viability. Researchers working on mice screened every available strain and discovered that the C57 BL/6 strain could develop *in vitro* through this stage (block occurs at the two-cell stage). Likely a result of a unique tolerance to hyperosmolarity (Hadi et al., 2005), the C57 BL/6 became the standard for embryo culture and manipulation experiments. For those of us who worked on other mammals, such as pigs, achieving development from the zygote to blastocyst stage required approaches whereby zygotes were placed in a co-culture environment (White et al., 1989), medium supplemented with oviductal fluid (Archibong et al., 1989), embedded in agar cylinders and cultured in a sheep oviduct (Prather et al., 1991), or *in vitro* in an organ culture system (Krisher et al., 1989).

Gradually, different formulations were described whereby embryos could be routinely cultured from the zygote to blastocyst stage, such as NCSU23 (Reed et al., 1992), Whitten’s Medium (Beckmann and Day, 1993), porcine zygote medium (PZM) (Yoshioka et al., 2002). One such culture medium was modified Tyrode’s Lactate without bicarbonate and buffered with HEPES (TL-HEPES) (Hagen et al., 1991). The discovery that TL-HEPES would support development from the zygote stage to the blastocyst stage was stumbled upon quite serendipitously. One Friday afternoon, zygotes were collected from a gilt that was supposed to provide 4-cell stage embryos. The zygotes were flushed from the oviduct by using TL-HEPES. Since the wrong stage of embryos was collected and there was no use for them, the zygotes were left in the flushing medium and placed in a humidified warm air incubator. The next week when old culture dishes were being removed from the incubator, it was noticed that the zygotes had become blastocyst stage embryos (RSP, personal observation). About that time, there was much debate regarding which components of the culture system were either inadequate or toxic to the developing embryo (reviewed by Petters and Wells, 1993). So, it was shown with TL-HEPES as the base medium that, contrary to some reports, neither glucose nor glutamine inhibited development (Hagen et al., 1991). Since TL-HEPES does not require a CO₂ atmosphere for buffering the pH, it has been widely used as a holding or embryo transfer medium for porcine embryos over the past 30 years (Redel et al., 2019). It was not until the 1990s that the first reports of embryo culture in pigs, whereby zygotes were rinsed from one sow’s oviduct, identified microscopically, cultured to the blastocyst stage, and then transferred to the oviduct of another sow with resulting pregnancies, were published (Beckmann and Day, 1993).

While the “block” to embryo development had been a major topic of scientific papers for over a half century, the “block” seems to have disappeared (Prather, 2010). In retrospect, the fine tuning of the culture system by altering the osmolarity (Baltz and Tartia, 2010) and using highly purified water may have caused the “block” to “disappear.” The “block” that was once a major hurdle to advancements in our understanding of early embryonic events. While culture systems are certainly not ideal, they have evolved

to the state of functionality that permit widespread repeatability and application of embryo related technologies.

Incorporation of powerful technologies such as RNA-sequencing now allow us to understand embryo requirements *in vitro* and improve standard medium formulations. For instance, the transcriptional profiles of *in vivo*-matured and -fertilized, *in vitro*-cultured blastocyst stage porcine embryos predominantly exhibited an upregulation of transcripts related to amino acid transport and metabolism pathways compared with their *in vivo*-derived counterparts (Bauer et al., 2010). Transcript abundance of the arginine transporter, solute carrier 7A1 (*SLC7A1*), was increased *in vitro*-cultured embryos, indicating a potential deficiency of arginine, and supplementation of arginine to embryo culture increased developmental parameters and resulted in the birth of live piglets (Redel et al., 2016b). Similarly, RNA-sequencing data led to supplementing glutamine to current culture system, and improved embryo development *in vitro* and led to successful term development after transfer (Chen et al., 2018). The same notion was used to supplement glycine in culture, and benefits of the supplementation was observed *in vitro*. (Redel et al., 2016a). However, 11 surrogates failed to become pregnant after receiving embryos cultured in supplemental glycine, indicating that *in vitro* measures of quality do not necessarily translate into *in vivo* developmental competence. These studies demonstrate that the adaptation of new technologies enable us to interpret the status of porcine embryos under culture conditions and assist us to continuously advance the culture systems.

PREGNANCY RATE WITH *IN VITRO*-PRODUCED EMBRYOS

In vitro-produced embryos are known to possess reduced quality and viability compared to their *in vivo*-derived counterparts. Before a reliable embryo culture system was developed, embryos were transferred at the one-cell stage into a gilt on days 0 or 1 after standing estrus (Lee et al., 2014; Whitworth et al., 2014). Improvements in the culture system have allowed for transfer of IVP blastocyst-stage embryos into a surrogate pig without compromising viability (Lee et al., 2013; Redel et al., 2016b; Chen et al., 2018). Morula- and blastocyst-stage embryos are typically transferred into a surrogate gilt on days 3, 4, or 5 after standing estrus. This asynchronized transfer strategy accounts for the developmental delay of IVP embryos (Bauer et al., 2010). Most term pregnancies are allowed to farrow; however, Cesarean section (C-section) can also be successfully performed. If a pregnancy is lost during gestation, the majority of loss occurs between days 25 to 45 by resorption, and in rare cases, the surrogate will abort the pregnancy after day 45 (Lai and Prather, 2003).

Since 2017, our group has performed four transfers of unmanipulated IVF embryos cultured *in vitro* that went to term with an 100% pregnancy rate and average of 10.3 ± 0.3 live piglets per litter. Based on the fact that 40–60 blastocyst-stage embryos were transferred in each case, ~1 in 5 embryos developed to term, indicating a significant improvement in the

culture system. However, reports of transfers by using *in vivo*-derived embryos indicate that ~1 in 3 blastocyst-stage embryos develop to term (Yoshioka et al., 2002; Martinez et al., 2015). Considering these rates, improvements in the developmental potential of IVP embryos can still be made to reduce the number of embryos that need to be transferred.

As discussed above, IVP embryos are essential for the application of ART. However, application of embryo manipulation or SCNT often compromises embryo viability, resulting in a low term development. The full-term pregnancy rate after transferring IVF embryos injected with CRISPR/Cas9 systems (microinjection) ranges from 40 to 47% (187 transfers since 2017) with an average of 5.7 ± 0.7 live piglets per litter. The full-term pregnancy rate after transferring SCNT-derived embryos generated from numerous cell lines was 20–30% (264 transfers since 2017) with an average of 3.2 ± 0.4 live piglets per litter. The full-term development from these embryos is typically lower compared to unmanipulated IVF embryos due to accumulated damage during embryo manipulation or gene editing itself. In addition, efficacy of SCNT also highly depends upon the type of donor cells used for the process (Hao et al., 2009; Richter et al., 2012; Whitworth et al., 2014). Examples between *in vivo*-, IVF-, and SCNT-derived embryo transfers and resulting litter sizes of live-born piglets can be found in **Table 1**, and the success rates of *in vivo*-derived pregnancies vs. pregnancies from different *in vitro* manipulations are depicted in **Figure 2**.

Previous embryo transfer reports certainly indicate that the current IVP system in pigs (both IVM and IVC) has been improved; blastocyst transfers can be practically applied, and the number of live piglets is closer to transferring *in vivo*-derived embryos. However, embryos and piglets originating from the IVP system can present abnormalities that are additional barriers to consider.

ABNORMALITIES OBSERVED IN *IN VITRO*-PRODUCED PREIMPLANTATION EMBRYOS

Development of the embryo culture system now successfully supports the production of porcine embryos *in vitro*, and the embryos are essential for studying early developmental processes as well as for the application of genetic engineering in pigs. However, the quality of embryos (e.g., cell number, apoptosis, epigenetic status) originating *in vitro* remains inferior to *in vivo*-derived embryos. Currently, development of IVF embryos to the blastocyst stage is ~40% on day 6 (Redel et al., 2016a; Yuan et al., 2017; Chen et al., 2020), with significant embryo loss or arrest at earlier stages. Blastocyst-stage embryos produced *in vitro* contain about 40 to 50 cells on day 6 (Redel et al., 2016a; Yuan et al., 2017; Chen et al., 2018, 2020), which is an improvement compared to the previous IVP systems (Macháty et al., 1998; Bauer et al., 2010). However, a significantly lower number of cells are detected in IVP day 7.5 blastocyst-stage embryos compared to *in vivo* counterparts, 50 vs. 80, respectively (Canovas et al., 2017), presumably contributing to the low term development. In

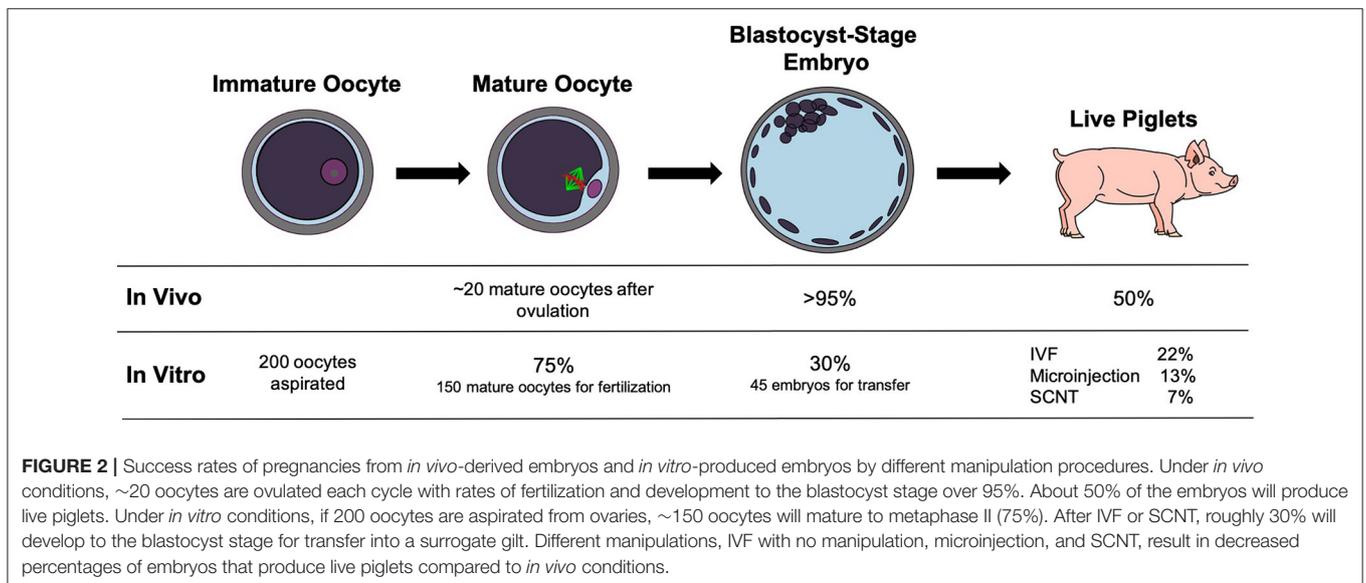
addition, the ratio of inner cell mass (ICM) to trophectoderm (TE) cell numbers, a predictive marker of embryo quality (Tao et al., 1995), is often skewed by IVP. The ICM:TE ratio of day 6 blastocyst-stage embryos cultured under the previous IVP system (~0.15–0.2) was lower than *in vivo* counterparts (~0.4–0.5) (Macháty et al., 1998; Yoshioka et al., 2002). Optimization of the culture system led to an improvement of the ICM:TE ratio of IVP embryos (~0.35–0.4) (Redel et al., 2016a; Jeong et al., 2017), which is close to the level of *in vivo*-derived blastocyst-stage embryos.

Regulation of DNA methylation is a major epigenetic event that occurs during preimplantation development in mammals and is critical for normal embryo development (Reik et al., 2001). Fertilized oocytes undergo a massive decrease in global DNA methylation during the transition from gametes to the blastocyst stage, except for imprinted genes and some repetitive elements (Messerschmidt et al., 2014). Artificial intrusions on natural embryo development, including *in vitro* maturation of oocytes, *in vitro* embryo culture, and varying oxygen tensions, are expected to influence to the epigenome of IVP embryos (El Hajj and Haaf, 2013; Gaspar et al., 2015; Sirard, 2017). Porcine IVP embryos are reported to possess a high level of DNA methylation compared to *in vivo*-derived counterparts (Deshmukh et al., 2011). The global level of 5-methylcytosine, measured through immunocytochemistry (ICC), was higher in IVF zygotes than *in vivo*-derived zygotes, and the higher DNA methylation level was maintained in cleavage and blastocyst-stage embryos, suggesting that IVP embryos may experience impaired post-fertilization demethylation. A recent study conducted an analysis of the whole genome DNA methylation status of *in vivo*-derived and IVP pig blastocyst-stage embryos at single nucleotide resolution (Canovas et al., 2017). Consistent with the results of the ICC assay, whole genome DNA methylation analyses revealed that global methylation levels, including CpG islands, promoters, intergenic regions, and repetitive elements, are higher in IVP blastocyst-stage embryos (~15%) than *in vivo*-derived blastocyst-stage embryos (~12%). *In vitro* culture also affects the expression and methylation of imprinted genes, which is critical for normal embryo development (Reik et al., 2001). Transcript abundance of imprinted genes, imprinting related genes, and X-linked genes differ between IVP and *in vivo*-derived blastocyst-stage embryos (Park et al., 2011, 2012; Canovas et al., 2017). Although the methylation status of differentially methylated regions (DMRs) is expected to be stable during embryogenesis, IVP pig blastocyst-stage embryos exhibit aberrant methylation patterns in several imprinted genes, such as *ZAC1*, *PEG10*, and *NNAT*, compared to their *in vivo* counterparts (Canovas et al., 2017). Interestingly, supplementation of oviductal or uterine fluid to fertilization and culture media induces shifts in the DNA methylation profiles of IVF embryos to become more similar to the epigenome of *in vivo*-derived embryos, indicating suboptimal culture conditions contributes to the epigenetic aberrations in porcine embryos produced *in vitro* (Canovas et al., 2017).

Incomplete epigenetic reprogramming of donor cell nuclei has been considered one of the main causes for developmental failure in cloned animals (Rideout et al., 2001; Mann and Bartolomei, 2002; Bonk et al., 2007). An early study on the DNA methylation

TABLE 1 | Live-born litter sizes after surgical transfer of embryos from different sources.

Embryo source	Genetically edited	Number embryos transferred (stage)	Number live piglets (average \pm SD)	Reference
<i>In vivo</i>	No	30 (Blastocyst)	9.4 \pm 0.8	Martinez et al., 2014
	No	30 (Blastocyst)	9.2 \pm 2.5	Martinez et al., 2015
	Yes (<i>NANOS2</i>)	30 (One-cell)	3.5 \pm 0.7	Park et al., 2017
IVF	No	40 (Blastocyst)	5.0 \pm 1.1	Redel et al., 2016a
	Yes (<i>NANOS2</i>)	52 (One-cell)	11 (One litter)	Park et al., 2017
	Yes (<i>CD163</i> , <i>CD1D</i> , <i>TMPRSS2</i> , <i>COL6A3</i> , <i>APC</i> , and <i>PAH</i>)	50 (Blastocyst)	8.6 \pm 1.1	Yuan et al., 2017
SCNT	No	40 (Blastocyst)	10.5 \pm 0.7	Chen et al., 2018
	Yes (<i>CMAH</i>)	192–257 (One-cell)	3.0 \pm 3.0	Kwon et al., 2013
	No	35–58 (Blastocyst)	3.2 \pm 2.7	Lee et al., 2013
	Yes (<i>RAG2</i>)	180–252 (One-cell)	2.0 \pm 1.1	Lee et al., 2014
		48 (Blastocyst)	6 (One litter)	
	Yes (<i>CD1D</i>)	201–239 (One-cell)	4.3 \pm 2.5	Whitworth et al., 2014
	Yes (<i>CD163</i>)	193–267 (One-cell)	4.6 \pm 3.1	Whitworth et al., 2014
	No	207–280 (One-cell)	2 (One litter)	Lee et al., 2015
	Yes (<i>NGN3</i>)	102 (One-cell)	3 (One litter)	Sheets et al., 2018
No	42 (Blastocyst)	2 (One litter)	Cecil et al., 2020	



status of cloned pig embryos found that genome-wide post-fertilization demethylation also occurs during preimplantation development (Kang et al., 2001). However, like IVF embryos, cloned embryos retain relatively higher DNA methylation levels compared to *in vivo*-derived embryos during preimplantation development (Kang et al., 2001; Bonk et al., 2008; Deshmukh et al., 2011). Suboptimal IVC could be a source for the incomplete epigenetic reprogramming of cloned embryos as shown in IVF embryos; however, the abnormal epigenetic status is likely related to the SCNT process because cloned embryos possess a higher global DNA methylation level compared to IVF embryos

(Kwon et al., 2008; Huan et al., 2014) and impairments in maintaining methylation imprints (Wei et al., 2011). Indeed, somatic driven DNA methyltransferase 1 (*DNMT1*) is known to interfere with DNA methylation reprogramming in cloned pig embryos (Song et al., 2017). Reducing *DNMT1* abundance by siRNA or epigenetic modification agents improves DNA methylation reprogramming and developmental competency of cloned pig embryos (Xu et al., 2013; Huan et al., 2014, 2015a,b; Song et al., 2017), supporting the fact that a high level of *DNMT1* is a barrier to the reprogramming of DNA methylation marks in cloned embryos. Furthermore, treatment of cloned

porcine embryos with histone deacetylase inhibitors promotes the DNA demethylation process by reducing *DNMT1* expression levels (Liang et al., 2015), leading to improved developmental competency (Zhang et al., 2007; Whitworth et al., 2011; Zhao et al., 2013).

Histone modifications establish molecular landmarks that differentiate active and inactive chromatin states (Rideout et al., 2001). Moreover, histone modifications are important factors that determine successful epigenetic reprogramming of cloned embryos as an open chromatin state must be achieved in the somatic donor cell to mimic the early embryo (Whitworth and Prather, 2010; Ogura et al., 2013). Acetylation and methylation of lysine residues on the amino-terminal tails of histone H3, such as H3K9ac, H3K27ac, and H3K4me3, are markers of an open chromatin state, correlating with the developmental potential of cloned embryos (Santos et al., 2003; Li et al., 2008). Cloned porcine embryos possess lower H3K9ac and H3K4me3 levels than normal fertilized embryos at early developmental stages, which represents a relatively inactive chromatin state (Zhai et al., 2018a,b). Conversely, the repressive histone marker, H3K9me3, is more abundant at early stages in cloned porcine embryos than those derived by IVF (Zhai et al., 2018a; Jeong et al., 2020). Patterns of histone modifications representing repressive chromatin structures serve as a barrier to successful epigenetic reprogramming of donor cells and correspond to a lower developmental potential. Indeed, induction of an open chromatin state in cloned porcine embryos by treatment with histone deacetylase inhibitors or histone-lysine methyltransferases of H3K9 leads to improvement of nuclear reprogramming and the developmental potential (Zhang et al., 2007; Whitworth et al., 2011; Song et al., 2014; Huang et al., 2016; Jeong et al., 2020; Weng et al., 2020).

ABNORMALITIES OBSERVED IN NEONATAL PIGLETS DERIVED FROM IVF OR SCNT

Different abnormalities have been detected in piglets derived by the IVP system and manipulation procedures, although predominantly the reports are related to cloned piglets. High rates of stillbirths have been noted for pregnancies carrying clones (Estrada et al., 2007; Schmidt et al., 2015; Ao et al., 2017; Cecil et al., 2020), and increased mortality within the first few days after farrowing is often observed with cloned pigs (Ao et al., 2017). Birth weight may be a predictor of survival probability for cloned piglets as those that died within 4 days after birth weighed ~28% less than those that survived for more than 4 days (Ao et al., 2017). As a contributing factor to perinatal mortality, abnormal umbilical cord vasculature and low placental weight as well as placental malformations have been observed for cloned fetuses and piglets, likely impairing nutrient transport and gas exchange (Lee et al., 2007; Ao et al., 2017). Regarding placentas of cloned neonatal piglets, presence of infarcts and villous hypoplasia may be the result of upregulation of apoptotic signaling pathways, which were detected by proteomic analyses (Lee et al., 2007). To overcome issues with stillbirths,

lack of mammary development in surrogates carrying clones, and delayed initiation of parturition, a C-section is typically performed on day 116 of gestation (Carter et al., 2002; Whyte et al., 2011). Delivery by C-section has been shown to decrease post-natal mortality over vaginal delivery by 20% (Schmidt et al., 2011).

Anatomical abnormalities can also be present in cloned piglets and are occasionally seen in piglets derived by IVF. Examples of abnormalities that can be observed by external examination include macroglossia, cleft palate, macrocephaly, flexor tendon contracture, polydactyly, kyphosis, and umbilical hernia, among others (Carter et al., 2002; Schmidt et al., 2015). Some of the abnormalities, such as flexor tendon contractures, can be corrected with physical therapy (Carter et al., 2002), but most abnormalities often lead to loss of the affected piglet. Internal abnormalities of the digestive tract include short or absent intestines and small or absent gall bladder (Schmidt et al., 2015). Cloned piglets can also have various heart and circulatory system defects that may lead to congestive heart failure (Carter et al., 2002; Schmidt et al., 2015). Lastly, issues of the reproductive system have been observed in cloned piglets, such as absent gonads, cryptorchidism, and enlarged gubernaculum (Schmidt et al., 2015).

Analysis of muscle tissues from normal and abnormal cloned piglets revealed numerous differentially expressed genes, such as those for the MAPK signaling pathway, the hypertrophic cardiomyopathy pathway, and imprinting, as well as altered DNA methylation patterns that could be involved in abnormality manifestation (Li et al., 2014). Although developmental abnormalities observed in SCNT-derived piglets are potentially linked to incomplete or faulty reprogramming of the donor cell nucleus after SCNT, the overall IVP system is suggested to contribute to the abnormalities as well.

IMPROVEMENTS REQUIRED TO EXPAND THE USE OF *IN VITRO*-PRODUCED PORCINE EMBRYOS

Application of ART, such as IVF, embryo transfer, and SCNT, facilitates rapid genetic improvements and are heavily used parts of the livestock industry. Regarding the cattle industry, the number of transfers by using IVP cattle embryos increased 7.3% from 2018 to 2019, pointing to a heavy reliance on these technologies (Viana, 2020). However, application of ART is limited in the swine industry, partially due to difficulties in embryo production. *In vitro* production of porcine embryos has not been adopted by this industry for several reasons. Since pigs have litters and a short generation interval, there is less incentive to incorporate ART to facilitate genetic improvements as compared to cattle who have single offspring and longer generation intervals. In pigs, establishment of pregnancy is only possible when at least two embryos are present in each uterine horn (Dziuk, 1968). Thus, a large number of IVP embryos must be transferred into a surrogate for a pregnancy to be established. Conventionally, a minimum of 150 one-cell stage embryos or 30 blastocyst-stage embryos are transferred into a single surrogate

pig. Other barriers hindering incorporation of IVP embryos in the swine industry include high rates of polyspermy, lack of non-surgical oocyte recovery or embryo transfer procedures, and unsuccessful cryopreservation.

Reducing Polyspermy

In vitro fertilization of porcine oocytes matured *in vitro* results in high rates of polyspermy that decreases development to the blastocyst stage and the number of cells in the ICM (Han et al., 1999b). The penetration of more than one spermatozoon often leads to developmental arrest of IVF embryos, thus impeding the use of IVF for pig production purposes. In pigs, the block to polyspermy is a two-step mechanism; the fast block occurs at the oocyte plasma membrane and the slow block occurs at the zona pellucida after cortical granule exocytosis. Specifically, the fast block is associated with shedding of folate receptor 4 (JUNO) from the oocyte plasma membrane which acts as the receptor for Izumo sperm-egg fusion 1 (IZUMO1) on the sperm head (Bianchi et al., 2014). The slow block is triggered by sperm-induced release of ions (i.e., calcium and zinc) followed by cortical granule exocytosis (Abbott and Ducibella, 2001; Tokuhira and Dean, 2018). The released cortical granules induce cleavage of zona protein 2 (ZP2), leading to increased zona hardening (Boccaccio et al., 2012; Burkart et al., 2012). Moreover, zinc released after sperm penetration can bind ZP1, which contributes to further crosslinking of the zona pellucida (Nishimura et al., 2019). Zona hardening has also been shown to be influenced by lectins and glycosidases in the mouse (Dolci et al., 1991), and treatment of porcine oocytes with different lectins decreased sperm binding to the zona pellucida (Hwang et al., 2002). Factors responsible for proper zona hardening are present within the oviductal fluid of pigs, partially explaining why this process is diminished during IVF (Mondéjar et al., 2013). Although polyspermy generally results in developmental arrest during preimplantation stages, accessory sperm heads have been detected within phagolysosomes of porcine embryos, suggesting that embryos possess a mechanism to degrade unnecessary spermatozoa for survival of the embryo (Xia et al., 2001). In fact, polyspermic porcine embryos could develop to term after transfer, and the piglets had the correct number of chromosomes (Han et al., 1999a).

Oocyte quality is known to be correlated with incidence of polyspermy and *in vitro*-matured oocytes present lower quality and higher rates of polyspermy. Components of porcine oviductal fluid, such as oviductal glycoprotein 1, osteopontin, and plasminogen, have been shown to reduce the number of penetrating sperm (Hao et al., 2006; Coy et al., 2008, 2012). A recently developed porcine IVF system increased the pH from 7.4 to 8.0, which is physiological for the oviductal ampulla, added oviductal fluid, and installed a barrier between gametes to promote sperm chemotaxis (Soriano-Úbeda et al., 2017). The system decreased sperm penetration by 21.8%; however, monospermy of the fertilized oocytes increased from 26.4 to 88.7% in the new system. Similarly, a porcine IVM system with the presence of C-X-C motif chemokine ligand 12 (CXCL12), vascular endothelial growth factor A (VEGFA), and Wingless-type MMTV integration site family member 5A

(WNT5A), increased nuclear maturation, development to the blastocyst stage, and retraction of transzonal projections with decreased incidence of polyspermy (Liu et al., 2020). Retraction of transzonal projections can promote cortical granule migration and exocytosis (Galeati et al., 1991).

As previously mentioned, addition of FLI to the porcine oocyte maturation system improved developmental competence of subsequent embryos. Moreover, oocytes matured in the presence of FLI presented decreased transzonal projections during IVM (Yuan et al., 2017). It is not clear whether the improved efficiency of IVM by using FLI is related to its ability to reduce polyspermy by decreasing transzonal projections as this has yet to be evaluated.

Non-surgical Embryo Transfer

The ability to conveniently transfer IVP embryos into surrogates is warranted for broad incorporation of IVF and SCNT technologies into the swine industry. However, unlike other livestock species, such as cows, routine non-surgical transfers in pigs have been limited due to the complexity of the female reproductive tract. Since the spiral shape of the cervix complements the male anatomy, passing a catheter through the cervix to reach the uterus is challenging as well as the fact that the uterine horns are long and narrow compared to other species. For transfer of IVP embryos in the research setting, a midventral laparotomy is performed to expose one ovary and oviduct, and the embryos are deposited in the ampullary-isthmic junction. Successful pregnancies have been established when morula and blastocyst-stage embryos are surgically transferred into a surrogate gilt (Chen et al., 2018, 2021; Koppes et al., 2020; Pfeiffer et al., 2020); however, this procedure is time-consuming and requires specialized skills, thus limiting the use of IVP embryos in the swine industry.

Non-surgical embryo transfers have been accomplished in pigs (averaging 2.5 min per procedure) by using an artificial insemination spirette to guide a modified flexible catheter for deposition of the embryos in the uterus (Li et al., 1996; Martinez et al., 2004). After 24 transfers of *in vivo*-derived embryos, 17 surrogates became pregnant (71%) and farrowed an average of 6.9 ± 0.7 piglets. The type of female, gilt or sow, did not impact success of the transfer, but more insertion force was needed to pass through the cervical canal of the gilts compared to the sows, as expected (Martinez et al., 2004). One major risk factor associated with this method is the possibility of puncturing the uterine wall with the catheter which could lead to infection. Furthermore, the number of embryos transferred is important as non-surgical transfer of 30 *in vivo*-derived blastocyst-stage embryos resulted in a farrowing rate of 38.9% (5.7 ± 2.4 piglets per litter) while transfer of 40 blastocyst-stage embryos resulted in a farrowing rate of 72.7% (9.9 ± 2.1 piglets per litter) (Martinez et al., 2015). Although the reports certainly encourage the use of non-surgical embryo transfer in pigs, only *in vivo*-derived embryos have been used in these studies. As pointed out previously, developmental potential of *in vivo*-derived embryos far surpasses IVP embryos, and the need for non-surgical transfers is primarily related to IVP embryos. Reports on the effectiveness of non-surgical embryo transfer using IVP embryos

should clarify whether the technology can be effectively used in the industry.

Cryopreservation

While cryopreservation of cells from pigs, such as fibroblast cells, is straightforward, cryopreservation of porcine gametes and embryos has been a challenge. Successful cryopreservation of boar sperm is season-, breed-, boar- and ejaculate-specific. Because of this variability, most artificial insemination in the swine industry is performed with fresh semen. For those ejaculates where sperm survive the cryopreservation, frozen semen results in a 20–30% decrease in farrowing rate with reduced litter sizes as compared to fresh semen (Silva et al., 2015).

Successful cryopreservation of embryos is even more challenging. The most consistent success is achieved after centrifugation of the early embryo and micromanipulation removal of the lipids prior to culture to the blastocyst stage and subsequent cryopreservation (Nagashima et al., 1995). Counterintuitively, removal of the lipids enhances development of the embryo rather than hinders it (Li et al., 2006), questioning the role of lipids in porcine embryo development. Transferring 163 delipidized, vitrified blastocyst-stage embryos into two surrogates resulted in a total of 10 piglets between the two pregnancies, confirming that these procedures are compatible with development *in vivo* (Li et al., 2006). Since the lipid removal procedure by micromanipulation is very labor intensive, a similar strategy was developed that separated the lipids through centrifugation but did not remove them from within the zona pellucida as a high-throughput method of cryopreservation for IVP embryos (Spate et al., 2013). An alternative method uses solid surface vitrification, whereby 50 oocytes or zygotes are moved through a series of equilibration and vitrification media and placed onto a cooling surface, such as aluminum foil, sitting on top of liquid nitrogen before being transferred to cryovials (Somfai and Kikuchi, 2021).

While there are a few reports of more conventional cryopreservation, those reports of successful cryopreservation of early pig embryos have not been widely repeatable, and the industry has not adopted the transfer of frozen embryos as a method of improving genetics or moving genetics around the world. *In vivo*-derived embryos survive cryopreservation better than IVP embryos. As discussed previously, this difference in survival may be a result of inherent developmental competence as oocytes derived from sexually mature animals are more developmentally competent than from sexually immature animals. Tajima et al. (2020) reported 35 embryo transfers that used 553 embryos resulting in 14 litters and 59 piglets (59/553 = 10%), and Hirayama et al. (2020) reported 12 embryo transfers with 180 embryos resulting in 8 sows farrowing and 37 piglets

(20%). However, in both cases, it is not clear how many embryos were cryopreserved as only the number transferred is reported. Thus, the percentages may overestimate the efficiency based on the number of embryos frozen. Maturation in the presence of FLI increased the quality of oocytes derived from prepubertal gilts, and similar improvements on the maturation and culture conditions may enhance the survivability of pig embryos after cryopreservation as has been reported for cattle (Stoecklein et al., 2021). Improvements in cryopreservation for porcine gametes and embryos will considerably enhance the applicability of *in vitro* technologies for the swine industry, allowing for rapid transfer of genetics at the national and international levels.

CONCLUSION

Over the years, considerable improvements have been made for IVP of porcine embryos. Pigs produced through these methods have been instrumental for advancements in biomedicine and animal agriculture. However, the success of porcine embryo IVP has not yet reached a level for this technology to be adopted by the swine industry as embryo viability is still decreased compared to *in vivo*-derived embryos. There have been several advances in porcine oocyte maturation and embryo culture, but certain areas require more attention to improve the IVP process as a whole. Lowering rates of polyspermy in IVP embryos, developing reliable methods of non-surgical transfer, and enhancing cryopreservation success of gametes and embryos are among some of the areas where improvements can be made and would be critical for use of IVP by the swine industry. Altogether, improvements in each step of IVP of porcine embryos will lead to the ultimate goal of increasing the pregnancy rate and number of live piglets from each pregnancy.

AUTHOR CONTRIBUTIONS

PC, RP, and KL created an outline for the review. PC, BR, KU, RP, and KL wrote the review. ER prepared the figure. All authors revised the manuscript. All authors contributed to the article and approved the submitted version.

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