

Figure S1. Stirred suspension bioreactor culture of porcine spermatogonia: (A) schematic of a 10 mL stirred suspension bioreactor; (B) maximum shear forces generated in the stirred suspension bioreactors; (C) proliferating spermatogonia index (number of EdU⁺ UCHL1⁺ cells) for 1-week-old porcine spermatogonia. $n = 3$, mean \pm SD. $p > 0.05$ (ns), $*p \leq 0.05$, $**p \leq 0.01$.

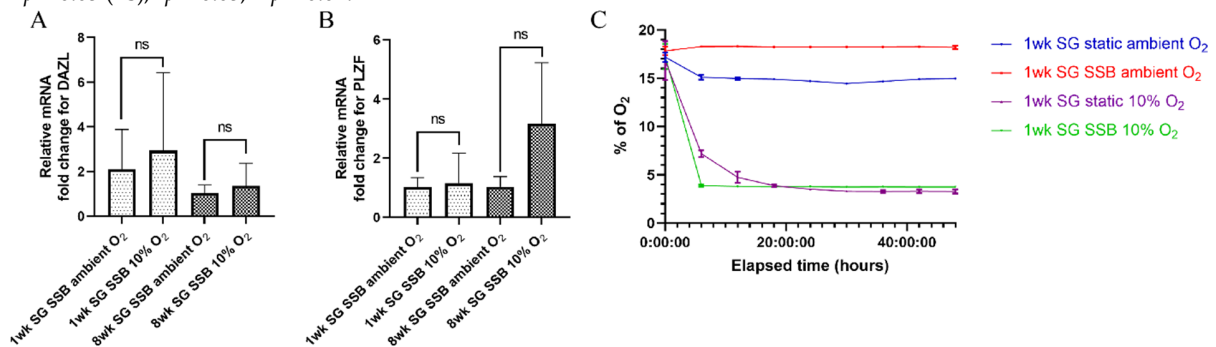


Figure S2. Gene expression of PLZF and DAZL: (A) relative mRNA fold change of DAZL for 1- and 8-week-old spermatogonia (SG); (B) relative mRNA fold change of PLZF for 1- and 8-week-old spermatogonia (SG). $n = 3$, mean \pm SD. $p > 0.05$ (ns), $*p \leq 0.05$, $**p \leq 0.01$, $***p \leq 0.001$, $****p \leq 0.0001$. (C) Percentage of O₂ measured via Oxygen Sensor Spots over the course of 48 h.

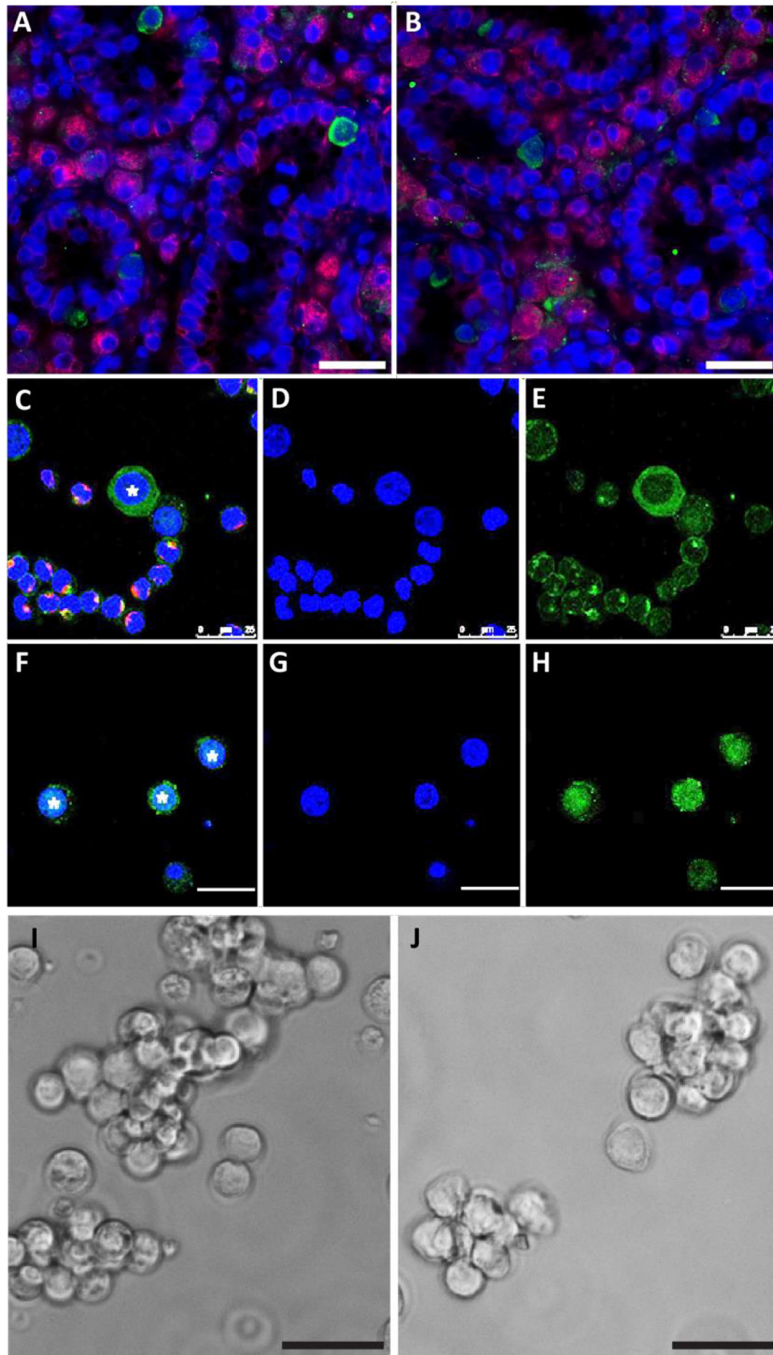


Figure S3. Immunofluorescence image of testis tissue and cells labeled for vimentin, UCHL1, and β -catenin, as well as brightfield images of spermatogonia clusters: (A,B) immunofluorescence image of UCHL1 (green) and vimentin (red) in 1-week-old (A) and 8-week-old testes (B); (C–H) immunofluorescence image of β -catenin (green) and vimentin (red): (C–E) static culture showing cytoplasmic β -catenin and (F–H) bioreactor culture showing nuclear β -catenin; (I,J) Brightfield image of 1-week-old (I) and 8-week-old (J) spermatogonia clusters after bioreactor culture. Scale bar 25 μ m.