



Figure S1. *LncPGCR* regulated the expression of the totipotent gene *Nanog* and PGC marker genes to promote PGCs development (**A**). qRT-PCR of *Nanog*, *Cvh*, and *C-kit* expression after overexpression or knockdown with *LncPGCR* in vitro. * $p < 0.05$, and ** $p < 0.01$ compared with untreated ESCs. Values are means \pm SEM. n=3. (**B**). qRT-PCR of *Nanog*, *Cvh*, and *C-kit* expression after overexpression or knockdown with *LncPGCR* in vivo. * $p < 0.05$, and ** $p < 0.01$ compared with the untreated genital ridge. Values are means \pm SEM. n=3. (**C**). The proportion of PGCs-like cells(*Cvh*⁺) in the genital ridge on E4.5d detected by flow cytometry analysis.

Table S1. *LncPGCR* promoter region different length PCR amplification primers.

| Name | Primer Sequence | Size | Enzyme Sites | Usage |
|------|--------------------------------|------|--------------|---------------|
| P1 | CCCTACGTACCTTGGGTTAGAGTCCTG | 1614 | XhoI | pLncPGCR-EGFP |
| P2 | CGGGGTACCCCTTGAGGGCTTCCTTGC | 1356 | KpnI | pGL3/1356 |
| P3 | CGGGGTACCGGTAGTCCCAGATGCCAGTCC | 1160 | KpnI | pGL3/1160 |
| P4 | CGGGGTACCGGTTCCACTGAACACCCTC | 1033 | KpnI | pGL3/1033 |
| P5 | CGGGGTACCATCCCAGCTTGTCTTGCT | 661 | KpnI | pGL3/661 |
| P6 | CGGGGTACCCCCAACCGAGGAGCGAAGA | 337 | KpnI | pGL3/337 |
| P0 | CCGCTCGAGATCAAAGCGATTCCCTGTA | | SnaBI | pLncPGCR-EGFP |

Table S2. Transcription factor TCF7L2 binding site mutation primer.

| Name | | Primer Sequence |
|--------|-----------|--|
| TCF7L2 | F-Kpn I | TGATCTTAATCAACCATGTTGCAAAGCTTCATACCGTAAGCACA |
| | TCF7L2-R1 | TGTGCTTACGGTATGAAGCTTGAAACATGGTTGATTAAGATCA |
| | TCF7L2-F1 | TGATCTTAATCAACCATGTTGCAAAGCTTCATACCGTAAGCACA |
| | R-Xho I | GCTTACTTAGATCGCAGATCTCGAGATCAAAGCGATTCCCTGTA |

Table S3. The transcript expression levels of PGCs lncRNA related to germ cell differentiation in ESCs, PGCs, and SSCs.

| lncRNA | ESCs_FPKM | PGCs_FPKM | SSCs_FPKM | P values |
|----------------|-----------|-----------|-----------|-------------|
| TCONS_00612668 | 0.304065 | 5.2481 | 2.91182 | 0.0268936 |
| TCONS_00765266 | 35.8417 | 0.0421958 | 0 | 2.90E-06 |
| TCONS_00874170 | 0 | 0.564523 | 0.0374735 | 0.00901045 |
| TCONS_00072199 | 7.20163 | 0.0787197 | 0 | 0.000343457 |
| TCONS_00362261 | 37.9311 | 1.1473 | 6.82561 | 0.0297688 |
| TCONS_00362344 | 22.9197 | 0.187924 | 0 | 0.000404435 |
| TCONS_00612668 | 0.304065 | 5.2481 | 2.91182 | 0.0268936 |
| TCONS_00659989 | 22.9598 | 0.270672 | 1.37532 | 0.00405561 |
| TCONS_00946310 | 1.28279 | 0.0733322 | 0 | 0.0338241 |

| | | | | |
|----------------|----------|-----------|----------|------------|
| TCONS_00580256 | 28.7449 | 0.379114 | 0 | 0.00588496 |
| TCONS_00612668 | 0.304065 | 5.2481 | 2.91182 | 0.0268936 |
| TCONS_00947438 | 5.34547 | 0.610657 | 0.114237 | 3.90E-06 |
| TCONS_00978804 | 0 | 0.105411 | 2.85304 | 0.0148691 |
| TCONS_00948124 | 0 | 0.524497 | 1.89062 | 0.00672651 |
| TCONS_00627195 | 32.6283 | 0.0533638 | 0 | 1.72E-08 |