

## Supplementary Material

### **Theophylline derivatives promote primordial follicle activation via cAMP-PI3K/Akt pathway and ameliorate fertility deficits in naturally aged mice**

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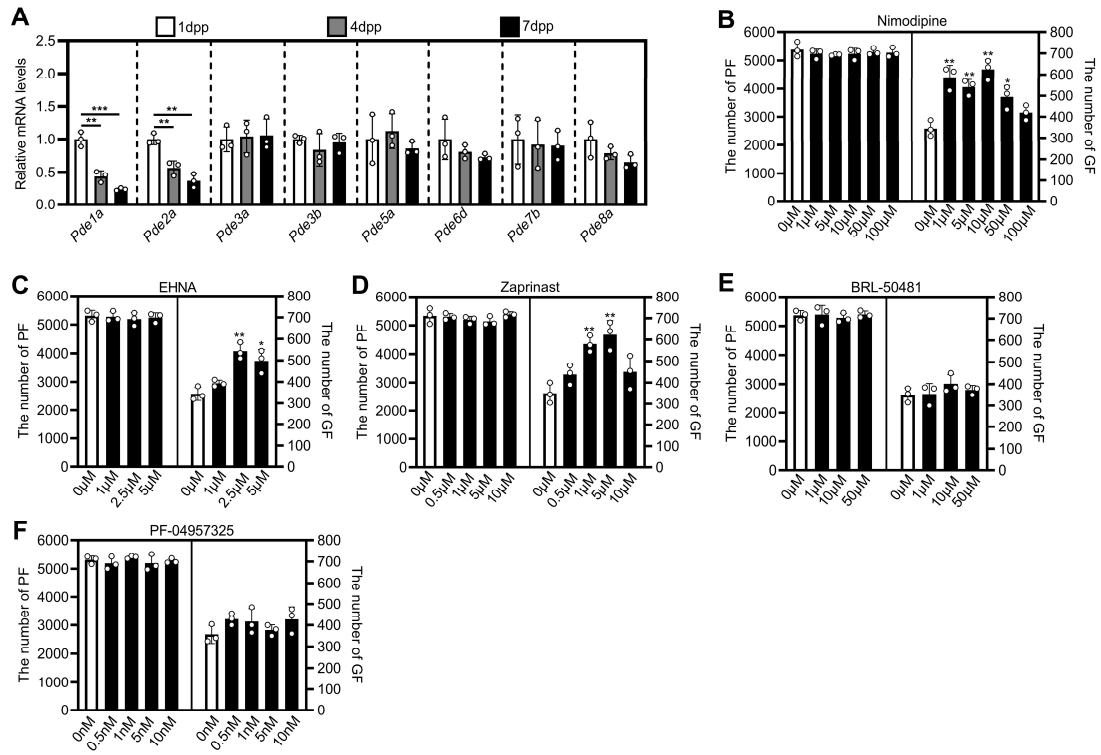
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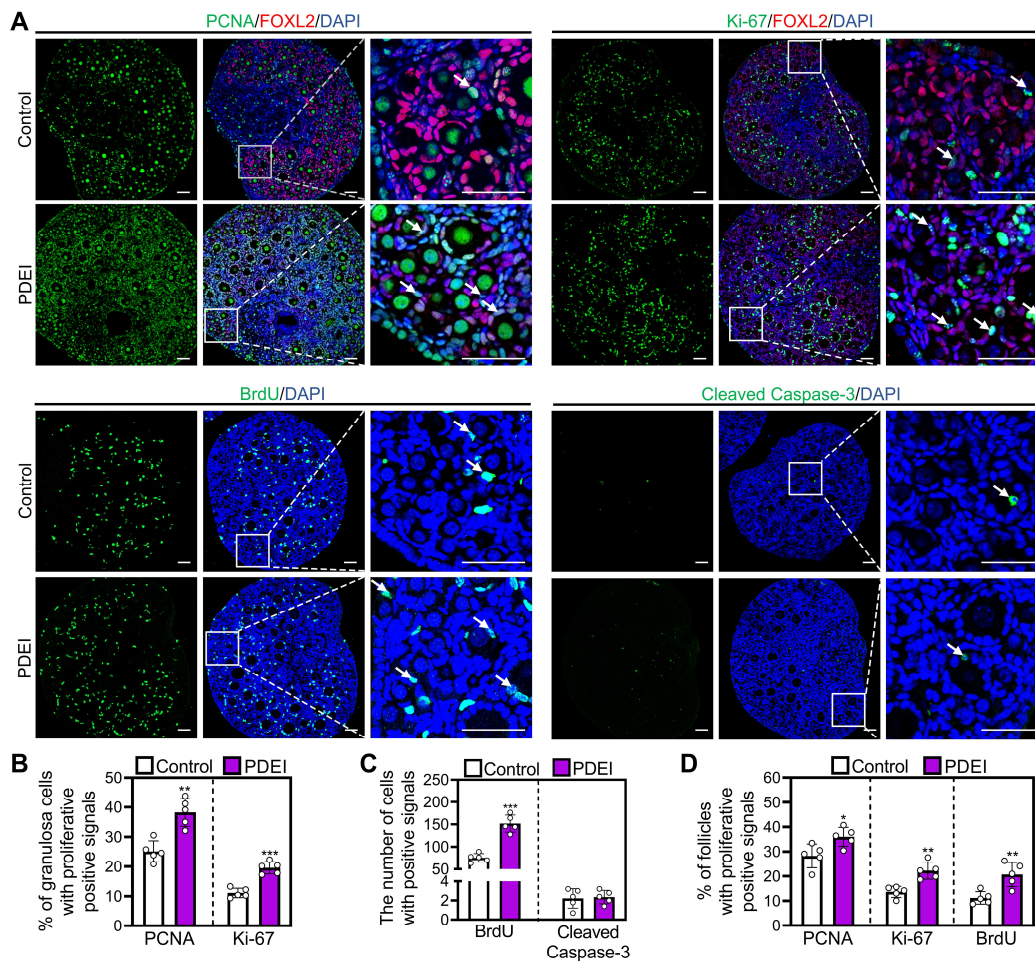
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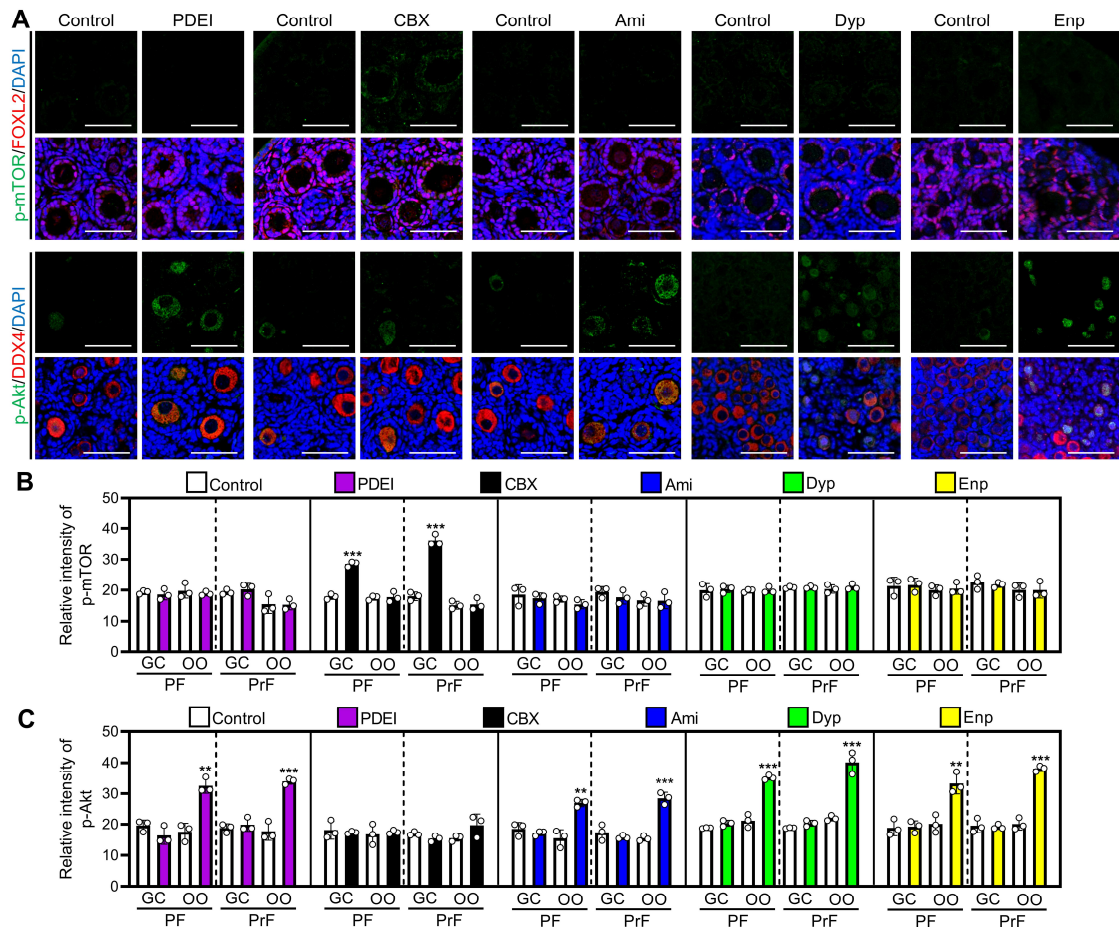
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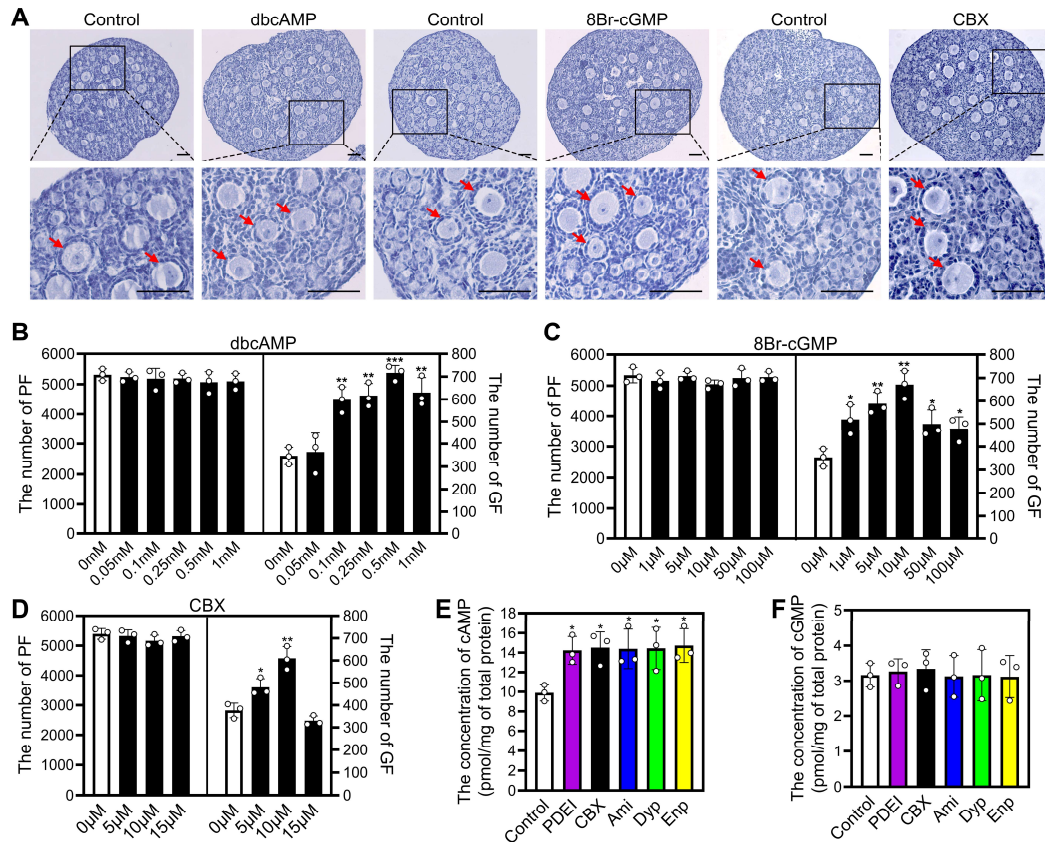
**Figure S1. Effects of specific phosphodiesterase inhibitors on mouse primordial follicle activation *in vitro*.** 1, 4, and 7 dpp mouse ovaries were collected for qRT-PCR analysis (A). 3 dpp mouse ovaries were cultured in medium alone or with nimodipine, EHNA, zaprinast, BRL-50481, or PF-04957325 in different concentrations for 4 days (B-F). A, The comparison of highly expressed phosphodiesterase subtype mRNA levels in 1,4 and 7dpp mouse ovaries (n = 3 biological replicates). \*\* $P < 0.01$ , \*\*\* $P < 0.001$  vs. 1 dpp group. B-F, The comparison of primordial and growing follicle number across various groups. PF, primordial follicle; GF, growing follicle. In each experiment, n = 3 biological replicates. Bars indicate the mean  $\pm$  SD. \* $P < 0.05$ , \*\* $P < 0.01$  and \*\*\* $P < 0.001$ . Two-tailed unpaired t-test was used to assess statistical significance.



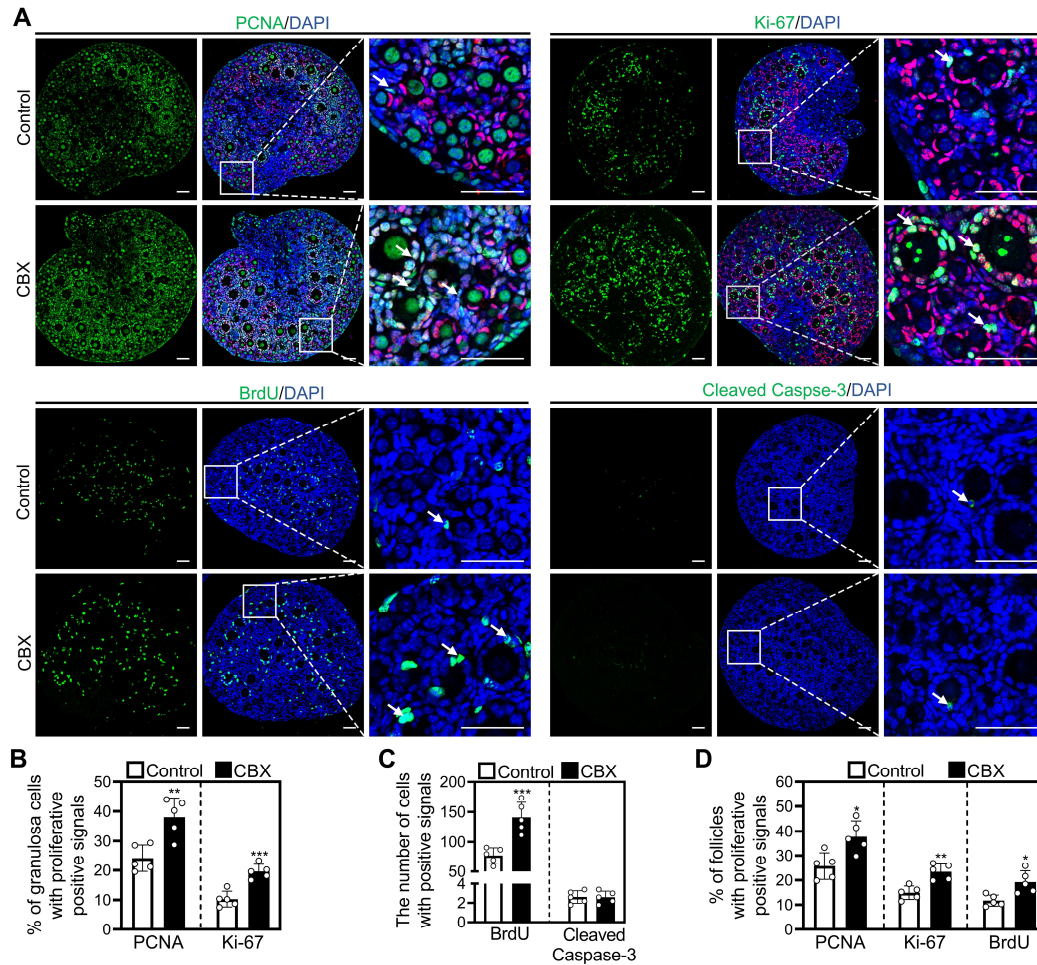
**Figure S2. Effects of PDEI on granulosa cell proliferation *in vitro*.** 3 dpp mouse ovaries were cultured in medium alone or with PDEI (the combination of 10  $\mu$ M nimodipine, 2.5  $\mu$ M EHNA, and 5  $\mu$ M zaprinast) for 2 days. **A**, PCNA, Ki-67, BrdU, and Cleaved Caspase-3 immunofluorescence stain (green) in control and PDEI group. FOXL2, red. **B-C**, The comparison of PCNA- and Ki-67-positive percentage in granulosa cells (**B**) and BrdU- and Cleaved Caspase-3-positive cell number (**C**) between control and PDEI groups. **D**, The comparison of PCNA-, Ki-67- and BrdU- positive percentage in primordial follicles between control and PDEI groups. The representative images were displayed. Scale bars, 50  $\mu$ m. In each experiment, n = 5 biological replicates. Bars indicate the mean  $\pm$  SD. \* $P$  < 0.05, \*\* $P$  < 0.01 and \*\*\* $P$  < 0.001. Two-tailed unpaired t-test was used to assess statistical significance.



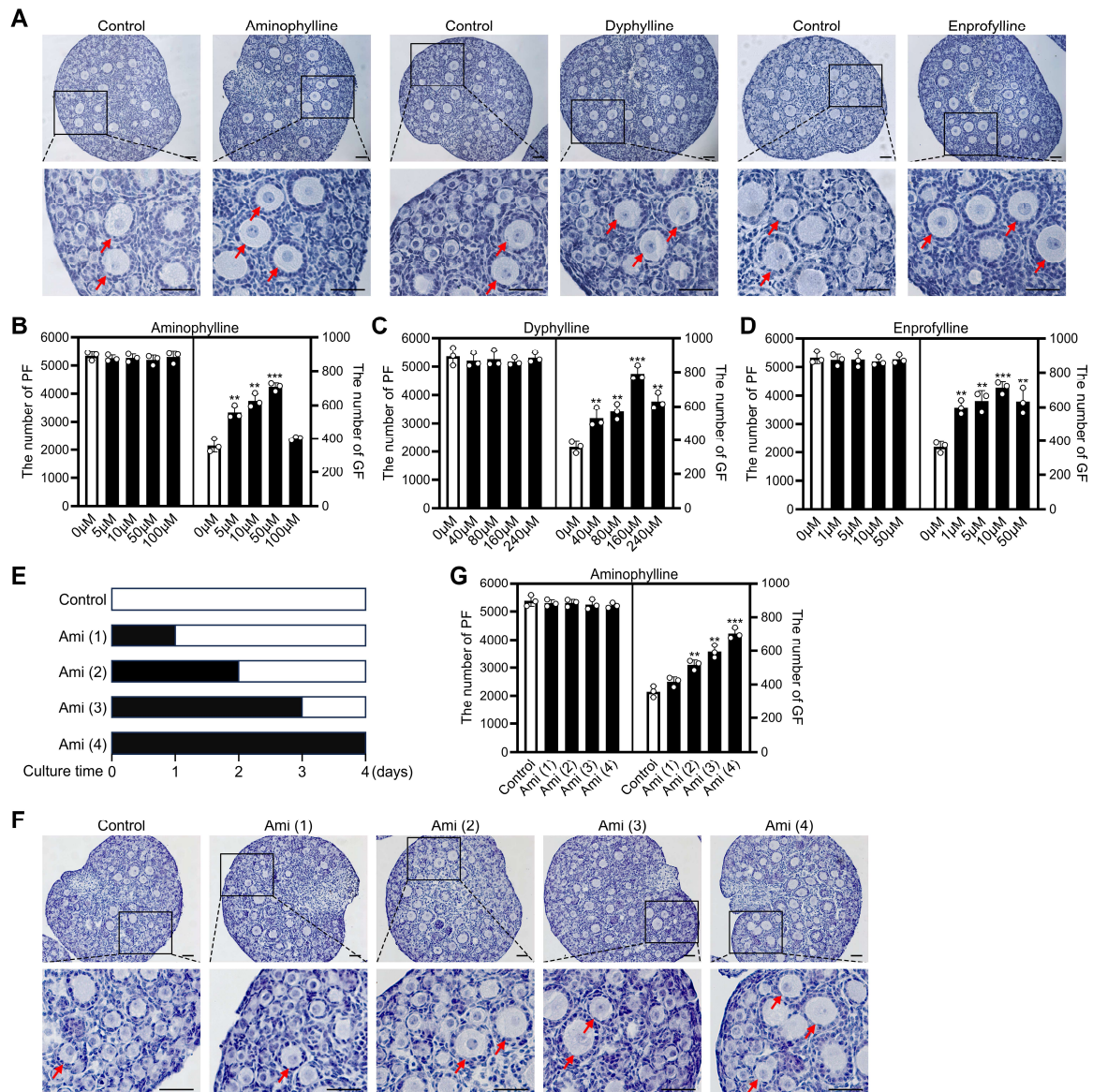
**Figure S3. Effects of CBX, PDEI, and theophylline derivatives on p-mTOR and p-Akt fluorescent intensities in cultured mouse ovaries.** 3 dpp mouse ovaries were cultured in medium alone or with PDEI (the combination of 10  $\mu\text{M}$  nimodipine, 2.5  $\mu\text{M}$  EHNA, and 5  $\mu\text{M}$  zaprinast), 10  $\mu\text{M}$  CBX, 50  $\mu\text{M}$  aminophylline (Ami), 160  $\mu\text{M}$  dyphylline (Dyp), or 10  $\mu\text{M}$  enprofylline (Enp) for 1 day. **A**, p-mTOR and p-Akt (green) immunofluorescence stain across various groups. FOXL2 or DDX4, red. **B-C**, The p-mTOR (**B**) and p-Akt (**C**) fluorescent intensities in granulosa cells and oocytes of primordial follicles and primary follicles. The representative images were displayed. GC, granulosa cell; OO, oocyte; PF, primordial follicle; PrF, primary follicle. Scale bars, 50  $\mu\text{m}$ . In each experiment,  $n = 3$  biological replicates. Bars indicate the mean  $\pm$  SD.  $**P < 0.01$  and  $***P < 0.001$ . Two-tailed unpaired t-test was used to assess statistical significance.



**Figure S4. Effects of dbcAMP, 8Br-cGMP and CBX on mouse primordial follicle activation *in vitro*.** 3 dpp mouse ovaries were cultured in medium alone or with dbcAMP, 8Br-cGMP, or CBX in different concentrations for 4 days (A-D), or cultured with PDEI (the combination of 10 μM nimodipine, 2.5 μM EHNA, and 5 μM zaprinast), 10 μM CBX, 50 μM aminophylline (Ami), 160 μM dyphylline (Dyp), or 10 μM enprofylline (Enp) for 1 day (E-F). A-D, The comparison of ovarian morphology (A) and primordial and growing follicle number (B-D) across various groups. The ovarian sections were hematoxylin-stained. E-F, The comparison of cAMP (E) and cGMP (F) concentration across various groups. The representative images were displayed. PF, primordial follicle; GF, growing follicle. Red arrows, growing follicles. In each experiment, n = 3 biological replicates. Bars indicate the mean ± SD. \* $P < 0.05$ , \*\* $P < 0.01$  and \*\*\* $P < 0.001$ . Two-tailed unpaired t-test was used to assess statistical significance.



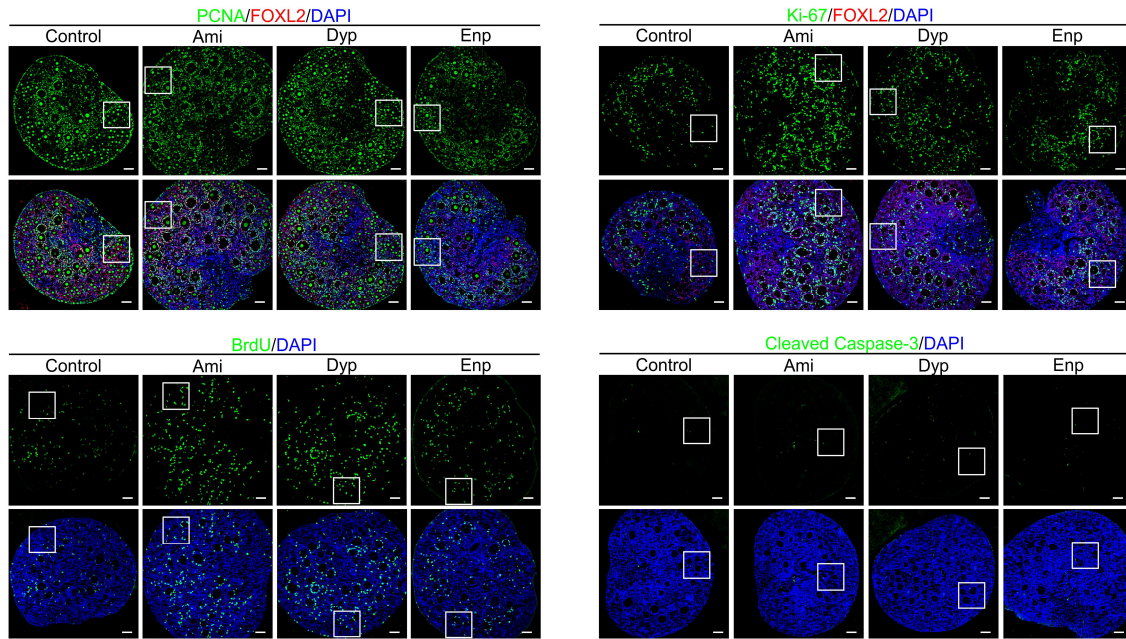
**Figure S5. Effects of CBX on granulosa cell proliferation *in vitro*.** 3 dpp mouse ovaries were cultured in medium alone or with 10  $\mu$ M CBX for 2 days. **A**, PCNA, Ki-67, BrdU, and Cleaved Caspase-3 immunofluorescence stain (green) in control and CBX groups. FOXL2, red. **B-C**, The comparison of PCNA- and Ki-67-positive percentage in granulosa cells (**B**) and BrdU- and Cleaved Caspase-3-positive cell number (**C**) between control and CBX groups. **D**, The comparison of PCNA-, Ki-67- and BrdU- positive percentage in primordial follicles between control and CBX groups. The representative images were displayed. Scale bars, 50  $\mu$ m. In each experiment, n = 5 biological replicates. Bars indicate the mean  $\pm$  SD. \* $P$  < 0.05, \*\* $P$  < 0.01 and \*\*\* $P$  < 0.001. Two-tailed unpaired t-test was used to assess statistical significance.



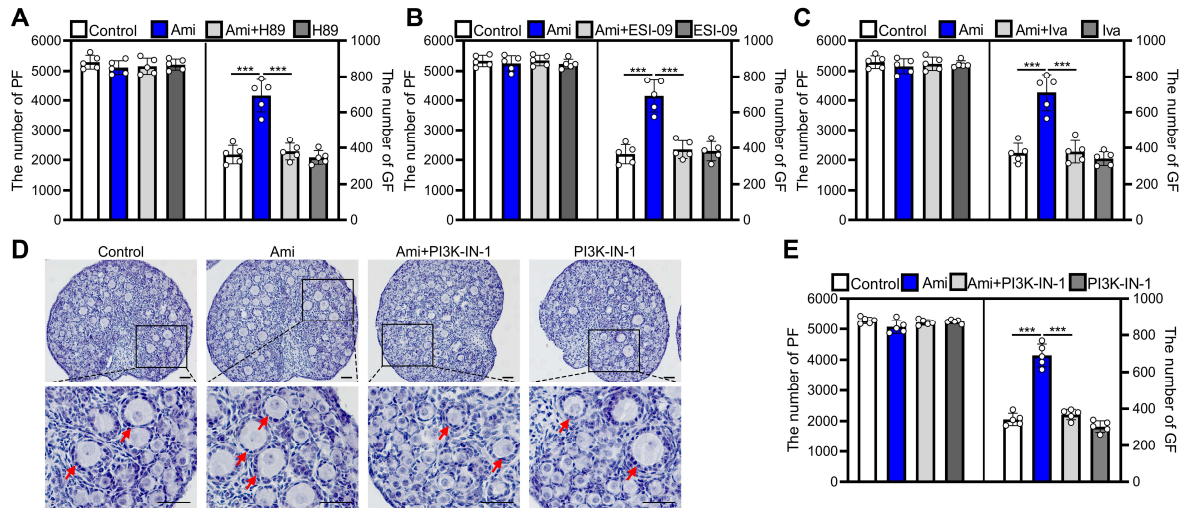
**Figure S6. Effects of the theophylline derivatives on mouse primordial follicles activation *in vitro*.** 3 dpp mouse ovaries were cultured in medium alone or with aminophylline, dyphylline, or enprofylline in different concentrations for 4 days (A-D) or cultured in medium alone or with aminophylline for indicated times (E-G). A-D, The comparison of ovarian morphology (A) and primordial and growing follicle number (B-D) across various groups. E, The timeline of the experiment was shown. 3 dpp mouse ovaries were cultured in medium alone or with aminophylline for indicated times and collected after 4 days of culture. F-G, The comparison of ovarian morphology (F) and primordial and growing follicle number (G) across various groups. Ami: aminophylline. The ovarian sections were hematoxylin-stained. Red arrows,

growing follicles. The representative images were displayed. Scale bars, 50  $\mu\text{m}$ . PF, primordial follicle; GF, growing follicle. In each experiment,  $n = 3$  biological replicates. Bars indicate the mean  $\pm$  SD.  $**P < 0.01$  and  $***P < 0.001$ . Two-tailed unpaired t-test was used to assess statistical significance.

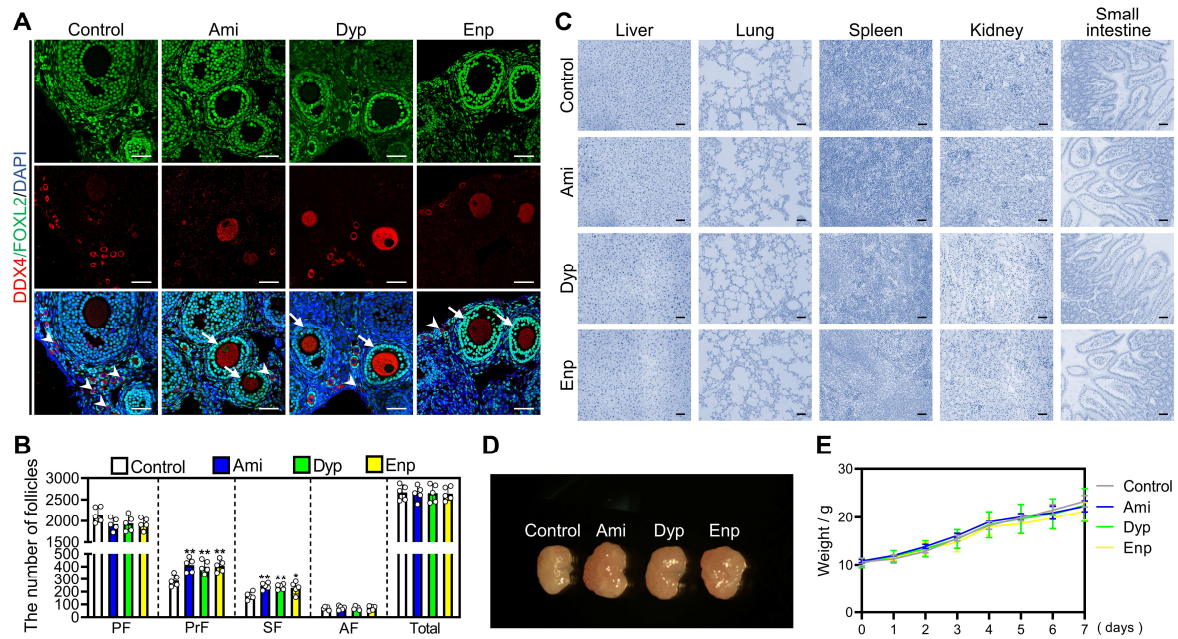




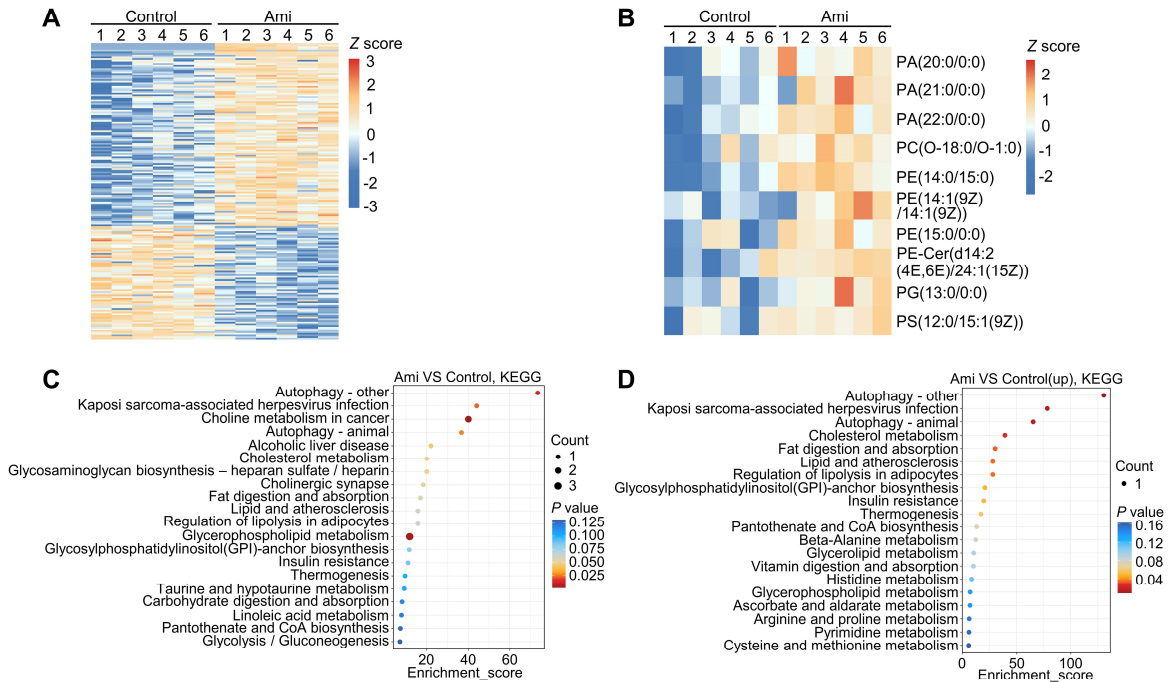
**Figure S7. Effects of the theophylline derivatives on granulosa cell proliferation *in vitro*** 3 dpp mouse ovaries were cultured in medium alone or with 50  $\mu\text{M}$  aminophylline (Ami), 160  $\mu\text{M}$  dyphylline (Dyp), or 10  $\mu\text{M}$  enprofylline (Enp) for 2 days. PCNA, Ki-67, BrdU, and Cleaved Caspase-3 (green) immunofluorescence stain across various groups. FOXL2, red. Figure 3H showed enlarged view of the boxed areas. Scale bars, 50  $\mu\text{m}$ .



**Figure S8. Effects of H89, ESI-09, ivabradine, and PI3K-IN-1 on aminophylline-induced mouse primordial follicle activation.** 3 dpp mouse ovaries were cultured in medium alone, supplemented with 50  $\mu$ M aminophylline (Ami), 5  $\mu$ M H89, 5 nM ESI-09, 2.5  $\mu$ M ivabradine (Iva), and/or 25  $\mu$ M PI3K-IN-1 for 4 days. **A-C**, The comparison of primordial and growing follicle number across various groups. **D-E**, The comparison of ovarian morphology (**D**) and primordial and growing follicle number (**E**) across various groups. The ovarian sections were hematoxylin-stained. Red arrows, growing follicles. The representative images were presented. Scale bars, 50  $\mu$ m. PF, primordial follicle; GF, growing follicle. In each experiment,  $n = 5$  biological replicates. Bars indicate the mean  $\pm$  SD. \*\*\* $P < 0.001$ . Two-tailed unpaired t-test was used to assess statistical significance.



**Figure S9. Effects of oral theophylline derivatives on adolescent mice.** The adolescent mice were administered water either with or without 4 mM aminophylline (Ami), 13 mM dyphylline (Dyp), or 0.8 mM enprofylline (Enp) for one week, followed by the collection of ovaries and organs. **A-B**, The comparison of ovarian morphology (**A**) and follicle number at different stages (**B**) across various groups. DDX4, red; FOXL2, green. **C**, Morphological comparison of liver, spleen, kidney and small intestine. **D**, The adolescent mice ovarian morphology across various groups. **e**, The body weight change across various groups. The representative images were displayed. Arrowheads, primordial follicles; arrows, growing follicles. PF, primordial follicle; PrF, primary follicle; SF, secondary follicle; AF, antral follicle. Scale bars, 50  $\mu$ m. In each experiment,  $n \geq 3$  for biological replicates. Bars indicate the mean  $\pm$  SD. \* $P < 0.05$  and \*\* $P < 0.01$ . Two-tailed unpaired t-test was used to assess statistical significance.



**Figure S10. Heatmap and KEGG analysis of differential metabolites in granulosa cells from naturally aged mice.** The naturally aged mice were administered water either with or without 4.0 mM aminophylline, and then the metabolome of granulosa cells was examined. **A-B**, Heatmap showed differential metabolites in granulosa cells from control and aminophylline (Ami) groups. **C-D**, The top twenty altered (**C**) and upregulated (**D**) pathways in granulosa cells from aminophylline group compared to those from control group by KEGG analysis.

**Table S1.** List of primary antibodies used in this study. IF: Immunofluorescence; WB: Western blotting.

Antibody	Catalog Code	Source	Host	Dilution	
				IF	WB
Akt	4691	Cell Signaling Technology	Rabbit	—	1:1000
p-Akt	4060	Cell Signaling Technology	Rabbit	1:200	1:1000
BrdU	ab1893	Abcam	Sheep	1:200	—
BAX	50599-2-Ig	Proteintech	Rabbit	—	1:1000
BCL2	26593-1-AP	Proteintech	Rabbit	—	1:1000
Cleaved Caspase-3	9664	Cell Signaling Technology	Rabbit	1:50	1:1000
DDX4	ab27591	Abcam	Mouse	1:200	1:1000
FOXL2	NB100-1277	Novus Biologicals	Goat	1:300	—
FOXO3a	12829	Cell Signaling Technology	Rabbit	1:100	1:1000
GDF9	ab254323	Abcam	Rabbit	—	1:1000
GLUT4	ab33780	Abcam	Rabbit	—	1:1000
HK1	ab150423	Abcam	Rabbit	—	1:1000
Ki-67	9129s	Cell Signaling Technology	Rabbit	1:100	—
mTOR	2972	Cell Signaling Technology	Rabbit	—	1:1000
p-mTOR	2971	Cell Signaling Technology	Rabbit	1:200	1:1000

PCNA	2586	Cell Signaling Technology	Mouse	1:100	1:1000
PFKL	ab181064	Abcam	Rabbit	—	1:1000
PKM2	4053	Cell Signaling Technology	Rabbit	—	1:1000
ZP3	sc-398359	Santa Cruz	Mouse		1:1000
$\alpha$ -Tubulin	ab195887	Abcam	Mouse	1:300	—
$\beta$ -actin	4967	Cell Signaling Technology	Rabbit	—	1:1000

**Table S2.** Primers for qRT-PCR used in this study.

Genes	Forwards (5'-3')	Backwards (5'-3')
<i>Aldoa</i>	CGTGTGAATCCCTGCATTGG	CAGCCCCTGGGTAGTTGTC
<i>Bax</i>	TTTCATCCAGGATCGAGCAGG	GCAAAGTAGAAGAGGGCAACCA C
<i>Bcl2</i>	CTACCGTCGTGACTTCGCA	TACCCAGCCTCCGTTATCC
<i>Caspase -3</i>	CCGGTTACTATTCCTGGAGA	TAACACGAGTGAGGATGTGC
<i>Eno1</i>	TGCGTCCACTGGCATCTAC	CAGAGCAGGCGCAATAGTTTTA
<i>Gdf9</i>	TCTTAGTAGCCTTAGCTCTCAGG	TGTCAGTCCCATCTACAGGCA
<i>Glut4</i>	AACTGGTCCTAGCTGTATTCT	CCAGCCACGTTGCATTGTA
<i>Hk1</i>	CGGAATGGGGAGCCTTTGG	GCCTTCCTTATCCGTTTCAATGG
<i>Ki-67</i>	ATCATTGACCGCTCCTTTAGGT	GCTCGCCTTGATGGTTCCT
<i>Ldhb</i>	CATTGCGTCCGTTGCAGATG	GGAGGAACAAGCTCCCGTG
<i>Pcna</i>	CGGCGTGAACCTGCAGAGCA	GGTTGCGGTCGCAGCGGTAT
<i>Pde1a</i>	CCAGACTGACTCCGTCCCAT	CCATTTTGCGTGTGAAAGTTGA
<i>Pde1b</i>	GATGCTGGAGTCGGATTGCC	TTCAGTGTCTAGGATTTGCCTTG

<i>Pde1c</i>	ACGTCCCAGAGGTTACGGT	GGCTGCATATTCCAGATTCTTCT
<i>Pde2a</i>	TGGCGTTGTGGACGATGAG	CGCGATAGAAAAGCGGATGG
<i>Pde3a</i>	CCTGGACTAGCGTGCTTAGGA	CAGGCGACCTTGAACCTCT
<i>Pde3b</i>	AAAGCGCAGCCGGTACTAT	CACCACTGCTTCAAGTCCCAG
<i>Pde4a</i>	AATGCCCTACAGACGCCTG	GACGGTGTGGCCCATTTT
<i>Pde4b</i>	TTCACGGTGGCTCATACATGC	CGCTGTCAAGATCGTAGAGGAA
<i>Pde4c</i>	TCCGAGAGCCAGTGGATTCT	CCTTGAGTTCCAATCGTGAAGA C
<i>Pde4d</i>	TTTTGCCAGTGCAATACATGATG	CAGAGCGAGTTCCGAGTTTGT
<i>Pde5a</i>	CGGCCTACCTGGCATTCTG	GCAAGGTCAAGTAACACCTGAT T
<i>Pde6a</i>	CAGCAACTACCACGATGTGAA	GTGGACATTGAAGAGCCTAGTG
<i>Pde6b</i>	GCAGCACTTTTTGAACTGGTG	CATTGCGCTGGCGGTACATA
<i>Pde6c</i>	GCGGCAGTTTGAAACGGTG	CATCATAGGCTGACTCTGCAC
<i>Pde6d</i>	CCCGTGTGCCCAAGAAAATC	CCACTCTTCTAGGCATTGTCCTT
<i>Pde6g</i>	AGGGTGAGATTCGGTCAGC	GCTCTTGAAGTGCCTTGTTTG
<i>Pde6h</i>	AGGGGTGAAAGGGTTTGGAGA	ATGATCCCGAACTGAGCAAGC
<i>Pde7a</i>	AGTGGATCACCTCTAAGAGACG	CGGACATCTCCTAGCATAACGAA T
<i>Pde7b</i>	TGCTAGGAGATGTACGACTAAG G	GGGCCTGCGGTATAATCCC
<i>Pde8a</i>	CCGAGCATCCACACTTCCG	TCAGCTACTGATACCTTCGAGG
<i>Pde8b</i>	AGAGCGGTGTGATCTACTGC	CGTCGGTCTGCACGAAGAG
<i>Pde9a</i>	CCACCATCTCCCTTTTAACCAC	CAGCACGCCCTGGATAAGT
<i>Pde10a</i>	GCAGGGGACAATCCTTGCC	CGTCAAACCTCTTGTGAACTCTT
<i>Pde11a</i>	AACAGGACCTACGATGAACAGG	TGAGGCAGATTCACCCTCGAT
<i>Pfkl</i>	GAACTACGCACACTTGACCAT	CTCCAAAACAAAGGTCCTCTGG

<i>Rpl19</i>	CTGAAGGTCAAAGGGAATGTGTT C	TGGTCAGCCAGGAGCTTCTTG
<i>Tpi</i>	CCAGGAAGTTCTTCGTTGGGG	CAAAGTCGATGTAAGCGGTGG
<i>Zp3</i>	CCTCAGGACTAACCGTGTGGA	CCATCAGGCGAAGAGAGAAAG