

C-phycoerythrin protects against low fertility by inhibiting reactive oxygen species in aging mice

Supplementary Material

Table S1-1. PC did not affect organ weight in D-gal-induced aging mice

Treatment	Organ weight (mg)			
	Mean \pm SEM (<i>n</i>)			
	Ovary	Liver	Spleen	Kidney
Ctrl	44.95 \pm 1.11 (6) ^a	1535.92 \pm 20.36 (6) ^a	104.33 \pm 3.37 (6) ^a	347.73 \pm 9.09 (6) ^a
D-gal	40.40 \pm 1.17 (6) ^a	1506.22 \pm 94.31 (6) ^a	99.47 \pm 4.26 (6) ^a	353.48 \pm 12.43 (6) ^a
D-gal+PC	41.77 \pm 3.16 (6) ^a	1427.40 \pm 76.66 (6) ^a	95.23 \pm 4.94 (6) ^a	323.30 \pm 12.73 (6) ^a

Within column, data without common superscript letters indicate significant difference ($P < 0.05$). *n* indicates the number of mice for each treatment.

Table S1-2. PC reversed some organ coefficients in D-gal-induced aging mice

Treatment	Organ coefficient (mg/g)			
	Mean \pm SEM (<i>n</i>)			
	Ovary/body	Liver/body	Spleen/body	Kidney/body
Ctrl	1.54 \pm 0.08 (6) ^a	52.31 \pm 1.28 (6) ^a	3.55 \pm 0.12 (6) ^a	11.82 \pm 0.25 (6) ^a
D-gal	1.23 \pm 0.05 (6) ^b	45.73 \pm 3.28 (6) ^a	3.01 \pm 0.14 (6) ^b	10.69 \pm 0.35 (6) ^b
D-gal+PC	1.50 \pm 0.11 (6) ^a	51.34 \pm 2.26 (6) ^a	3.42 \pm 0.13 (6) ^a	11.61 \pm 0.15 (6) ^a

Within column, data without common superscript letters indicate significant difference ($P < 0.05$). *n* indicates the number of mice for each treatment.

Table S2-1. PC did not increase oocyte number in D-gal-induced aging mice

Treatment	Oocyte number per mouse	
	Mean \pm SEM (<i>n</i>)	
	0 h after hCG	14 h after hCG
Ctrl	57.83 \pm 6.89 (6) ^a	54.17 \pm 1.80 (6) ^a
D-gal	50.67 \pm 3.92 (6) ^a	55.17 \pm 4.56 (6) ^a
D-gal+PC	52.00 \pm 8.12 (6) ^a	51.67 \pm 4.96 (6) ^a

Within column, data without common superscript letters indicate significant difference ($P < 0.05$).
n indicates the number of mice for each treatment

Table S2-2. PC increased oocyte PB1 extrusion in D-gal-induced aging mice

Treatment	PB1 extrusion (%)	
	Mean \pm SEM (<i>n</i>)	
	<i>in vivo</i> maturation	<i>in vitro</i> maturation
Ctrl	90.03 \pm 2.05 (6) ^a	83.61 \pm 1.13 (6) ^a
D-gal	69.72 \pm 1.29 (6) ^b	47.45 \pm 4.13 (6) ^b
D-gal+PC	85.15 \pm 3.09 (6) ^a	71.11 \pm 2.82 (6) ^c

Within column, data without common superscript letters indicate significant difference (ab, $P < 0.001$;
ac, $P < 0.05$; bc, $P < 0.01$).
n indicates the number of mice for each treatment.

Table S2-3. PC inhibited oocyte fragmentation in D-gal-induced aging mice

Treatment	Fragmentation (%)	
	Mean \pm SEM (<i>n</i>)	
	<i>in vivo</i> maturation	<i>in vitro</i> maturation
Ctrl	5.78 \pm 0.52 (7) ^a	9.67 \pm 0.58 (6) ^a
D-gal	13.08 \pm 1.60 (6) ^b	36.57 \pm 3.02 (6) ^b
D-gal+PC	7.01 \pm 0.48 (7) ^a	23.34 \pm 1.96 (6) ^c

Within column, data without common superscript letters indicate significant difference (ab, $P < 0.001$;
ac, $P < 0.001$; bc, $P < 0.01$).
n indicates the number of mice for each treatment.

Table S2-4. PC inhibited oocyte aneuploidy in D-gal-induced aging mice

Treatment	Aneuploidy (%)	
	Mean \pm SEM (<i>n</i>)	
	<i>in vivo</i> maturation	<i>in vitro</i> maturation
Ctrl	3.41 \pm 0.52 (9) ^a	5.41 \pm 0.94 (9) ^a
D-gal	14.57 \pm 0.35 (9) ^b	18.71 \pm 0.88 (9) ^b
D-gal+PC	7.77 \pm 0.43 (9) ^c	10.19 \pm 0.44 (9) ^c

Within column, data without common superscript letters indicate significant difference ($P < 0.001$).
n indicates the number of mice for each treatment.

Table S3. PC normalized spindle chromosome complex in D-gal-induced aging mice

Treatment	Abnormal SCCs (%)	
	Mean \pm SEM (<i>n</i>)	
Ctrl	16.31 \pm 3.68 (6) ^a	
D-gal	43.32 \pm 2.28 (6) ^b	
D-gal+PC	21.40 \pm 3.01 (6) ^a	

Data without common superscript letters indicate significant difference ($P < 0.001$).
n indicates the number of mice for each treatment.

Table S4. D-gal and PC did not influence ovary telomere length or telomerase activity

Treatment	T/S ratio	Telomerase activity
	Mean \pm SEM (<i>n</i>)	Mean \pm SEM (<i>n</i>)
Ctrl	1.22 \pm 0.03 (8) ^a	1.60 \pm 0.06 (6) ^a
D-gal	1.20 \pm 0.02 (8) ^a	1.49 \pm 0.03 (6) ^a
D-gal+PC	1.18 \pm 0.02 (8) ^a	1.62 \pm 0.09 (6) ^a

Within column, data without common superscript letters indicate significant difference ($P < 0.05$).
n indicates the number of mice for each treatment.

Table S5-1. PC rescued the expression of some antioxidant genes

Gene name	Treatment (<i>n</i>)			superscript letters annotation
	Ctrl (6)	D-gal (6)	D-gal+PC (6)	
<i>Gclm</i>	1.02 ± 0.01 ^a	1.66 ± 0.19 ^b	1.22 ± 0.20 ^{ab}	ab, <i>P</i> < 0.05
<i>Gclc</i>	1.02 ± 0.00 ^a	1.58 ± 0.16 ^b	1.02 ± 0.15 ^{ac}	ab, <i>P</i> < 0.05; bc, <i>P</i> < 0.01
<i>Gpx1</i>	1.02 ± 0.01 ^a	1.60 ± 0.22 ^a	1.21 ± 0.17 ^a	<i>P</i> > 0.05
<i>Gpx3</i>	1.02 ± 0.00 ^a	1.49 ± 0.19 ^b	0.63 ± 0.10 ^c	ab, <i>P</i> < 0.05; ac, <i>P</i> < 0.05; bc, <i>P</i> < 0.001
<i>Gsr</i>	1.02 ± 0.01 ^a	1.27 ± 0.14 ^a	1.12 ± 0.16 ^a	<i>P</i> > 0.05
<i>Gsta4</i>	1.02 ± 0.01 ^a	1.35 ± 0.14 ^a	1.19 ± 0.13 ^a	<i>P</i> > 0.05
<i>Gstm1</i>	1.02 ± 0.00 ^a	1.30 ± 0.19 ^a	1.31 ± 0.22 ^a	<i>P</i> > 0.05
<i>Gstm2</i>	1.02 ± 0.01 ^a	1.29 ± 0.14 ^a	1.23 ± 0.13 ^a	<i>P</i> > 0.05
<i>Gstt1</i>	1.02 ± 0.00 ^a	1.36 ± 0.13 ^a	1.19 ± 0.20 ^a	<i>P</i> > 0.05
<i>Mgst1</i>	1.02 ± 0.00 ^a	1.33 ± 0.22 ^{ab}	1.75 ± 0.25 ^b	ab, <i>P</i> < 0.05
<i>Sod1</i>	1.02 ± 0.00 ^a	1.37 ± 0.12 ^a	1.01 ± 0.14 ^a	<i>P</i> > 0.05
<i>Sod2</i>	1.02 ± 0.00 ^a	1.30 ± 0.14 ^{ab}	1.73 ± 0.29 ^b	ab, <i>P</i> < 0.05
<i>Cat</i>	1.03 ± 0.01 ^a	0.76 ± 0.08 ^b	1.33 ± 0.11 ^c	ab, <i>P</i> < 0.05; ac, <i>P</i> < 0.05; bc, <i>P</i> < 0.001
<i>Glx1</i>	1.01 ± 0.00 ^a	1.35 ± 0.08 ^a	0.84 ± 0.09 ^a	<i>P</i> > 0.05
<i>Glx2</i>	1.02 ± 0.00 ^a	1.67 ± 0.21 ^b	1.26 ± 0.18 ^{ab}	ab, <i>P</i> < 0.05
<i>Prdx3</i>	1.02 ± 0.00 ^a	1.32 ± 0.13 ^a	1.24 ± 0.19 ^a	<i>P</i> > 0.05
<i>Txn2</i>	1.03 ± 0.01 ^a	1.85 ± 0.15 ^b	1.39 ± 0.22 ^{ab}	ab, <i>P</i> < 0.01
<i>Txnr1</i>	1.01 ± 0.00 ^a	1.30 ± 0.13 ^a	1.47 ± 0.37 ^a	<i>P</i> > 0.05
<i>Txnr2</i>	1.02 ± 0.01 ^a	1.48 ± 0.18 ^a	1.27 ± 0.24 ^a	<i>P</i> > 0.05
<i>Ccs</i>	1.01 ± 0.01 ^a	1.08 ± 0.14 ^a	1.16 ± 0.12 ^a	<i>P</i> > 0.05

Within row, data without common superscript letters indicate significant difference (see superscript letters annotation).

n indicates the number of mice for each treatment.

Table S5-2. PC increased SOD activity in D-gal-induced aging mice

Treatment	SOD activity (U/mg)
	Mean ± SEM (<i>n</i>)
Ctrl	144.30 ± 3.41 (10) ^a
D-gal	67.92 ± 1.63 (10) ^b
D-gal+PC	89.21 ± 1.49 (10) ^c

Data without common superscript letters indicate significant difference (*P* < 0.001).

n indicates the number of mice for each treatment.

Table S5-3. D-gal or PC did not affect GSH-Px activity

Treatment	GSH-Px activity (U/mg)
	Mean \pm SEM (<i>n</i>)
Ctrl	47.78 \pm 2.84 (10) ^a
D-gal	48.92 \pm 3.39 (10) ^a
D-gal+PC	44.23 \pm 4.59 (10) ^a

Data without common superscript letters indicate significant difference ($P < 0.05$).
n indicates the number of mice for each treatment.

Table S5-4. PC did not reverse the increased CAT activity in D-gal-induced aging mice

Treatment	CAT activity (U/mg)
	Mean \pm SEM (<i>n</i>)
Ctrl	25.92 \pm 2.05 (10) ^a
D-gal	33.21 \pm 1.83 (10) ^b
D-gal+PC	38.01 \pm 2.74 (10) ^{bc}

Data without common superscript letters indicate significant difference (ab, $P < 0.05$; ac, $P < 0.01$).
n indicates the number of mice for each treatment.

Table S5-5. PC decreased MDA content in D-gal-induced aging mice

Treatment	MDA content (mmol/g)
	Mean \pm SEM (<i>n</i>)
Ctrl	0.27 \pm 0.02 (9) ^a
D-gal	0.36 \pm 0.03 (9) ^b
D-gal+PC	0.25 \pm 0.03 (9) ^a

Data without common superscript letters indicate significant difference ($P < 0.01$).
n indicates the number of mice for each treatment.

Table S6-1. PC did not influence relative ATP level in cumulus cells or oocytes in D-gal-induced aging mice

Treatment	Relative ATP level	
	Mean \pm SEM (<i>n</i>)	
	cumulus cells	oocytes
Ctrl	1.07 \pm 0.14 (8) ^a	1.55 \pm 0.09 (8) ^a
D-gal	1.23 \pm 0.13 (8) ^a	1.40 \pm 0.08 (8) ^a
D-gal+PC	1.33 \pm 0.17 (8) ^a	1.41 \pm 0.09 (8) ^a

Within column, data without common superscript letters indicate significant difference ($P < 0.05$). *n* indicates the number of mice for each treatment.

Table S6-2. Mitochondrial distribution in D-gal-induced oocytes was normalized by PC

Treatment	Mitochondrial distribution (%)	
	Mean \pm SEM (<i>n</i>)	
	even	aggregated
Ctrl	94.24 \pm 1.39 (6) ^a	5.76 \pm 1.39 (6) ^a
D-gal	62.03 \pm 6.96 (6) ^b	37.97 \pm 6.96 (6) ^b
D-gal+PC	88.95 \pm 1.57 (6) ^a	11.05 \pm 1.57 (6) ^a

Within column, data without common superscript letters indicate significant difference ($P < 0.001$). *n* indicates the number of mice for each treatment.

Table S7. PC reduced high ROS levels induced by D-gal in MII oocytes

Treatment	ROS relative fluorescence
	Mean \pm SEM (<i>n</i>)
Ctrl	123.44 \pm 10.54 (6) ^a
D-gal	370.14 \pm 56.83 (6) ^b
D-gal+PC	181.25 \pm 28.22 (6) ^{ac}

Data without common superscript letters indicate significant difference (ab, $P < 0.001$; bc, $P < 0.01$). *n* indicates the number of mice for each treatment.

Table S8. PC inhibited D-gal-induced early apoptosis in MII oocytes

Treatment	Oocytes with early stage apoptosis (%)
	Mean \pm SEM (<i>n</i>)
Ctrl	6.19 \pm 0.62 (6) ^a
D-gal	19.89 \pm 1.07 (6) ^b
D-gal+PC	6.39 \pm 1.14 (7) ^a

Data without common superscript letters indicate significant difference ($P < 0.001$). *n* indicates the number of mice for each treatment.

Table S9-1. PC rescued litter size in D-gal-treated mice

Treatment	Litter size
	Mean \pm SEM (<i>N</i>)
Ctrl	8.69 \pm 0.47 (16) ^a
D-gal	6.15 \pm 0.73 (20) ^b
D-gal+PC	8.68 \pm 0.67 (19) ^{ac}

Data without common superscript letters indicate significant difference (ab, $P < 0.05$; bc, $P < 0.01$). *N* indicates the number of female mice with plugs and that gave birth for each treatment.

Table S9-2. Offspring birth weight was not significantly different among control, D-gal and D-gal+PC groups

Treatment	Birth weight (g)
	Mean \pm SEM (<i>N</i> , <i>n</i>)
Ctrl	1.57 \pm 0.05 (16, 139) ^a
D-gal	1.54 \pm 0.06 (20, 123) ^a
D-gal+PC	1.39 \pm 0.05 (19, 165) ^a

Data without common superscript letters indicate significant difference ($P < 0.05$). *N* indicates the number of female mice with plugs and that gave birth for each treatment. *n* indicates the total number of offspring for each treatment.

Table S9-3. Postnatal growth of female pups from weeks 1 to 8

week	Treatment (<i>N</i> , <i>n</i>)		
	Mean \pm SEM		
	Ctrl (16, 69)	D-gal (19, 62)	D-gal+PC (19, 82)
1	5.28 \pm 0.06 ^a	5.22 \pm 0.06 ^a	5.02 \pm 0.07 ^a
2	8.94 \pm 0.06 ^a	8.91 \pm 0.07 ^a	8.37 \pm 0.08 ^a
3	13.07 \pm 0.06 ^a	12.78 \pm 0.07 ^a	12.46 \pm 0.05 ^a
4	19.03 \pm 0.06 ^a	18.73 \pm 0.06 ^a	18.26 \pm 0.13 ^a
5	21.06 \pm 0.06 ^a	20.81 \pm 0.08 ^a	20.42 \pm 0.07 ^a
6	22.37 \pm 0.06 ^a	22.26 \pm 0.08 ^a	21.91 \pm 0.16 ^a
7	23.36 \pm 0.06 ^a	23.30 \pm 0.08 ^a	23.14 \pm 0.13 ^a
8	24.34 \pm 0.06 ^a	24.34 \pm 0.08 ^a	23.97 \pm 0.18 ^a

Within row, data without common superscript letters indicate significant difference ($P < 0.05$).

N indicates the number of female mice with plugs and that gave birth for each treatment. *n* indicates the total number of offspring for each treatment.

Table S9-4. Postnatal growth of male pups from weeks 1 to 8

week	Treatment (<i>N</i> , <i>n</i>)		
	Mean \pm SEM		
	Ctrl (16, 70)	D-gal (18, 61)	D-gal+PC (18, 83)
1	5.21 \pm 0.05 ^a	5.01 \pm 0.07 ^a	4.74 \pm 0.07 ^a
2	11.77 \pm 0.09 ^a	11.37 \pm 0.07 ^a	11.51 \pm 0.04 ^a
3	17.15 \pm 0.09 ^a	17.51 \pm 0.06 ^a	17.93 \pm 0.07 ^a
4	20.98 \pm 0.09 ^a	21.44 \pm 0.06 ^a	21.77 \pm 0.07 ^a
5	24.71 \pm 0.09 ^a	25.08 \pm 0.06 ^a	25.43 \pm 0.07 ^a
6	27.12 \pm 0.09 ^a	27.65 \pm 0.06 ^a	28.05 \pm 0.08 ^a
7	28.54 \pm 0.09 ^a	28.91 \pm 0.06 ^a	29.10 \pm 0.08 ^a
8	30.01 \pm 0.11 ^a	30.19 \pm 0.07 ^a	30.27 \pm 0.08 ^a

Within row, data without common superscript letters indicate significant difference ($P < 0.05$).

N indicates the number of female mice with plugs and that gave birth for each treatment. *n* indicates the total number of offspring for each treatment.