

Figure S1

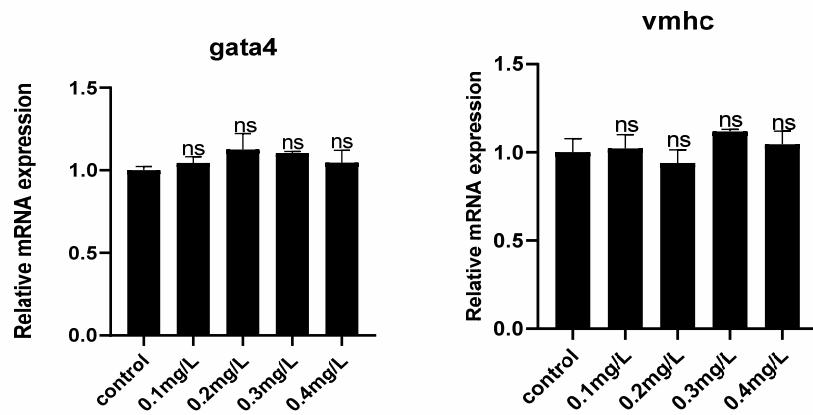
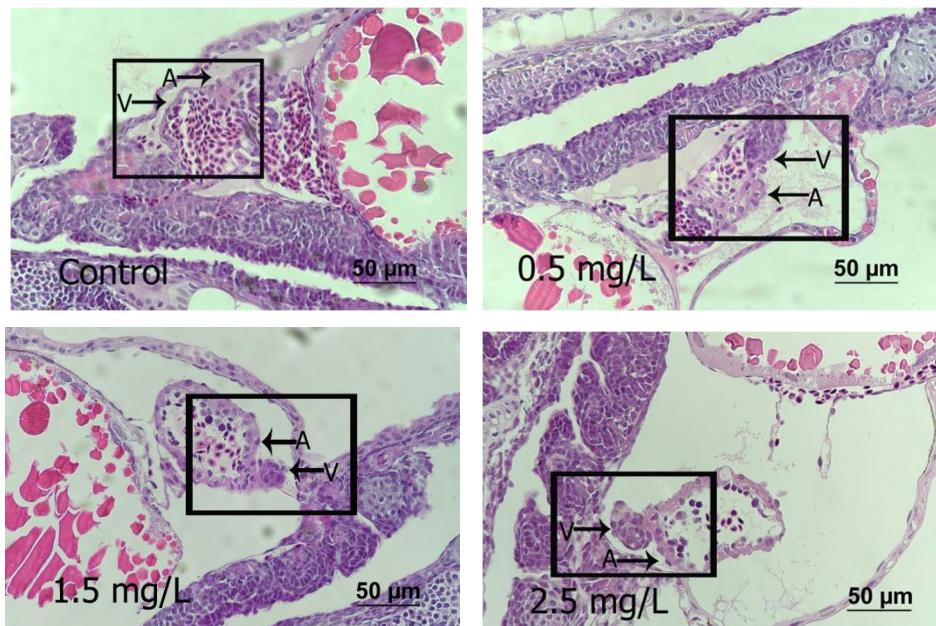


Figure S1 Effect of diflubenzuron exposure on mRNA expression of the genes related to heart development

The transcription of *gata4* and *vmhc* in zebrafish embryos after exposure to various concentrations of diflubenzuron (0, 0.1, 0.2, 0.3 and 0.4 mg/L) at 96 hpf. Each bar represents mean \pm SD ($n=24$) of three independent experiments. *ns*: no significant difference.

Figure S2

(A)



(B)

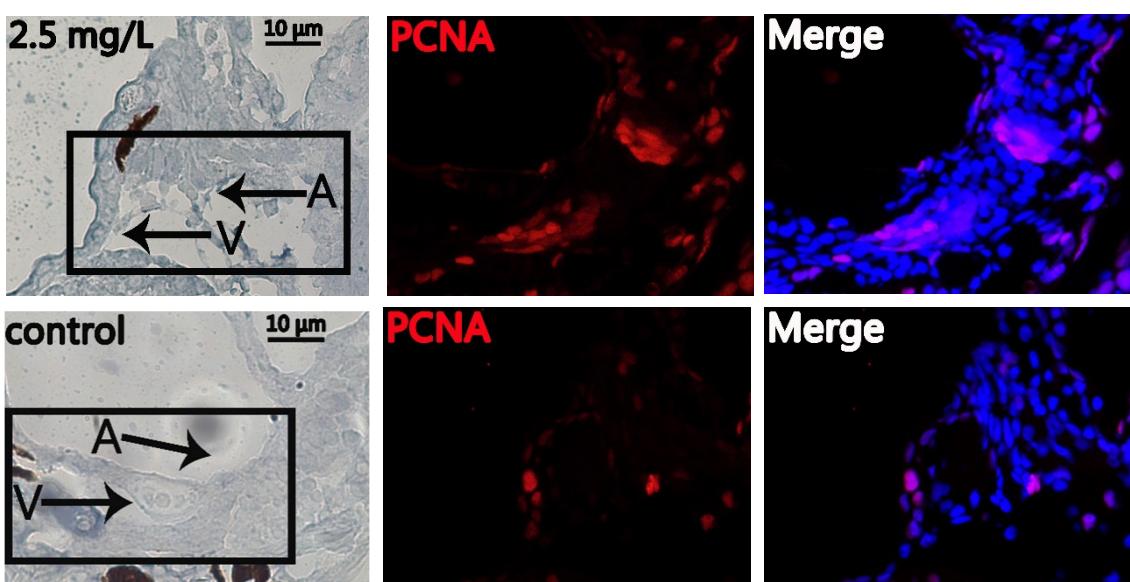


Figure S2 Exposure to diflubenzuron induces cardiac developmental toxicity in zebrafish embryos.

H&E staining of heart sections in control and 0.5, 1.5 and 2.5 mg/L diflubenzuron-exposed zebrafish at 96 hpf (A), The heart slices were prepared, incubated with PCNA

primary antibodies and stained with cy3-tagged second antibodies (**B**). DAPI was used to stain the nucleus. Each of experiment was repeated at least three times.

Figure S3

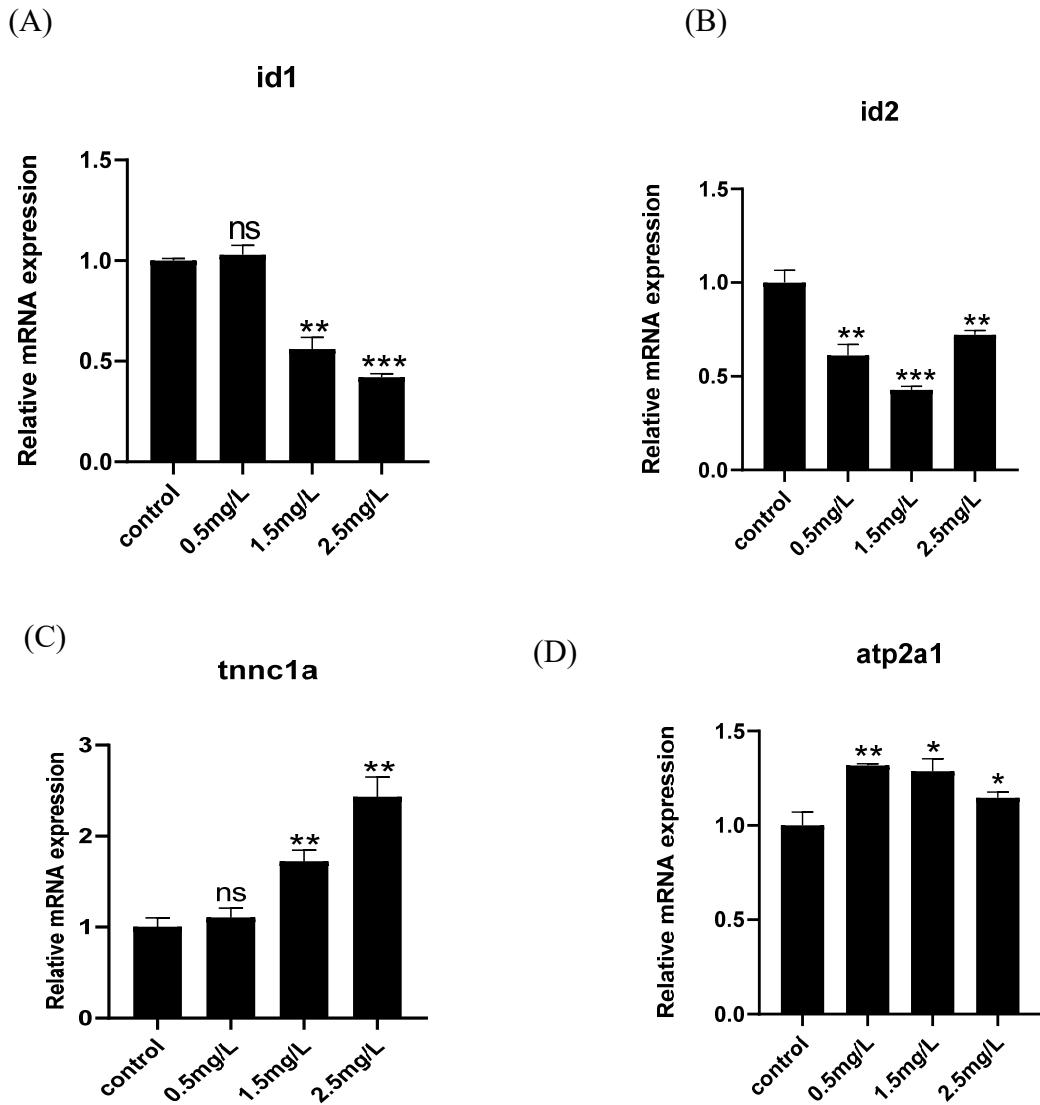


Figure S3 Exposure of diflubenzuron affects the gene expression of embryos in BMP signaling pathway and calcium dependent signaling pathway

Q-PCR was used to detect the genes in BMP signaling pathway (*id1* and *id2*) (**A and B**) and calcium dependent signaling pathway (*tnnc1a* and *atp2a1*) (**C and D**). Each bar represents mean \pm SD (n=24) of three independent experiments. Asterisks indicate significant differences from control, * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$.

Table S1
Sequences of the qRT-PCR primers used in this study.

Target gene	Prime sequence (5'-3')
β-actin-F	AGCACGGTATTGTGACTAACTG
β-actin-R	TCGAACATGATCTGTGTCATC
nppa-F	GCTCCTGGTTGGCAGCAG
nppa-R	GAGCTGCTGCTCCTCTCGG
gata4-F	CTACAGGCACCCCAGCAGA
gata4-R	AGAGCCCGAGACCCGAAAT
vmhc-F	GAGCTTGATGAGGCAGAAG
vmhc-R	CAGCATAACGGAGACACAG
tbx5-F	AGTGACAGCGAACCAAAAG
tbx5-R	GTCTGGATGGACATAAAGC
myh6-F	CACCAGCAGACACTGGATG
myh6-R	GCTCCAAGTCCATTCTGAC
tbx2b-F	GGGGAAACAATGGATGGCTAA
tbx2b-R	CCTAAGTGGCTGGAAACC
klf2a-F	CCGTCTATTCCACATTTCG
klf2a-R	TCCAGTTCATCCTCACCT
bax-F	TCGAACATGATCTGTGTCATC
bax-R	TATGGCTGGGTCACTTTCTC
bcl2-F	TGGCGTCCCAGGTAGATAAT
bcl2-R	ACCGTACATCTCCACGAAGG
p53-F	CCCGGATGGAGATAACTG
p53-R	CACAGTTGCCATTCAGCAC
casepase3-F	GAGACCGCTGCCCATCACTAG
casepase3-R	ATCCTTCACGACCACATCT
casepase9-F	GCCTTCTTGATTCCCTGCG
casepase9-R	TCTTGGCCTGGTTGGTCTCA
id1-F	CAGCAAAGTTGGAGGAGAGG
id1-R	AGCCGTTCTCCACAGAGATGCT
id2-F	GAGTGTAAACGACGACGGAGC
id2-R-F	TGATGCAGGCTGGTTATCGC
bmp4-F	GGTCATTATTATGCCAAGT
bmp4-R	AAACGAATCGCAGAGGAGT
atp2a1-F	CCGACAAAATGGCACCTTG
atp2a1-R	CTGCAGTCAACTTGGCACC
tnncl1a-F	GGCAGAGCAACTCACCGAT
tnncl1a-R	GTAGGGTTCTGGCCAACAT