

Supplementary Materials: Molecular Effects of Irradiation (Cobalt-60) on the Control of *Panonychus citri* (Acari: Tetranychidae)

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Table S1. Primers were used for the qPCR.

Gene ID	Protein	qPCR Primers (5' to 3')
Unigene0009307	apolipoprotein D	F: TGGTCAACGAGCTACCGAAT R: GTCGTTGAGTTGTATCCGCA
Unigene0019519	calmodulin-like	F: TGAGGTTGATGCCGATGGTA R: ACATGTCTCAATT CAGCAGCA
Unigene0022395	Breakpoint cluster region protein	F: TCCAAGCCAAACAAAACCCCT R: AGGAATCTCATGGTGCTCA
Unigene0017154	serine protease	F: CAGGGCATTACTGTGGAGGA R: AGCGAGTCTGTTCA GTTCCA
Unigene0015558	Intestinal-type alkaline phosphatase 1	F: CAAAGAGTCGC ACTGGTACA R: GTTCCGTTCCCTGGTCCAC
Unigene0054633	Cu/Zn superoxide dismutase	F: AGGAAACATTGAAGCCGATCA R: GTCCACCAGCATT CCTGTG
Unigene0021115	microtubule-associated proteins 1A/1B light chain 3A-like	F: TTGTACCCCCAACACCTTCCA R: ATICCCACCAGCGAGAAGAA
Unigene0022207	DNA repair endonuclease XPF-like	F: ACCATGCCACACTTT CAG R: TGAGGTCAATAAGGT CAGCGA
Unigene0024951	autophagy-related 4	F: GGTTGTATGCTACGCTGTGG R: TACCTTCAGTCGCACCCATT

COG Function Classification of *Panonychus citri*-Unigene.fa Sequence

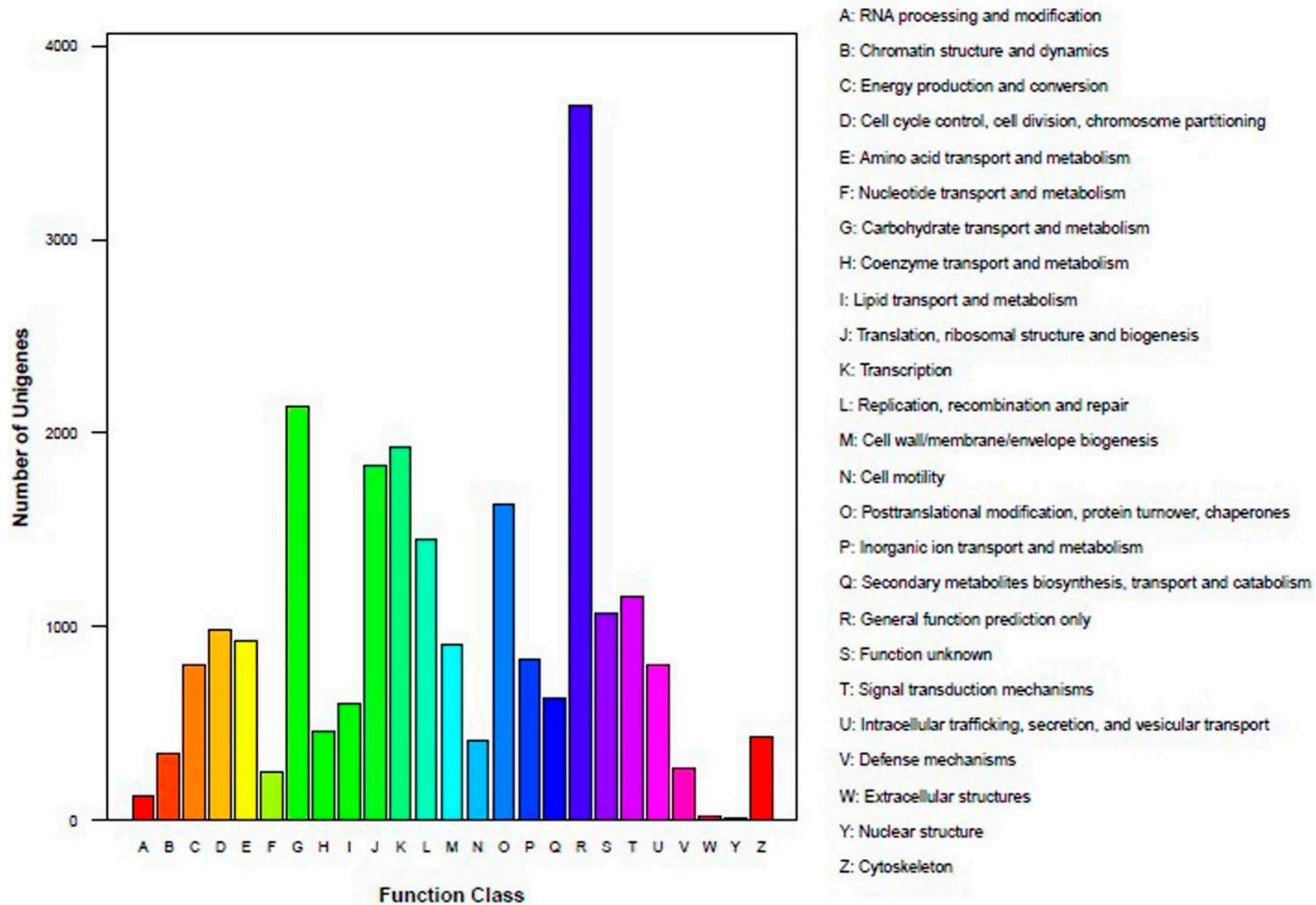


Figure S1. COG (Clusters of Orthologous Groups) function classification annotation of unigenes from *P. citri* transcriptome.

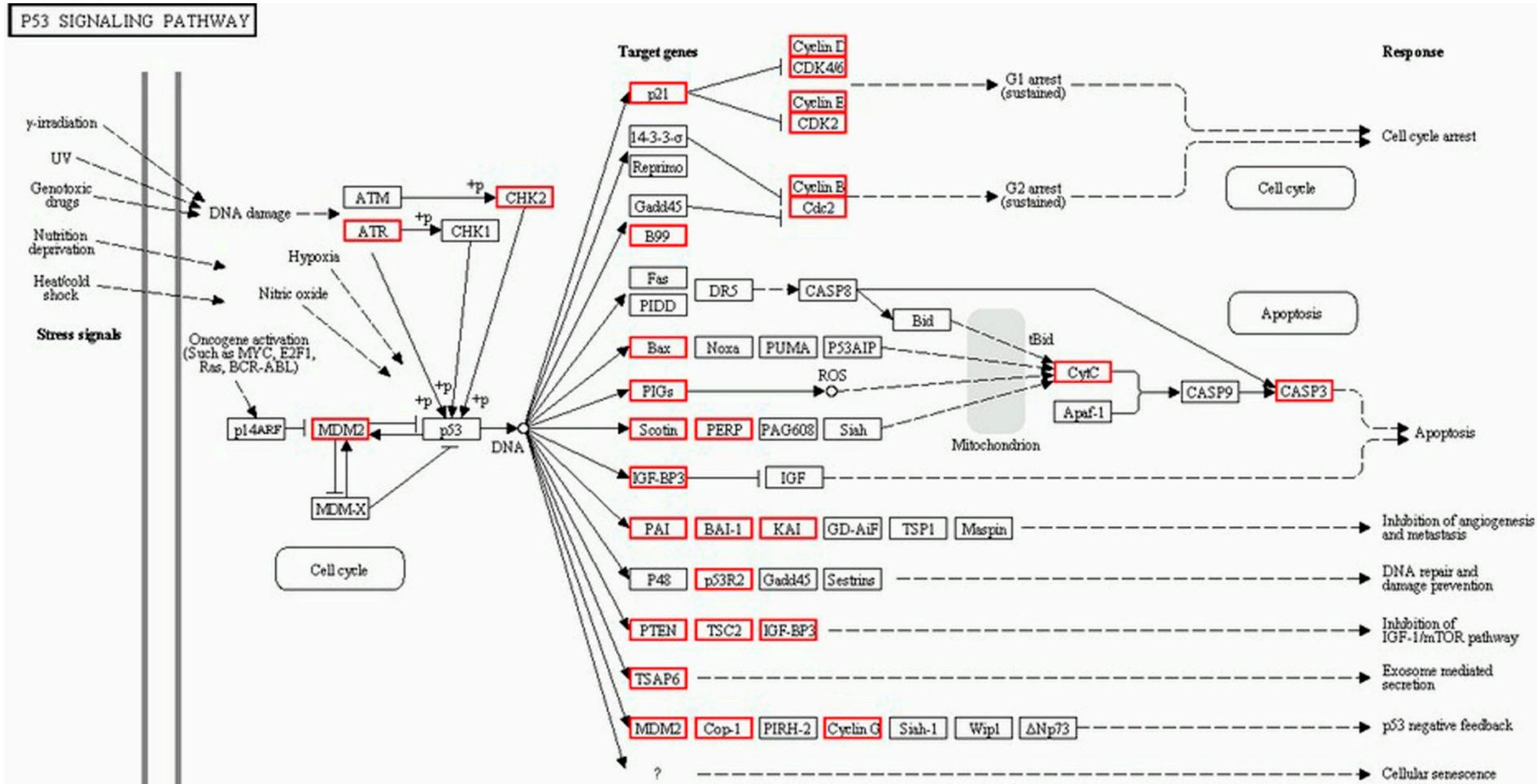


Figure S2. P53 signaling pathway for unigene annotated by KEGG. The unigenes with significant different expression are marked in red box. More information can refer to KEGG pathway: map04115 (Available online: http://www.genome.jp/dbget-bin/www_bget?pathway+map04115).