Caffeine-induced food-avoidance behavior is mediated by neuroendocrine signals in Caenorhabditis elegans

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High-dose caffeine uptake is a developmental stressor and causes food-avoidance behavior (aversion phenotype) in C. elegans, but its mode of action is largely unknown. In this study, we investigated the molecular basis of the caffeineinduced aversion behavior in C. elegans. We found that aversion phenotype induced by 30 mM caffeine was mediated by JNK/MAPK pathway, serotonergic and dopaminergic neuroendocrine signals. In this process, the dopaminergic signaling appears to be the major pathway because the reduced aversion behavior in cat-2 mutants and mutants of JNK/MAPK pathway genes was significantly recovered by pretreatment with dopamine. RNAi depletion of hsp-16.2, a cytosolic chaperone, and cyp-35A family reduced the aversion phenotype, which was further reduced in cat-2 mutants, suggesting that dopaminergic signal is indeed dominantly required for the caffeine-induced food aversion. Our findings suggest that aversion behavior is a defense mechanism for worms to survive under the high-dose caffeine conditions. [BMB Reports 2017; 50(1): 31-36]

INTRODUCTION

Caffeine is one of the most widely used drugs in the world as a part of consumed food. Caffeine uptake induces both beneficial and adverse effects on a body in a dosage- and developmental stage-dependent manner (1, 2). A moderate dose of caffeine uptake has a neuroprotective power and is also beneficial to physical performance in human as previously described (3). However, overconsumption of caffeine can be harmful to human health. Although the effects of caffeine can be lethal, the mode of its action in the body is still

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unclear.

In our previous study, we investigated effects of caffeine on Caenorhabditis elegans. C. elegans is an excellent model organism to analyze the effects of bioactive molecules on animal behaviors. We found that an overdose (30 mM) of caffeine treatment inhibited the early larval development and induced a stress response by up-regulating heat shock proteins, suggesting that high-dose caffeine is a developmental stressor (4). Interestingly, we also found that the food-aversion phenotype (food-avoidance behavior) was induced by the caffeine treatment, which was abolished by RNAi depletion of hsp-16.2 (4). HSP-16.2 is a small heat shock inducible cytosolic chaperone protein in C. elegans (5). The above finding suggests that the aversion phenotype is a defensive behavior for animal survival in response to an overdose of caffeine.

In this study, we further investigated whether the caffeineinduced aversion behavior is mediated by a neuronal or non-neuronal tissue response. Here we report that the food aversion phenotype observed after treatment with 30 mM of caffeine is mediated by neuroendocrine signals, in which dopamine serves as a dominant signaling and serotonin serves as a supplementary signaling. Further we show that these neuroendocrine signals are associated with cellular stress responses mediated by jnk-1, hsp-16.2, cyp-35A2, and cyp-35A4.

RESULTS

Caffeine induced food-avoidance behavior (aversion phenotype) in a dosage and developmental stage-dependent manner

We found that the level of food avoidance (% aversion) was caffeine-dose dependent. Among them, 30 mM caffeine treatment at the L1 larval stage caused 18.0 \pm 4.5% of food aversion (Fig. 1A). However, caffeine treatment at the later larval stages, L2 and L3 showed more reduced levels of aversion than L1, and caffeine treatment at the L4 larval stage did not cause significant food aversion phenotype (Fig. 1A). In our previous study, we reported that high-dose caffeine induced developmental growth retardation when treated at early larval stages (2). As in caffeine-induced growth retardation, we observed that early-stage larvae were more sensitive to caffeine for aversion behavior than later-stage larvae (Fig. 1A). We also reported that cvp-35A gene family,

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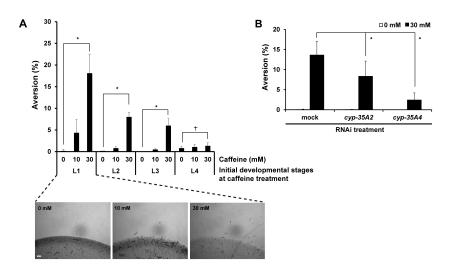


Fig. 1. Caffeine induced aversion phenotype in *C. elegans*. (A) Wild-type worms synchronized at the L1 larval stage (0 mM, n = 673; 10 mM, n = 535; 30 mM, n = 565), L2 larval stage (0 mM, n = 649; 10 mM, n = 531; 30 mM, n = 536), L3 larval stage (0 mM, n = 847; 10 mM, n = 348; 30 mM, n = 487), and L4 larval stage (0 mM, n = 249; 10 mM, n = 322; 30 mM, n = 306) were cultured on NGM agar plates containing 0, 10, or 30 mM caffeine, and their aversion phenotype was observed 24 h after caffeine treatment at 20° C. Representative images of worms treated with 0, 10 or 30 mM caffeine when they were at the L1 larval stage are shown under the graph. (B) The caffeine-induced aversion phenotype after RNAi depletion of *cyp-35A2* and *cyp-35A4* in wild type. After RNAi depletion, the synchronized L1 larvae were cultured on NGM agar plates containing 0 or 30 mM caffeine, and the aversion phenotype was observed at h after the caffeine treatment (mock RNAi, n = 763; *cyp-35A2* RNAi, n = 823; *cyp-35A4* RNAi, n = 654). In (A, B), values are shown as average % food aversion from three independent experiments. Bar, 200 μm. Error bars represent s.d. *P < 0.05. TP > 0.05.

the detoxification pathway genes were highly induced after 30 mM caffeine treatment, and that RNAi depletion of these genes showed partial recovery from growth retardation caused by caffeine treatment (2). Therefore, we examined whether RNAi depletion of *cyp-35* genes also affects the aversion phenotype. In fact, we found that RNAi of *cyp-35A2* and *cyp-35A4* caused lower levels of aversion than mock RNAi control (Fig. 1B). This result suggests that growth retardation induced by 30 mM caffeine was affected by food aversion.

Caffeine-induced food-avoidance behavior is mediated by INK/MAPK pathway

To further investigate the correlation between food avoidance and growth retardation, we next examined MAPK signaling pathway, which is required for microbial aversion (6), and RNAi depletion of MAPK pathway genes was previously reported to cause growth retardation (wormbase.org). We examined food aversion behavior under 0 or 30 mM caffeine condition, after RNAi depletion of genes involved in the MAPK pathway. We found that the aversion phenotype was suppressed when tir-1, sek-1, pmk-1, mlk-1, mek-1, vhp-1, kgb-1, jkk-1 and jnk-1 genes were depleted under 30 mM caffeine conditions compared to mock RNAi control (Fig. 2A: mock RNAi, $11.0 \pm 0.4\%$; tir-1 RNAi, $4.4 \pm 0.4\%$; sek-1 RNAi, $6.3 \pm 1.1\%$; pmk-1 RNAi, $5.6 \pm 1.4\%$; mlk-1 RNAi, $7.5 \pm 1.2\%$; mek-1 RNAi, $3.5 \pm 1.6\%$; vhp-1 RNAi, $5.1 \pm 0.6\%$; kbg-1 RNAi, $3.9 \pm 0.3\%$; jkk-1 RNAi, $2.9 \pm 1.0\%$;

jnk-1 RNAi, 3.8 ± 0.4%). Among them, RNAi depletion of tir-1, mek-1, kgb-1, ikk-1, and ink-1 reduced the aversion phenotype more effectively than the other genes (Fig. 2A). We also tested caffeine-induced aversion phenotype using genetic loss-of-function mutants of mlk-1, mek-1, kgb-1, and jnk-1 genes. The results obtained using the loss-of-function mutants were similar to those obtained by RNAi depletion (Fig. 3A). We judge that efficiency of RNAi depletion in this study was high enough to evaluate deficiency of tested gene activities. These results suggest that activation of MAPK pathway genes is collectively required for caffeine-induced food aversion. Among the MAPK pathway genes, JNK kinase pathway genes appear to have a major role for the control of caffeine-induced food aversion behavior. Furthermore, the reduced level of food avoidance we observed in loss-of-function mutants of MAPK pathway genes was unlikely related to growth retardation.

Caffeine-induced food-avoidance behavior is mediated by both serotonin and dopamine signals, in which dopamine has a major role

To investigate a possible correlation between caffeine-induced food-avoidance behavior and a neuronal signaling, we observed the aversion phenotype after caffeine treatment in dopamine-synthesis defective *cat-2*, serotonin-synthesis defective *tph-1*, and both serotonin- and dopamine-synthesis defective *bas-1* mutants. We found that the aversion phenotype was reduced significantly in *cat-2*, *tph-1*, and more significantly in

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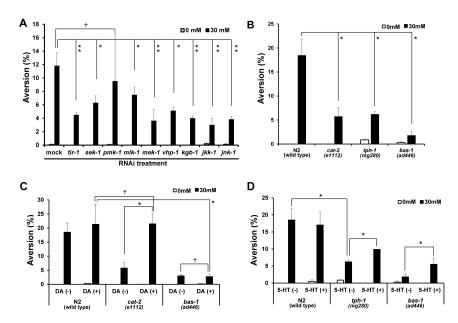


Fig. 2. The JNK/MAPK pathway and neuronal signaling were required for caffeine-induced aversion phenotype. (A) The JNK/MAPK pathway mediates caffeine-induced aversion phenotype demonstrated by RNAi depletion of *tir-1*, *sek-1*, *pmk-1*, *mlk-1*, *mek-1*, *vhp-1*, *kgb-1*, *jkk-1* and *jnk-1* in N2 worms. After RNAi depletion, synchronized L1 larvae were cultured on NGM agar plates containing 0 or 30 mM caffeine, and the aversion phenotype was observed at 24 h after caffeine treatment (mock RNAi, n = 894; *tir-1* RNAi, n = 821; *sek-1* RNAi, n = 797; *pmk-1* RNAi, n = 877; *mlk-1* RNAi, n = 676; *mek-1* RNAi, n = 874; *vhp-1* RNAi, n = 809; *kgb-1* RNAi, n = 781; *jkk-1* RNAi, n = 1020; *jnk-1* RNAi, n = 898). (B) Dopamine-deficient *cat-2*, serotonin-deficient *tph-1*, serotonin and dopamine-deficient *bas-1* mutants showed defects in food aversion at 30 mM caffeine. *cat-2*, *tph-1* and *bas-1* mutant worms synchronized at the L1 larval stage were cultured on NGM agar plates containing 0 or 30 mM caffeine, and the aversion phenotype was observed at 24 h after the caffeine treatment (N2, n = 589; *cat-2*, n = 380; *tph-1*, n = 966; *bas-1*, n = 623). (C, D) N2, *cat-2*, *tph-1* and *bas-1* worms synchronized at the L1 larval stage were pre-treated for 10 min with dopamine (DA) or serotonin (5-HT), cultured on NGM agar plates containing 0 or 30 mM caffeine, and the aversion phenotype was monitored at 24 h after caffeine treatment (N2, n = 793; *cat-2*, n = 511; *tph-1*, n = 763; *bas-1*, n = 524). Values are shown as average % food aversion. Error bars represent s.d. *P < 0.05. **P < 0.005. *P > 0.05.

bas-1 mutants when worms were exposed to 30 mM caffeine (Fig. 2B: N2, 18.4 \pm 3.2%; cat-2, 5.8 \pm 1.8%; tph-1, 6.2 \pm 0.6%; bas-1, 1.8 \pm 0.8%). These results suggest that both dopamine and serotonergic signaling pathways are required for the food avoidance after caffeine treatment. The defect of food-avoidance behavior in cat-2 mutants, but not bas-1 mutants, was sufficiently rescued by acute-dopamine treatment (Fig. 2C). However, the caffeine-induced aversion phenotype was not sufficiently rescued by acute-serotonin treatment in neither tph-1 nor bas-1 mutants (Fig. 2D). These results suggest that although both dopaminergic and serotonergic signals are required for the aversion phenotype, dopaminergic signaling is the major pathway for controlling the aversion phenotype in the presence of high-dose caffeine. To confirm that these signaling pathways are active at the L1 larval stage, we monitored expression of dat-1::GFP dopaminergic transgene and tph-1::GFP serotonergic transgene in the L1 stage transgenic worms. We indeed confirmed that both dat-1::GFP and tph-1::GFP were highly expressed at the L1 larval stage (Fig. S1).

JNK kinase signaling possibly functions upstream of dopaminergic neuronal signaling to control caffeine-induced food-avoidance behavior

To test whether the reduced caffeine-induced aversion phenotype in mutants of JNK/MAPK pathway genes are restored by acute-dopamine and serotonin treatment, we measured the caffeine-induced aversion phenotype in wild-type and mutants of JNK/MAPK pathway genes with or without DA or 5-HT treatment. We hypothesized that if JNK/MAPK signaling pathway is acting upstream of dopaminergic or serotonergic signaling the reduced aversion phenotype in mutants of JNK/MAPK signaling pathway genes is rescued by either dopamine or serotonin treatment. We found that aversion phenotype was rescued by neither DA nor 5-HT treatment in kgb-1 mutant and mek-1 mutants while the aversion phenotype was partially rescued by DA, but not noticeably by 5-HT, in mlk-1 and jnk-1 mutants (Fig 3A: mlk-1: control, 3.5 \pm 0.3%, with DA, 5.5 \pm 0.8%; jnk-1: control, 0%, with DA, 6.8 \pm 1.1%). Remarkably, the caffeine-induced aversion phenotype that was almost completely suppressed in jnk-1

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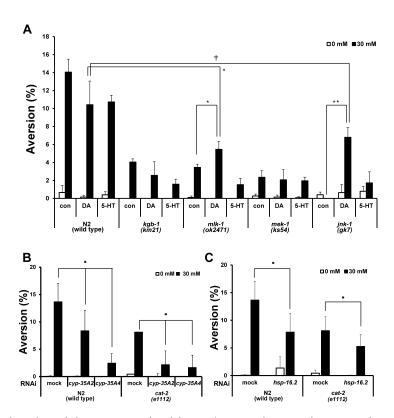


Fig. 3. Caffeine-induced food-avoidance behavior was mediated by JNK/MAPK and neuroendocrine signaling. (A) Aversion phenotype in JNK/MAPK mutants (kgb-1, mlk-1, mek-1, jnk-1) after 0 mM or 30 mM caffeine treatment with or without acute-dopamine or serotonin treatment. Worms synchronized at the L1 larval stage were cultured on NGM agar plates containing 0 or 30 mM caffeine, and the aversion phenotype was observed at 24 h after the caffeine treatment (N2, n = 4099; kgb-1, n = 1559; mlk-1, n = 1638; mek-1, n = 1704; jnk-1, n = 929). Values are shown as average % food aversion \pm s.d. (B) Aversion phenotype after RNAi depletion of cyp-35A2, cyp-35A4 in N2 and cat-2 mutants. After the RNAi depletion, the synchronized L1 larvae were cultured on NGM agar plates containing 0 or 30 mM caffeine, and the aversion phenotype was observed at 24 h after caffeine treatment (individual RNAi, n > 654). (C) Aversion neon NGM agar plates containing 0 or 30 mM caffeine, and the aversion phenotype was observed at 24 h after caffeine treatment (individual RNAi, n > 723). Values are shown as average % food aversion. Error bars represent s.d. *P < 0.05.

mutants was significantly recovered by dopamine, but not by serotonin, treatment (Fig. 3A). These results indicate that JNK pathway has a critical role for controlling caffeine-induced aversion phenotype, and that this pathway is possibly acting upstream of dopaminergic signaling to control caffeine-induced food-avoidance behavior.

We previously reported that expression of *cyp-35A* family genes and *hsp-16.2*, a cytosolic chaperone, was induced by caffeine treatment and that RNAi depletion of *hsp-16.2* reduced the caffeine-induced aversion phenotype (4). In this study, we also showed that expression of *cyp-35::GFP* transgene was induced by caffeine treatment in the L1 larval stage of transgenic worms (Fig. S2). In addition, we showed that RNAi depletion of *cyp-35A2* and *cyp-35A4* also reduced the aversion phenotype (Fig. 1B). These findings suggest that high expression of *cyp-35A* family and *hsp-16.2* genes induces the aversion phenotype in response to a toxic environment

induced by high-dose caffeine. We further examined whether the reduced level of food aversion after RNAi depletion of *cyp-35A2*, *cyp-35A4*, or *hsp-16.2* is further reduced in *cat-2* mutants. We found that the reduction of food aversion behavior after RNAi depletion of *cyp-35A2*, *cyp-35A4*, and *hsp-16.2* was more substantial in 30 mM-caffeine treated *cat-2* mutants than in 30 mM-caffeine treated wild-type N2 animals (Fig. 3B, 3C). These findings suggest two possibilities: the caffeine-induced aversion phenotype which is enhanced by expression of *cyp-35A2*, *cyp-35A4*, and *hsp-16.2* genes, requires dopaminergic signal pathway; or the stress-responding *cyp-35A2*, *cyp-35A4*, and *hsp-16.2* genes and the dopaminergic signal pathway are acting in parallel, so that when both pathways were depleted, the defect in food aversion was observed additively.

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DISCUSSION

A food-avoidance behavior is considered as one of defense mechanisms from harsh environments. Our previous study revealed that high-dose caffeine treatment induces HSP-16.2, a small heat shock inducible cytosolic chaperone protein and RNAi depletion of hsp-16.2 abolished the food-aversion behavior (4). This result suggests that HSP-16.2 is required for the food-avoidance behavior and that this behavior is a program for survival. HSP-16.2 was expressed under stressful conditions, and the expression of HSP-16.2 was associated with nuclear translocation of DAF-16, which is a transcriptional activator of HSP-16.2 (7). This finding suggests that survival pathway via HSP-16.2 and the aversion behavior is possibly coupled with life-span control of worms. In this study we also revealed that RNAi of cyp-35A2 (cytochrome P450) and cyp-35A4, which are highly induced by caffeine treatment (2), reduced the food-avoidance behavior, suggesting that a detoxification pathway mediated by cyp-35A2 and cyp-35A4 is also required for inducing the food-avoidance behavior after caffeine treatment.

Based on these findings, we further investigated underlying molecular mechanism that controls the caffeine-induced food-avoidance behavior. We hypothesized that high-dose caffeine uptake triggers changes in normal cellular processes and stimulates the food-avoidance behavior. It was previously reported that a disruption of cellular activities induced a food-avoidance behavior which was controlled by serotonergic and JNK/MAPK kinase signaling pathways (6). Similarly, our results revealed that JNK/MAPK pathway, and serotonergic and dopaminergic neuroendocrine signals are required for food-avoidance behavior induced by caffeine treatment, in which dopaminergic signaling seems to be the major pathway to respond to caffeine treatment. Lack of neuroendocrine signals abolished or significantly suppressed the food-avoidance behavior, which was substantially recovered by providing the signals exogenously. These results indicate that production of the neuroendocrine signals is essential for the food-avoidance behavior induced by caffeine treatment. In addition, it was reported that eating behaviors involved dopaminergic neural circuitry which affected the food- and drug-related cues in the brain (8, 9). This idea is supported by the findings that well conserved dopamine biosynthesis and transport pathways are required to maintain homeostasis in humans, Drosophila and C. elegans (10). Recently, it was reported that caffeine suppresses nighttime sleep through dopamine signaling in Drosophila (11). Nevertheless, thus far, transcription factors, which are involved in dopamine signaling after caffeine treatment, remain to be determined. Taken together, high-dose caffeine treatment to C. elegans has adverse effects on the development, and induces a food-avoidance behavior for the survival. For this response the JNK/MAPK pathway is required. In addition, the neuroendocrine signals, serotonin and dopamine are also required. Caffeine has various beneficial effects on the body when appropriate doses are administered. However, overdose of caffeine is a stressor to the body and induces a survival pathway, which results in changes of various cellular activities. Especially, caffeine consumption at early developmental stages can be lethal through inducing growth retardation and severe food avoidance.

MATERIALS AND METHODS

C. elegans strains and maintenance

All strains were maintained at either 15°C or 20°C on Nematode Growth Medium (NGM) agar plates seeded with *Escherichia coli* OP50 as previously described (12). The following strains were used: CB1112: cat-2(e1112) *II*, MT7988: bas-1(ad446) *III*, CY573: vbls5[cyp-35B1p::GFP + gcy-7p::GFP]?, FK171: mek-1(ks54) X, KU21: kgb-1(km21) *IV*, MT15434: tph-1(mg280) *II*, RB1908: mlk-1(ok2471) V, VC8: jnk-1(gk7) *IV*, BZ555: egls1 [dat-1p::GFP]?, GR1366: mgls42 [tph-1::GFP + rol-6(su1006)]?. N2 was used as a wild type. For caffeine treatment, the appropriate amount of caffeine (Sigma, St. Louis, MO, USA) was included in the NGM agar plates before autoclaving to achieve final concentration of 30 mM.

RNA interference (RNAi) analysis

RNAi analysis was performed using the "RNAi-by-soaking" method as described previously (13). Wild-type worms were synchronized at the L1 larval stage in absence of food, and soaked for 24 h in each of the double-strand RNA solutions transcribed *in vitro* from respective cDNA templates, which were PCR-amplified from corresponding L4440 feeding RNAi clones using T7 primer, 5'-GTAATACGACTCACTATAGGCC-3', and L4440 T7 primer, 5'-ATTAATACGACTCACTATAG GGA-3'. Worms were then transferred to either NGM plates containing caffeine or control NGM plates at 20°C. Aversion phenotype was observed at 24 h after the transfer of the worms. Efficiency of RNAi was examined by final phenotype of individual test genes.

Food-avoidance behavior (aversion phenotype) assay

To check for food avoidance after caffeine treatment, synchronized L1 worms were transferred to *E. coli* OP50-seeded NGM plates with or without 30 mM caffeine. Food avoidance of worms was calculated by dividing the number of worms that stayed outside of food area with the total number of worms on the plate as described previously (6). Each RNAi or drug treatment was conducted in triplicate to evaluate statistical significance of the data, unless otherwise indicated.

Acute dopamine and serotonin treatment

Dopamine (Sigma) and serotonin (Sigma) solutions were prepared in M9 buffer (3.0 g KH_2PO_4 , 6.0 g Na_2HPO_4 , 0.5 g NaCl, 1.0 g NH_4Cl in 1 L H_2O). Synchronized L1 worms were treated with dopamine (3 mg/ml, 15 mM) or serotonin (3

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mg/ml, 7.4 mM) for 10 min, based on previous reports (14, 15). The optimal conditions for acute dopamine and serotonin treatments were determined by finding the concentration for the most effective rescue from the food aversion defect (Fig. S3). After the drug treatments, worms were transferred to either caffeine-containing or control NGM plates at 20°C. Control worms were treated with M9 buffer.

Statistical analysis

All experiments were repeated more than three times for statistical evaluation of data. P values were calculated by Student's t-test. We considered P < 0.05 as being significant.

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CONFLICTS OF INTEREST

The authors have no conflicting financial interests.

REFERENCES

- 1. Sutphin GL, Bishop E, Yanos ME, Moller RM and Kaeberlein M (2012) Caffeine extends life span, improves healthspan, and delays age-associated pathology in *Caenorhabditis elegans*. Longev Healthspan 1, 1-9
- Min H, Kawasaki I, Gong J and Shim YH (2015) Caffeine induces high expression of cyp-35A family genes and inhibits the early larval development in *Caenorhabditis* elegans. Mol Cells 38, 236-242
- 3. Cabrera C, Artacho R and Giménez R (2006) Beneficial

- effects of green tea—a review. J Am Coll Nutr 25, 79-99
- Al-Amin M, Kawasaki I, Gong J and Shim YH (2016) Caffeine induces the stress response and up-regulates heat shock proteins in *Caenorhabditis elegans*. Mol Cells 39, 164-168
- Fonte V, Kipp DR, Yerg J et al (2008) Suppression of in vivo beta-amyloid peptide toxicity by overexpression of the HSP-16.2 small chaperon protein. J Biol Chem 283, 784-791
- Melo JA and Ruvkun G (2012) Inactivation of conserved C. elegans genes engages pathogen- and xenobioticassociated defenses. Cell 149, 452-466
- Hartwig K, Heidler T, Moch J, Daniel H and Wenzel U (2009) Feeding a ROS-generator to Caenorhabditis elegans leads to increased expression of small heat shock protein HSP-16.2 and hormesis. Genes Nutr 4. 59-67
- 8. Baik JH (2013) Dopamine signaling in food addiction: role of dopamine D2 receptors. BMB Rep 46, 519-526
- Min S, Chung J (2016) Identification of a neural pathway governing satiety in Drosophila. BMB Rep 49, 137-138
- Caldwell KA, Shu Y, Roberts NB, Caldwell GA and O'Donnell JM (2013) Invertebrate models of dystonia. Curr. Neuropharmacol 11, 16-29
- Nall AH, Shakhmantsir I, Cichewicz K, Birman S, Hirsh J and Sehgal A (2016) Caffeine promotes wakefulness via dopamine signaling in *Drosophila*. Sci Rep 12, 20938
- 12. Brenner S (1974) The genetics of Caenorhabditis elegans. Genetics 77, 71-94
- Maeda I, Kohara Y, Yamamoto M and Sugimoto A (2001) Large-scale analysis of gene function in *Caenorhabidits* elegans by high-throughput RNAi. Curr Biol 11, 171-176
- Sharfer WR and Kentyon CJ (1995) A calcium-channel homologue required for adaptation to dopamine and serotonin in Caenorhabdits elegans. Nature 375, 73-78
- Benedetto A, Au C, Avilla DS, Milatovic D and Aschner M (2010) Extracellular dopamine potentiates Mn-induced oxidative stress, lifespan reduction, and dopaminergic neurodegeneration in a BLI-3-dependent manner in Caenorhabditis elegans. PLos Genet 6, e1001084

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