Akt signaling-associated metabolic effects of dietary gold nanoparticles in Drosophila

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Gold nanoparticles (AuNPs) are often used as vehicles to deliver drugs or biomolecules, due to their mild effect on cell survival and proliferation. However, little is known about their effect on cellular metabolism. Here we examine the in vivo effect of AuNPs on metabolism using Drosophila as a model. Drosophila and vertebrates possess similar basic metabolic functions, and a highly conserved PI3K/Akt/mTOR signaling pathway plays a central role in the regulation of energy metabolism in both organisms. We show that dietary AuNPs enter the fat body, a key metabolic tissue in Drosophila larvae. Significantly, larvae fed with AuNP show increased lipid levels without triggering stress responses. In addition, activities of the PI3K/Akt/mTOR signaling pathway and fatty acids synthesis are increased in these larvae. This study thus reveals a novel function of AuNPs in influencing animal metabolism and suggests its potential therapeutic applications for metabolic disorders.

Gold nanoparticles (AuNPs) possess many attractive optical and electronic properties that have proven to be of high utility in biomedical applications1. Biomolecular and cellular interactions of AuNPs are particularly interesting and versatile, which have led to numerous AuNPs-based diagnostic, imaging and therapeutic tools2–7. More recently, AuNPs were found to be cell permeable8,9 in addition to their well known low cytotoxicity and high biocompatibility10,11. Consequently, several groups including ours have shown that AuNPs can serve as nanoscale vehicles for drug delivery2,8,12. Despite the well-documented progress, direct dietary use of AuNPs in living organisms and possible consequences on cellular metabolism are little explored.

Living organisms can sense nutritional conditions to coordinate cellular metabolism. Interestingly, Drosophila and vertebrates are very similar in their basic metabolic functions, i.e., sugar, lipid and amino acid homeostasis13,14. In both organisms, a highly conserved phosphatidylinositol 3-kinase (PI3K) pathway has a pivotal role in nutrition uptake and storage15–18. Aberrant PI3K signaling in human results in metabolic disorders and often occurs in type-2 diabetes or cancers19–21. As a response to dietary nutrient, the upstream signaling molecule PI3K is activated, leading to re-localization and activation of Akt19. The Akt kinase has two major metabolic target molecules, FOXO and mTOR22,23. Phosphorylation by Akt inactivates FOXO, a transcription factor that controls expression of key enzymes of lipogenesis including Fatty Acid Synthase (FAS) and Acetyl-CoA Carboxylase (ACC)24.

In this work, we aimed to explore whether dietary AuNPs affect cellular metabolism and used Drosophila larvae as an in vivo model. We fed larvae with several concentrations of AuNPs and found that they were absorbed by the fat body, a key metabolic tissue in Drosophila larvae. In these larvae, stress responses including expression of heat shock genes, activation of the JNK pathway or induction of autophagy were not induced compared to those in control larvae. Despite their little effects on growth or stress responses, lipid levels were increased in larvae fed with AuNPs. By examining metabolic pathways that control lipid anabolism, we found that dietary AuNPs increased activities of the PI3K/Akt/mTOR signaling pathway and fatty acids synthesis in these larvae. In addition, inhibition of mTOR abolished these effects of AuNPs, suggesting that dietary AuNPs could affect Drosophila lipid metabolism by promoting the PI3K/Akt/mTOR signaling pathway.
Figure 1 | Uptake of AuNPs by fly larva tissues. (A) Hydrated diameter of AuNPs (23 nm) were measured by dynamic light scattering (DLS). (B) 15 nm AuNPs characterized by transmission electron microscopy (TEM). (C) Uptake of AuNPs by fly larva tissues, as measured by quantitative inductively coupled plasma mass spectrometry (ICP-MS). This bar graph shows representative data of four individual sample analyses.
Results

AuNPs of 15 nm (Fig. 1A,B, supplementary Fig. S1) were synthesized and formulated in the diet. *Drosophila* larvae fed with AuNPs were dissected to determine the distribution of AuNPs among different tissues. Quantitative analysis with inductively coupled plasma mass spectrometry (ICP-MS) detected localization of AuNPs in the fat body (equivalent to mammalian white adipose tissue and liver), but not in brain, salivary gland or imaginal discs (Fig. 1C). Further analysis by transmission electron microscopy (TEM) showed that AuNPs localized to lipid droplets of fat body cells (supplementary Fig. S4C, D).

We first examined whether cellular stress was up-regulated by dietary AuNPs. Cellular stress could increase expression of heat shock proteins, activate the c-Jun N-terminal Kinase (JNK) pathway, and trigger autophagy. In larvae fed with 0.5 or 2 nM AuNPs, expression of *hsp83* or *hsp70* was unchanged (Fig. 2A, B, supplementary Fig. S2A, B); activity of the JNK pathway was unchanged or slightly reduced, as measured by expression of its downstream gene *puckered* (Fig. 2C, supplementary Fig. S2C); autophagy was also unchanged or slightly reduced, as measured by lysotracker staining (Fig. 2D, supplementary Fig. S2D). However, 5 nM dietary AuNPs did slightly increase expression of *hsp70* and *puckered*, but not that of *hsp83* (supplementary Fig. S2A-C). As a conclusion, up to 2 nM dietary AuNPs did not trigger cellular stress responses in larvae, although higher concentration of AuNPs might have a mild effect. For this reason we used 2 nM AuNPs for the rest of experiments.

The localization of AuNPs to lipid droplets of fat body cells triggered our interest to examine its possible consequences on lipid metabolism. *Drosophila* and mammals regulate lipid metabolism via the highly conserved SREBP pathway, which activates expression of key enzymes of fatty acid synthesis including FAS and ACC. In larvae fed with AuNPs, triacylglycerol (TAG, the main stored form of lipid) levels were slightly higher (~10%, Fig. 3A). In addition, expression of FAS and ACC were increased, suggesting that enhanced fatty acid synthesis was responsible for the change of lipid levels (Fig. 3B,C). The fly food provided excessive nutrition, and lipid anabolism in larvae already reached a high level. Thus we also fed larvae in the condition of calorie restriction (CR), where lipid anabolism was limited. Under this background, larvae fed with AuNPs showed even higher TAG levels (~20%) compared to the CR control (Fig. 4A).

The SREBP pathway is a downstream effector of the PI3K/Akt/mTOR pathway. To test whether this signaling pathway was activated by dietary AuNPs, we examine the activity of PI3K, Akt, and expression of their downstream genes. In standard diet, the activity of

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Akt in larvae fed with AuNPs was comparable to that in control, consistent with the result that AuNPs only had a mild effect on lipid metabolism under this condition. In the presence of CR, dietary AuNPs significantly enhanced Akt activity (Fig. 4B). We also measured membrane bound tGPH in the fat body, whose levels reflected the activity of PI3K. Larvae fed with AuNPs showed higher activity of PI3K in the fat body than the control (Fig. 4C,D). These results confirmed that the upstream signaling molecules (PI3K and Akt) were activated by dietary AuNPs. Activated Akt can enhance expression of FAS and ACC via the mTOR/SREBP cascade, and repress expression of InR and 4EBP via FOXO (supplementary Fig. S3A), suggesting that rapamycin blocked the stimulation of fatty acid synthesis by AuNPs. Therefore, dietary AuNPs possibly affect *Drosophila* fat metabolism via the PI3K/Akt/mTOR pathway (Fig 5E).

**Discussion**

There are several ways that AuNPs can affect PI3K/Akt signaling. PI3K/Akt signaling generally senses nutritional conditions to coordinate metabolism. Although larvae fed with or without AuNPs consume the same amount of food (supplementary Fig. S4A), it is possible that AuNPs may promote nutrient uptake at cellular levels. To test this possibility, we examined total sugar levels as sugar metabolism provided acetyl-CoA precursors (pyruvate and citrate) for fatty acid synthesis. We found that larvae fed with AuNPs had higher

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**Figure 3 | Dietary AuNPs increases lipid levels and fatty acids synthesis in larvae.** (A) Normalized TAG levels in larvae fed with AuNPs were 10% higher than that in control. *, *P* < 0.05. (B), (C) Expression of key enzymes for fatty acids synthesis FAS (B) and ACC (C) were increased by dietary AuNPs. **, *P* < 0.01.
sugar levels than the control in CR food, while this difference was insignificant in standard food (supplementary Fig. S4B). These data suggest that AuNPs enhance nutrient uptake in CR food and may at least partially explain our observations. Alternatively, AuNPs may affect activities of signaling molecules of the PI3K/Akt pathway. Activations of both PI3K and Akt occur on the cell membrane. It is also likely that endocytosis of AuNPs facilitates recruitment of PI3K and Akt on the cell membrane and promotes their activations.

Consistent with this model, higher levels of tGPH are detected at the cell membrane (Fig. 4C,D), reflecting that higher levels of PI3K are localized to the cell membrane. Therefore this mechanism may also contribute to the effect of AuNPs.

AuNPs have shown great promise in biomedicine, e.g. they may serve as vehicles to deliver chemical drugs or biomolecules. While it is generally considered that AuNPs are biocompatible and non-toxic for cell growth, other biochemical activities of AuNPs are little

Figure 4 | Calorie restriction enhances effects of AuNPs on fat metabolism and Akt signaling. (A) Dietary AuNPs increased larva TAG levels by 10% in standard food, and by 20% under the condition of calorie restriction (CR). *, P<0.05. (B) Under CR, larvae fed with AuNPs showed increased Akt activity compared to the control (lane 4 vs. lane 3). (C) (D) Under CR, larvae fed with AuNPs (D) had higher PI3K activity in the fat body compared to the control (C).
Figure 5 | The effect of AuNPs on larva fat metabolism is dependent on Akt/mTOR signaling. (A) (B) In standard diet (A) and CR (B), the increase in larva TAG levels by dietary AuNPs was abolished when the Akt/mTOR signaling pathway was blocked by rapamycin. *, P<0.05. ns, not significant. (C) (D) Increased expression of FAS (C) and ACC (D) by dietary AuNPs was abolished when the Akt/mTOR signaling pathway was blocked. **, P<0.01. ns, not significant. (E) A diagram for the effect of dietary AuNPs on larva fat metabolism.
studied. In this work, we focus on the metabolic effects of dietary AuNPs in Drosophila larvae. We find that dietary AuNPs promote the Akt signaling pathway and alter lipid metabolism, an unexpected in vivo effect of nanomaterials on cellular metabolism. Given that metabolic diseases such as diabetes have become worldwide epidemic, our new observations shed new light on using AuNPs, either by themselves or coupled with other functional drugs, as nanomedicine against metabolic disorders.

Methods
Preparation and characterization of AuNPs. AuNPs of 15 nm were synthesized by reduction of HAuCl4, as previously reported32, and their sizes were characterized with transmission electron microscopy and dynamic light scattering.

Fly strain and culture. yw D. melanogaster was used in this study. Flies were raised in cages. 0–1 hour embryos were collected from fruit juice agar plates, counted and distributed on SY food (10% sucrose, 10% yeast, 1.5% agar)33 with or without AuNPs. AuNPs were coated with 2 ug/ml BSA for 10 minutes and mixed with the food before solidification. Same amount of BSA is also included in the control food. Larvae were harvested 108 hours later for analysis. To control the fly food contained 25% nutrients of standard food, and larvae were harvested 156 hours after egg laying. To prepare larval tissues for detection of AuNPs distribution, larvae fed with AuNPs were transferred to control food for 8 hours, allowing evacuation of AuNP food. Larvae were then transferred to 6 cm dishes and washed 10 times in 1×PBST before dissection. Dissected tissues were transferred to microcentrifuge tubes and washed 10 times in 1×PBST.

Measurement of mRNA, protein and lipid levels. Total RNAs were extracted from larvae with TRIzol reagent (Invitrogen). The first-strand cDNAs were generated by iScriptTM cDNA Synthesis Kit (Bio-Rad). Quantitative RT-PCR assays were performed with SYBR Green PCR Master Mix and the Applied Biosystems ABI Prism 7000 Sequence Detection System (Applied Biosystems; Foster City, CA). The average expression levels of actin, tubulin and TRP were used as internal controls for normalization of mRNA levels. For western blotting, anti-phospho-Akt and anti-Tubulin antibodies (Cell Signaling) were used. For measurement of TAG, larvae were suspended in 400 ul chilled lysis buffer (0.01 mM KH2PO4, 1 mM EDTA, protease inhibitors cocktail, 0.05% Tween 20), lysed by Tissueyzer II (Qiagen, set at 30 Hz for 3 min), and heated at 70°C for 5 min. Cleared supernatant was used to measure TAG with TAG determination kits (Sigma,TR0100). Lipid levels were normalized to total protein levels.

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Author contributions
CF and HS designed the project, BW, NC and YW performed experiments, JL, LS, JW, QH and CF wrote the main manuscript text and BW prepared figures 1-5. All authors reviewed the manuscript.

Additional information
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