



# Inhibition of HIPK3 by AST487 Ameliorates Mutant HTT-Induced Neurotoxicity and Apoptosis *via* Enhanced Autophagy

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## Dear Editor,

Accumulation of misfolded and aggregation-prone proteins is the common hallmark of many neurodegenerative disorders, and lowering the levels of these proteins may provide promising strategies for the potential treatment of some of these diseases [1, 2]. Among them, Huntington's disease (HD) is a monogenic disease caused by mutation of the *HTT* (huntingtin) gene [3], which encodes the mutant HTT protein (mHTT) with an expanded polyglutamine tract (polyQ). The monogenic nature of HD provides high confidence for the causal relationship between mHTT and disease pathology, making HD suitable for testing the potential beneficial effects of reducing disease-causing proteins.

The gain of toxic function of mHTT is the major cause of HD, and lowering mHTT protein levels has been shown to effectively alleviate its toxicity. mHTT is known to be degraded by autophagy [4], and lowering its protein levels may ameliorate its downstream toxicity and treat HD [5]. Much evidence supports the role of apoptosis in HD; mHTT can induce apoptosis and then promote neuronal

death [6], so the modulation of autophagy and apoptosis may be a potential means of reducing neuronal death. Turning off the transgene in a transgenic HD mouse model that expresses inducible mHTT N-terminal fragments reverses the neuropathology and motor deficits [7]. Other genetic strategies such as delivering short-hairpin RNAs, small interfering RNAs, antisense oligonucleotides [8], and CRISPR/Cas9-mediated genome editing [9] can attenuate the neuropathology in HD mouse models. We identified the kinase HIPK3 (homeodomain interacting protein kinase 3) as a novel modulator of mHTT protein levels from an unbiased genetic screen [10]. Knocking-down *HIPK3* or loss of its kinase activity by mutagenesis lowers mHTT levels *via* autophagy [10]. Meanwhile, whether HIPK3 contributes to neurotoxicity and whether inhibiting its kinase function rescues HD-relevant phenotypes remained unknown.

In this study, we elucidated the potential role of HIPK3 in HD pathogenesis and the possible therapeutic effects of the small-molecule HIPK3 inhibitor AST487 in HD models, including HD mouse primary neurons, human induced pluripotent stem cell (iPSC)-derived neurons, and HD fly models.

We first investigated the potential pathological role of HIPK3 in HD. mHTT-induced cytotoxicity under stressed culture conditions occurred in neurons from a knock-in mouse model (HD<sup>Q7/Q140</sup>, Q indicates polyglutamine) expressing endogenous mHTT protein from its original locus [11]. To induced apoptosis phenotypes, neurons were cultured in a medium without supplements N2 and B27, and their shrinkage was measured by the Tuj1<sup>+</sup> (neuron-specific class III beta-tubulin) area in each neuron [12]. Knocking down *HIPK3* rescued the apoptotic phenotype (Fig. 1A). To further confirm this in a human neuronal model, we cultured neurons derived from human iPSCs

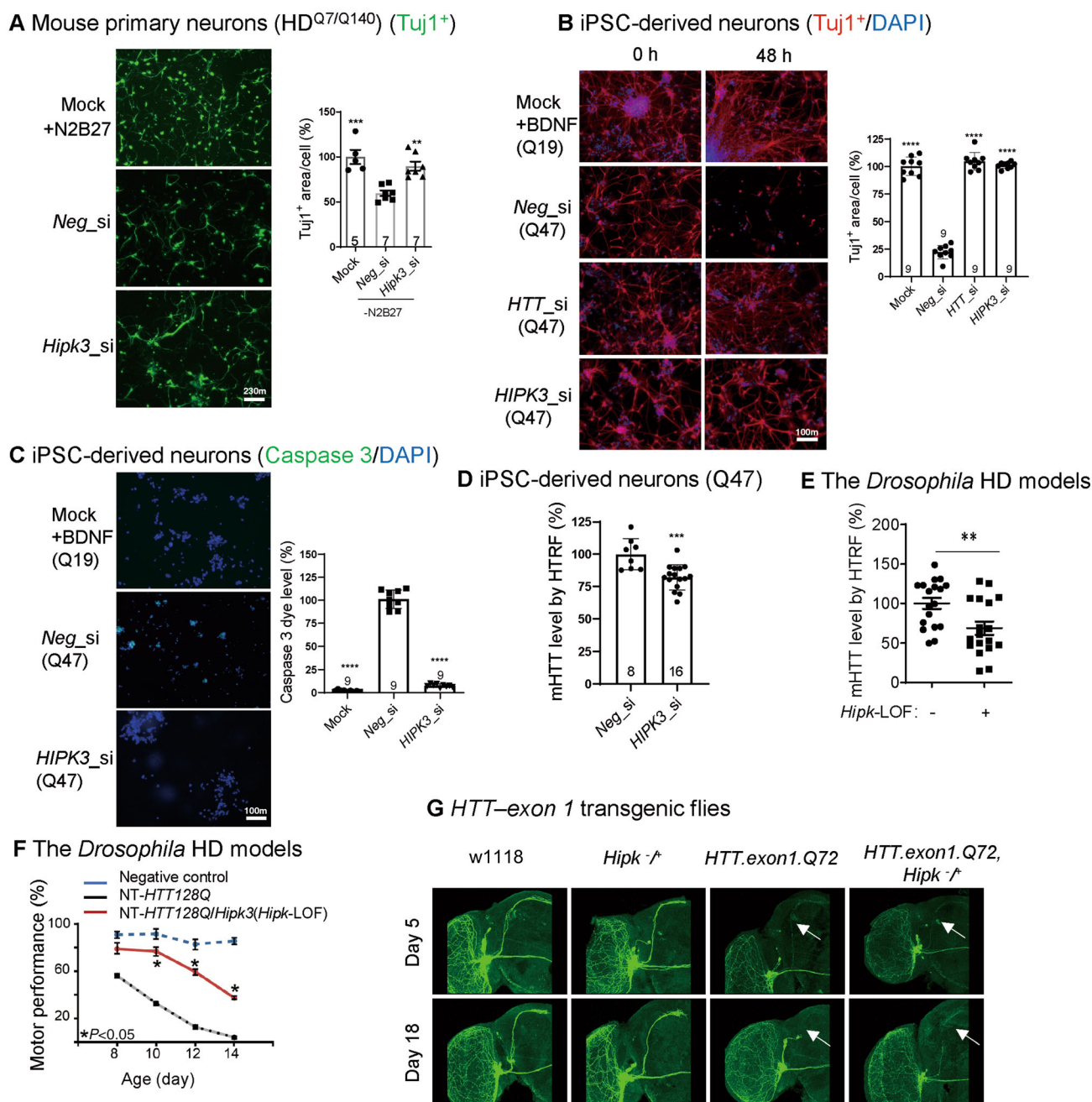
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**Fig. 1.** The potential pathological role of HIPK3 in HD. **A** Representative images and quantification of the Tuj1<sup>+</sup> area in mouse primary neurons (HD<sup>Q7/Q140</sup>). Data are normalized to controls with N2 and B27. Note that the mock group was the same for both the Fig. 1A and Fig. 2E quantification (scale bar, 10 µm; *n*, number of mice of each genotype). **B** Representative immunostaining results of the neuron-specific tubulin marker Tuj1 and DAPI showing the morphology of human iPSC-derived neurons (HD: polyQ = 47; WT: polyQ = 19). Note that the mock group was the same for both the Fig. 1B and Fig. 2F quantification [scale bar, 10 µm; data are normalized to the wild-type (WT)]. **C** Representative images and quantification of the caspase signal in human iPSC-derived neurons (HD: polyQ = 47; WT: polyQ = 19), note that the mock group was the same for both the Fig. 1C and Fig. 2G quantification, scale bar, 10

µm. **D** mHTT levels in HD iPSC-derived neurons (HD: polyQ = 47) treated with *HIPK3* RNAi and controls assessed by the pair of antibodies 2B7/3B5H10 using HTRF (homogeneous time-resolved fluorescence). **E** mHTT levels in fly heads assessed by 2B7/3B5H10 using HTRF (fly model: NT-*HTT128Q*). **F** *Hipk3* depletion rescues the climbing deficit in HD flies. **G** Neurodegeneration in axons of small ventral lateral clock neurons (sLNv) labeled by mCD8GFP protein using GMR61G12-GAL4 in a fly model. Mock, N2 and B27 were not removed from the medium for mouse primary neuron culture, BDNF was not removed from the medium for iPSC-derived neuron culture. *Neg\_si*, siScramble; *Hipk3\_si*, si*Hipk3*. Data are presented as the mean ± SD. \**P* < 0.05; \*\**P* < 0.01; \*\*\**P* < 0.001; ns, *P* > 0.05; one-way ANOVA and *post hoc* Dunnett's tests (A–C, F), and unpaired *t*-test (D, E).

with or without brain-derived neurotrophic factor (BDNF). Knocking down *HIPK3* rescued the neuronal shrinkage (Fig. 1B) and apoptotic phenotype (Fig. 1C), and decreased the mHTT level in the human iPSC-derived neurons (Q47) (Fig. 1D). Taken together, targeting the kinase HIPK3 lowered the mHTT level and ameliorated mHTT neurotoxicity and apoptosis. To further confirm the role of *Hipk3* *in vivo*, we tested two HD *Drosophila* models. A loss-of-function (LOF) mutation of the *Drosophila* homolog of *Hipk3* (*Hipk-LOF*) significantly decreased the mHTT level in the HD transgenic model expressing the N-terminal fragment of human mHTT with 128Q (NT-*HTT-128Q*, Fig. 1E). Consistent with this, the LOF mutation also rescued the climbing deficits in this model (Fig. 1F). We next investigated another HD fly model expressing the human mHTT exon 1 fragment (*HTT-exon1-Q72*) to validate the role of *Hipk3* *in vivo*. In this model, the small ventral lateral clock neurons are labeled by mCD8GFP protein [13], so neurodegeneration *in vivo* is clearly indicated by the neuronal morphology, such as axon shrinkage and loss. *Hipk-LOF* rescued the mHTT-induced neuronal degeneration, while the same group of neurons in the wild-type flies were not influenced by *Hipk-LOF* (Fig. 1G), suggesting that the effects were HD-specific. Taken together, the effects of *Hipk3* are likely to play a role *in vivo* and to be evolutionarily conserved.

We then investigated the possibility of targeting HIPK3 using small molecular inhibitors, which may provide potential entry points to HD therapeutics. AST487 (Fig. 2A) has been reported to be a potent HIPK3 inhibitor based on a kinase screen [14]. We confirmed this by showing that AST487 blocked HIPK3 activity *in vitro* in a dose-dependent manner with an  $IC_{50}$  of 276.15 nmol/L (Fig. 2B). We then tested the effects of AST487 on mouse striatal HD cells (STHdh<sup>Q7/Q111</sup>), and found that mHTT toxicity was significantly rescued by AST487 at 2  $\mu$ mol/L, as measured by an indicator dye for active caspase 3 (Fig. 2C). We further confirmed that mHTT toxicity was significantly rescued in an STHdh<sup>Q7/Q7</sup>-transfected model (Fig. 2D). AST487 was also sufficient to rescue the outgrowth deficits in mouse primary neurons (HD<sup>Q7/Q140</sup>) (Fig. 2E). Consistent with the results in mouse cells, AST487 rescued the disease-relevant phenotypes in the iPSC-derived neurons from HD patients after BDNF removal. Treatment with AST487 at 5  $\mu$ mol/L rescued the neuronal shrinkage (Fig. 2F) and apoptosis (Fig. 2G). We further demonstrated that AST487 treatment decreased

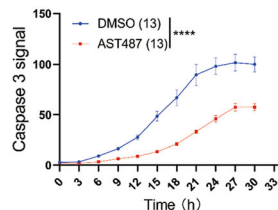
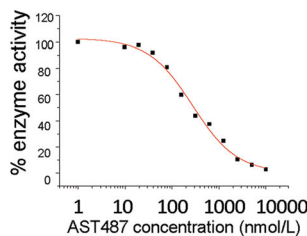
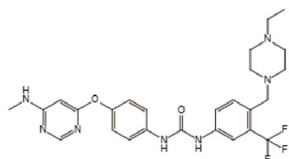
mHTT levels (Fig. 2H). Finally, to confirm the effects of AST487 *in vivo*, we examined the mHTT level in HD transgenic flies expressing full-length mHTT, and showed that AST487 significantly reduced the mHTT levels in fly transgenic models (Fig. 2I). Collectively, these results confirmed the significant role of HIPK3 and the rescue effect of inhibitors both *in vitro* and *in vivo*.

We have demonstrated that HIPK3 modulates autophagy in a previously study. Here, we further showed that *HIPK3* RNAi (RNA interference) or AST487 treatment enhanced the autophagosome marker protein LC3B-II levels in mouse primary neurons (Fig. 2J) and human iPSC-derived neurons (Fig. 2K), indicating that the autophagy level was up-regulated. *HIPK3* RNAi decreased the caspase signal (Figs 1C and 2G) and significantly decreased the cleaved caspase 3 in mouse primary neurons (Fig. 2L) and human iPSC-derived neurons (Fig. 2M), suggesting that HIPK3 is involved in the pathogenesis in HD neurons. Collectively, inhibition of the HIPK3 kinase ameliorates mHTT-induced neurotoxicity and apoptosis.

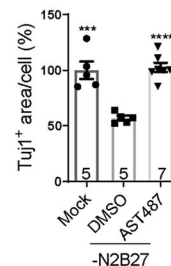
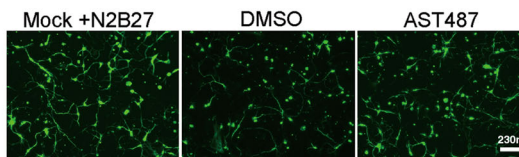
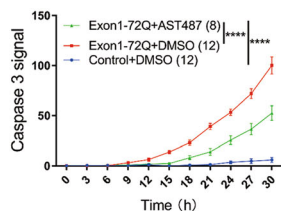
Gain of toxic function of mHTT is the major cause of HD, so lowering mHTT protein levels may ameliorate its downstream toxic effects and treat HD. In neurodegenerative disorders, neuronal death occurs *via* one of two mechanisms: apoptosis or necrosis [15]. So far, there is no strong evidence to support that HD is correlated with apoptosis per se, however, the processes involved in initiating apoptosis happens before the onset of HD [15]. Autophagy and apoptosis are basic physiological processes, so the modulation of some regulators of autophagy and apoptosis may be an ideal way to treat neurodegenerative disorders.

The HTT protein is very large (348 kDa), its entire crystal structure and function are still unknown, so it is very difficult to develop therapeutic drugs that target it. So, protein kinases and receptors are ideal drug targets for HD and target-discovery approaches for similar diseases. The current study provides both genetic and chemical biology evidence demonstrating that HIPK3 is a potential therapeutic target of HD: knocking down *HIPK3* or treatment with the HIPK3 inhibitor AST487 (Figs 1 and 2). Thus, targeting the kinase HIPK3 to lower the mHTT level would be a potential direction for developing HD drugs. Collectively, targeting HIPK3 using small-molecule inhibitors could be beneficial to HD patients, while further optimization of the compounds and further *in vivo* studies in the HD mouse models are needed.

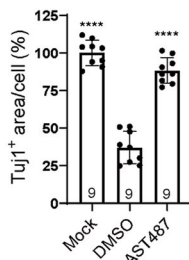
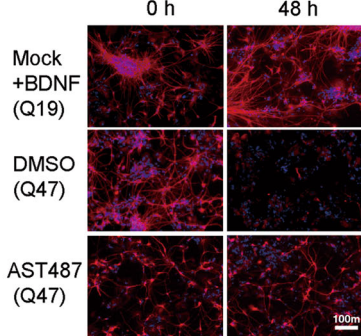
**A** AST487 chemical structure    **B** HIPK3 kinase activity    **C** Mouse striatal cells (STHdh<sup>Q7/Q111</sup>)



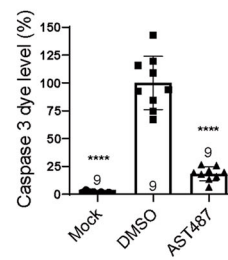
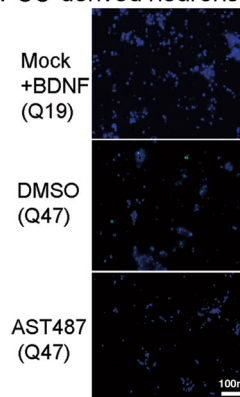
**D** Mouse striatal cells (STHdh<sup>Q7/Q7</sup>)    **E** Mouse primary neurons (HD<sup>Q7/Q140</sup>) (Tuj1<sup>+</sup>)



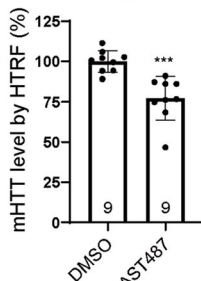
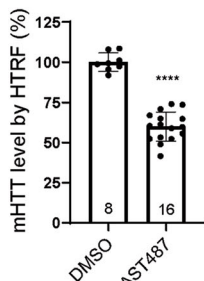
**F** iPSC-derived neurons (Tuj1<sup>+</sup>/DAPI)



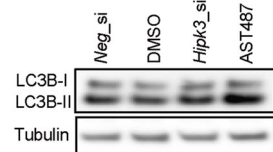
**G** iPSC-derived neurons (Caspase 3/DAPI)



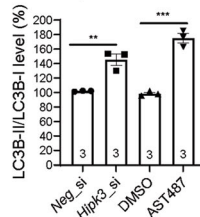
**H** iPSC-derived neurons (Q47)    **I** fl-HTT transgenic flies



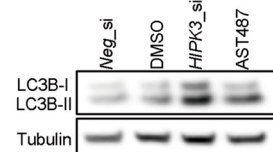
**J** Mouse primary neurons (HD<sup>Q7/Q140</sup>)



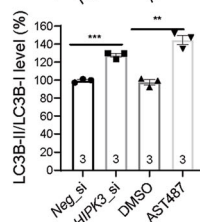
LC3B-II/Tubulin ratio : 1.0 1.0 1.33 1.66



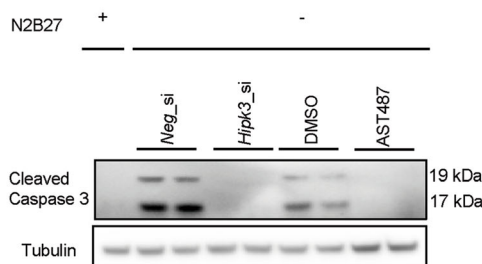
**K** iPSC-derived neurons (Q47)



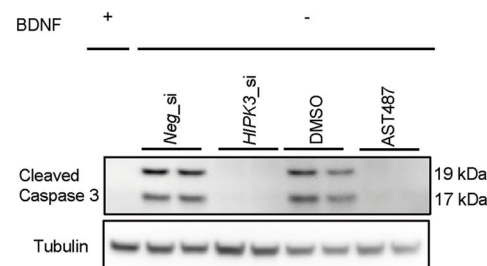
LC3B-II/Tubulin ratio : 1.0 1.0 2.10 1.83



**L** Mouse primary neurons (HD<sup>Q7/Q140</sup>)



**M** iPSC-derived neurons (Q47)



**Fig. 2.** The potential pathological role of AST487 in HD. **A** Chemical structure of AST487. **B** AST487 regulates HIPK3 kinase activity. The plots are fitted with a logistic curve. **C** Apoptosis in STHdh<sup>Q77/Q111</sup> cells at different time points with or without AST487 treatment (active caspase 3 was measured using NucView 488 dye). Data are presented as the mean  $\pm$  SD. **D** As in C, but using STHdh<sup>Q77/Q7</sup> cells transfected with exon 1 *HTT-Q72* plasmid. **E** Representative images and quantification of primary neurons (HD<sup>Q77/Q140</sup>), note that the mock group was the same for both the Fig. 1A and Fig. 2E quantification, scale bar, 10  $\mu$ m. **F** Representative immunostaining results of the neuron-specific tubulin marker Tuj1 and DAPI showing the morphology of human iPSC-derived neurons (HD: Q47; WT: Q19) [note that the mock group was the same for both the Fig. 1B and Fig. 2F quantification; scale bar, 10  $\mu$ m; data are presented as the mean  $\pm$  SD (normalized to the WT)]. **G** Representative images and quantification of the caspase 3 signal in human iPSC-derived neurons (HD: polyQ = 47; WT: polyQ = 19), note that the mock group was the same for both the Fig. 1C and Fig. 2G quantification, scale bar, 10  $\mu$ m. **H** mHTT levels in human iPSC-derived neurons (HD: polyQ = 47; WT: polyQ = 19). **I** mHTT levels in fly heads assessed by 2B7/3B5H10 using HTRF (fly model: fl-*HTT-Q128*). **J** Representative images and quantification of LC3B Western blots for *Hipk3* RNAi or *Hipk3* inhibitor AST487 treatment in mouse primary neurons (HD<sup>Q77/Q140</sup>) ( $n = 3$  replicates). **K** As in J, but in human iPSC-derived neurons (HD: polyQ = 47) ( $n = 3$  replicates). **L** Representative caspase 3 Western blots for *Hipk3* RNAi or *Hipk3* inhibitor in mouse primary neurons (HD<sup>Q77/Q140</sup>). **M** As in L, but in human iPSC-derived neurons (HD: polyQ = 47). Mock, N2 and B27 were not removed from the medium for mouse primary neuron culture, BDNF was not removed from the medium for iPSC-derived neuron culture. Data are presented as the mean  $\pm$  SD. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ ; ns,  $P > 0.05$ ; two-way ANOVA (C, D), one-way ANOVA and *post hoc* Dunnett's tests (E–G), or the unpaired *t*-test (H–K).

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**Conflict of interest** The authors claim no conflicts of interest.

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