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## Abstract

Cellular layers enter the lysosome compartment by endocytosis, phagocytosis, or autophagy. Inside the lysosomal compartment, flm components of complex structure are corrupted into their building squares. These are

proteins is required.

, E e higii<sup>4</sup>Shigii<sup>4</sup>aia, , ei

4 4 .....

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he he ef legalai l'he iag h lii, a hag ha e a abaa ee. Sa ai la ia a ae a hag iee ea fe e ia i e e . hag i le e e a ee el i era e. Bi fr / elge i he e a ah/a fa hag / ee i a i irel . b he el a a he i ea a lefe i e i a hag. ... i fa hag b lie i ea i he a e.

I he e 4 I hee a Jiae, aa he ia / ih ai<sup>n</sup>i . V-. iaeae fhጚ e e e а, a, ei, a haite, ei ait, g h h i it.Te e hei ite fhe e f gae, a he e<sup>¶</sup>e,i a i e he i a, he i egi fhe e i i g. eet be e et ate a it he eht f a ... i . i a bea . i het b a hi g a . et f he ab h tae. i f a f ta e a ae... ei (LIMpS) a t a a f<sup>4</sup>a e a a e, , ei (LLMAPS) a<sup>4</sup> e a e<sup>4</sup>, , ei, e he i a , e , i e<sup>4</sup> f, i i<sup>4</sup>, , i a bee, e e<sup>4</sup>, ea h hei, b, a e h, gh hig a hihi el fg ei e eelig -g ael/ih a a ie i Si e heelge i e eeli degalai, a e i a abe f e baeV beli a/ihi he a/ a .a. e [-10]. he Hiele eblae HegaHai e ale, a/eH. i/a he a ei e f gaie /ih.higiiłaai aalie ha Mulgagiiłi . bi el, hi g i il a i a ... ei (Sa) a, , he e he gahe a vi e i a ... ageb lie. e. al, he ha ebee ile i el a MM B. ... e e i g he elia f Sa ... e. ... le ie b b a / ih a a e .ai f. e. el Sa ... e. c. el a l'heb . a agg.egai f i e i a ... e, a Lee ab i het he e a a i Hobaeg , hig i it.

L a., e i e e, he ahe i , aa e he h l i f., ei . Fer f he, ei ae / , a a i - , ab , e i ae, rhe, ea a, ee l e i ae l a e l a e l a e i ha e a, a e he a, a i, ei e, ei e, ei ae fa i e f h l i e e. e ae i ae l i ai e- , e e ei , a a l ae a a, e e giel i i le a, ei e f he e l i, ahra. I, a i a e , eh e a bee i el a l igh f. e, a l i he li ae, ei e a i g a ai i e e, a ige , e ai g i i le ea e e e i i le e, a ige , e ai g i i le ea e e e , . . i ha li g, . h e ha li g, a l legalai f fa er , i e i he e ae a , a e. I allii,

a., eae haebee ... e<sup>4</sup> .e.e. .b.e he a. fa. if. ii<sup>4</sup>e he . I e.a i hei e ai / ..., ee f a... ei ig he ahe i ea<sup>4</sup> i... e<sup>4</sup> ifei e f he... ei / ihi he a e.i. e., a i he ae f ahe i , e.a i i<sup>4</sup>ae,

a e, i, e, a i he a e f a he i , e, a i i a e, a h, -ga a i ha e, T i i h- h, e, e a e 0.0 ei la a he -1.2  $\Gamma_D(T/0-1.2 \Gamma_D(\cdot)$  hag b hiele a b 18211 f21101012113130.2.83720.537 f, i, Mh ( Ma a i ) 0.5( be Lybie h e a lei e) 0.5( f)] TBL018 T/TBL hee e - he e f he e e i p 1-h of 0 I0 · L FX BBC FX