

Table 5. Bioinformatic associations among gene candidates identified by RNAi

<i>C. elegans</i> gene ID	Association	Component of:
K11G12.4 (<i>smf-1</i>)	PINK1 and DJ-1 *	
R05D11.6	PINK1 and DJ-1 *	
F26E4.11 (<i>hrdl-1</i>)	PINK1 and DJ-1 *	ERAD
C35D10.2	DJ-1 [†]	Autophagy
M7.5 (<i>atgr-7</i>)		UPS and autophagy
F32A6.3 (<i>vps-41</i>)	UCHL-1 *	UPS and autophagy

*Coexpressed in microarray topology map, as per Kim *et al.* (1).

[†]Identified within same protein interactome network, as per Li *et al.* (2).

1. Kim SK, *et al.* (2001) A gene expression map for *Caenorhabditis elegans*. *Science* 293:2087-2092.
2. Li S, *et al.* (2004) A map of the interactome network of the metazoan *C. elegans*. *Science* 303:540-543.