

<i>H. sapiens</i> Gene	Variant	Ensemble Gene	<i>C. elegans</i> Locus ID	<i>C. elegans</i> Gene	Animals Defective (% ± SD) [*] - NC1687	n	Animals Defective (% ± SD) [*] - HML429	n
<i>HSP90AA1</i>	SYN-NET ^{LGD}	ENSG00000080824	C47E8.5	<i>daf-21</i>	72.0 ± 24.4	25	Emb	n/a
<i>HSPA8</i>	SYN-NET ^{LGD} rdNS-SCZ	ENSG00000109971	F26D10.3	<i>hsp-1</i>	28.8 ± 14.2	66	Emb	n/a
<i>YWHAZ</i>	SYN-NET ^{MIS}	ENSG00000164924	M117.2	<i>par-5</i>	15.4 ± 10.5	26	71.0 ± 27.2	31
<i>YWHAZ</i>	SYN-NET ^{MIS}	ENSG00000164924	F52D10.3	<i>ftt-2</i>	3.3 ± 2.4	60	35.2 ± 6.8	54
<i>GIT1</i>	SYN-NET ^{MIS}	ENSG00000108262	F14F3.2	<i>git-1</i>	3.3 ± 2.4	60	32.4 ± 12.1	37
pPD129.36 empty vector					7.9 ± 6.7	101	0.0 ± 0.0	20

(1) ^{*} = % and standard deviation (SD) denote the weighted % and weighted standard deviation, respectively.