

<b>Genotype</b>	<b>mean RS± std. error</b>	<b>% change</b>	<b>P-value</b>	<b>N=</b>
<b>Experiment 1:</b>				
wild type	<b>3.5±0.2</b>	--	--	30
<i>daf-7(e1372)</i>	<b>5.4±0.1</b>	<b>+54%</b>	<0.0001	30
<b>Experiment 2:</b>				
wild type	<b>3.6±0.2</b>	--	--	30
<i>daf-1(m40)</i>	<b>4.9±0.1</b>	<b>+36%</b>	<0.0001	30
<i>daf-14(m77)</i>	<b>5.3±0.2</b>	<b>+47%</b>	<0.0001	30
<b>Experiment 3:</b>				
wild type	<b>3.8 ±0.2</b>	--	--	45
<i>daf-4(e1364)</i>	<b>7.7 ±0.3</b>	<b>+103%</b>	<0.0001	26
<b>Experiment 4:</b>				
wild type	<b>3.9±0.3</b>	--	--	12
<i>daf-7(e1372)</i>	<b>5.6±0.2</b>	<b>+44%</b>	<0.0001	17
<b>Experiment 5:</b>				
wild type	<b>3.3±0.1</b>	--	--	30
<i>daf-7(e1372)</i>	<b>4.1±0.2</b>	<b>+24%</b>	<0.0001	30
<i>daf-14(m77)</i>	<b>4.4±0.2</b>	<b>+33%</b>	<0.0001	30