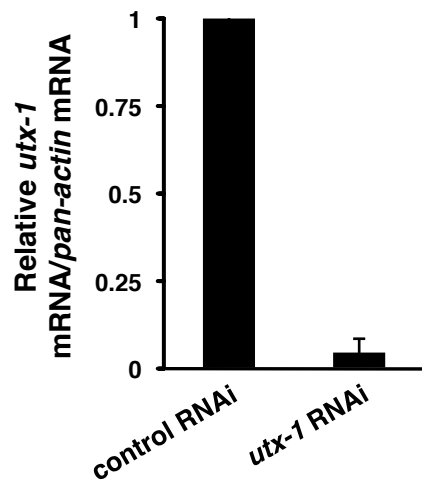


Demethylase Screen			
	Initial screen		
RNAi construct	Mean +/- SD	p values	# worms
control	19.760 +/- 0.568		86/90
rbr-2	17.767 +/- 0.805	0.0319	30/30
<i>tag-279</i>	18.000 +/- 0.701	0.0323	30/30
<i>F23D12.5</i>	18.225 +/- 1.079	0.2650	28/30
<i>C06H2.3</i>	18.833 +/- 0.716	0.2005	30/30
<i>psr-1</i>	19.100 +/- 1.052	0.7186	30/30
<i>C29F7.6</i>	19.756 +/- 1.047	0.8629	22/30
<i>F29B9.2</i>	20.495 +/- 0.851	0.6823	28/30
F43G6.6	22.456 +/- 0.758	0.0586	29/30
utx-1	23.560 +/- 1.455	0.0012	27/30
T26A5.5	24.104 +/- 1.259	0.0003	28/30
control	18.419 +/- 0.473		52/60
<i>amx-1</i>	18.954 +/- 0.604	0.5803	29/30
lsd-1	22.393 +/- 0.589	<0.0001	27/30
Repeat of targets with >10% change in lifespan			
control	17.981 +/- 0.422		78/90
<i>F43G6.6</i>	16.761 +/- 0.431	0.0581	81/90
<i>lsd-1</i>	19.025 +/- 0.496	0.0560	85/90
control	19.185 +/- 0.397		89/90
rbr-2	16.111 +/- 0.432	<0.0001	90/90
control	18.852 +/- 0.487		88/90
utx-1	21.652 +/- 0.584	<0.0001	82/90
T26A5.5 #1	21.144 +/- 0.567	<0.0009	86/90
T26A5.5 #2	21.374 +/- 0.477	<0.0005	88/90

	1			2			
N2 + RNAi	Mean +/- SD	p values	# worms	Mean +/- SD	p values	# worms	Combined p values
N2 control	18.158 +/- 0.525		68/90				
N2 <i>T26A5.5 (Adult)</i>	21.173 +/- 0.624	0.0005	64/90				
N2 control	15.716 +/- 0.343		79/90	16.989 +/- 0.317		73/95	
N2 <i>utx-1 (L1)</i>	17.697 +/- 0.386	<0.0001	95/116	19.662 +/- 0.649	<0.0001	66/93	<0.0001
N2 control	16.540 +/- 0.417		80/88	17.214 +/- 0.451		73/95	
N2 <i>utx-1 (Adult)</i>	21.385 +/- 0.584	<0.0001	60/73	20.813 +/- 0.564	<0.0001	66/93	<0.0001
Strain	Mean +/- SD	p values	# worms	Mean +/- SD	p values	# worms	Combined p values
N2	21.736 +/- 0.529		63/90	20.550 +/- 0.570		76/90	
<i>T26A5.5(ok2364)</i>	21.140 +/- 0.430	0.1782	67/90	20.427 +/- 0.514	0.7993	88/90	
stDp2/+	11.771 +/- 0.385		45/90	12.594 +/- 0.349		63/92	
stDp2/ <i>utx-1(tm3118)</i>	17.684 +/- 0.355	<0.0001	64/86	16.054 +/- 0.378	<0.0001	71/92	<0.0001
N2 control	13.717 +/- 0.509		66/88	16.932 +/- 0.607		73/90	
N2 <i>utx-1</i>	15.779 +/- 0.477	0.0040	72/85	20.931 +/- 0.610	<0.0001	71/91	<0.0001
<i>glp-1(e2141)</i> control	16.369 +/- 0.442	0.0002	86/92	20.674 +/- 0.608	<0.0001	89/92	<0.0001
<i>glp-1(e2141) utx-1</i>	18.191 +/- 0.432	0.0088	84/92	23.960 +/- 0.666	0.0006	89/92	<0.0001
Statistical comparison							
N2 <i>utx-1</i> / <i>glp-1 utx-1</i>		0.0005			<0.0001		<0.0001
Two-way ANOVA N2/ <i>glp-1 +/- utx-1</i>		0.0244			0.5740		0.0738
N2 control	15.378 +/- 0.453		75/93				
N2 <i>utx-1</i>	18.940 +/- 0.666	<0.0001	68/92				
<i>eat-2(ad1116)</i> control	17.025 +/- 0.463	0.0219	80/88				
<i>eat-2(ad1116) utx-1</i>	19.875 +/- 0.524	<0.0001	85/93				
Statistical comparison							
N2 <i>utx-1</i> / <i>eat-2 utx-1</i>		0.5274					
Two-way ANOVA N2/ <i>eat-2 +/- utx-1</i>		<0.0001					
N2 control	19.627 +/- 0.659		71/93	21.369 +/- 0.479		74/90	
N2 <i>utx-1</i>	25.162 +/- 0.785	<0.0001	64/92	24.452 +/- 0.641	<0.0001	75/90	<0.0001
<i>daf-16(mu86)</i> control	15.596 +/- 0.284	<0.0001	78/92	16.224 +/- 0.370	<0.0001	85/90	<0.0001
<i>daf-16(mu86) utx-1</i>	16.429 +/- 0.255	0.0195	69/92	16.614 +/- 0.337	<0.6034	88/90	<0.0640
Statistical comparison							
N2 <i>utx-1</i> / <i>daf-16(mu86) utx-1</i>		<0.0001			<0.0001		<0.0001
Two-way ANOVA N2/ <i>daf-16 +/- utx-1</i>		<0.0001			0.0036		<0.0001
N2 control	21.369 +/- 0.479		74/90	19.627 +/- 0.659		71/93	
N2 <i>utx-1</i>	24.452 +/- 0.641	<0.0001	75/90	25.162 +/- 0.785	<0.0001	64/92	<0.0001
<i>daf-2(e1370)</i> control	39.792 +/- 1.056	<0.0001	81/90	27.431 +/- 1.535	<0.0001	74/90	<0.0001
<i>daf-2(e1370) utx-1</i>	39.226 +/- 1.212	0.5367	59/92	28.893 +/- 1.302	0.6484	63/91	0.7153
Statistical comparison							
N2 <i>utx-1</i> / <i>daf-2(e1370) utx-1</i>		<0.0001			<0.0001		<0.0001
Two-way ANOVA N2/ <i>daf-2 +/- utx-1</i>		0.0393			0.0777		0.0207

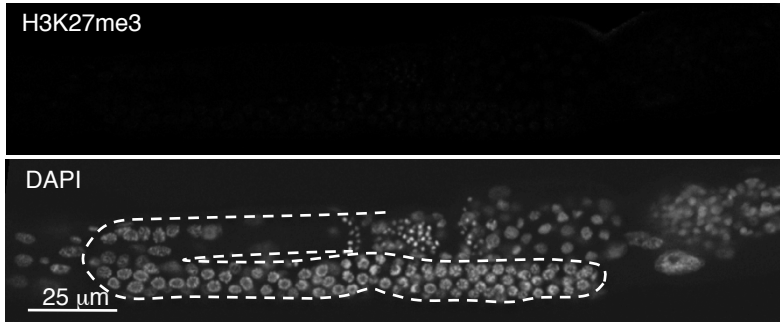
Supplementary Table 2



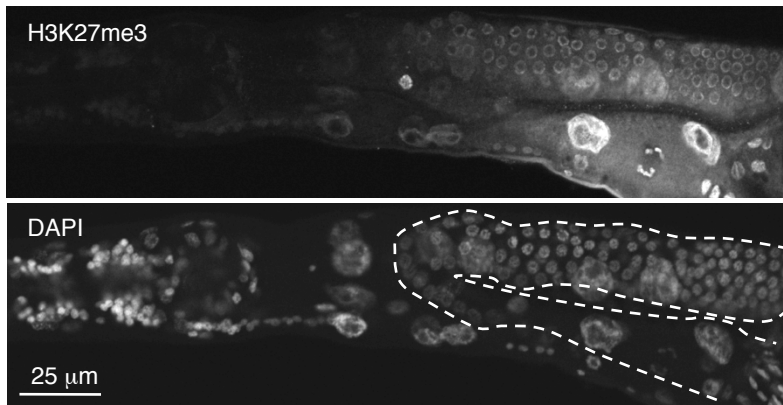
Supplementary Fig. 1

A

control RNAi

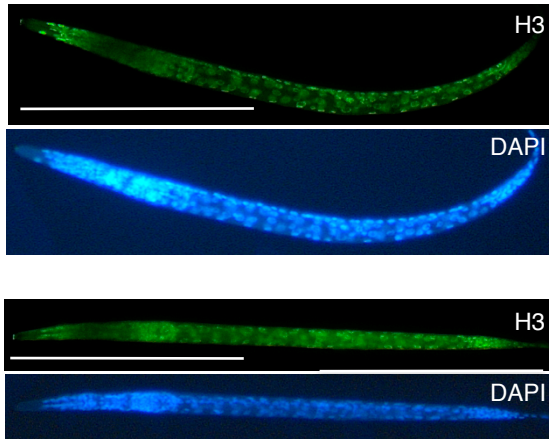


***utx-1* RNAi**

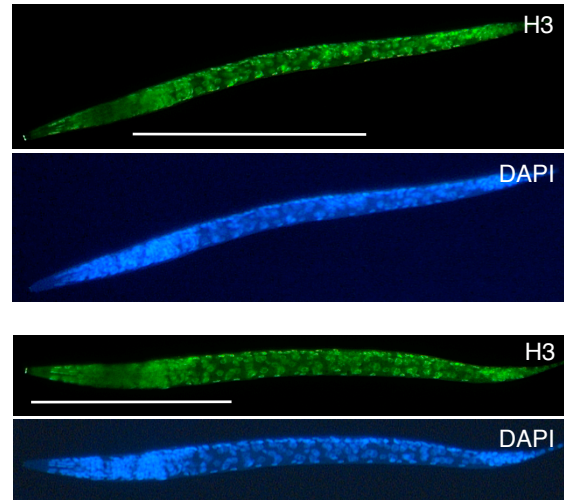


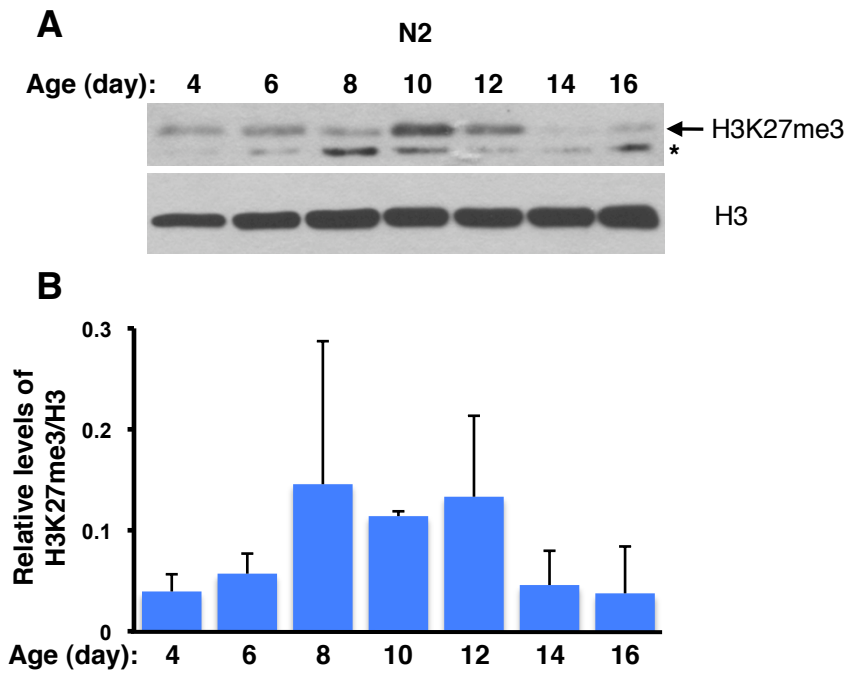
B

control RNAi



***utx-1* RNAi**





Supplementary Fig. 3