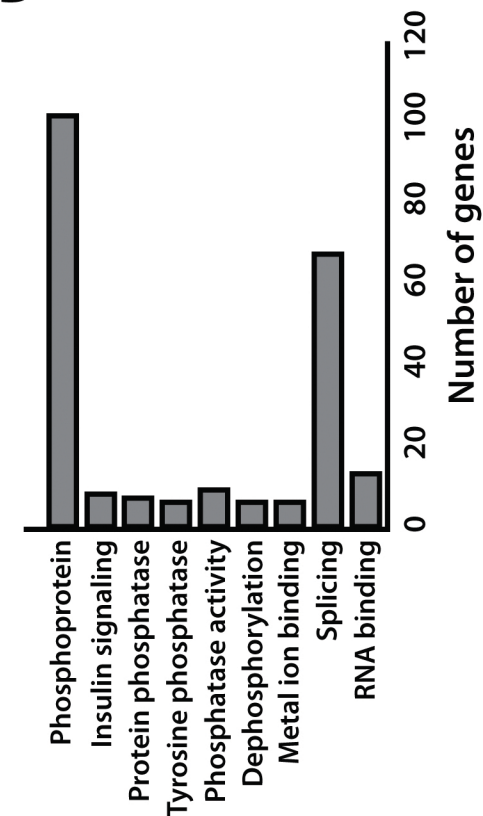


Supplemental Fig. 3

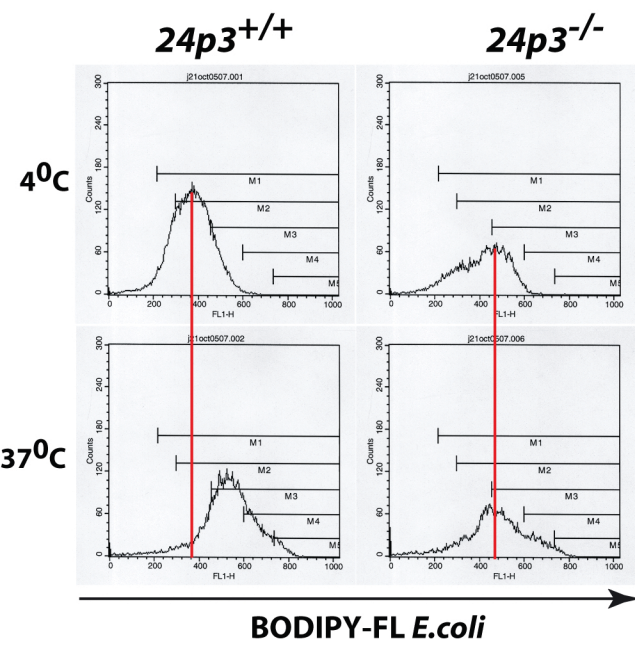
A

Transcript ID(Array Design)	24p3 ^{-/-} /wt	Sequence	Alignments	Sequence Length
mmu-mir-362	3.39	AACACACCUGUUCAAGGAUUCA	X:6819108-6819172 (-)	22
mmu-mir-31	3.04	UGCUAUGCCAACAUAUUGCCAUC	4:88556461-88556566 (-)	23
mmu-mir-1958	2.13	UAGGAAAGUGGAAGCAGUAAGU	4:48225991-48226087 (-)	22
mmu-mir-1936	2.04	UAACUGACCUGCUGUGAACUGGC	12:103923138-103923230 (-)	23
mmu-mir-363	1.88	AAUUGCACGGUAUCCAUCUGUA	X:50094870-50094944 (-)	22
mmu-mir-423	1.65	UGAGGGGCAGAGAGCGAGACUUU	11:78891566-78891674 (-)	23
mmu-mir-483	1.72	AAGACGGGAGAAGAGAAGGGAG	7:149840829-149840901 (-)	22
mmu-mir-18a	0.58	ACUGCCCUAAGUGCUCCUUCUG	14:115443073-115443168 (+)	22
mmu-mir-878	0.54	GCAUGACACCACACUGGGUAGA	X:64054683-64054760 (-)	22
mmu-mir-1903	0.45	CCUUCUUCUUCUUCUGAGACA	8:130883141-130883220 (+)	22
mmu-mir-1966	0.43	AAGGGAGCUGGCUCAGGAGAGAGUC	8:108139366-108139473 (+)	25
mmu-mir-669b	0.49	AGUUUUGUGUGCAUGUGCAUGU	2:10389417-10389513 (+)	22
mmu-mir-335	0.58	UUUUUCAUUAUUGCUCCUGACC	6:30691299-30691396 (+)	22
mmu-mir-654	0.36	UAUGUCUGCUGACCAUCACCUU	12:110961428-110961511 (+)	22

B



C



D

