

Table S2. PLLP migration distance is not affected by *lef1* loss of function

A						
	Siblings			<i>lef1</i>^{u767}		
	Average	s.d.	<i>n</i>	Average	s.d.	<i>n</i>
27 hours	393	68.5	15	344	92.6	7
30 hours	493	80	15	409	108.6	7
33 hours	649	80.7	15	555	119.2	7
36 hours	815	77.8	15	668	117.9	7

B						
	Siblings			<i>lef1</i> morphants		
	Average	s.d.	<i>n</i>	Average	s.d.	<i>n</i>
27 hours	412	57.1	5	350	32.4	38
30 hours	580	47.2	19	564	31.6	37
33 hours	642	71.1	19	620	52.8	40
36 hours	861	61.6	17	799	84.9	48

Table shows average distance (μm) from the caudal limit of the otic vesicle and the leading edge of the PLL primordium.

s.d., standard deviation; *n*, number of embryos counted per point.

(A) Sibling versus *lef1*^{u767} mutant embryos.

(B) Sibling versus *lef1* morphant embryos.