

Table S3. BrdU incorporation index in siblings and *lef1^{u767}* mutant embryos

	Sibling total count			<i>lef1^{u767}</i> total count		
	Average	s.d.	<i>n</i>	Average	s.d.	<i>n</i>
1	0.510	0.094	8	0.228	0.105	10
2	0.524	0.098	8	0.268	0.069	8
3	0.540	0.061	8	0.249	0.062	11
4	0.529	0.098	8	0.279	0.074	8

	Sibling leading zone			<i>lef1^{u767}</i> leading zone			Sibling trailing zone			<i>lef1^{u767}</i> trailing zone		
	Average	s.d.	<i>n</i>	Average	s.d.	<i>n</i>	Average	s.d.	<i>n</i>	Average	s.d.	<i>n</i>
1	0.484	0.134	8	0.189	0.118	10	0.327	0.142	10	0.198	0.116	10
2	0.541	0.150	8	0.332	0.087	8	0.348	0.105	8	0.307	0.137	8
3	0.513	0.108	8	0.170	0.116	11	0.322	0.091	8	0.268	0.168	11
4	0.582	0.161	8	0.226	0.096	8	0.310	0.143	8	0.198	0.079	8

Table shows average fraction of the number of BrdU-positive cells over the total number of cells in the PLL primordium after 1, 2, 3 and 4 deposited neuromasts in the whole primordium, leading and trailing zones.

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