

Supplementary Table 1. Effects of morpholino oligonucleotides

Type of embryo	Gene tested	Stage	Number of embryos			Figure	Description
			Down	No effect	Up		
<i>jam1a</i> MOatg	<i>runx1</i>	26hpf	37	0	0	Fig. 1g	
<i>jam1a</i> MOatg	<i>efnb2a</i>	26hpf	0	14	0	Fig. 1j	
<i>jam1a</i> MOatg	<i>cmyb</i>	36hpf	36	7	0	Fig. 1l	
<i>jam1a</i> MOatg	<i>rag1</i>	4dpf	22	0	0	Fig. 1m	
<i>jam1a</i> MOatg	<i>scl-a/β</i>	30hpf	16	0	0	Fig. 1n	Full-length <i>scl</i> probe
<i>jam1a</i> MOatg	<i>scl-a</i>	30hpf	0	17	0	Fig. 1o	5' <i>scl</i> probe
<i>jam1a</i> MOatg	<i>fli1</i>	12hpf	0	20	0	Fig. 2b	
<i>jam1a</i> MOatg	<i>fli1</i>	17hpf	27*	3	0	Fig. 2e	
<i>jam1a</i> MOatg	<i>dlc</i>	14hpf	0	20	0	Fig. 4o	
<i>jam1a</i> MOatg	<i>dld</i>	14hpf	0	24	0	Fig. 4q	
<i>jam1a</i> MOatg	<i>runx1</i>	28hpf	24	3	0	Fig. 4s	<i>jam1a</i> MOatg only
<i>jam1a</i> MOatg	<i>runx1</i>	28hpf	4	20**	0	Fig. 4t	<i>jam1a</i> MOatg + <i>dlc</i>
<i>jam1a</i> MOatg	<i>runx1</i>	28hpf	5	26***	0	Fig. 4u	<i>jam1a</i> MOatg + <i>dld</i>
<i>jam1a</i> MOatg	<i>gata1</i>	24hpf	0	18	0	E.D. Fig. 4d	
<i>jam1a</i> MOatg	<i>l-plastin</i>	24hpf	0	13	0	E.D. Fig. 4e	
<i>jam1a</i> MOatg	<i>l-plastin</i>	4dpf	33	2	0	E.D. Fig. 4f	Expression in the CHT
<i>jam1a</i> MOatg	<i>kdrl</i>	26hpf	0	20	0	E.D. Fig. 4g	Expression in the trunk
<i>jam1a</i> MOatg	<i>kdrl</i>	26hpf	20	0	0	E.D. Fig. 4h	Expression in the CHT
<i>jam1a</i> MOatg	<i>tbx20</i>	26hpf	0	13	0	E.D. Fig. 4i	
<i>jam1a</i> MOatg	<i>flt4</i>	26hpf	0	14	0	E.D. Fig. 4j	
<i>jam1a</i> MOatg	<i>cdh17</i>	24hpf	0	24	0	E.D. Fig. 4k	
<i>jam1a</i> MOatg	<i>desma</i>	26hpf	0	21	0	E.D. Fig. 4l	
<i>jam1a</i> MOatg	<i>nkx3.1</i>	26hpf	0	24	0	E.D. Fig. 4m	
<i>jam1a</i> MOatg	<i>shha</i>	16hpf	0	17	0	E.D. Fig. 4n	
<i>jam1a</i> MOatg	<i>shha</i>	26hpf	0	17	0	E.D. Fig. 4o	
<i>jam1a</i> MOatg	<i>vegfa</i>	17hpf	0	23	0	E.D. Fig. 4p	
<i>jam1a</i> MOatg	<i>vegfa</i>	26hpf	0	17	0	E.D. Fig. 4q	
<i>jam1a</i> MOatg	<i>runx1</i>	26hpf	47	4	0	E.D. Fig. 7b	<i>hsp:Gal4/UAS:NICD</i> (-)
<i>jam1a</i> MOatg	<i>runx1</i>	26hpf	1	4	15	E.D. Fig. 7b	<i>hsp:Gal4/UAS:NICD</i> (+)
<i>jam1a</i> MOatg	<i>runx1</i>	26hpf	59	1	0	E.D. Fig. 7d	<i>fli1:Gal4/UAS:NICD</i> (-)
<i>jam1a</i> MOatg	<i>runx1</i>	26hpf	0	0	22	E.D. Fig. 7d	<i>fli1:Gal4/UAS:NICD</i> (+)
<i>jam1a</i> MOatg	<i>notch1b</i>	26hpf	0	15	0	E.D. Fig. 9b	
<i>jam1a</i> MOatg	<i>notch3</i>	26hpf	0	17	0	E.D. Fig. 9d	
<i>jam1a</i> MOatg	<i>dlc</i>	26hpf	0	12	0	E.D. Fig. 9f	
<i>jam1a</i> MOatg	<i>dll4</i>	26hpf	0	17	0	E.D. Fig. 9h	
<i>jam1a</i> MOatg	<i>Tp1:GFP</i>	28hpf	7	1	0	E.D. Fig. 9j	<i>jam1a</i> MOatg only
<i>jam1a</i> MOatg	<i>Tp1:GFP</i>	28hpf	3	5	0	E.D. Fig. 9k	<i>jam1a</i> MOatg + <i>dlc</i>
<i>jam1a</i> MOatg	<i>Tp1:GFP</i>	28hpf	2	6	0	E.D. Fig. 9l	<i>jam1a</i> MOatg + <i>dld</i>
<i>jam1a</i> MOex7	<i>runx1</i>	26hpf	24	5	0	Fig. 1h	
<i>jam1a</i> MOex7	<i>efnb2a</i>	26hpf	0	13	0	Fig. 1k	
<i>jam1a</i> MOex7	<i>fli1</i>	12hpf	0	20	0	Fig. 2c	
<i>jam1a</i> MOex7	<i>fli1</i>	17hpf	27*	4	0	Fig. 2f	

<i>jam1a</i> MOex7	<i>cmyb</i>	36hpf	13	3	0	E.D. Fig. 3g	
<i>jam1a</i> MOex7	<i>rag1</i>	4dpf	25	0	0	E.D. Fig. 3i	
<i>jam2a</i> MOatg	<i>runx1</i>	26hpf	33	2	0	Fig. 3h	
<i>jam2a</i> MOatg	<i>efnb2a</i>	26hpf	1	29	0	Fig. 3j	
<i>jam2a</i> MOatg	<i>fli1</i>	17hpf	26*	1	0	Fig. 3k	
<i>jam2a</i> MOatg	<i>runx1</i>	26hpf	41	0	0	E.D. Fig. 7e	<i>hsp:Gal4/UAS:NICD</i> (-)
<i>jam2a</i> MOatg	<i>runx1</i>	26hpf	1	14	0	E.D. Fig. 7e	<i>hsp:Gal4/UAS:NICD</i> (+)
<i>jam2a</i> MOatg	<i>runx1</i>	26hpf	35	12	0	E.D. Fig. 7f	<i>fli1:Gal4/UAS:NICD</i> (-)
<i>jam2a</i> MOatg	<i>runx1</i>	26hpf	0	0	14	E.D. Fig. 7f	<i>fli1:Gal4/UAS:NICD</i> (+)
<i>jam2a</i> MOex5	<i>runx1</i>	26hpf	23	3	0	E.D. Fig. 5f	
<i>jam2a</i> MOex5	<i>efnb2a</i>	26hpf	0	24	0	E.D. Fig. 5h	
<i>jam2a</i> MOex5	<i>cmyb</i>	36hpf	17	4	0	E.D. Fig. 5j	
<i>jam2a</i> MOex5	<i>rag1</i>	4dpf	17	0	0	E.D. Fig. 5l	

'Down' and 'Up' denote downregulation and upregulation of each gene, respectively. *The migration of *fli1*⁺ cells is delayed. **The expression of *runx1* is partially restored. ***The expression of *runx1* is almost fully restored. E.D. Fig., Extended Data Figure.

Supplementary Table 2. Primer sequences for cDNA cloning, RT-PCR, and qPCR

Gene symbol	GenBank Accession #	Primers for cDNA cloning	
		Forward	Reverse
<i>jam1a (f11r)</i>	NM_001004667	TCGTCACAGGCATACATGGT	GAATCCGTTAGCCTCATCCA
<i>jam2a</i>	NM_001098264	AGTGGAGGTGCACGAGTTCT	CAGAACCACAGCTGAGACCA
<i>dll4</i>	NM_001079835	GCATTTTTCACGACCGTTTT	ATGATCCGTGCAAAGACCTC
<i>scl-5'</i>	NM_213237	AAACCTCAGACATGGATGACC	CTTGAGCTCCCTGCCTTTTA
<i>cdh17</i>	NM_194422	ATGAAAGCTGAAGCCAAGGA	AATGACGCCGAATACTCCTG
Primers for RT-PCR			
		Forward	Reverse
<i>jam1a (f11r)</i>	NM_001004667	GGTGTCTGGAAGTGGTGGAT	AAAAGCCAATATGCAAAATGAA
<i>jam2a</i>	NM_001098264	GATCGCGCAGATATTGAAGG	TCGGCTGAAATATCCCTGTC
<i>ef1a</i>	NM_131263	CAAGGAAGTCAGCGCATACA	TGATGACCTGAGCGTTGAAG
Primer for qPCR			
		Forward	Reverse
<i>jam1a (f11r)</i>	NM_001004667	ATTTGGAGTTCCCGAGTGTG	TGGCAGATATCCCTTTTTCG
<i>jam2a</i>	NM_001098264	GCATCGAGTGGAAGAGGAAA	GGTTTCTCCAGTGAGATGG
<i>ef1a</i>	NM_131263	ACCGGCCATCTGATCTACAA	CAATGGTGATACCACGCTCA
<i>notch1a</i>	NM_131441	AGCATGATGTGAACGAGTGC	CCGTTCTTACAGGGAGACGA
<i>notch1b</i>	NM_131302	AAACGACGCCACCTGTTTAG	GCACTCGCATTGGTAGTTGTT
<i>notch3</i>	NM_131549	TGTGACCTAGACCGAAACGA	CACACAGGTGCCCTGATTTA
<i>runx1</i>	NM_131603	CAAGTGCCACATATCAGAG	CTGCTCAGAGAAAGCTAACG