

Table S1. Putatively conserved lincRNAs between *Homo sapiens* and *Branchiostoma lanceolatum*. Analysis of the genes surrounding the lincRNA focusing on three genes (upstream2, upstream1 and lincRNA). Some lincRNAs can be adscribed to more than one hypothetically conserved microsyntenic clusters.

Orthologous lincRNAs ¹	State of the cluster in human ²	Human orthologous lincRNA ³
BL02382 Sc0000007 7 + linc	Synteny conserved in the coding genes	AC005258.2
BL57655 Sc0000007 114 + linc	*Conserved microsynteny	LOC100505666
BL58800 Sc0000008 5 - linc	One gene with the strand inverted	LINC01124 AC007405.1
BL62558 Sc0000008 145 - linc	*Conserved microsynteny	TCONS_I2_00003921
BL42493 Sc0000014 4 + linc	*Conserved microsynteny	TCONS_00016098
BL39635 Sc0000017 32 - linc	Two genes with the strand inverted	LOC101448202
BL76639 Sc0000022 35 + linc	One gene with the strand inverted	SULT1C2P1
BL81207 Sc0000033 4 - linc	Synteny conserved in the coding genes	TCONS_00019766
BL63533 Sc0000034 21 + linc	*Conserved microsynteny but extra gene in human	LINC01877
BL79865 Sc0000034 22 - linc	*Conserved microsynteny but extra gene in human	TCONS_00002678
BL42364 Sc0000034 23 - linc	*Conserved microsynteny but extra gene in human	AC097717.1
BL34463 Sc0000054 53 - linc	Correct order but strands inverted	AL355338.2
BL56140 Sc0000081 24 - linc	Synteny conserved in the coding genes	AC012309.2
BL88002 Sc0000083 10 - linc	*Conserved microsynteny	AC007298.2
BL70460 Sc0000095 4 + linc	Two genes with the strand inverted	TCONS_00018032
BL35238 Sc0000141 19 - linc	One gene with the strand inverted	TCONS_00025555
BL43288 Sc0000247 8 + linc	Two genes with the strand inverted	NM_001302493.1
BL83080 Sc0000266 1 - linc	*Conserved microsynteny	AC115284.2
BL66360 Sc0000273 1 - linc	*Conserved microsynteny	AC012354.3
BL89644 Sc0000273 2 - linc	*Conserved microsynteny	LINC01833
BL26716 Sc0000273 3 - linc	*Conserved microsynteny	SIX3-AS1
BL68659 Sc0000384 9 - linc	*Conserved microsynteny	TCONS_00011917

¹GenID, Scaffold, virtual coordinates and strand separated by “|”. ²Description of the synteny of the cluster status in human. ³ID of the putative human orthologous lincRNA. * indicates a perfect match in strand and order of the three core genes.

Table S2. Putatively conserved lincRNAs between *Homo sapiens* and *Branchiostoma lanceolatum*. Analysis of the genes surrounding the lincRNA focusing on three genes (lincRNA, downstream1 and downstream2). Some lincRNAs can be adscribed to more than one hypothetically conserved microsyntenic clusters.

Orthologous lincRNAs ¹	State of the cluster in human ²	Human orthologous lincRNA ³
BL48403 Sc00000000 10 - linc	Syntenly conserved in the coding genes	AC004160.1
BL91553 Sc00000000 32 + linc	*Conserved microsyntenly	HOXB-AS1
BL83549 Sc00000000 42 - linc	Syntenly conserved in the coding genes	TCONS_00003443
BL49148 Sc00000004 180 + linc	Two genes with the strand inverted	HOXB-AS4
BL88007 Sc00000005 228 + linc	*Conserved microsyntenly	TCONS_00016695
BL88764 Sc00000006 154 + linc	Correct order but strands inverted	AC009159.2
BL21777 Sc00000012 28 + linc	*Conserved microsyntenly	TCONS_l2_00025400
BL43485 Sc00000012 77 + linc	*Conserved microsyntenly but extra gene in human	KCNJ2-AS1
BL38852 Sc00000015 95 + linc	Two genes with the strand inverted	TCONS_00025331
BL23696 Sc00000017 69 + linc	One gene with the strand inverted	TCONS_00003896
BL78928 Sc00000022 40 - linc	One gene with the strand inverted	AC100825.1
BL61801 Sc00000024 6 + linc	Two genes with the strand inverted	Z98257.1
BL41653 Sc00000050 7 + linc	Two genes with the strand inverted	AC092364.1
BL64814 Sc00000066 36 - linc	Syntenly conserved in the coding genes	TCONS_00014852
BL70524 Sc00000080 28 + linc	*Conserved microsyntenly	Z98880.1
BL84269 Sc00000083 14 + linc	*Conserved microsyntenly	AC114812.2
BL66046 Sc00000083 15 - linc	*Conserved microsyntenly	AC090001.1
BL62840 Sc00000091 33 + linc	*Conserved microsyntenly	LINC02410
BL90962 Sc00000097 13 - linc	Syntenly conserved in the coding genes	AL158013.1
BL67055 Sc00000174 7 - linc	Two genes with the strand inverted	AC012254.3
BL33122 Sc00000193 3 - linc	Syntenly conserved in the coding genes	AC006398.1
BL49481 Sc00000194 3 - linc	One gene with the strand inverted	LINC00210
BL57370 Sc00000219 7 + linc	*Conserved microsyntenly	AC092364.1
BL52347 Sc00000273 6 - linc	Syntenly conserved in the coding genes	LINC01438
BL52353 Sc00000273 7 - linc	*Conserved microsyntenly	TCONS_00022526
		AC093702.1

¹GenID, Scaffold, virtual coordinates and strand separated by “|”. ²Description of the syntenly of the cluster status in human. ³ID of the putative human orthologous lincRNA. * indicates a perfect match in strand and order of the three core genes.