nairpin	263 (20%)	53	2567	(44%)	1139	10945	(20%)	2217	8965	(41%)	3668	19007	(42%)	0538	38407	(35%)	13014	
Y-shaped	3 (67%)	2	49	(11%)	5	327	(30%)	99	120	(50%)	60	160	(33%)	52	659	(33%)	219	
clover-shaped	0 (n.a.)	0	0	(n.a.)	0	6	(80%)	5	1	(0%)	0	1	(100%)	1	8	(75%)	6	
complex shapes	0 (n.a.)	0	0	(n.a.)	0	0	(n.a.)	0	1	(0%)	0	0	(n.a.)	0	1	(0%)	0	
any shape	266 (21%)	55	2616	(44%)	1144	11278	(21%)	2321	9087	(41%)	3728	15828	(42%)	6591	39075	(35%)	13839	
The fold counts, estimated true positive rate (in parenthesis), and estimated true																		

intronic

intergenic

any location

coding

Table S2: Count statistics for short fold classes

positive counts are given for each location/shape-class of short folds. The "any shape" row and the "any location" column gives the marginalized counts for each set of fold classes. The entry at the lower right corner thus holds the overall counts for the set

5'UTR

3'UTR

type\location

of long folds.