

Table 5. Length of transcripts <10 kb

miRNA/cluster	Chromosome (Strand)	Genomic coordinates of pre-miRNA/cluster	Approximate 5' boundary of the primary transcript	Approximate 3' boundary of the primary transcript	Approximate length (bp) of the primary transcript	Supporting evidences
hsa-mir-200c~141	12 (+)	6943123-6943615	6942737	6943865	1128	TSS,ESTs,Ditags,polyA
hsa-mir-497-195	17 (-)	6861658-6862065	6861211	6863698	2487	ESTs,5'CAGE
hsa-mir-34b~34c	11 (+)	11088873-110889450	110888630	110889820	1190	TSS,CpG,5'CAGE,ESTs,polyA
hsa-mir-29b-2~29c	1 (-)	206041820-206042491	206041489	206046102	4613	ESTs,5'CAGE,Ditags,polyA
hsa-mir-572	4 (+)	10979549-10979643	10979348	10984460	5112	TSS,CpG,5'CAGE,Ditags,ESTs,polyA
hsa-mir-124a-1	8 (-)	9798308-9798392	9794986	9800634	5648	TSS,CpG,5'CAGE,ESTs,polyA
hsa-mir-99b~let-7e~125a	19 (+)	56887677-56888404	56885133	56888521	3388	TSS,CpG,5'CAGE,ESTs
hsa-mir-424~503	X (-)	133508407-133508094	133508008	133511322	3314	TSS,CpG,5'CAGE,ESTs
hsa-mir-200b~200a~429	1 (+)	1092347-1094330	1088033	1094500	6467	TSS,CpG,5'CAGE,Ditags,ESTs
hsa-mir-223	X (+)	65155437-65155546	65152025	65156338	4313	ESTs,5'CAGE,polyA
hsa-mir-23a~27a~24~2	19 (-)	13808101-13808473	13804510	13808884	4374	ESTs,polyA
hsa-mir-219-2	9 (-)	130194718-130194814	130193109	130195318	2209	TSS,CpG,5'CAGE,Ditags,ESTs
hsa-mir-106a~20b~19b-2~92-2~363	X (-)	133131074-133131974	133129103	133133925	4822	TSS,CpG,polyA
hsa-mir-802	21 (+)	36014883-36014976	36009993	36018276	8283	Ditags
hsa-mir-142	17 (-)	53763592-53763678	53763264	53764936	1672	TSS,5'CAGE,ESTs,polyA
hsa-let-7i	12 (+)	61283733-61283816	61283559	61284132	573	TSS,CpG,ESTs,Ditags
hsa-mir-493	14 (+)	100405150-100405238	100401271	100409002	7731	ESTs, Ditags
hsa-mir-210	11 (-)	558089-558198	555660	558587	2927	TSS,CpG,5'CAGE,ESTs,polyA
hsa-mir-92b	1 (+)	153431592-153431687	153431072	153432003	931	TSS,CpG,5'CAGE,ESTs,Ditags