

Table 6. 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-193b~365-1 | 16 (+) | 14305325-14310729 | 14303596 | 14315047 | 11451 | TSS,CpG,5'CAGE,ESTs,polyA |
| hsa-mir-374~545 | X (-) | 73423664-73423917 | 73416950 | 73429102 | 12152 | TSS,5'CAGE,ESTs,polyA |
| hsa-mir-15a-16-1 | 13 (-) | 49521110-49521338 | 49516048 | 49554190 | 38142 | TSS,CpG,ESTs,polyA |
| hsa-mir-181c~181d | 19 (+) | 13846513-13846825 | 13837066 | 13848576 | 11510 | TSS,CpG,ESTs,polyA |
| hsa-let-7a-1~7f-1~7d | 9 (+) | 95978060-95981023 | 95968554 | 95981591 | 13037 | TSS,CpG,5'CAGE,ESTs,polyA |
| hsa-mir-183~96~182 | 7 (-) | 129197459-129202090 | 129191947 | 129207520 | 15573 | TSS,CpG,5'CAGE,ESTs |
| hsa-mir-9-2 | 5 (-) | 87998427-87998513 | 87996020 | 88006038 | 10018 | TSS,CpG,5'CAGE,ESTs,polyA |
| hsa-mir-222~221 | X (-) | 45490529-45491474 | 45489851 | 45595877 | 106026 | TSS,CpG,5'CAGE,ESTs,polyA |