

Additional file 4.

Figure S1. The length distribution of unannotated sequences in non-pregnant and pregnant libraries

Figure S2. The stem loop structures of precursors of predicted miRNA candidates.
The red line parts indicate the mature sequences. (A)Non-pregnant (B) Pregnant

Figure S3. Analysis workflow of the Solexa sequencing results

Figure S1

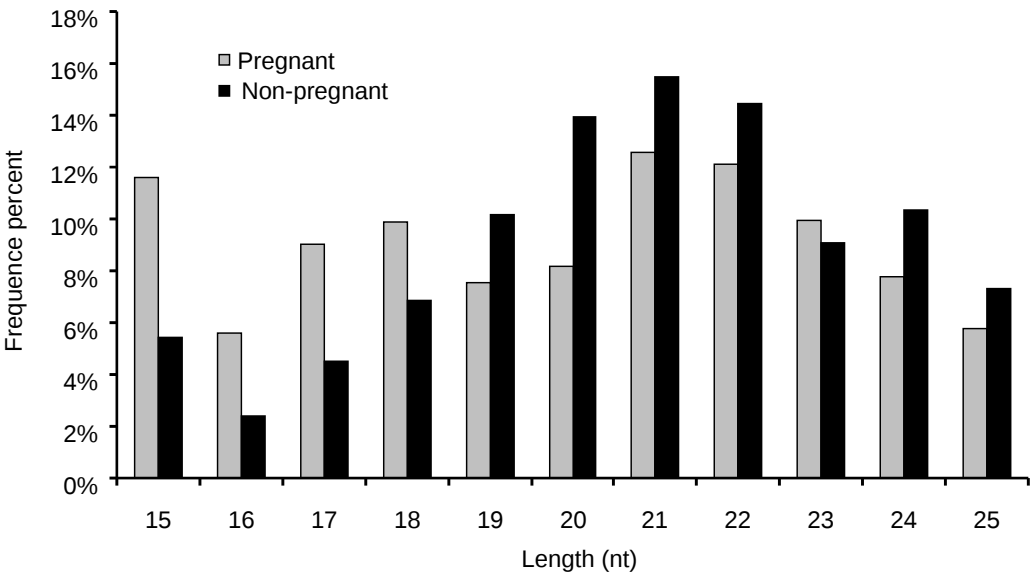
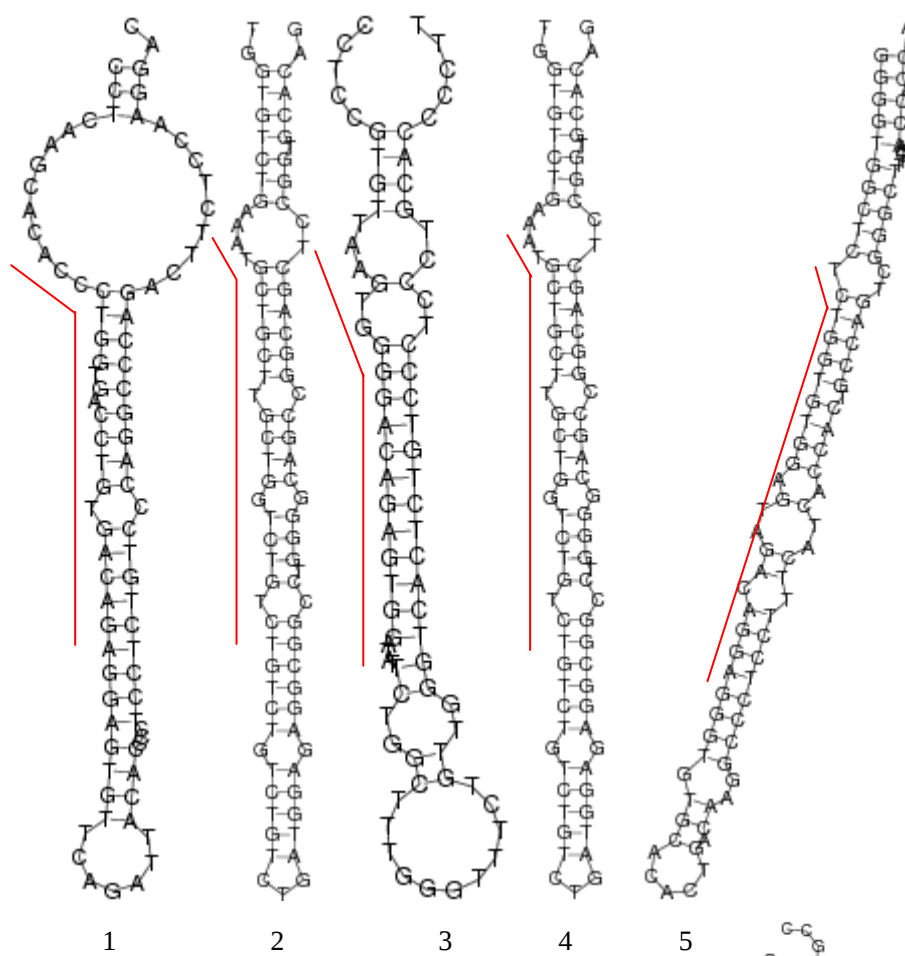


Figure S2

A



B

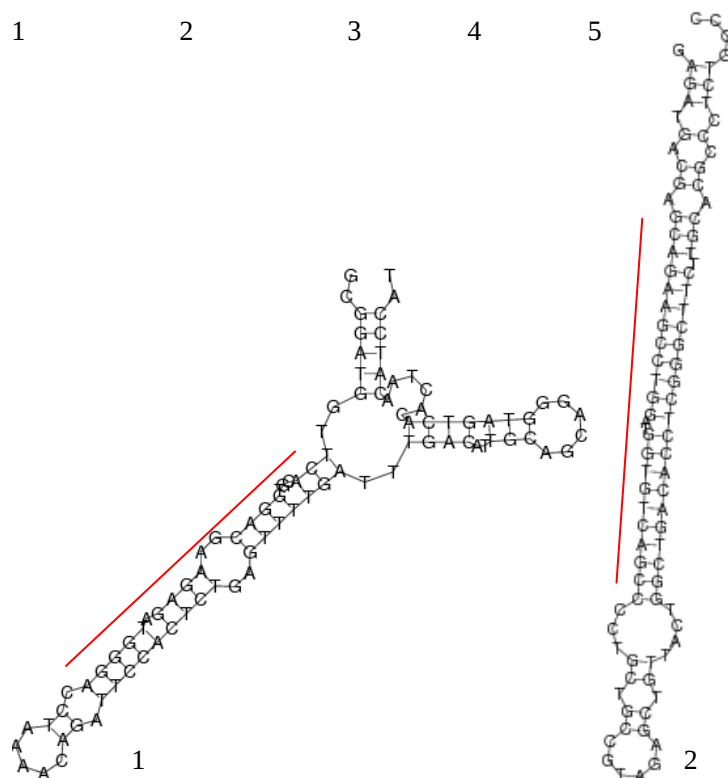


Figure S3

