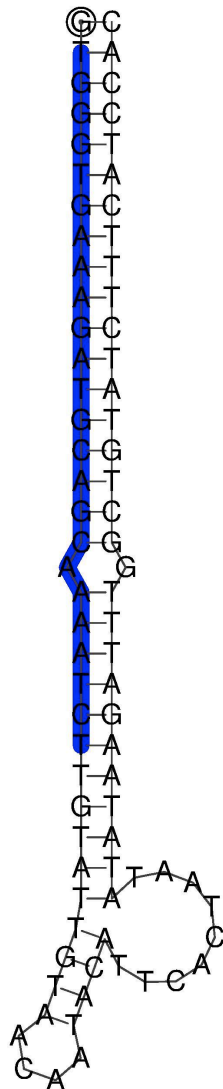


Figure S1. Predicted secondary stem-loop structures of the litchi-specific miRNAs.

Name: lch-miRC1

Sequence: TGGGTGAAAGATGCAGCAAAATCT

Normalized abundance: 414.24



Name: lch-miRC2

Sequence: TTCGATTTCGAACCCAGAGATGTCT

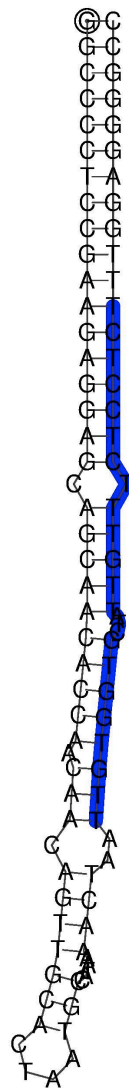
Normalized abundance: 216.73



Name: lch-miRC3

Sequence: TTGTGGTGCTATTGTTTCTCCTCT

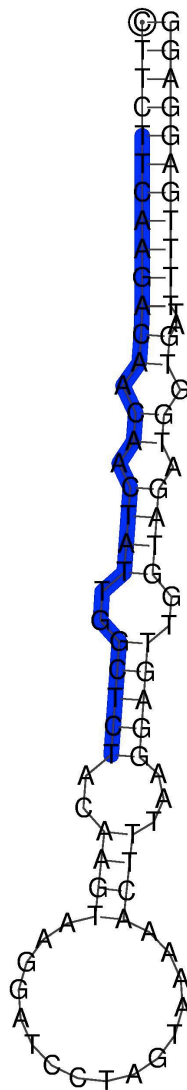
Normalized abundance: 115.11



Name: lch-miRC4

Sequence: TTCAAGACAACAACACTATTGGCTCT

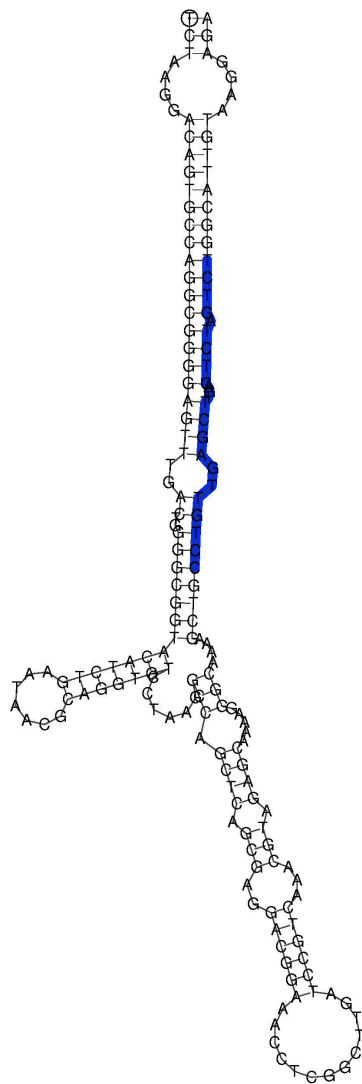
Normalized abundance: 270.65



Name: lch-miRC5

Sequence: CCTGTTGAGCTTGACTCTAGTCT

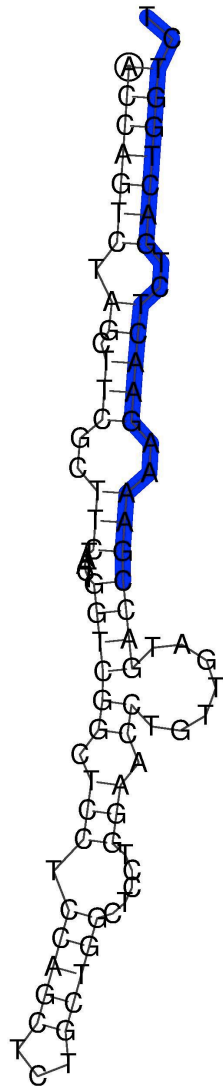
Normalized abundance: 66.62



Name: lch-miRC6

Sequence: CGAAAAGAACTCTGACTGGTCT

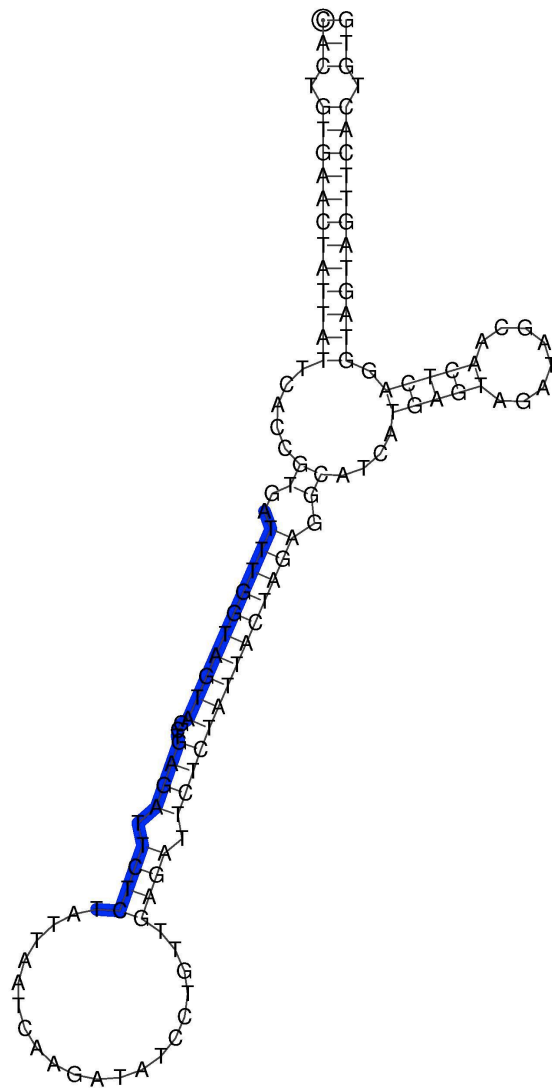
Normalized abundance: 107.62



Name: lch-miRC7

Sequence: ATTTGGTAGTAGCTGAGATTCTCT

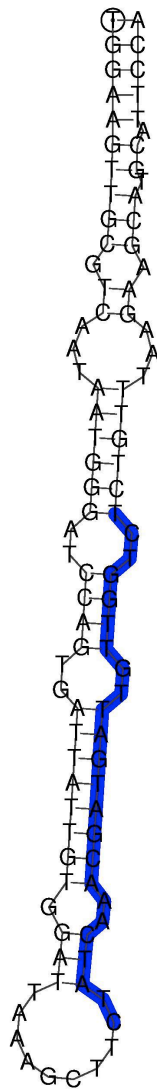
Normalized abundance: 46.58



Name: lch-miRC8

Sequence: CTATCAAACGATGATTGTTGGTCT

Normalized abundance: 23.58

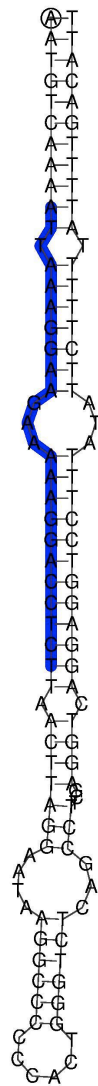




Name: lch-miRC9

Sequence: ATTAAAGGAAGAAAAGGACCTCT

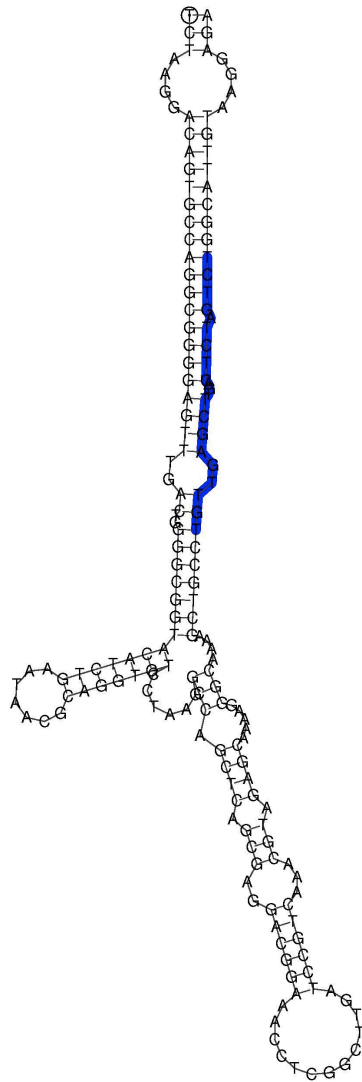
Normalized abundance: 19.18



Name: lch-miRC10

Sequence: TGTTGAGCTTGACTCTAGTCT

Normalized abundance: 38.60



Name: lch-miRC11

Sequence: CGGAGAAGGGCAATTACTCATTCT

Normalized abundance: 7.87

