Supporting Information

Logic-signal-based Multiplex Detection of MiRNAs with High Tension Hybridization and Multiple Signal Amplification

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Figure S1. Electrophoresis analysis for the formation of Y-DNA. Lane 1: Y_{1-210} , Lane 2: Y_{2-210} , Lane 3: Y_{3-210} , Lane 4: Y_{210} -DNA, Lane 5: Y_{1-21} , Lane 6: Y_{2-21} , Lane 7: Y_{3-21} , Lane 8: Y_{21} -DNA, Lane 9: Y_{1-155} , Lane 10: Y_{2-155} , Lane 11: Y_{3-155} , Lane 12: Y_{155} -DNA, Lane 13: Marker

2. Simultaneously Detection of miR-155 and miR-21



Figure S2. (a) Emission spectra before and after the addition of miR-155 and miR-21. (b) Emission intensity of the output at 570 nm from the four input combinations. (d) Logic scheme and truth table of the integrated INH and OR gates. The excitation wavelength is 380 nm.



Figure S3. (a) Emission spectra before and after the addition of miR-21 and miR-210. (b) Emission intensity of the output at 610 nm from the four input combinations. (d) Logic scheme and truth table of the integrated INH and OR gates. The excitation wavelength is 380 nm.

- 4. DNAs and MiRNAs sequences
- 1. **AM**₁₅₅: 5'-NH₂-GCGAGTACCCCTATCACGATTAGCATTAATTTCGACACTCGC-NH₂- 3'
- 2. AM₂₁₀: 5'-NH₂-GCGAGTTCAGCCGCTGTCACACGCACAGTTTCGAGACTCGC-NH₂- 3'
- 3. AM₂₁: 5'-NH₂-GCGAGTTCAACATCAGTCTGATAAGCTATTTCGAGACTCGC-NH₂- 3'
- 4. 17ES₁₅₅: 5'-CATCTCTTCTCCGAGCCGGTCGAAATTAATGCTAATC -3'
- 5. 17ES₂₁₀: 5'-TCATCTCTCCGAGCCGGTCGAAACTGTGCGTGTGA-3'
- 6. 17ES₂₁: 5'-CATCTCTTCTCCGAG CCGGTCGAAATAGCTTATCAGA -3'
- 7. **BF-SS**₁₅₅:
- 8. 5'-Biotin-TTTTTTTTTTTTTTTTAATrAGGAAGAGATGTACATAGTAGTG-Cy3-3'
- 9. **BF-SS**₂₁₀: 5'-Biotin-TTTTTTTTACAGTrAGGAAGAGATGGTTAAAGCTTG-FAM- 3'
- 10. BT-SS₂₁:

5'-Biotin-TTTTTTTTTTTTGCTATrAGGAAGAGATGATTGAGTACATG-Tex red-3'

- 11. Y₁₋₁₅₅: 5'-GCGAGTCACTACTATGTACATCTCTTCCACTCGCTTGCTGGATCCGCATGACA TTCGCCGTAAGT-3'
- 12. Y₂₋₁₅₅: 5'-GCGAGTCACTACTATGTACATCTCTTCCACTCGCACTTACGGCGAATGACCGA ATCAGCCT-3'
- 13. Y₃₋₁₅₅: 5'-GCGAGTCACTACTATGTACATCTCTTCCACTCGCAGGCTGATTCGGTTCATGC GGATCCAGCAA-3'
- 14. Y₁₋₂₁₀: 5'-GCGAGTCAAGCTTTAACCATCTCTTCCACTCGCTTGCTGGATCCGCATGACAT TCGCCGTAAGT-3'
- 15. Y₂₋₂₁₀: 5'-GCGAGTCAAGCTTTAACCATCTCTTCCACTCGCACTTACGGCGAATGACCGAA TCAGCCT-3'
- 16. Y₃₋₂₁₀: 5'-GCGAGTCAAGCTTTAACCATCTCTTCCACTCGCAGGCTGATTCGGTTCATGCG GATCCAGCAA-3'
- 17. Y₁₋₂₁: 5'-GCGAGTCATGTACTCAATCATCTCTTCCACTCGCTTGCTGGATCCGCATGACA TTCGCCGTAAGT-3'
- 18. Y₂₋₂₁: 5'-GCGAGTCATGTACTCAATCATCTTCTCCACTCGCACTTACGGCGAATGACCGA

ATCAGCCT-3'

- 19. Y₃₋₂₁: 5'-GCGAGTCATGTACTCAATCATCTCTTCCACTCGCAGGCTGATTCGGTTCATGC GGATCCAGCAA-3'
- 20. miR-155: UUAAUGCUAAUCGUGAUAGGGGU
- 21. 1-mis-155: UUAAUGCUAAUCGUGAUAGCGGU
- 22. miR-210: CUGUGCGUGUGACAGCGGCUGA
- 23. miR-21: UAGCUUAUCAGACUGAUGUUGA