

Electronic Supplementary Material

Rapid selection of aptamers based on Protein Microarray

Fang Yu ^a, Hui Li^{*a}, Wei Sun^b, Danke Xu^{*a}, Fuchu He^{*a,b}

^a *State Key Laboratory of Analytical Chemistry for Life Science, School of Chemistry and Chemical Engineering, Nanjing University, Nanjing, Jiangsu 210046, China*

^b *State Key Laboratory of Proteomics, National Center for Protein Sciences Beijing, Beijing Institute of Radiation Medicine, Beijing 102206, China*

Contents

Fig. S1 The fluorescence images of enriched aptamers on negative microarrays in every round of the first selection.

Table S1 The selected aptamers candidates for the first circle selection of Lac.

Fig. S2 The fluorescence images of enriched aptamers on negative microarrays in every round of the second selection.

Table S2 The selected aptamers candidates for the second circle selection of Lac.

Fig. S1 The fluorescence images of enriched aptamers on negative microarrays in every round of the first selection.

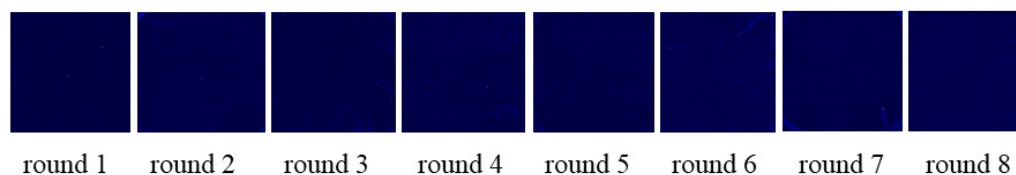


Table S1 The selected aptamers candidates for the first circle selection of Lac.

Aptamer-ID	Sequences of N40 from 5' to 3'	ΔG (kcal mol ⁻¹)	The sixth round	Truncated sequences of library from 5' to 3'	ΔG (kcal mol ⁻¹)
YFL-1	CGGTGCATCTATGGCTACTAG CTTTCTGCCTATACTAC	-2.27	2	AGGCAGGACACCGTAACCGGTGCATCTATGGCTACTAGCT TTTCTGCCT	-11.92
YFL-2	AATACTCCTGTTACCGTGCAT CTATGGCCATTGGCTTTTC	-2.26	2	GGCAGGACACCGTAACAATACTCCTGTTACCGTGCATCTAT GGCCATTGGCTTTTCTGCT	-9.35
YFL-3	CCTAACACGTACGGGGCATT ATGGCATAGCTCTTCTCC	-0.77	6	GGCAGGACACCGTAACCTAACACGTACGGGGCATTATG GCATAGCTCTTCTCCCTGCT	-7.16
YFL-4	ACGGGCTGATGCTCTCTTTAT TTTACCTAAATAAAGTGTC	-7.06	7	GCAGGACACCGTAACACGGGCTGATGCTCTCTTTATTTTAC CTAAATAAAGTGTCTGCTG	-14.3
YFL-5	CGGGCATTGCTCTCAATTTA GTCTCAAATGTGGCCTGC	-5.01	2	CAGGCAGGACACCGTAACCGGGCATTGCTCTCAATTTAG TCTCAAATGTGGCCTGCCTG	-14.04
YFL-6	CGGTGCATCTATGGCTACTAG CTTCTGCCTATACTAC	-2.27	18	AGGCAGGACACCGTAACCGGTGCATCTATGGCTACTAGCT CTTCTGCCT	-11.21
YFL-7	GGGTGCACACTCTTATTTTTA CACGAGCCAAAAATATGTC	-3.53	3	GCAGGACACCGTAACGGGTGCACACTCTTATTTTTACACG AGCCAAAAATATGTCTGCTG	-9.51
YFL-8	TCATCCCAAGTCCGGTGCCA TCTATGGGCTTCGCTTTTC	-2.47	3	GGCAGGACACCGTAACTCATCCCAAGTCCGGTGCCATCT ATGGGCTTCGCTTTTCTGCT	-8.19
YFL-9	CCCTAGTTCCTGGTGCATTTAT GGCAAAGCTTTTCTGCC	-2.75	3	AGGCAGGACACCGTAAACCCTAGTTCCTGGTGCATTTATG GCAAAGCTTTTCTGCC	-9.26
YFL-10	ACTGCTTTATCCCCGTCGGCTT GGCTCTTCGACAGTGTGG	-3.33	4	GCAGGACACCGTAAACTGCTTTATCCCCGTCGGCTTGGCT CTTCGACAGTGTGGCTGCTACC	-6.06
YFL-11	ACTGCTTTATCCCCGTCGGCTT GGCTCTTCGACAGTGTGG	-3.33	2	GGCAGGACACCGTAAACTGCTTTATCCCCGTCGGCTTGGC TCTTCGACAGTGTGGCTGCT	-5.83
YFL-12	ATTGAGCAGACGGGAGACCT TTAGAGTTGTAAGTTGAGT	-2.85	4	GAGCAGACGGGAGACCTTTAGAGTTGTAAGTTGAGTCTGC TA	-7.12
YFL-13	GGTGCATCCATGGCTTTTAGC TCTTCTGAACTGTACAC	-2.68	2	GACAGGACAGGACACCGTAAACCGGTGCATCCATGGCTTTAG CTTCTTCTGAACTGTACA	-10.7
YFL-14	GGTGCATCTATGGCTTTGCTC TTCTACCTGTTCTACGAG	-1.61	2	GACAGGACAGGACACCGTAAACCGGTGCATCTATGGCTTTGCT CTTCTACCTGTTCTACGAGCTGCTACCTCCCTCTCT	-8.58
YFL-15	GACCGGTTCTCGTAGGTGCGC GTATGCGGCTATGCTTTTC	-4.83	2	GGCAGGACACCGTAAACGACCGGTTCTCGTAGGTGCGCGTA TGCGGCTATGCTTTTCTGCT	-10.1
YFL-16	GGTGCCTGTACACGGCTTTGC TTTTCTGCCTCTGTTTAC	-4.19	2	CAGGCAGGACACCGTAAACGGTGCCTGTACACGGCTTTGCT TTTTCTGCCTC	-11.76

Fig. S2 The fluorescence images of enriched aptamers on negative microarrays in every round of the second selection.

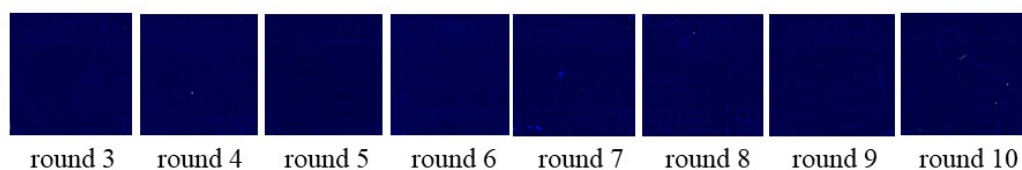


Table S2 The selected aptamers candidates for the second circle selection of Lac.

Aptamer-ID	Sequences of N40 from 5' to 3'	ΔG (kcal mol ⁻¹)	Truncated sequences of library from 5' to 3'	ΔG (kcal mol ⁻¹)	Repeat times		
					The sixth round	The seventh round	The eighth round
YFL-1	CGGTGCATCTATGGCTACTA GCTTTTCCTGCCTATACTAC	-2.27	AGGCAGGACACCGTAACCGGTGCATCTATGGC TACTAGCTTTTCCTGCCT	-11.92	2	1	2
YFL-4	ACGGGCTGATGCTCTCTTTA TTTACCTAAATAAAGTGTC	-7.06	GCAGGACACCGTAACACGGGCTGATGCTCTCT TTATTTTACCTAAATAAAGTGCTCTGC	-14.3	2	1	7
YFL-5	CGGGCATTGCTCTCAATTT AGTCTCAAATGTGGCCTGC	-5.01	CAGGCAGGACACCGTAACCGGGCATTGCTCT CAATTTAGTCTCAAATGTGGCCTGCCTG	-14.04	1	1	2
YFL-6	CGGTGCATCTATGGCTACTA GCTCTTCCTGCCTATACTAC	-2.27	AGGCAGGACACCGTAACCGGTGCATCTATGGC TACTAGCTCTTCCTGCCT	-11.21	0	4	4
YFL-17	GGGCTATGCTCTAAATCTT CCTACTGACGCAATTTGGGA	-1.17	CCGTAACGGGCTATGCTCTAAATCTTCCTACT GACGCAATTTGGACTGCTACCTCCCTCC	-2.66	2	2	1
YFL-7	GGGTGCACACTCTTATTTTT ACACGAGCCAAAAATATGT C	-3.53	GCAGGACACCGTAACGGGTGCACACTCTTATT TTTACACGAGCCAAAAATATGCTCTGC	-9.51	2	2	4
YFL-10	ACTGCTTTATCCCCGTCGGC TTGGCTCTTCGACAGTGTTG	-3.33	GCAGGACACCGTAACACTGCTTTATCCCCGTC GGCTTGGCTCTTCGACAGTGTTGGCTGCTACC	-6.06	11	5	17
YFL-18	AACAACCTCGTATCCGGTGC ATTTATGGCGAATGCTTTTC	-4.71	GCAGGACACCGTAACAACAACCTCGTATCCGG TGCATTTATGGCGAATGCTTTTCCTGC	-7.96	1	3	3
YFL-11	ACTGCTTTATCCCCGTCGGC TTGGCTCTTCGACAGTGTTG	-3.33	GCAGGACACCGTAACACTGCTTTATCCCCGTC GGCTTGGCTCTTCGAC	-5.12	4	3	10
YFL-19	ACTGCTTTATCCCCGTCGGC TTGTCTCTTCGACAGTGTTG	-3.33	GCAGGACACCGTAACACTGCTTTATCCCCGTC GGCTTGTCTCTTCGACAGTGTTGGCTGCTACC	-6.06	17	23	43
YFL-14	GGTGCATCTATGGCTTTGCT CTTCTACCTGTTCTACGAG	-1.61	ACAGGCAGGACACCGTAACGGTGCATCTATGG CTTTGCTCTTCCTACCTGTTCTACGAGCTGCTA CCTCCCTCCTC	-8.14	11	10	7
YFL-20	ACTGCTTTATCCCCGTCGGC TTGTCTCTTCGACAGTGTTG	-3.33	GCAGGACACCGTAACACTGCTTTATCCCCGTC GGCTTGTCTCTTCGAC	-5.21	5	5	12
YFL-21	GGGCGCACGCTCTGAATTTT TCATATGAAACAATTCTGTC	-5.97	GCAGGACACCGTAACGGGCGCACGCTCTGAAT TTTTCATATGAAACAATTCTGCTCTGC	-11.61	2	4	8
YFL-22	TCGCATCGGTGCATCTATGG CGCACGCTCTTCCTGCCTTC	-4.81	AGGCAGGACACCGTAACCTCGCATCGGTGCATC TATGGCGCACGCTCTTCCTGCCT	-10.71	1	0	3

YFL-23	TTCCGGATAATCAGACGATA TATGAGGAATTCTGAGGGTA	-2.05	CCGTAACTTCCGGATAATCAGACGATATATGA GGAATTCTGAGGGTACTGCTACCTCCCTC	-6.44	1	1	2
YFL-24	CTTTCAGGGTGGTGCCGTTT CGGGCTTTGTCTCTTCTGTC	-2.75	CAGGCAGGACACCGTAACCTTTCAGGGTGGTG CCGTTTCGGGCTTTGTCTCTTCTGCTG	-11.37	0	1	2
YFL-25	ACTGCTTTATCCCCGTCGGC TTGCTCTTCGACAGTGTG	-3.33	GCAGGACACCGTAACACTGCTTTATCCCCGTC GGCTTTGCTCTTCGAC	-5.12	1	1	1
YFL-26	CGGGCTTTAGCTCTAAAAAA TGTCACCATCTTTTTTGTCTGTC	-4.37	GCAGGACACCGTAACCGGGCTTTAGCTCTAAA AAATGTCCACCATCTTTTTTGTCTGTC	-10.7	0	1	2

Fig. S3 The secondary structures of six selected aptamers were simulated with OligoAnalyzer 3.1 at the minimum ΔG . (a) YFL-13, (b) YFL-16, (c) YFL-21, (d) YFL-22, (e) YFL-24, (f) YFL-26.

