

Supporting information for

Characterization and application of a DNA aptamer binding to Tryprophan

Xiaojuan Yang^{1,2}, Tao Bing^{1,2}, Hongcheng Mei^{1,2}, Canliang Fang^{1,2}, Zehui Cao¹, Dihua
Shangguan¹*

¹ Beijing National Laboratory for Molecular Sciences, Key Laboratory of Analytical Chemistry for Living Biosystems, Institute of Chemistry, Chinese Academy of Sciences, Beijing 100190, China

² Graduate School of the Chinese Academy of Science, Beijing 100039, China

Tel & Fax: +86 10 62528509 E-mail address: sgdh@iccas.ac.cn

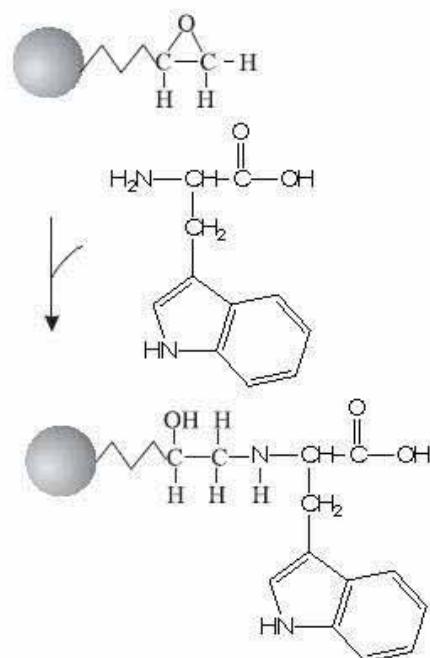


Figure S1. Immobilizton procedure of L-tryptophan to the Epoxy-activated Sepharose 6BTM

Table S1. The changes of the selection conditions during SELEX procedure

Round	positive beads (μL)	beads negative beads (μL)	Incubating time (min)	Washing (μL) × times	buffer
1th	200	0	60	100×2	
2th	60	20	60	100×2	
3th	20	20	30	100×2	
4th	20	40	30	150×3	
5th	20	60	30	150×3	
6th	15	60	30	150×3	
7th	15	60	30	150×4	
8th	10	60	5	150×4	
9th	10	60	5	200×4	
10th	10	60	5	200×4	
11th	10	60	5	200×4	
12th	10	60	5	200×4	

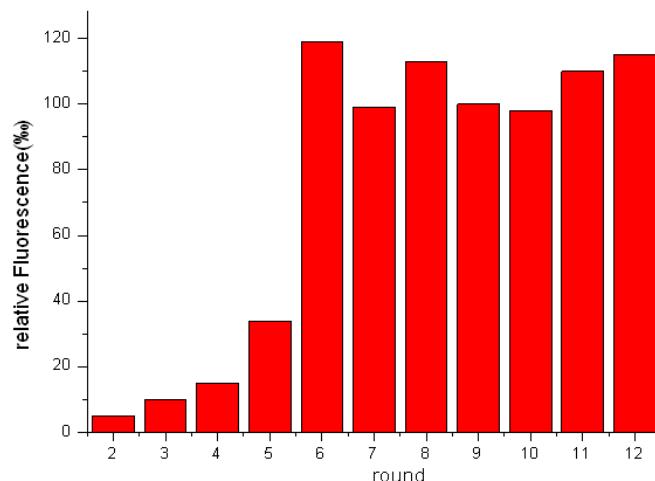


Figure S2. The progress of the selection. The bar represents the percentage of eluted ssDNA to the total amounts of ssDNA in each selection round. Relative Fluorescence=(F_e) $\times 1000\%/(F_0)$, where, F_e is the fluorescence intensity of the eluate. F_0 is the initial fluorescence intensity before binding.

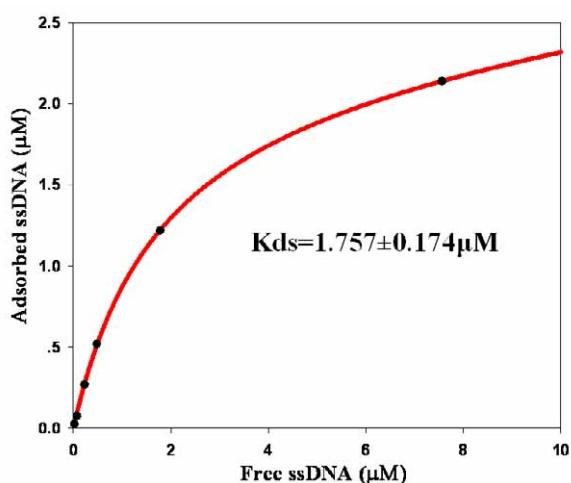


Figure S3. Binding curve of Trp3a-1 to L-tryptophan immobilized beads. Binding studies were performed with a fixed amount of L-tryptophan immobilized beads and a concentration series(0.05-10 μM) of the FAM-labelled aptamers. Binding curve was obtained and the K_d was calculated by sigmaplot 11 software