

**Cy5/BHQ dye–quencher pairs in fluorogenic qPCR probes: effects of charge and hydrophobicity**

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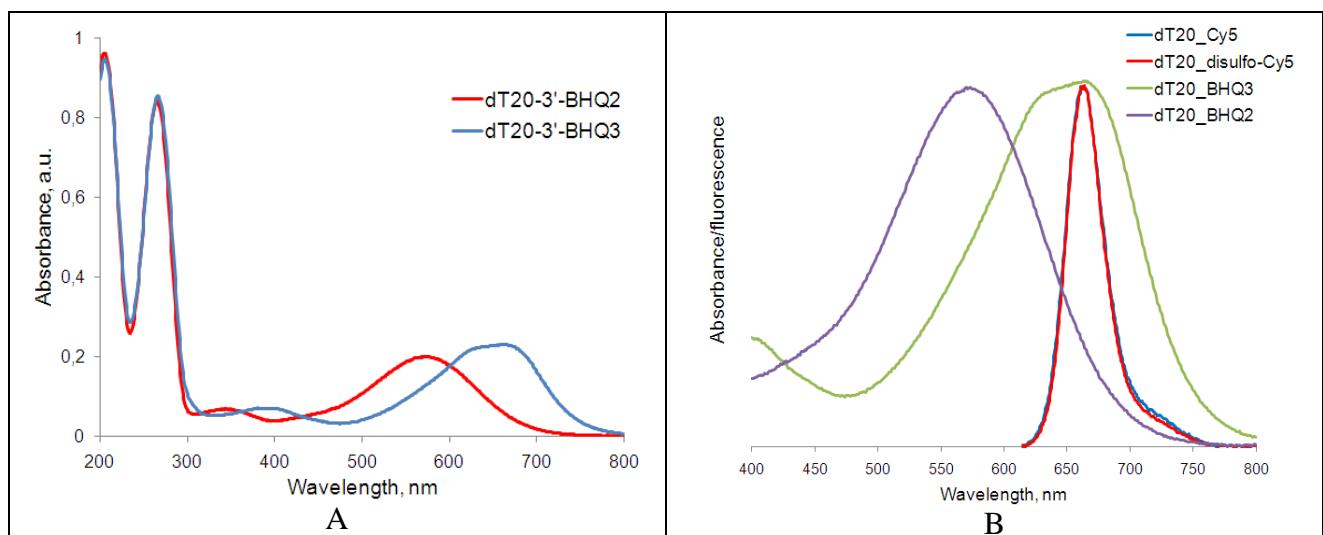
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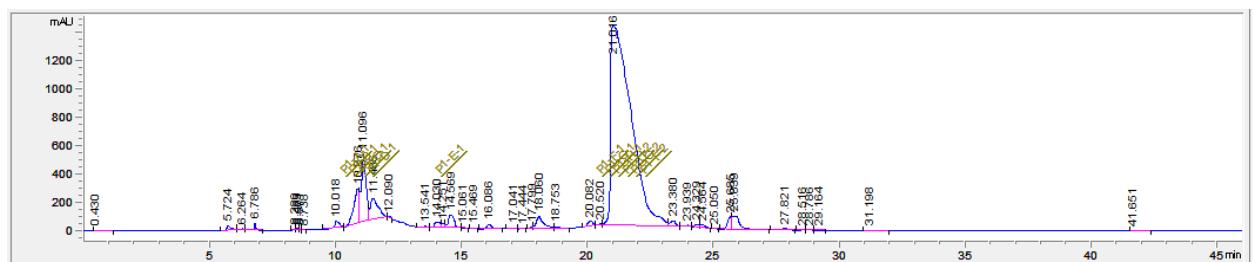
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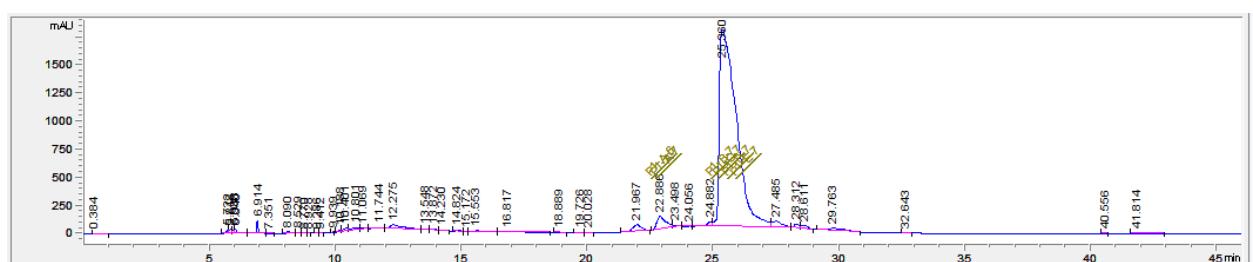
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**Fig. S1.** A) UV/Vis spectra of model T<sub>20</sub> oligonucleotides modified with BHQ2 and BHQ3 quenchers; B) normalized BHQ2 and BHQ3 absorbance vs Cy5 and disulfo-Cy5 fluorescence on model T<sub>20</sub> oligonucleotides.



**Fig. S2-1.** RP-HPLC traces of reaction mixture - sCy5-azide conjugation with unpurified G3z-BHQ2 oligonucleotide

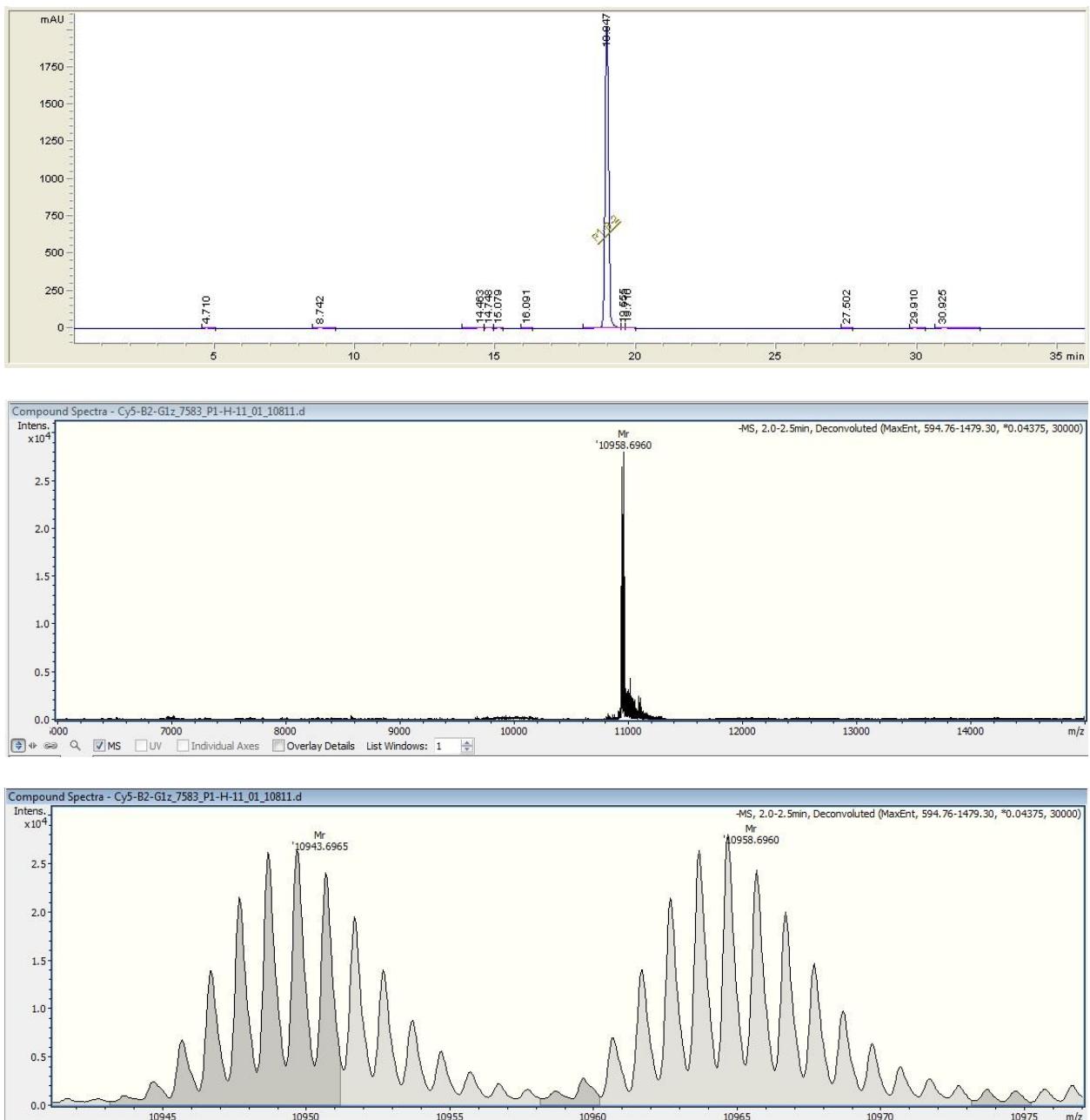


**Fig. S2-2.** RP-HPLC traces of reaction mixture - Cy5-azide conjugation with purified G3z-BHQ3 oligonucleotide

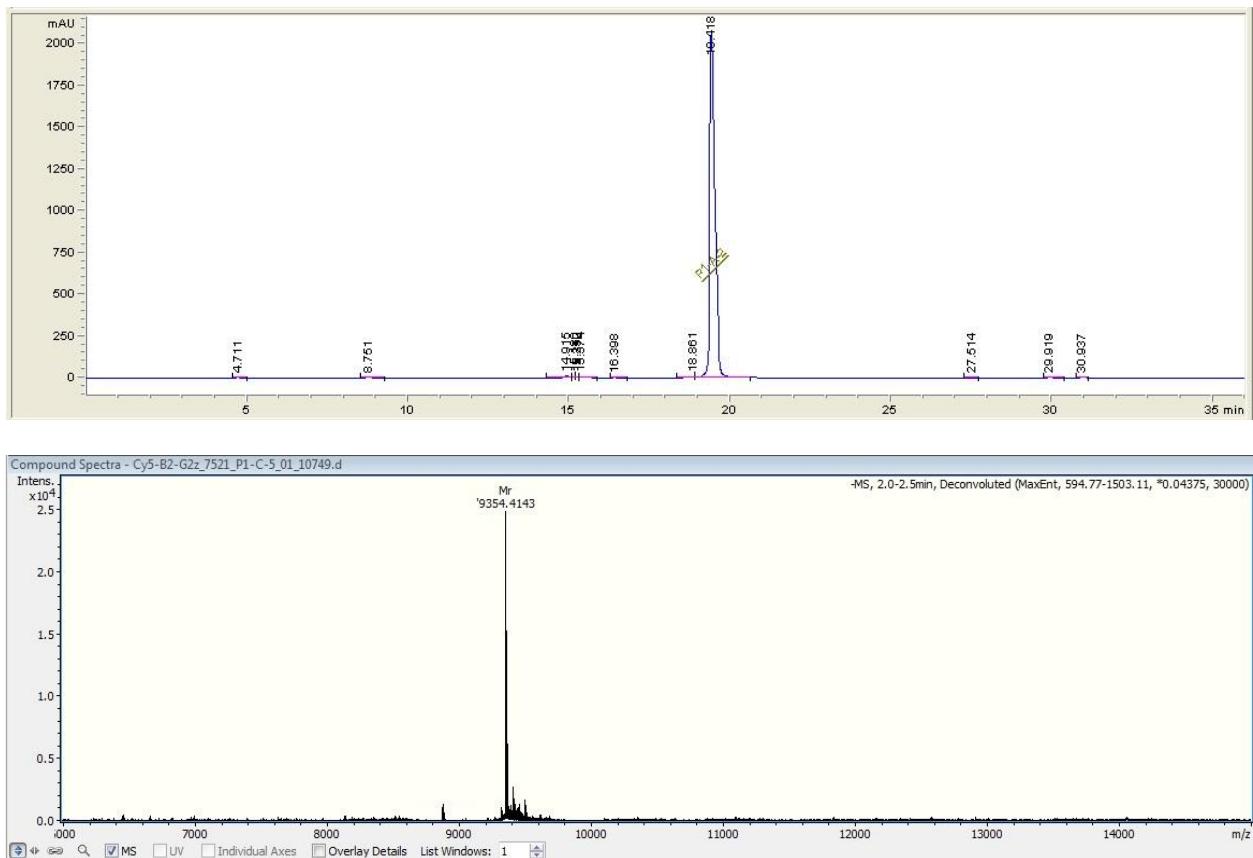
**Table S1.** Taqman probes and their ESI-MS data.

#	Name	Sequence, 5'→3'	Calculated mass	Detected mass
1	Cy5-B2-G1z	<b>Cy5</b> -CTGAGCTTAGTYAAGGCAAATAATGCTCAG- <b>BHQ2</b>	10936.2 10951.3	10943.7 10958.7
2	Cy5-B2-G2z	<b>Cy5</b> -CAGCGTCTAGTGATCCCGTTATTGGC- <b>BHQ2</b>	9346.2	9354.4
3	Cy5-B2-G3z	<b>Cy5</b> -TACCCAACGTAAAGCAGCAACAGGGTA- <b>BHQ2</b>	9375.3	9383.5
4	Cy5-B2-G4z	<b>Cy5</b> -CTGAACYTGTCTGGCCATCCTTGGTT- <b>BHQ2</b>	9297.2 9312.2	9305.4 9320.4
5	Cy5-B2-P4z	<b>Cy5</b> -CACTCTGACTACTACCTTAAACAGAGCG- <b>BHQ2</b>	10198.8	10206.5
6	Cy5-B2-P8z	<b>Cy5</b> -GGACTGCAGTCR TTGCTGTTAAC- <b>BHQ2</b>	8776.8 8792.8	8785.3 8801.3
7	sCy5-B2-G1z	<b>sCy5</b> -CTGAGCTTAGTYAAGGCAAATAATGCTCAG- <b>BHQ2</b>	11094.1 11109.2	11089.6 11104.6
8	sCy5-B2-G2z	<b>sCy5</b> -CAGCGTCTAGTGATCCCGTTATTGGC- <b>BHQ2</b>	9504.1	9500.3
9	sCy5-B2-G3z	<b>sCy5</b> -TACCCAACGTAAAGCAGCAACAGGGTA- <b>BHQ2</b>	9533.2	9529.3
10	sCy5-B2-G4z	<b>sCy5</b> -CTGAACYTGTCTGGCCATCCTTGGTT- <b>BHQ2</b>	9455.1 9470.1	9451.3 9466.3
11	sCy5-B2-P4z	<b>sCy5</b> -CACTCTGACTACTACCTTAAACAGAGCG- <b>BHQ2</b>	10356.7	10352.5
12	sCy5-B2-P8z	<b>sCy5</b> -GGACTGCAGTCR TTGCTGTTAAC- <b>BHQ2</b>	8934.7 8950.7	8931.2 8947.2
13	Cy5-B3-G1z	<b>Cy5</b> -CTGAGCTTAGTYAAGGCAAATAATGCTCAG- <b>BHQ3</b>	11196.7 11211.7	11203.9 11218.9
14	Cy5-B3-G2z	<b>Cy5</b> -CAGCGTCTAGTGATCCCGTTATTGGC- <b>BHQ3</b>	9606.6	9614.7
15	Cy5-B3-G3z	<b>Cy5</b> -TACCCAACGTAAAGCAGCAACAGGGTA- <b>BHQ3</b>	9635.7	9643.6
16	Cy5-B3-G4z	<b>Cy5</b> -CTGAACYTGTCTGGCCATCCTTGGTT- <b>BHQ3</b>	9557.6 9572.6	9565.5 9580.5
17	Cy5-B3-P4z	<b>Cy5</b> -CACTCTGACTACTACCTTAAACAGAGCG- <b>BHQ3</b>	10459.2	10466.8
18	Cy5-B3-P8z	<b>Cy5</b> -GGACTGCAGTCR TTGCTGTTAAC- <b>BHQ3</b>	9037.3 9053.3	9045.5 9061.5
19	sCy5-B3-G1z	<b>sCy5</b> -CTGAGCTTAGTYAAGGCAAATAATGCTCAG- <b>BHQ3</b>	11354.6 11369.6	11349.9 11364.9
20	sCy5-B3-G2z	<b>sCy5</b> -CAGCGTCTAGTGATCCCGTTATTGGC- <b>BHQ3</b>	9764.5	9760.4
21	sCy5-B3-G3z	<b>sCy5</b> -TACCCAACGTAAAGCAGCAACAGGGTA- <b>BHQ3</b>	9793.6	9789.5
22	sCy5-B3-G4z	<b>sCy5</b> -CTGAACYTGTCTGGCCATCCTTGGTT- <b>BHQ3</b>	9715.5 9730.5	9711.4 9726.4
23	sCy5-B3-P4z	<b>sCy5</b> -CACTCTGACTACTACCTTAAACAGAGCG- <b>BHQ3</b>	10617.1	10612.7
24	sCy5-B3-P8z	<b>sCy5</b> -GGACTGCAGTCR TTGCTGTTAAC- <b>BHQ3</b>	9195.2 9211.2	9191.4 9207.4

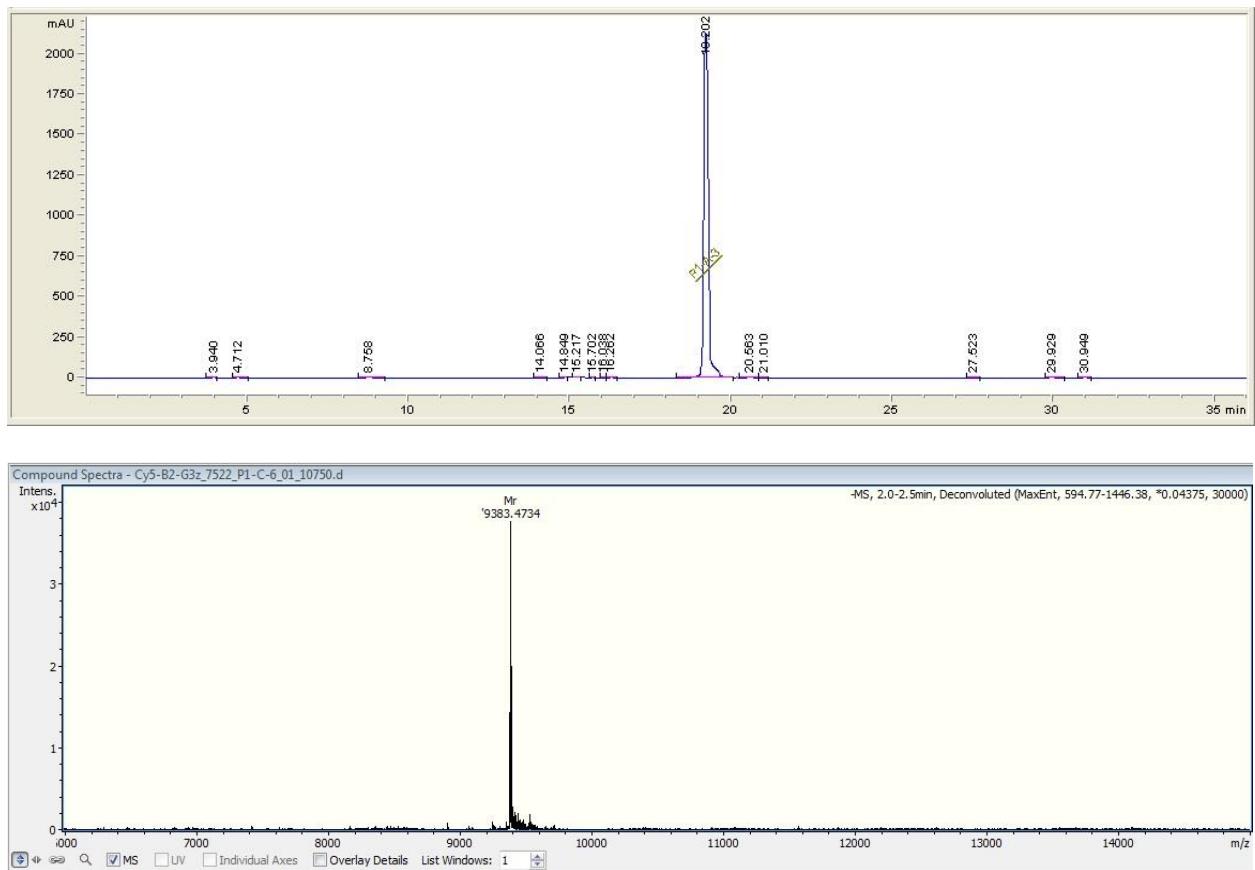
**Fig. S3-1.** HPLC profile and ESI MS of **Cy5-B2-G1z** probe.



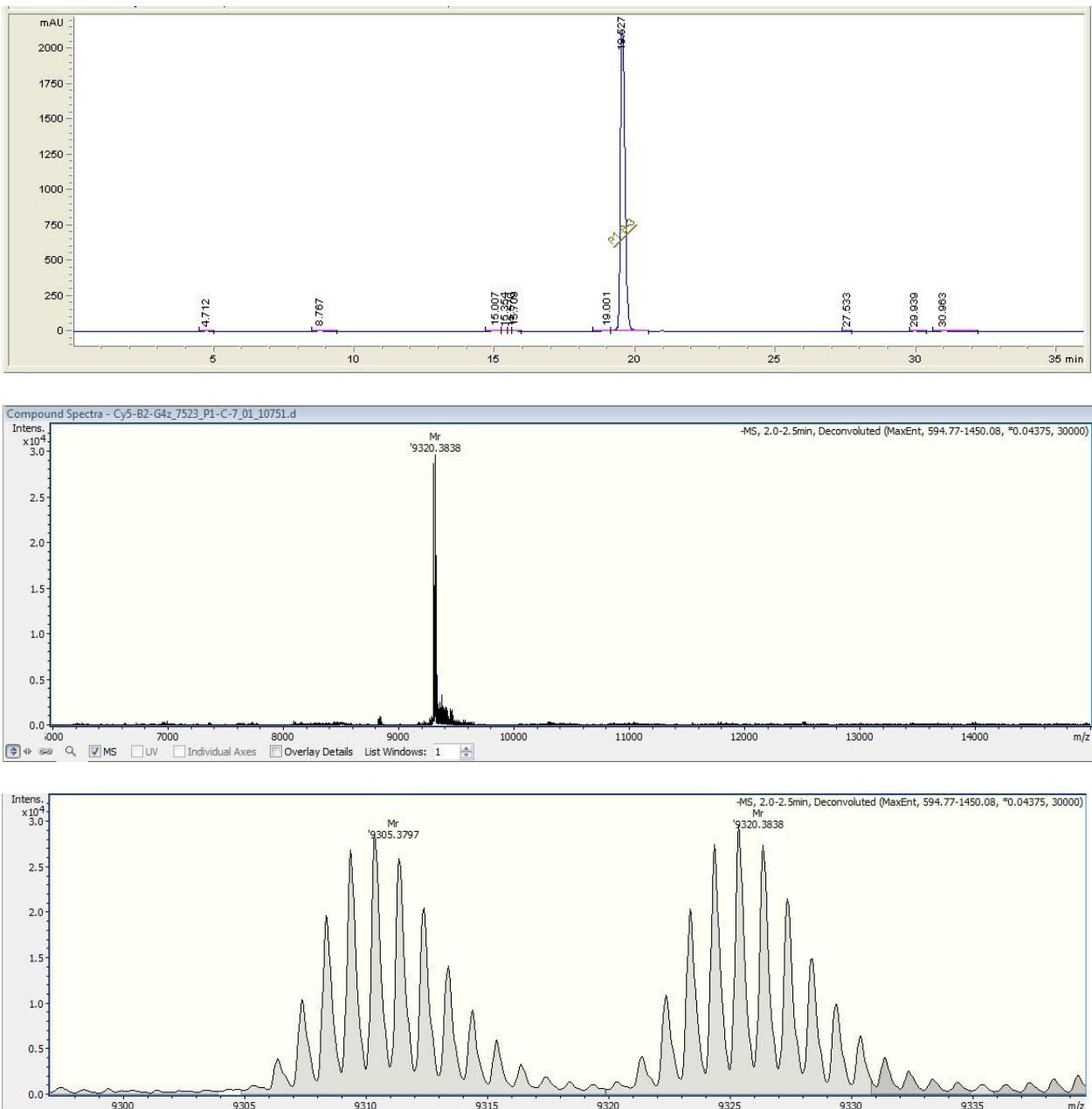
**Fig. S3-2.** HPLC profile and ESI MS of **Cy5-B2-G2z** probe.



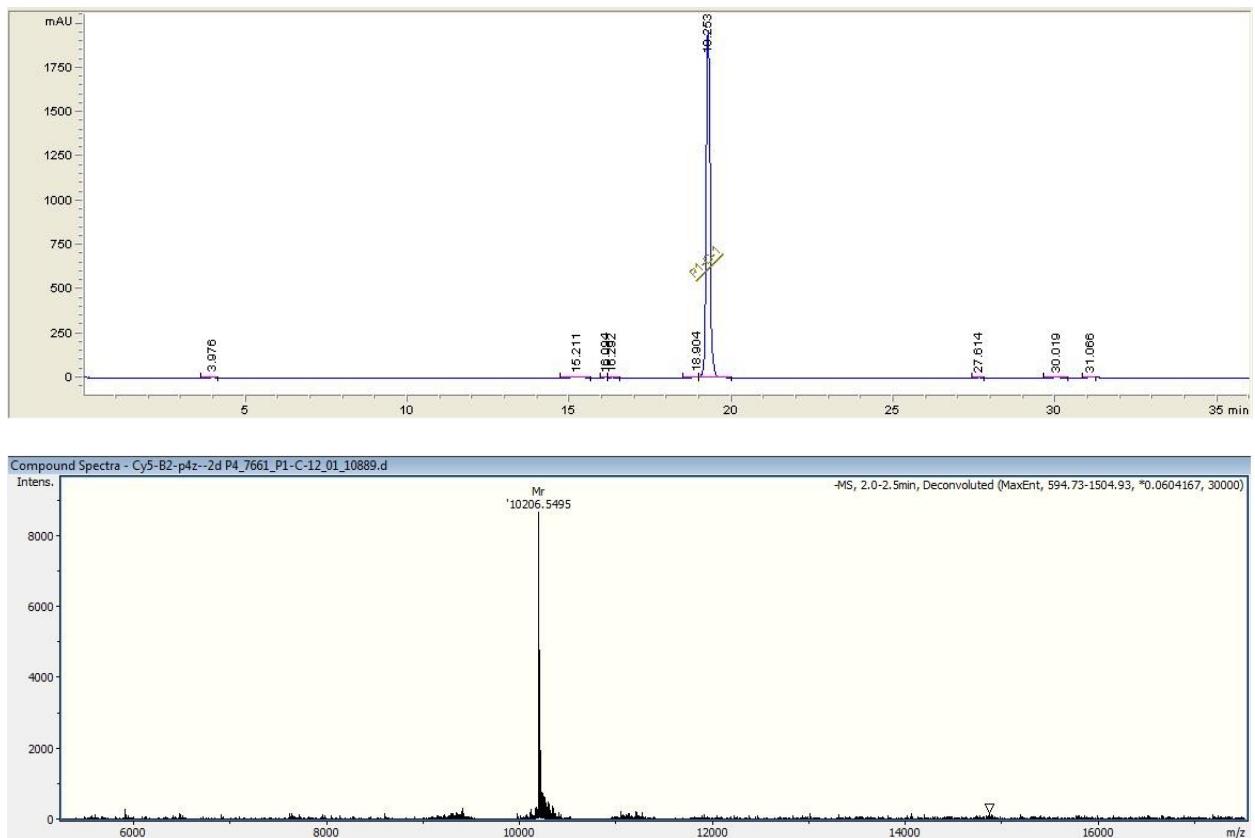
**Fig. S3-3.** HPLC profile and ESI MS of **Cy5-B2-G3z** probe.



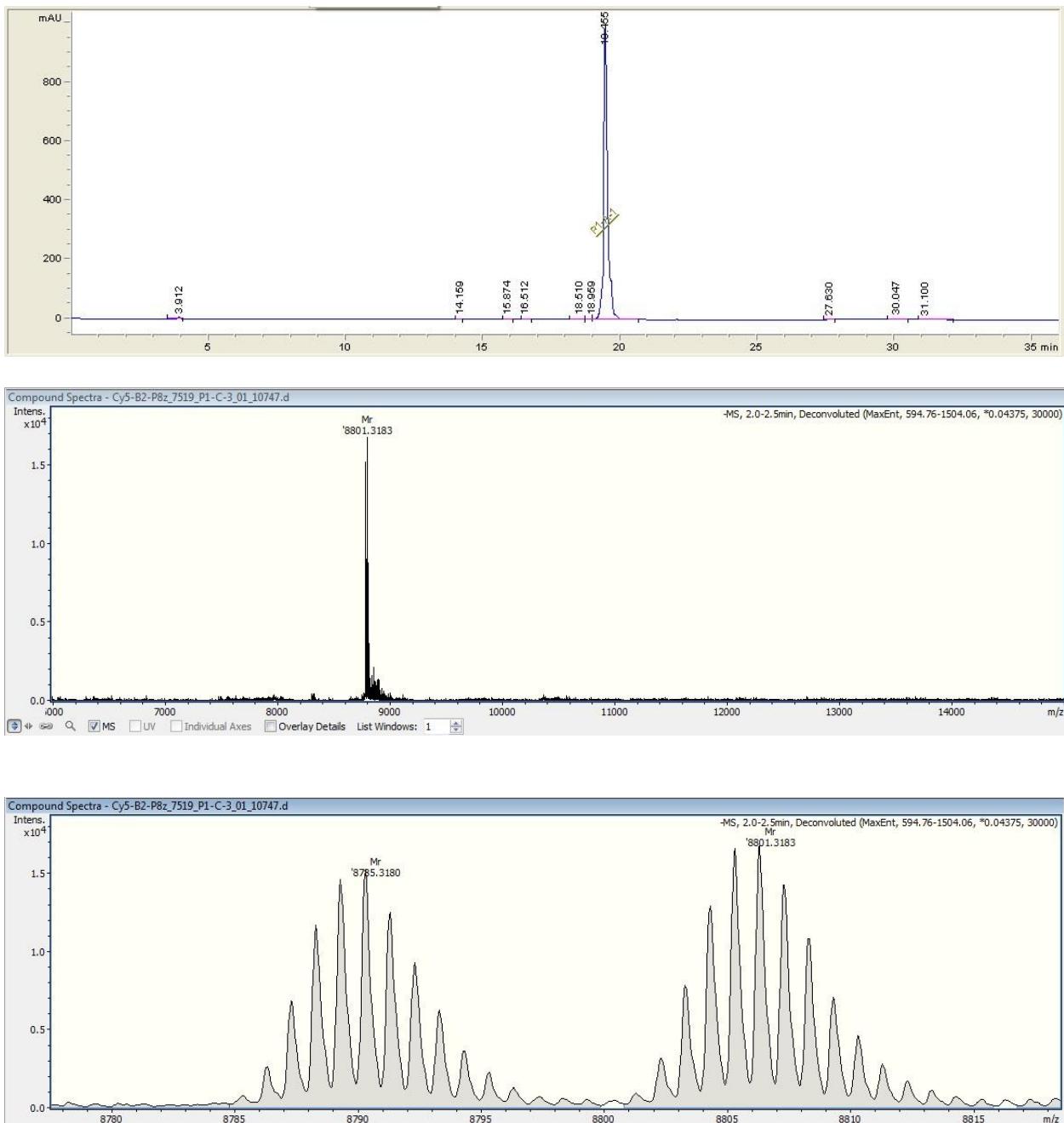
**Fig. S3-4.** HPLC profile and ESI MS of **Cy5-B2-G4z** probe.



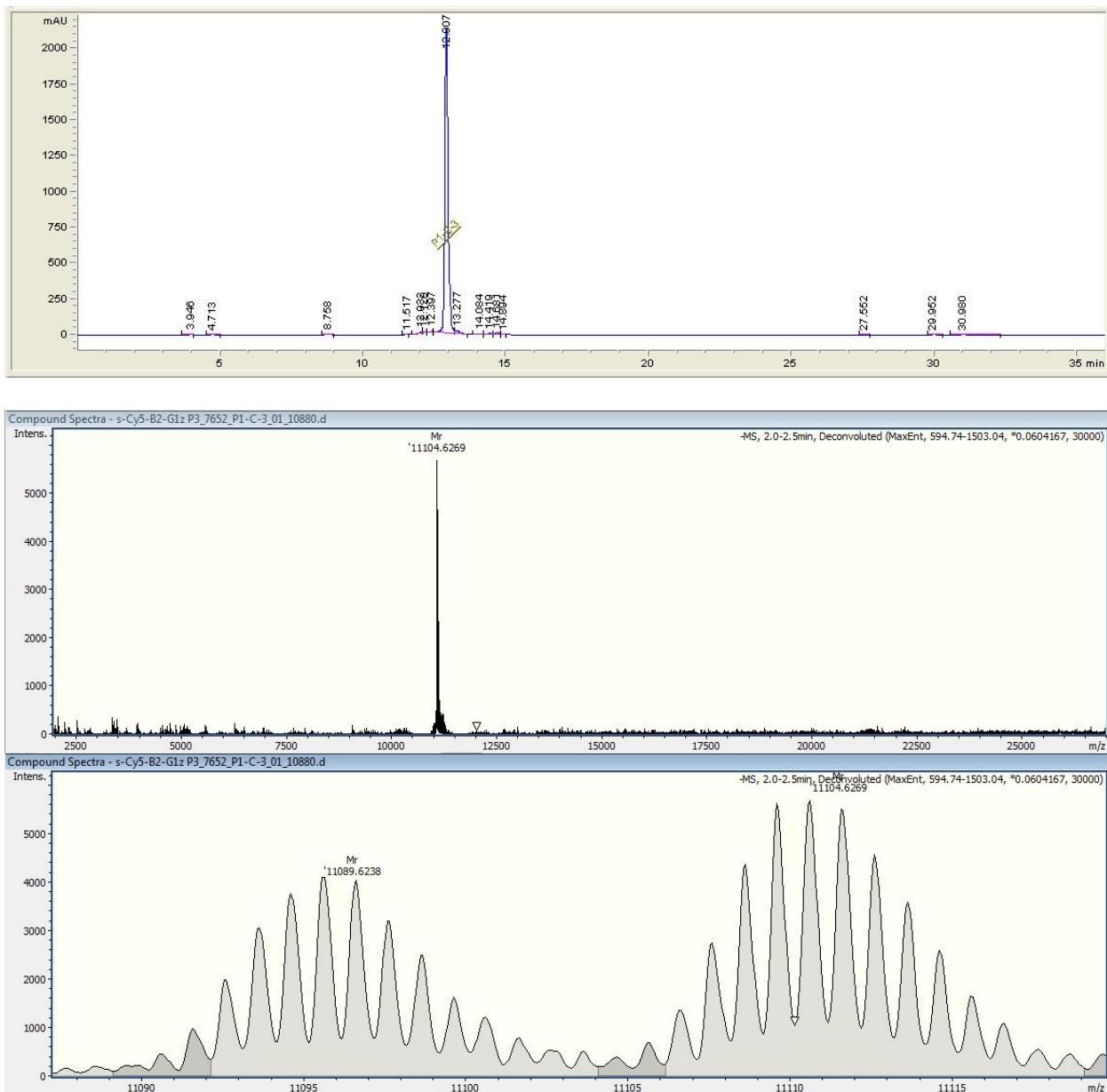
**Fig. S3-5.** HPLC profile and ESI MS of **Cy5-B2-P4z** probe.



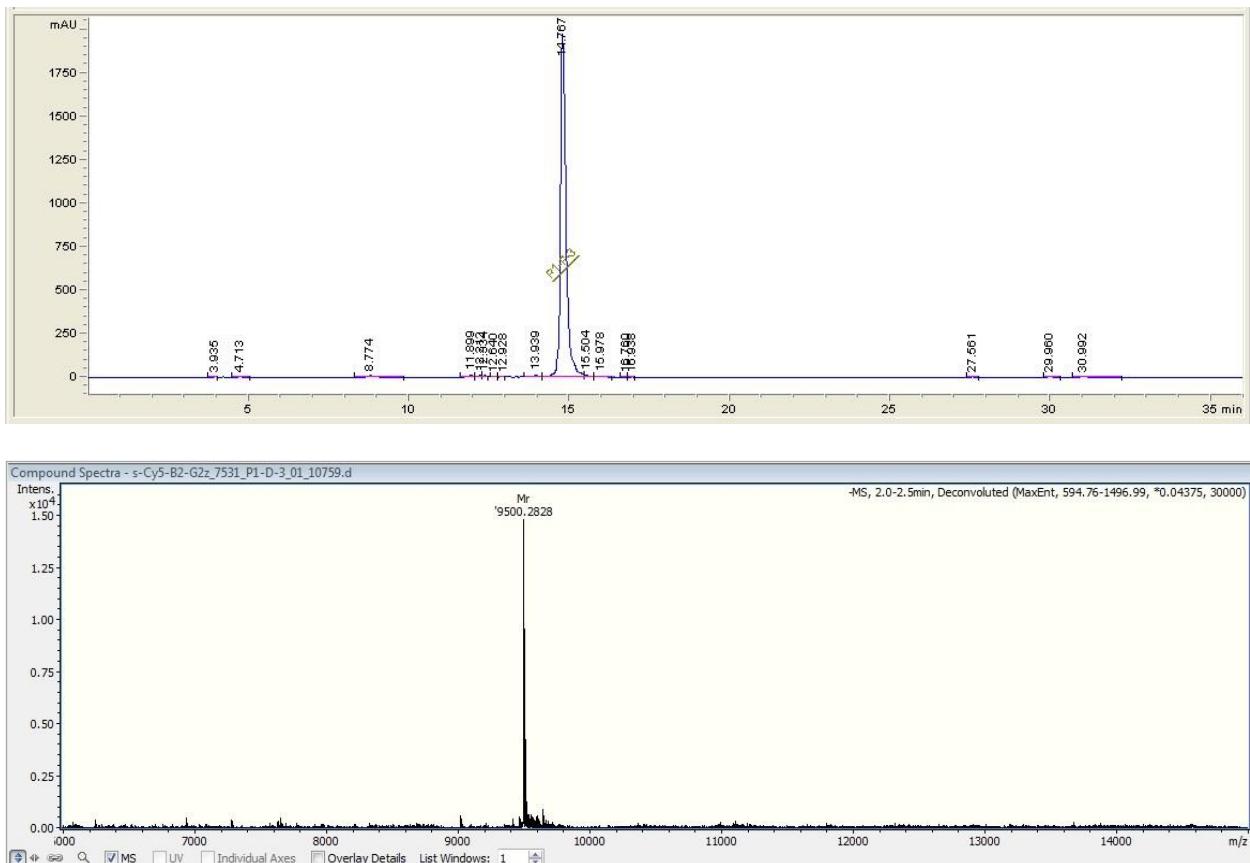
**Fig. S3-6.** HPLC profile and ESI MS of Cy5-B2-P8z probe.



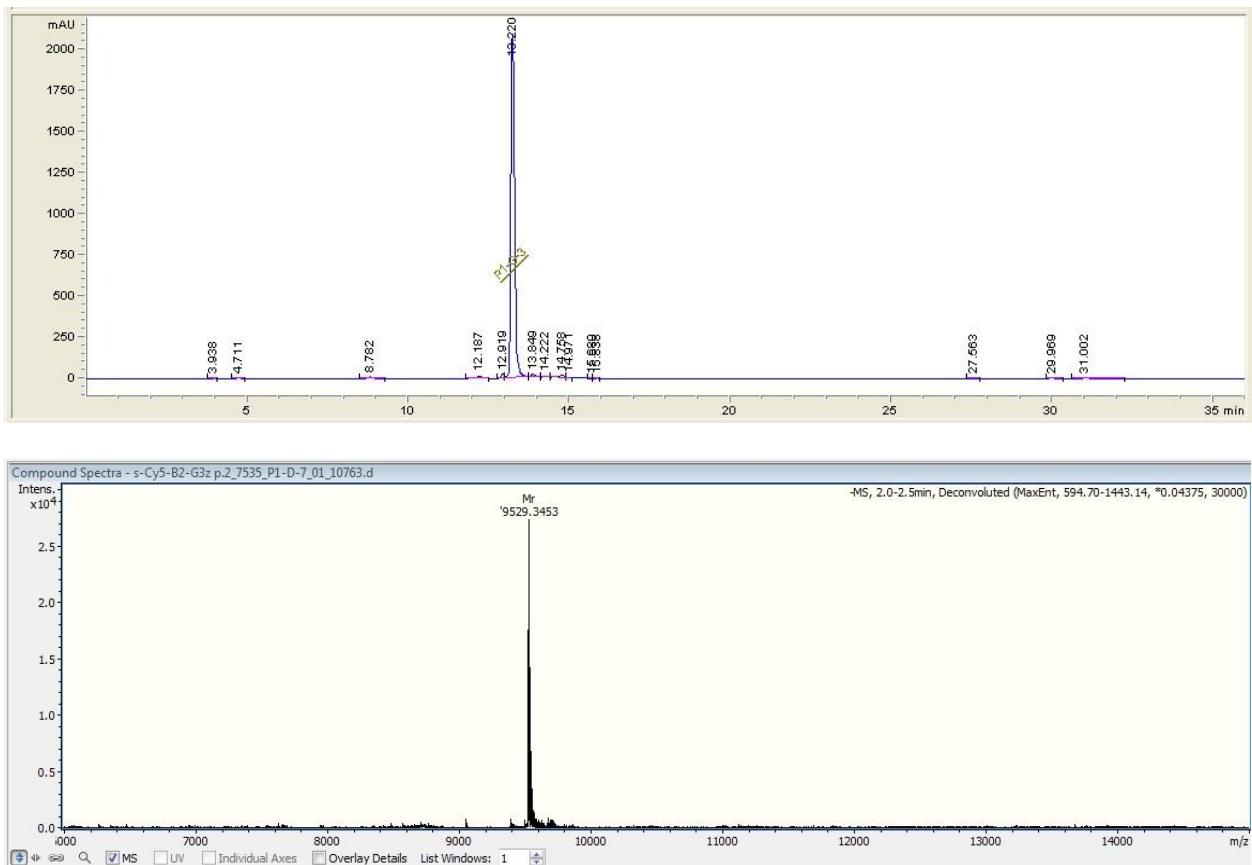
**Fig. S3-7.** HPLC profile and ESI MS of sCy5-B2-G1z probe.



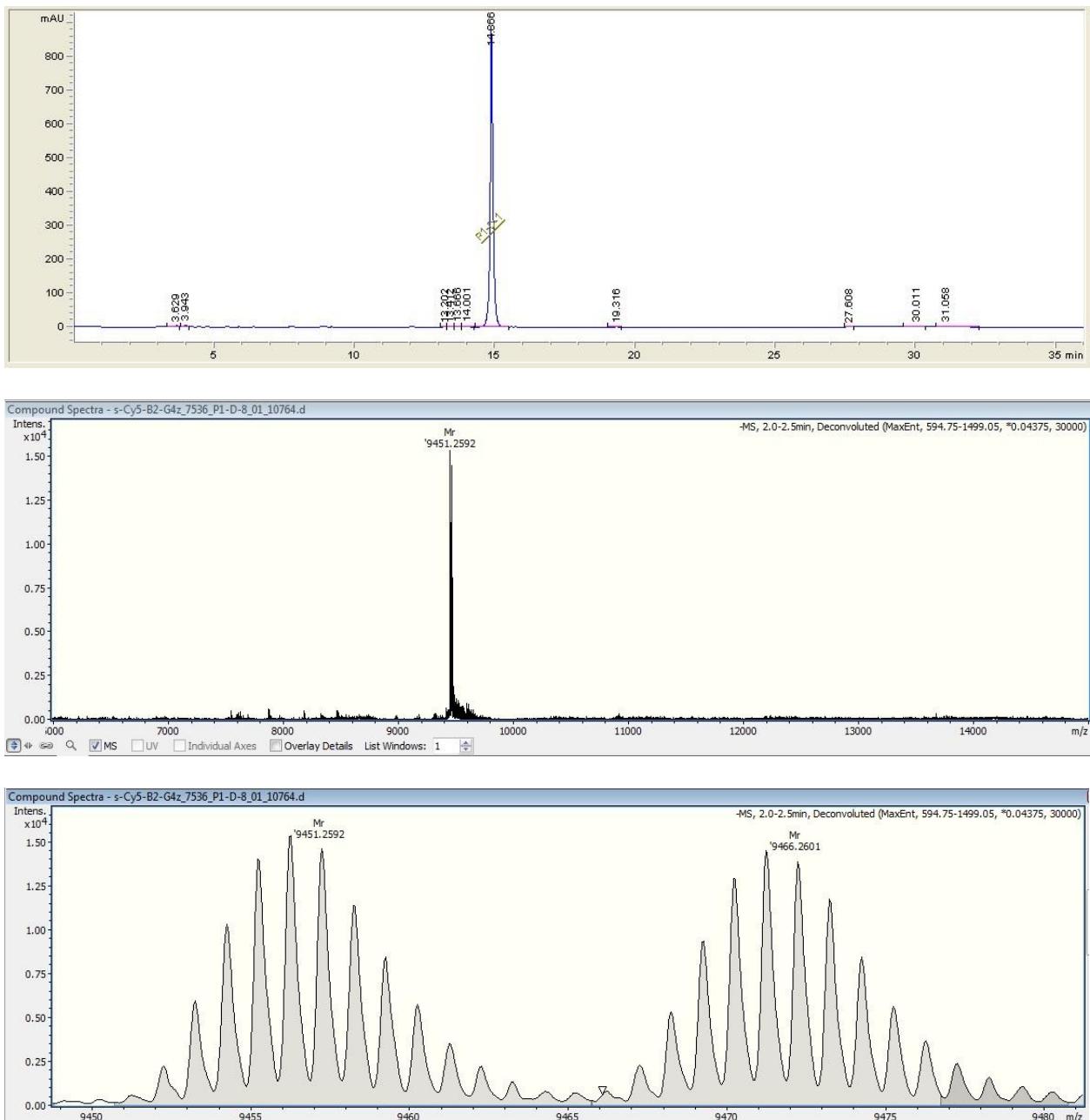
**Fig. S3-8.** HPLC profile and ESI MS of sCy5-B2-G2z probe.



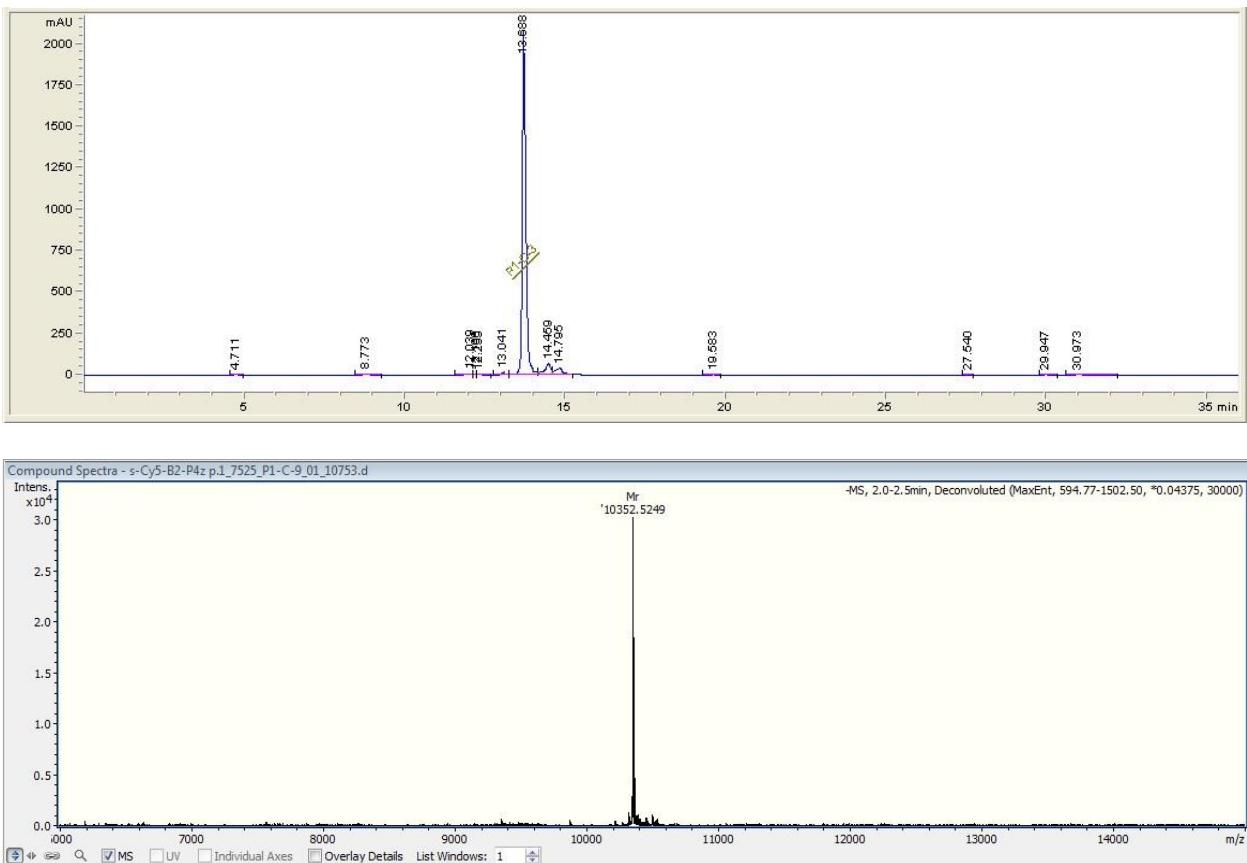
**Fig. S3-9.** HPLC profile and ESI MS of sCy5-B2-G3z probe.



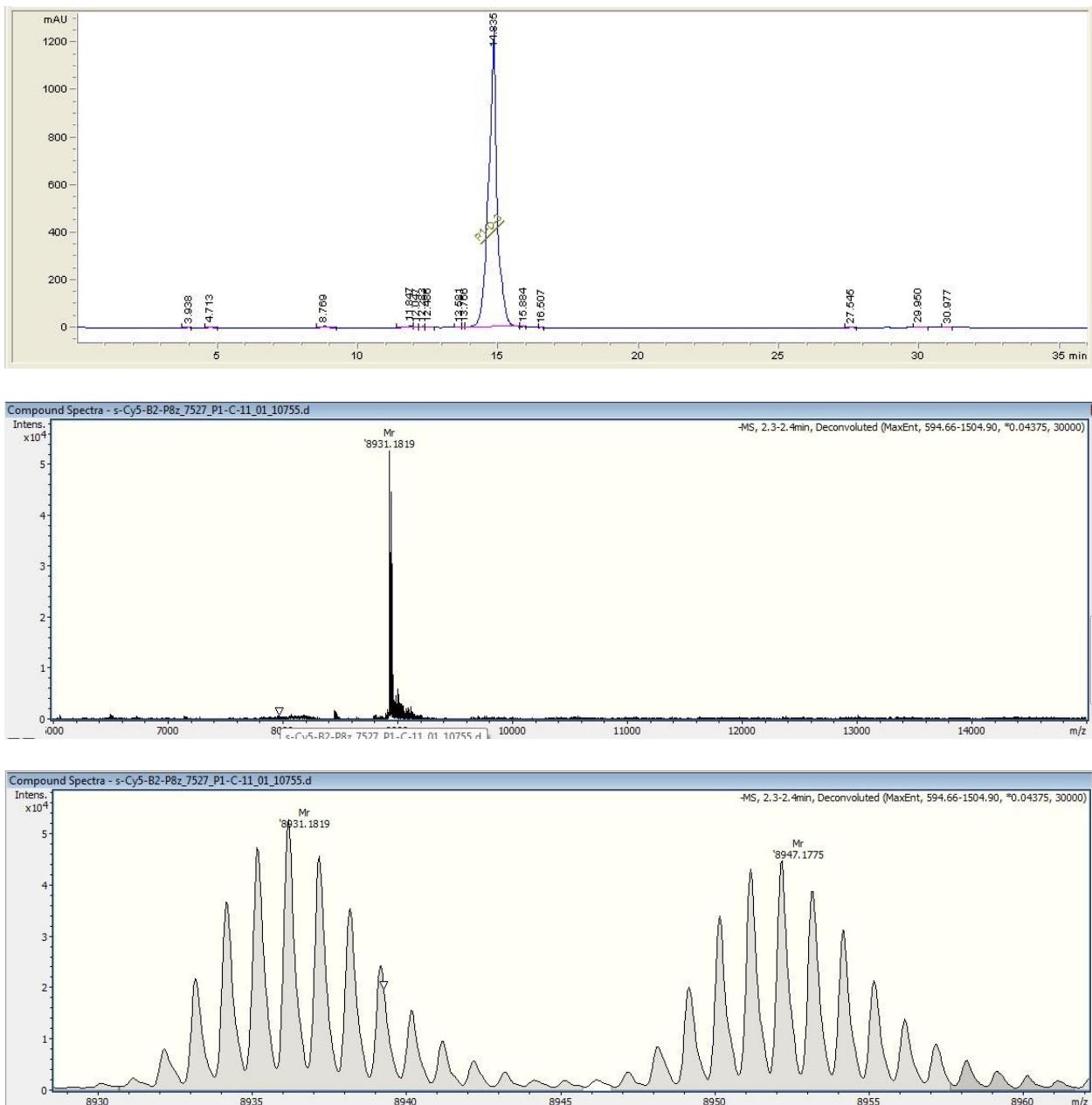
**Fig. S3-10.** HPLC profile and ESI MS of sCy5-B2-G4z probe.



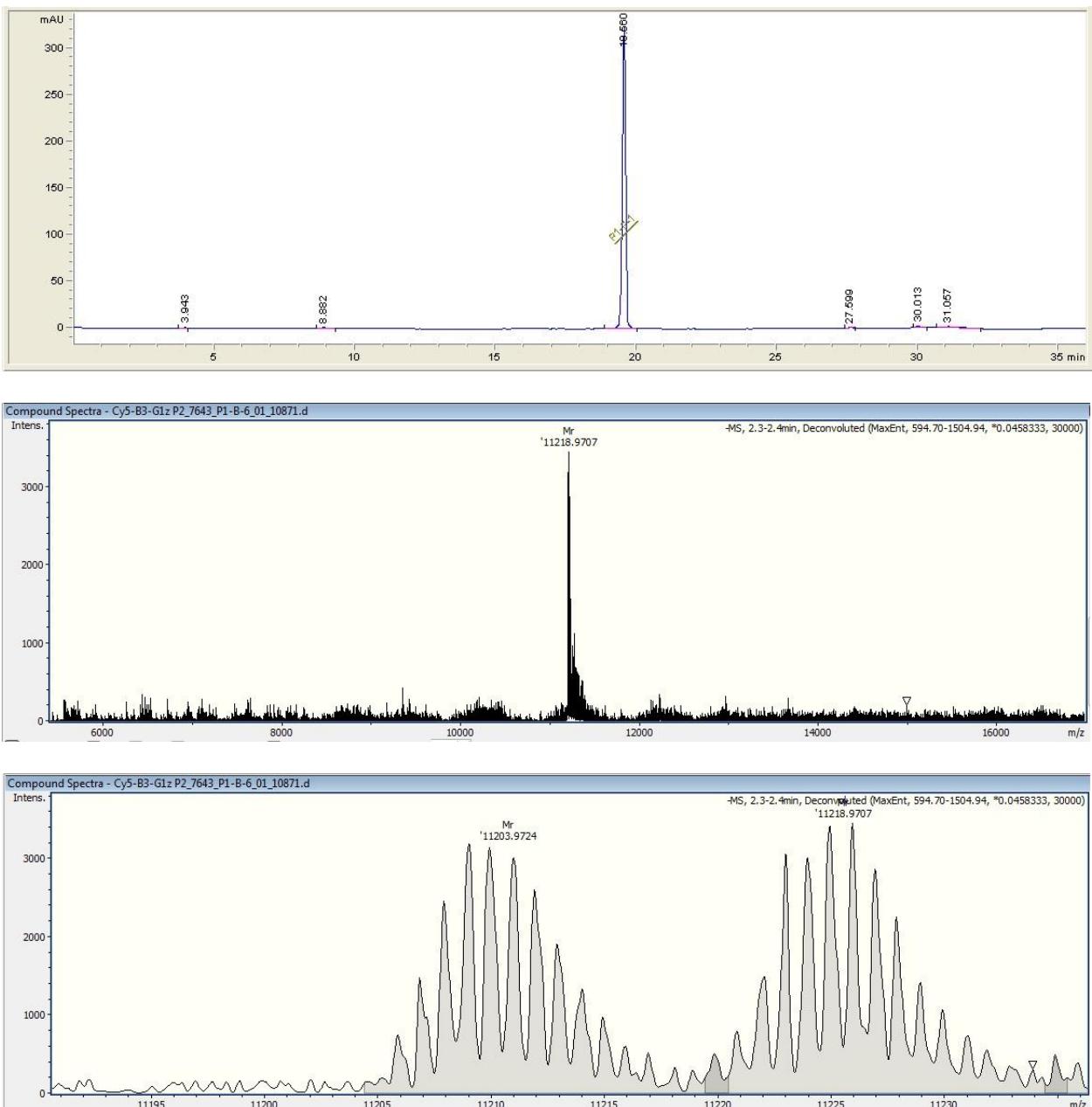
**Fig. S3-11.** HPLC profile and ESI MS of sCy5-B2-P4z probe.



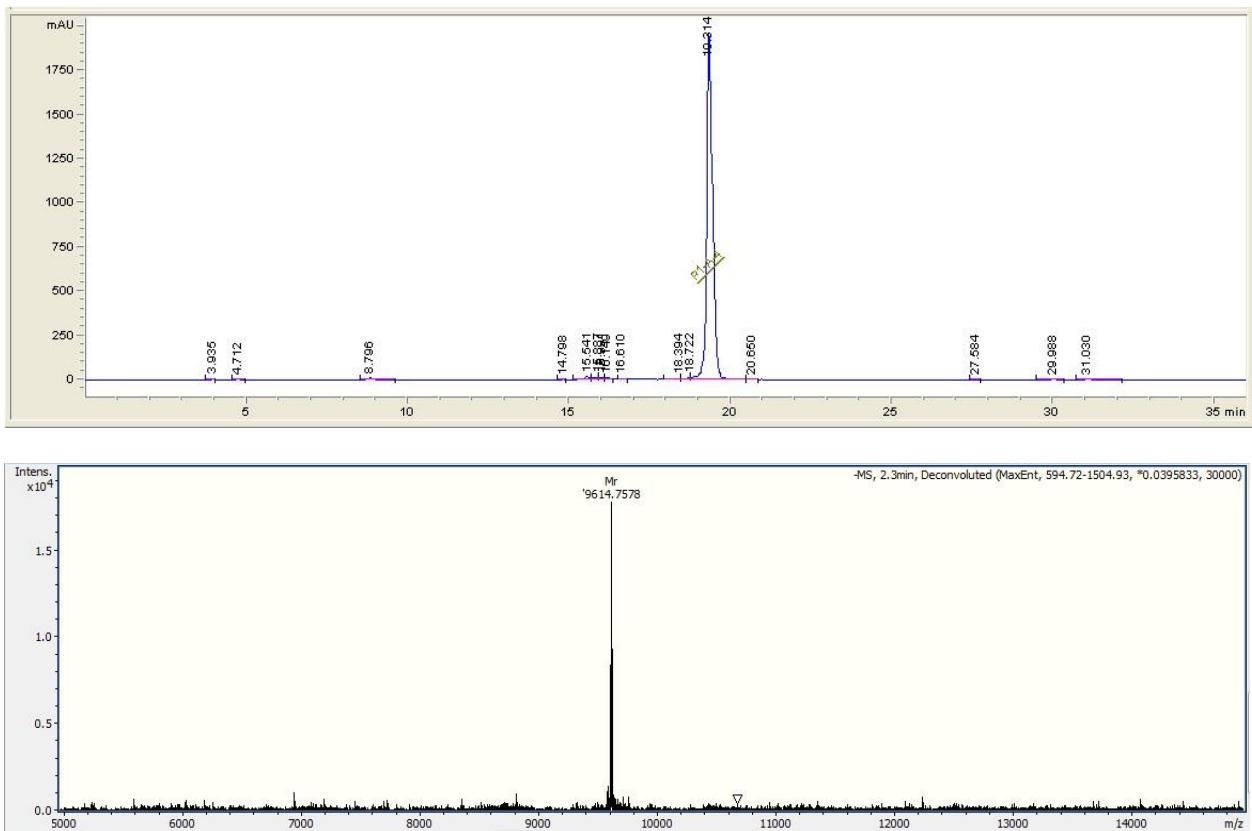
**Fig. S3-12.** HPLC profile and ESI MS of sCy5-B2-P8z probe.



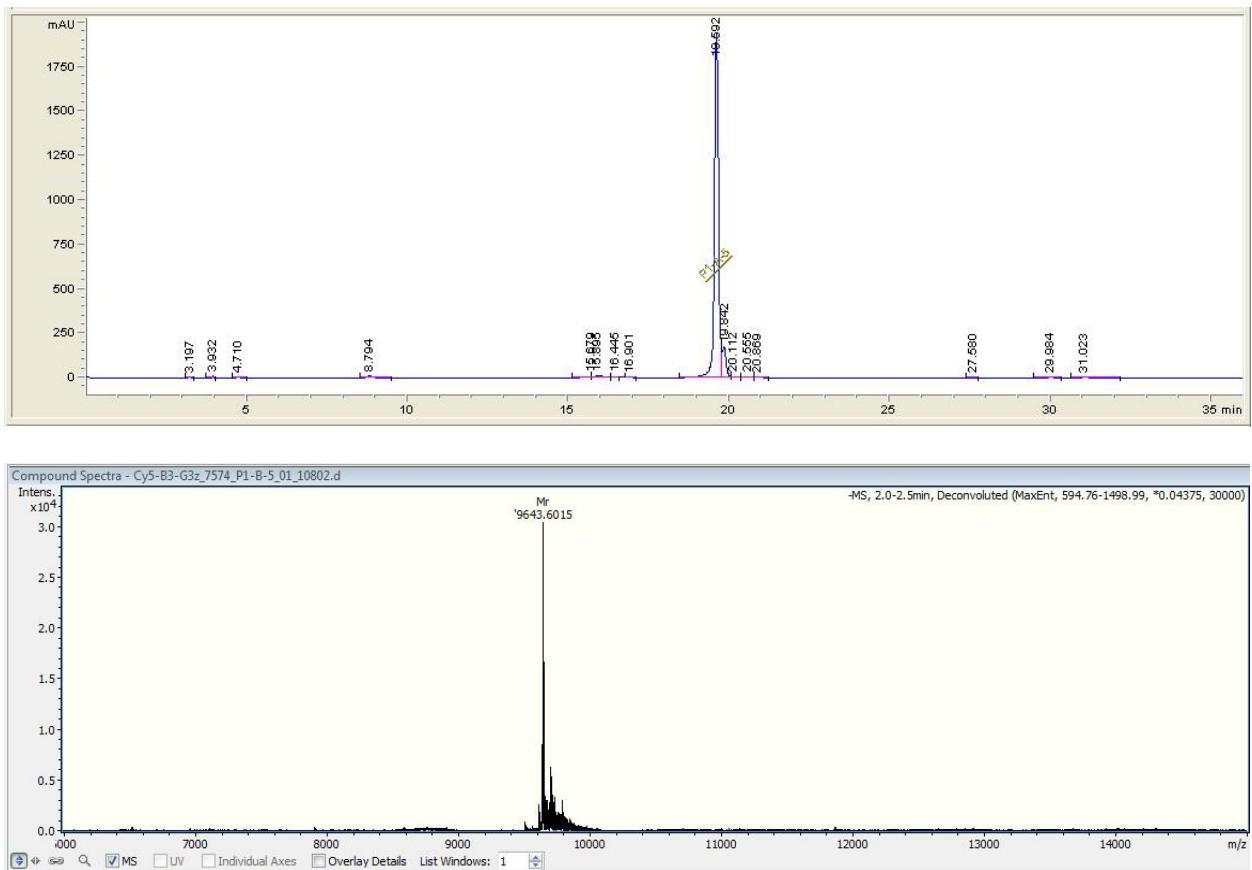
**Fig. S3-13.** HPLC profile and ESI MS of Cy5-B3-G1z probe.



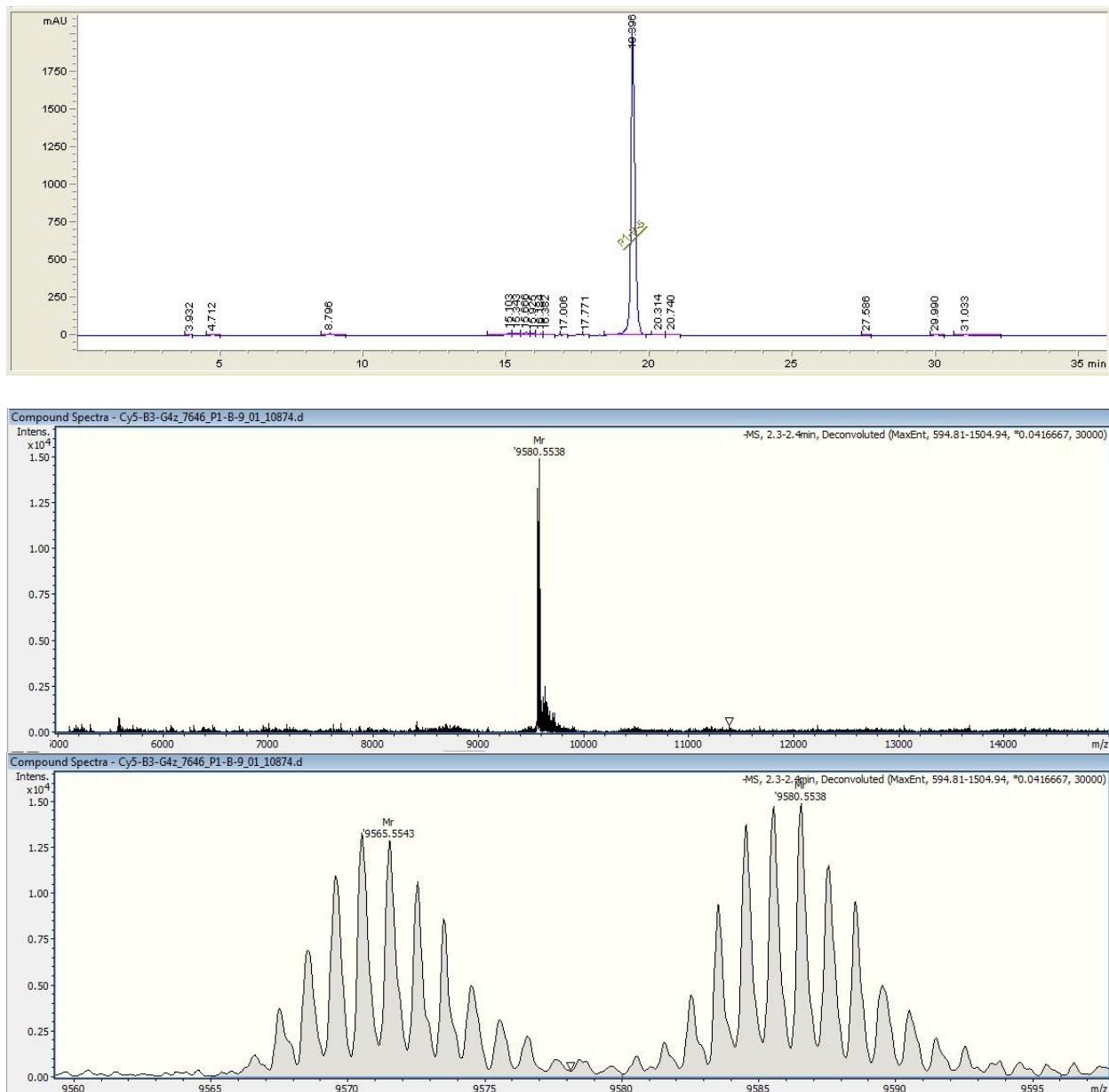
**Fig. S3-14.** HPLC profile and ESI MS of Cy5-B3-G2z probe.



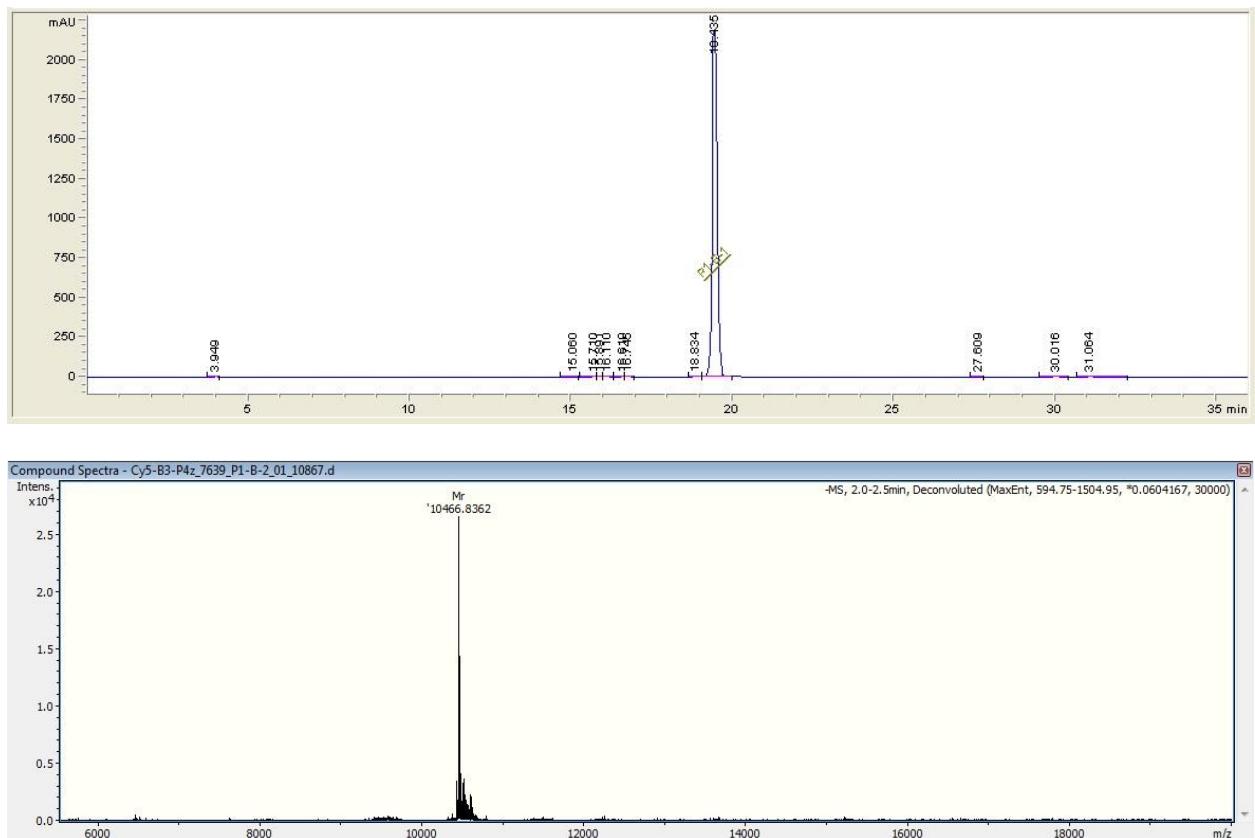
**Fig. S3-15.** HPLC profile and ESI MS of Cy5-B3-G3z probe.



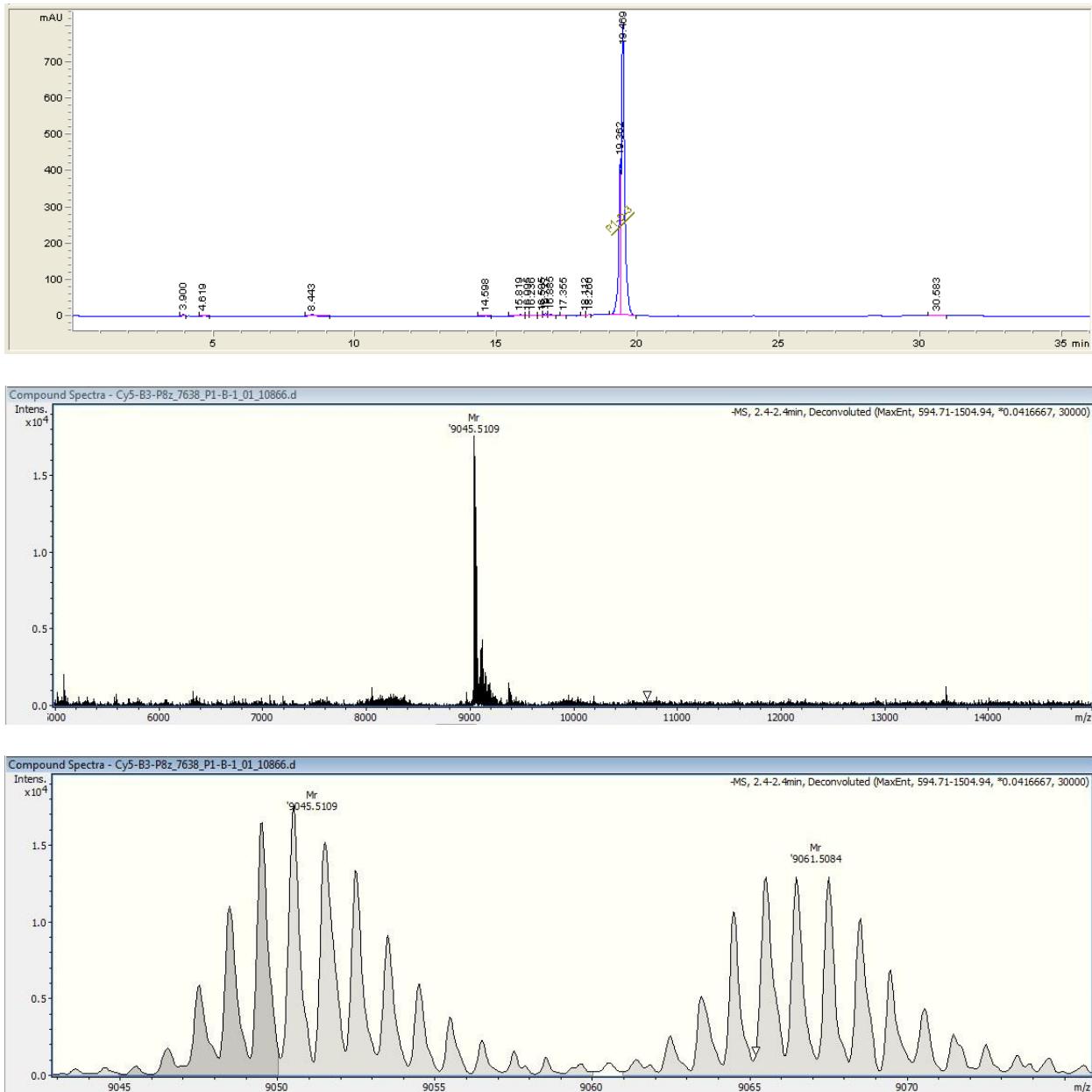
**Fig. S3-16.** HPLC profile and ESI MS of Cy5-B3-G4z probe.



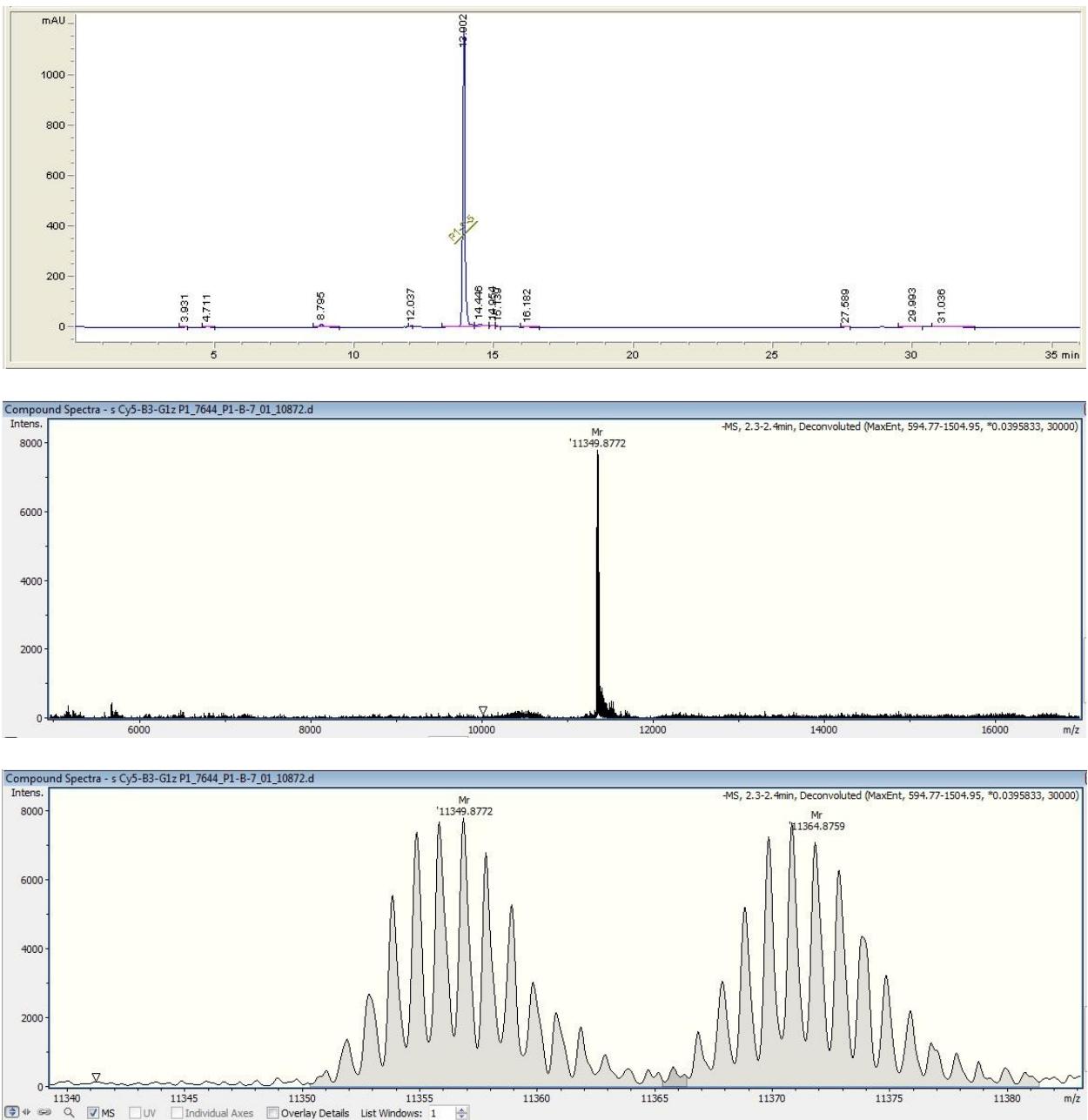
**Fig. S3-17.** HPLC profile and ESI MS of Cy5-B3-P4z probe.



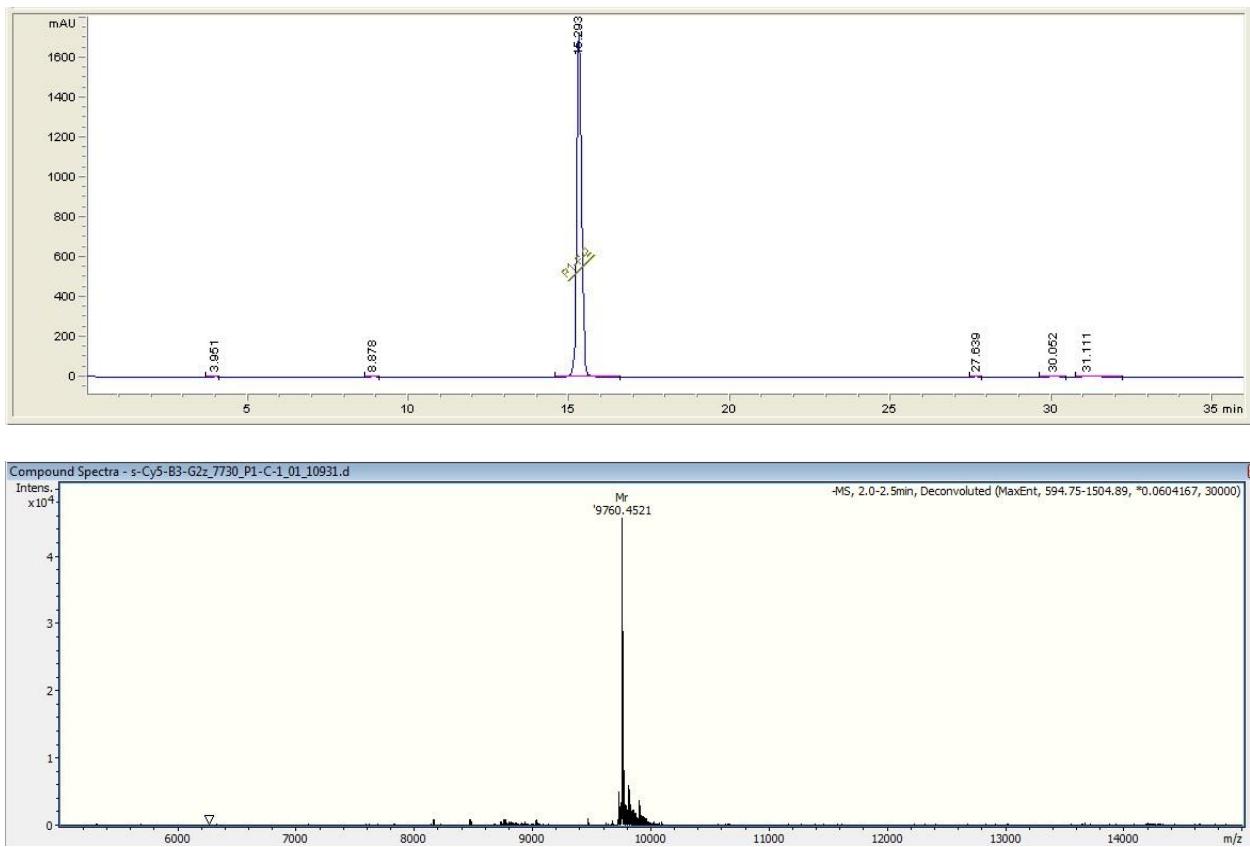
**Fig. S3-18.** HPLC profile and ESI MS of Cy5-B3-P8z probe.



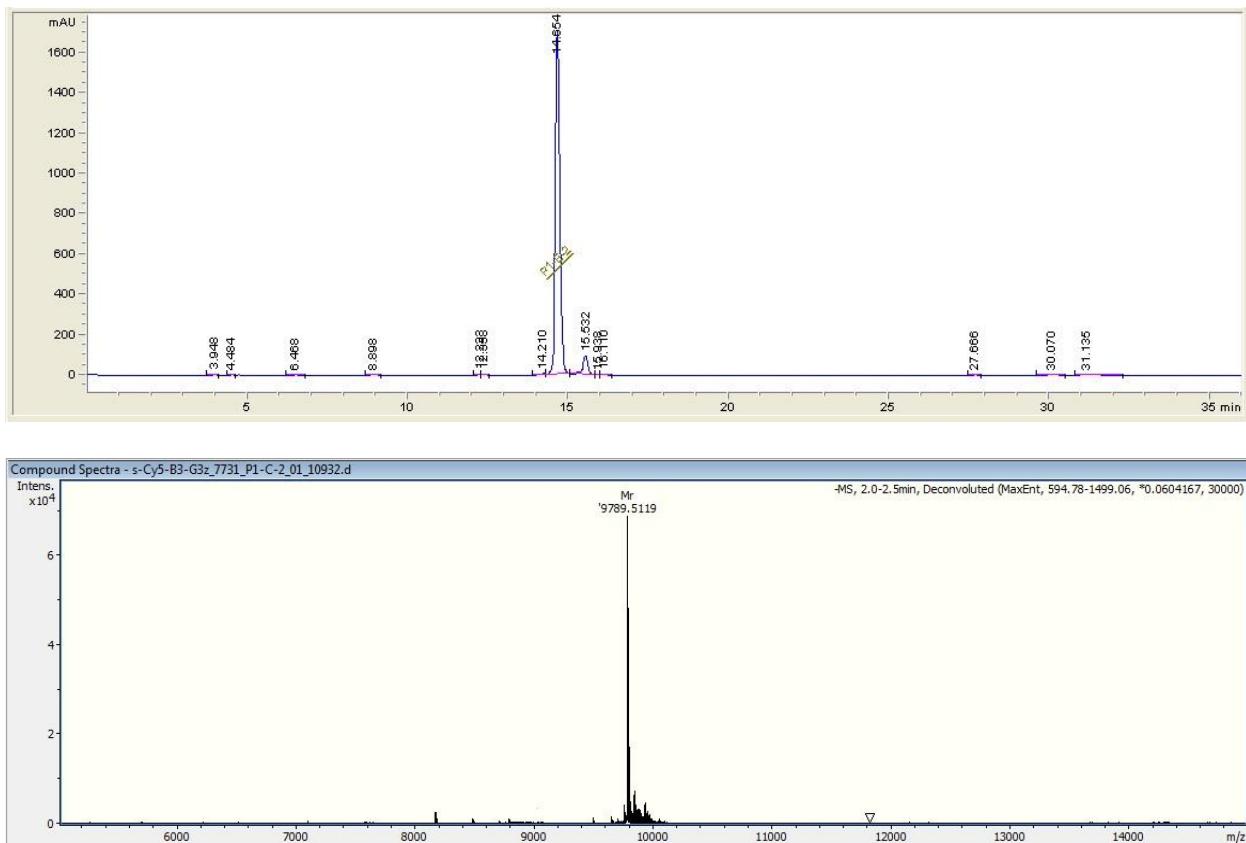
**Fig. S3-19.** HPLC profile and ESI MS of sCy5-B3-G1z probe.



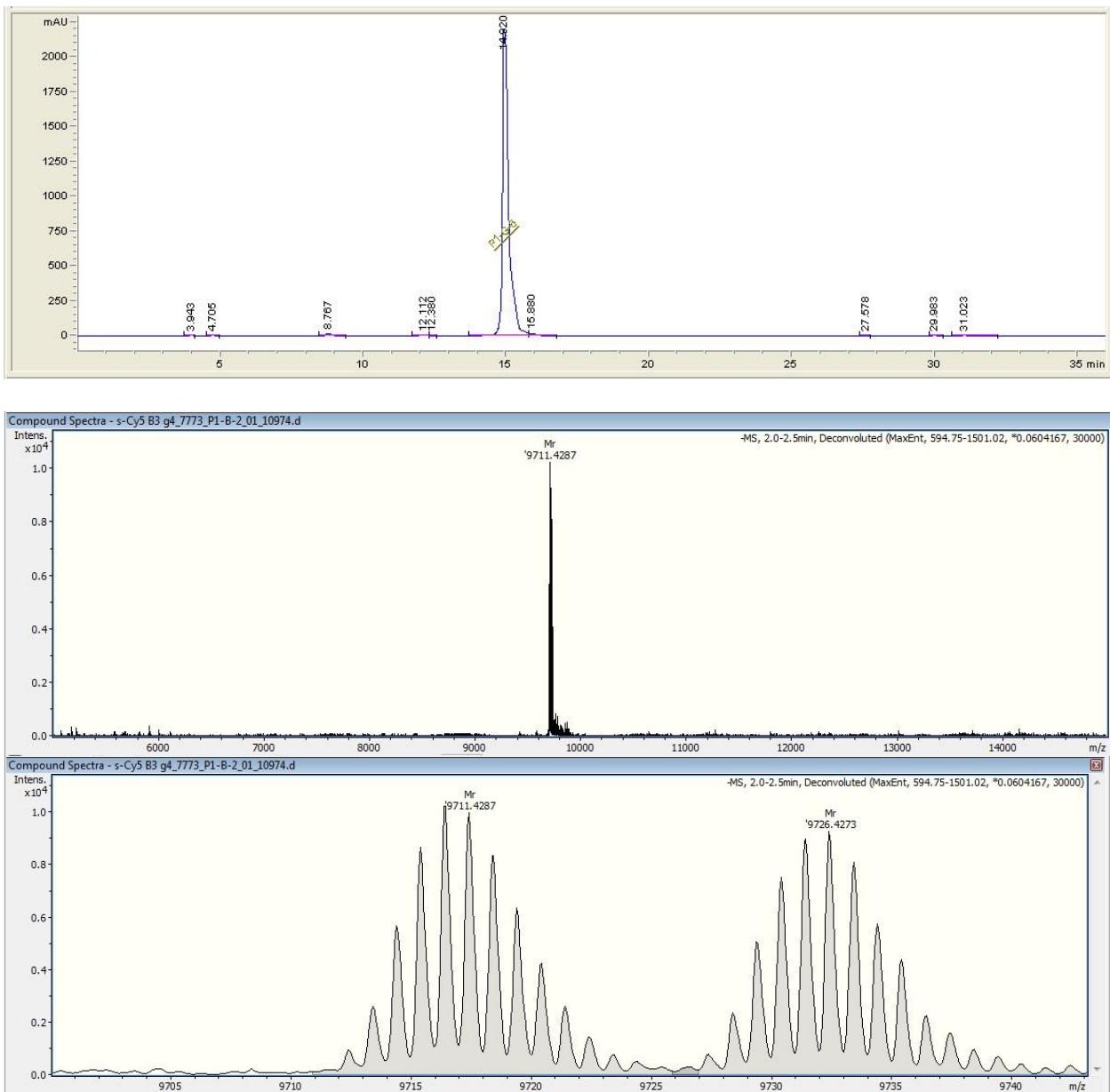
**Fig. S3-20.** HPLC profile and ESI MS of sCy5-B3-G2z probe.



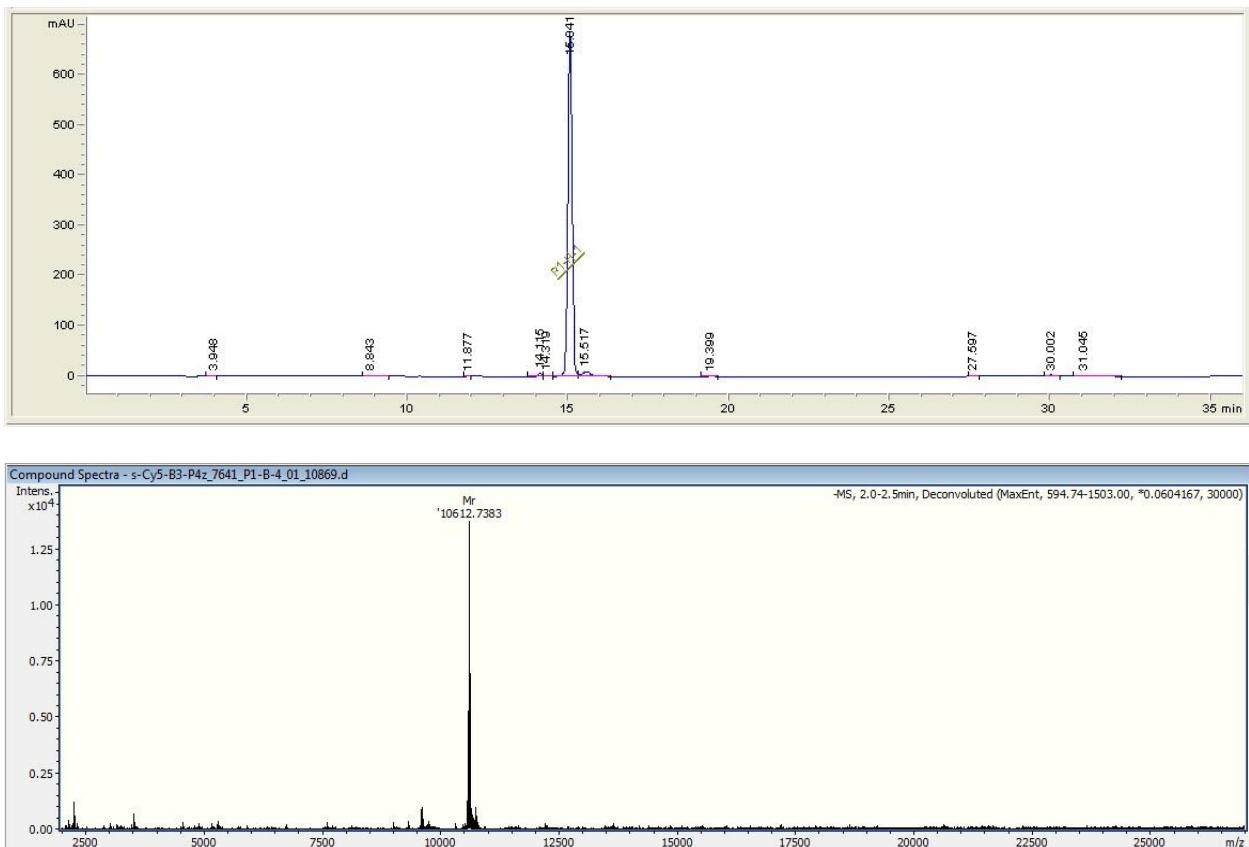
**Fig. S3-21.** HPLC profile and ESI MS of sCy5-B3-G3z probe.



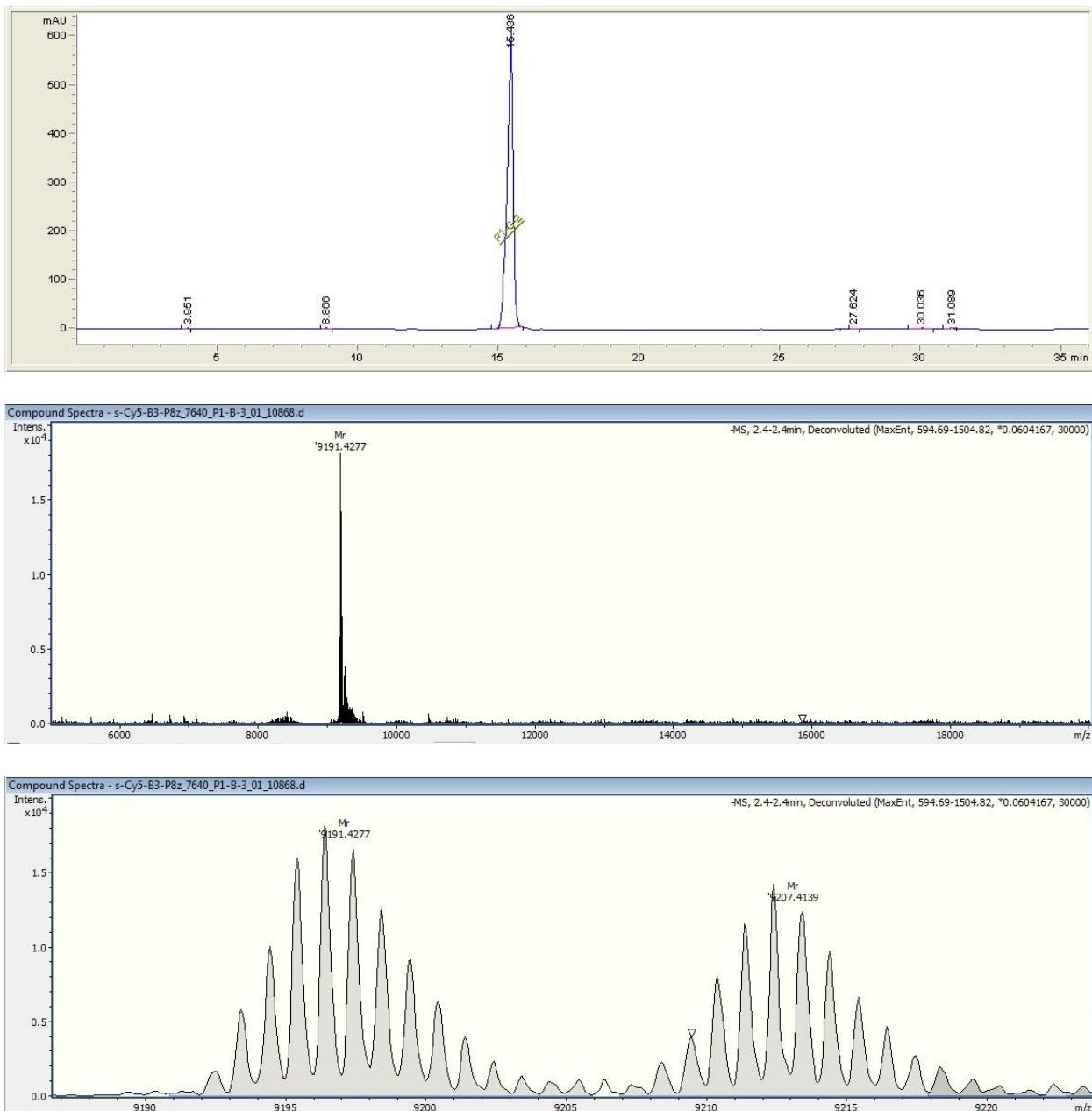
**Fig. S3-22.** HPLC profile and ESI MS of sCy5-B3-G4z probe.



**Fig. S3-23.** HPLC profile and ESI MS of sCy5-B3-P4z probe.

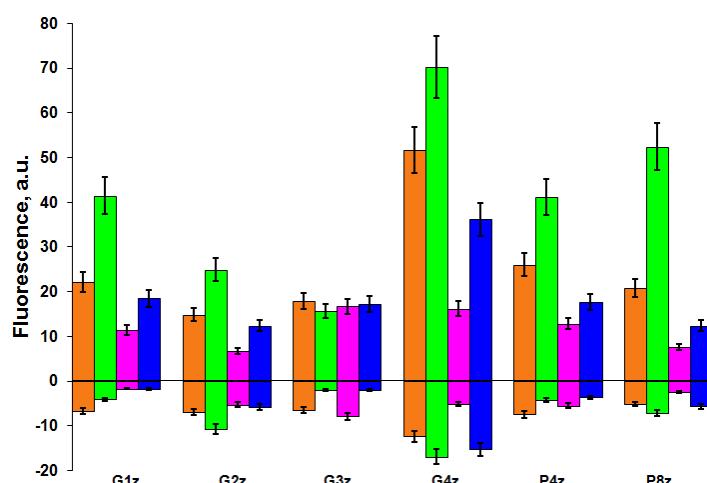


**Fig. S3-24.** HPLC profile and ESI MS of sCy5-B3-P8z probe.



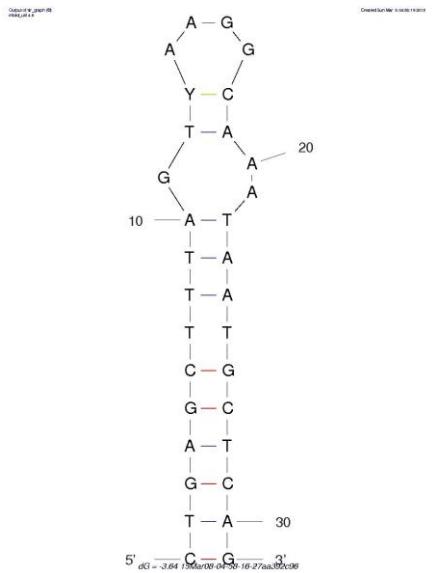
**Table S2.** Melting of Cy5/sCy5–BHQ2/BHQ3 labeled probes with fluorescence detection.

Probe	Dye/quencher	Tm, °C
G1	Cy5/BHQ2	64.3
	sulfo Cy5/BHQ2	68.9
	Cy5/BHQ3	62.3
	sulfo Cy5/BHQ3	67.4
G2	Cy5/BHQ2	50.2
	sulfo Cy5/BHQ2	46.2
	Cy5/BHQ3	51.7
	sulfo Cy5/BHQ3	48.2
G3	Cy5/BHQ2	58.8
	sulfo Cy5/BHQ2	62.8
	Cy5/BHQ3	54.2
	sulfo Cy5/BHQ3	58.8
G4	Cy5/BHQ2	50.7
	sulfo Cy5/BHQ2	46.2
	Cy5/BHQ3	50.2
	sulfo Cy5/BHQ3	44.1
P4	Cy5/BHQ2	48.7
	sulfo Cy5/BHQ2	52.7
	Cy5/BHQ3	48.7
	sulfo Cy5/BHQ3	51.2
P8	Cy5/BHQ2	58.3
	sulfo Cy5/BHQ2	55.3
	Cy5/BHQ3	62.3
	sulfo Cy5/BHQ3	59.8

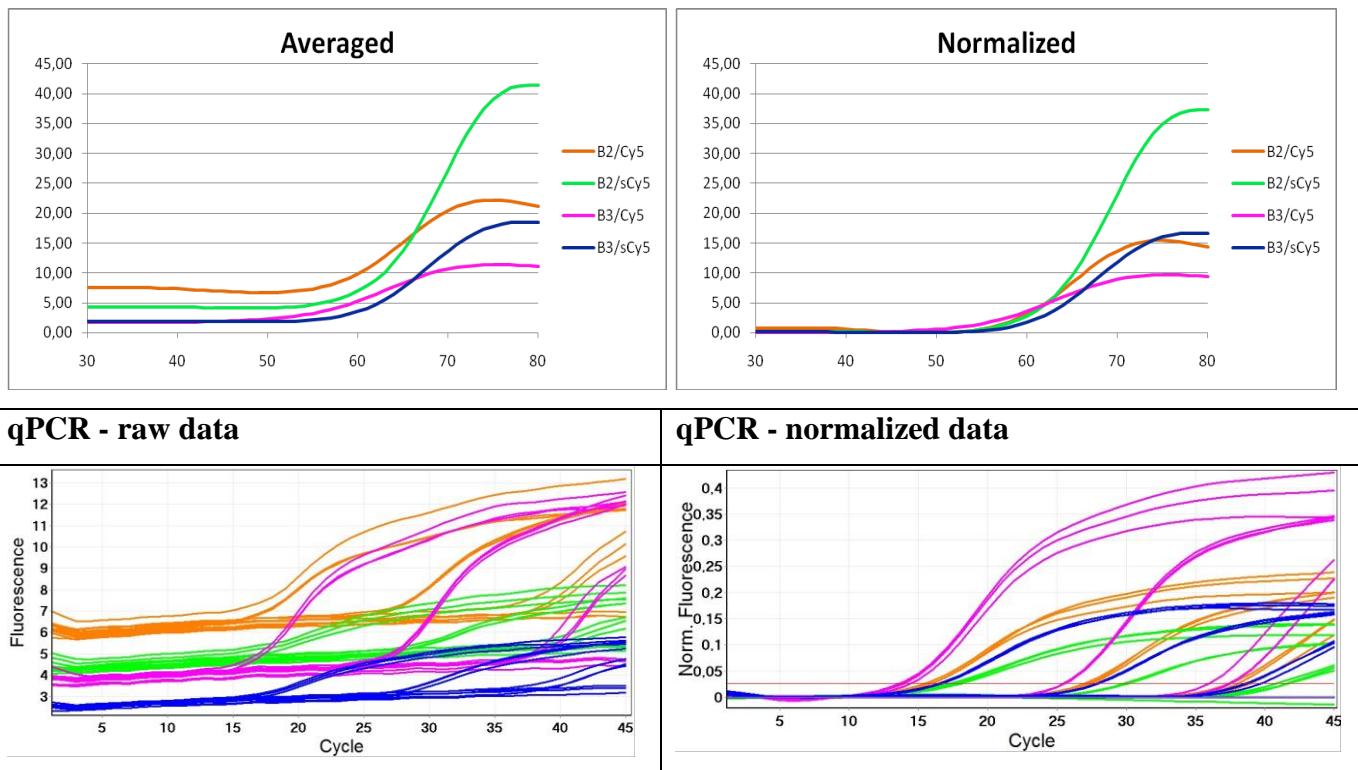


**Fig S4.** Fluorescence background (negative) and light-up (positive) of structured and molten probes; █ Cy5-B2, █ sCy5-B2, █ Cy5-B3, █ sCy5-B3.

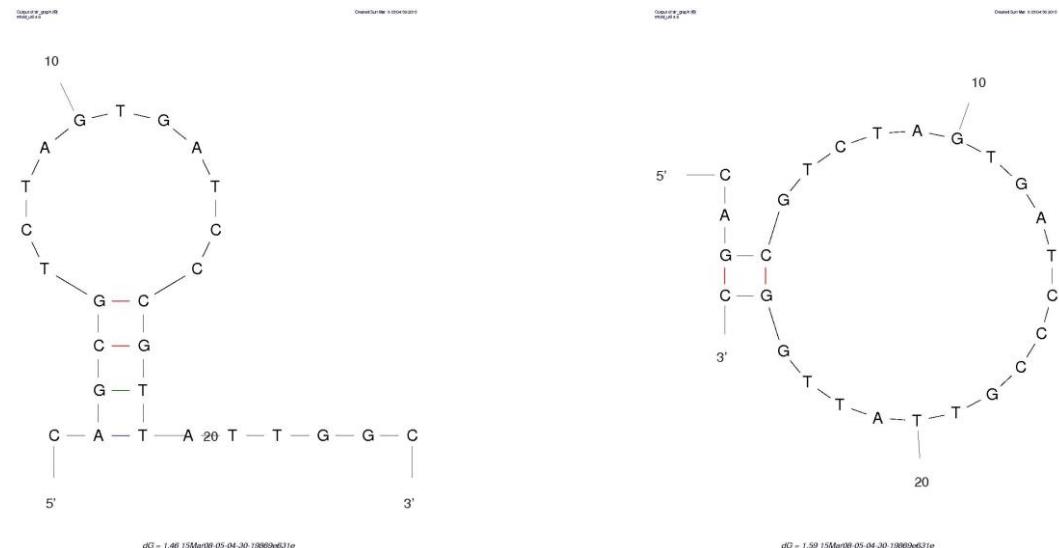
**Fig. S5-1.** Mfold secondary structure, raw and processed melting and qPCR data for G1z probes.



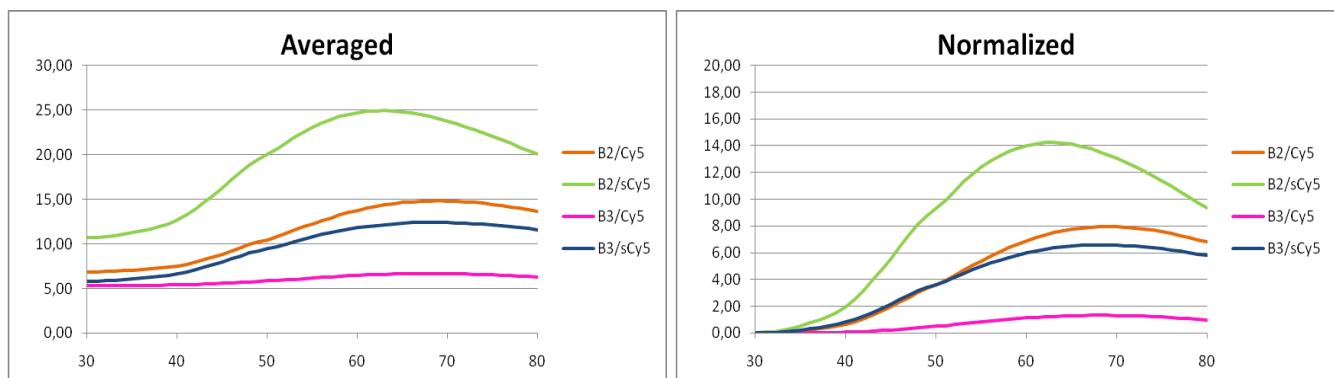
Melting



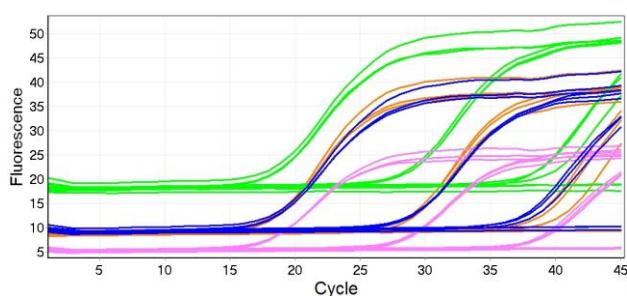
**Fig. S5-2.** Mfold secondary structure, raw and processed melting and qPCR data for **G2z** probes.



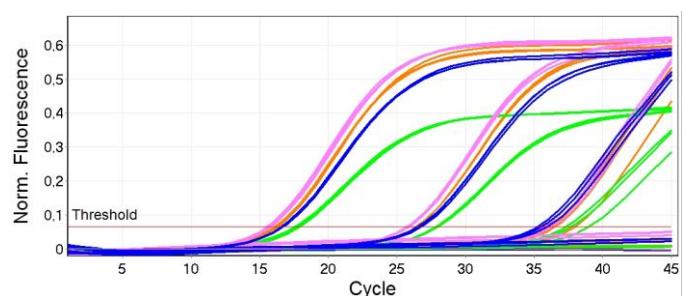
## Melting



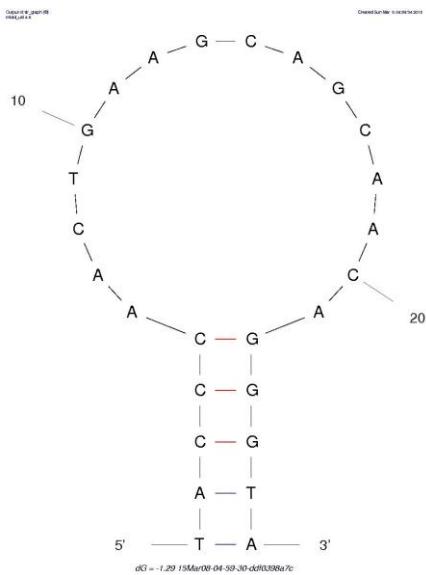
## qPCR - raw data



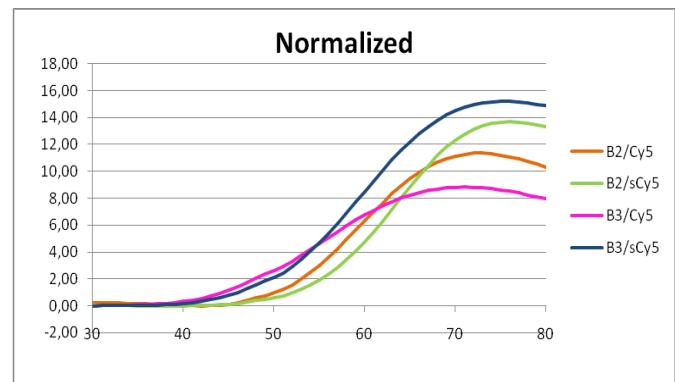
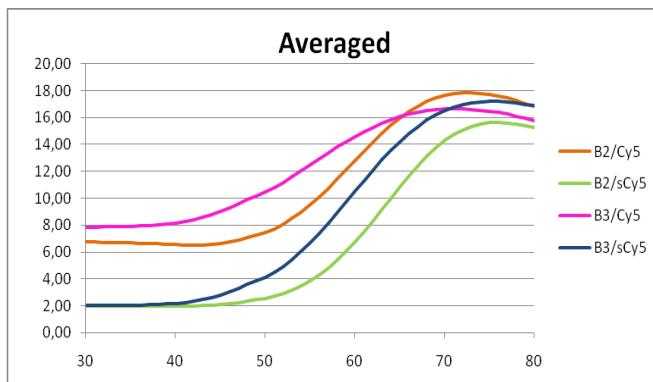
## qPCR - normalized data



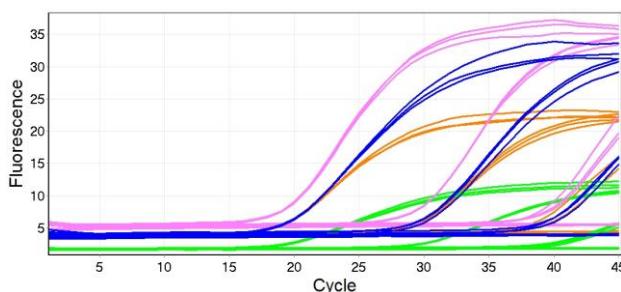
**Fig. S5-3.** Mfold secondary structure, raw and processed melting and qPCR data for **G3z** probes.



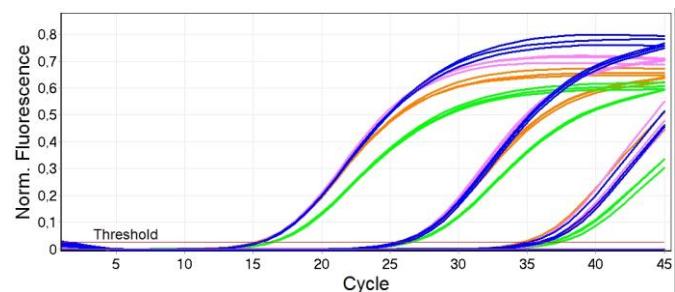
### Melting



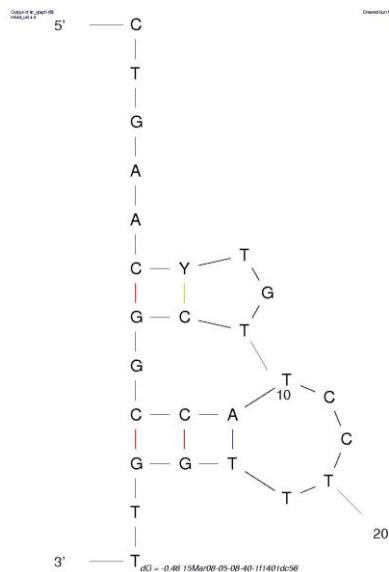
### qPCR - raw data



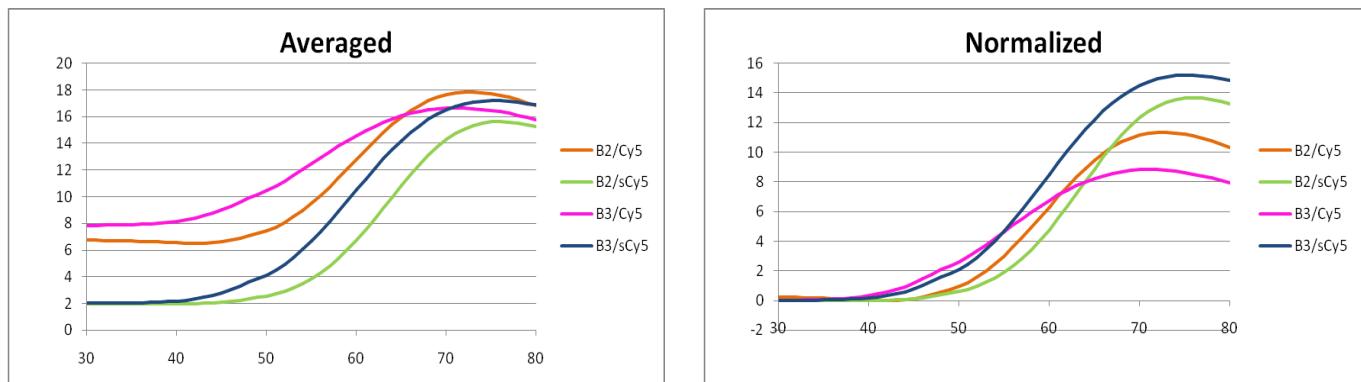
### qPCR - normalized data



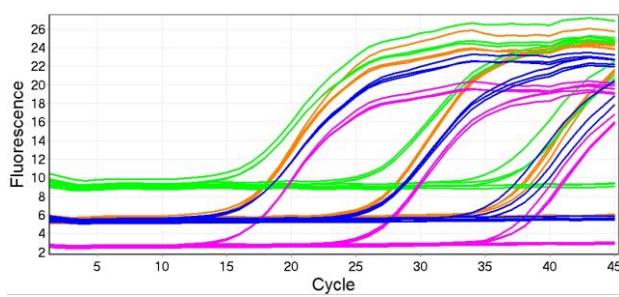
**Fig. S5-4.** Mfold secondary structure, raw and processed melting and qPCR data for **G4z** probes.



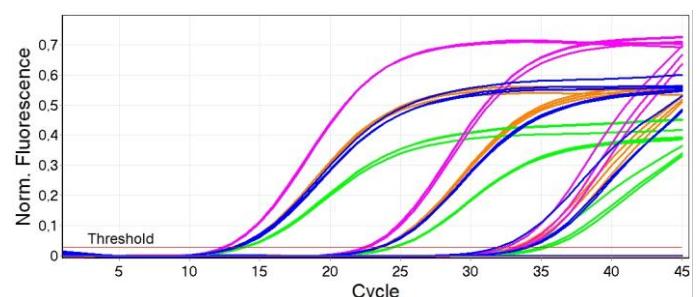
### Melting



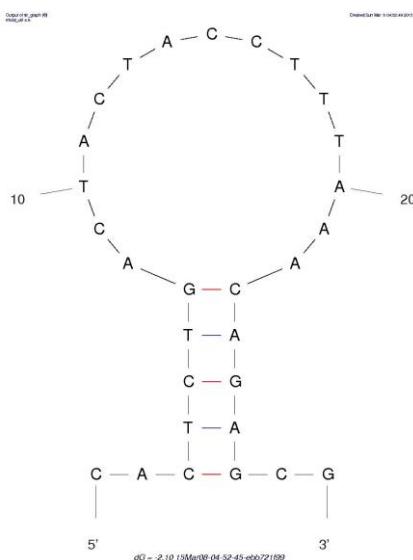
### qPCR - raw data



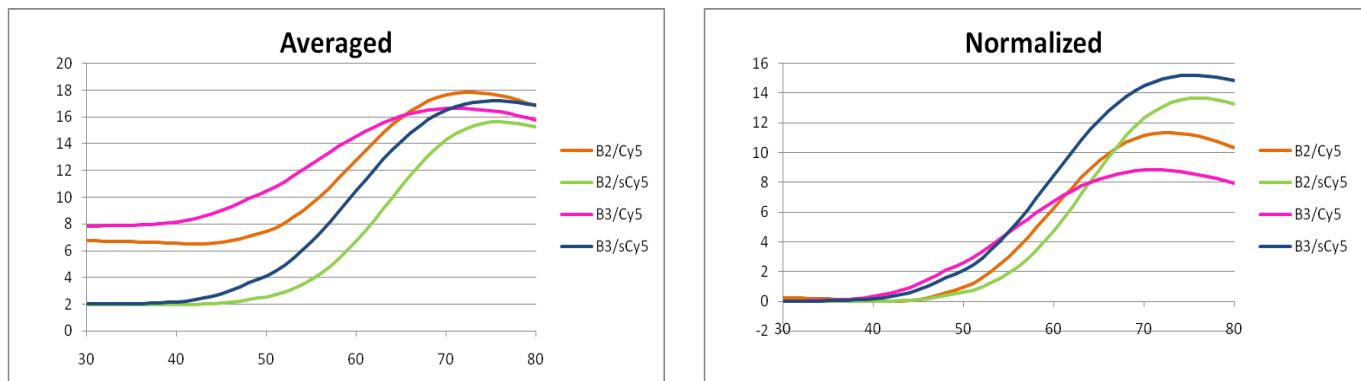
### qPCR - normalized data



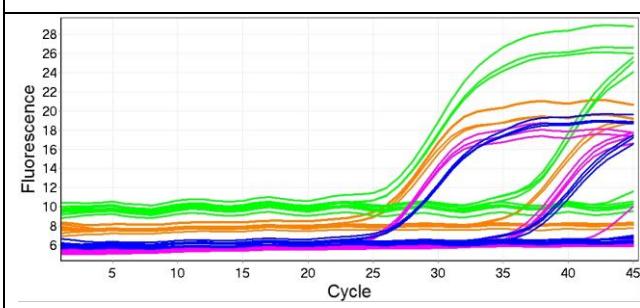
**Fig. S5-5.** Mfold secondary structure, raw and processed melting and qPCR data for **P4z** probes.



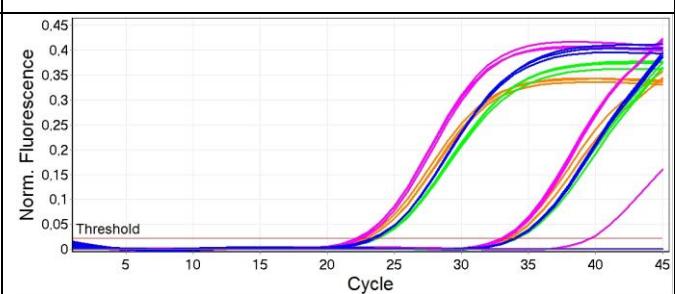
### Melting



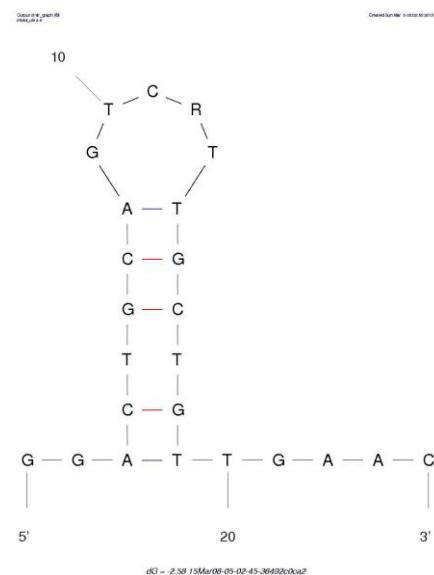
**qPCR - raw data**



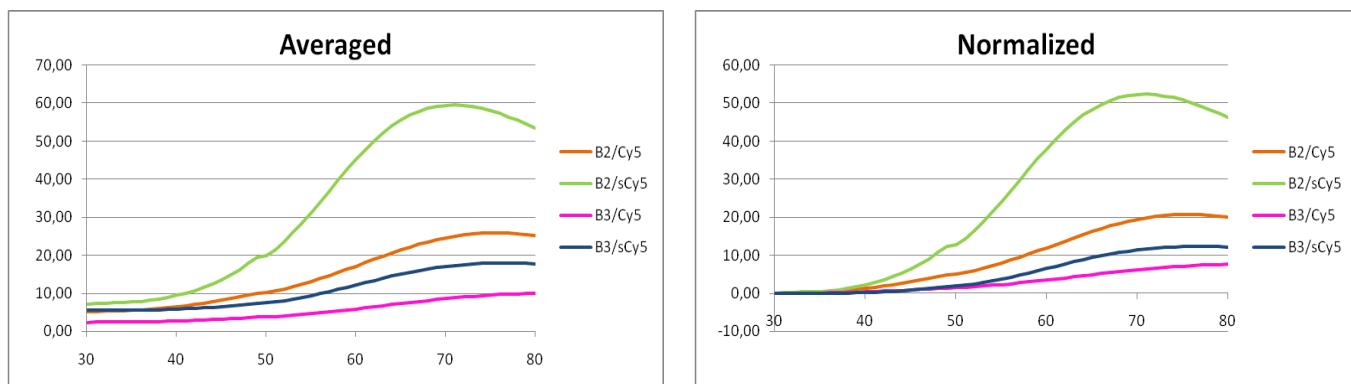
**qPCR - normalized data**



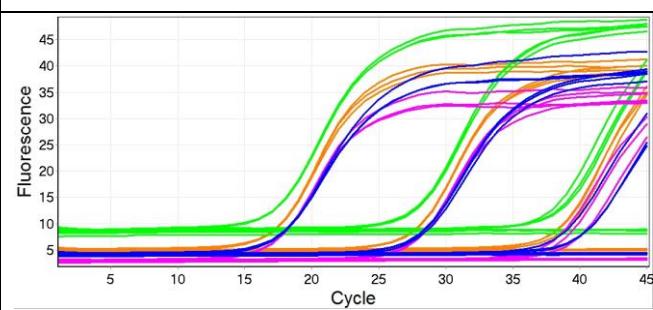
**Fig. S5-6.** Mfold secondary structure, raw and processed melting and qPCR data for **P8z** probes.



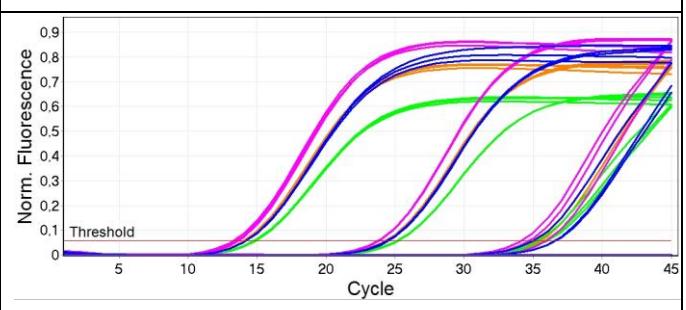
### Melting

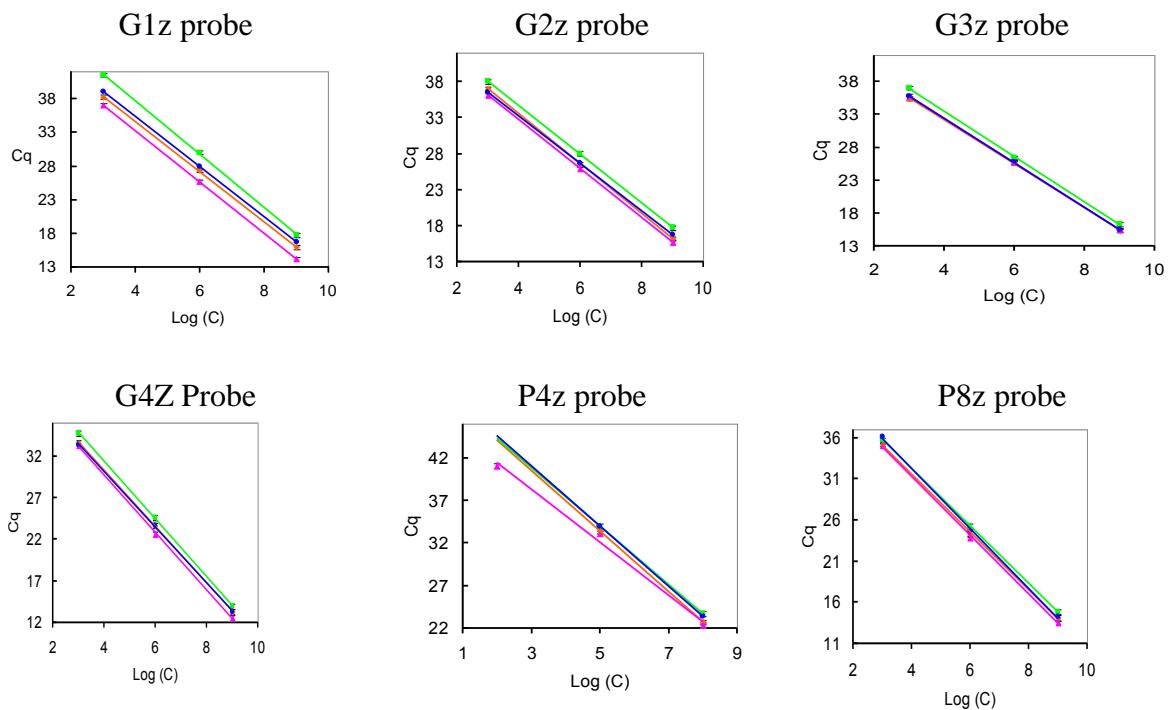


### qPCR - raw data



### qPCR - normalized data

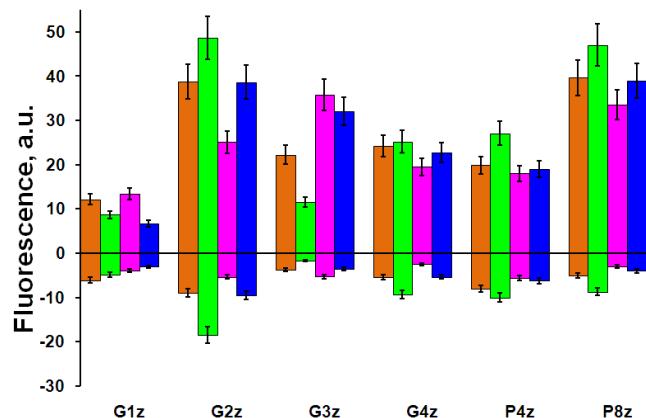




**Fig. S6.** qPCR data for **Cy5-B2**, **sCy5-B2**, **Cy5-B3**, and **sCy5-B3** dual labeled probes; plot: dependence of quantification cycle number on the logarithm of DNA target concentration. For numerical values see Table S3†.

**Table S3.** qPCR data for **Cy5-B2**, **sCy5-B2**, **Cy5-B3**, and **sCy5-B3** dual labeled probes.

Probe	Probe sequence, 5'-3'	qPCR data				
		Target DNA, copies/reaction	Cq, cycles			
			Cy5-B2	sCy5-B2	Cy5-B3	sCy5-B3
<b>G1z</b>	<b>Dye</b> -CTG AGC TTT AGT YAA GGC AAA TAA TGC TCA <b>G-Quencher</b>	1.0·10 <sup>9</sup>	15.9±0.2	17.7±0.3	14.1±0.2	16.7±0.1
		1.0·10 <sup>6</sup>	27.3±0.3	28.0±0.3	25.6±0.3	28.0±0.2
		1.0·10 <sup>3</sup>	38.2±0.4	38.0±0.4	37.0±0.3	39.0±0.4
<b>G2z</b>	<b>Dye</b> -CAG CGT CTA GTG ATC CCG TTA TTG GC- <b>Quencher</b>	1.0·10 <sup>9</sup>	16.2±0.2	17.7±0.3	15.7±0.3	16.7±0.2
		1.0·10 <sup>6</sup>	26.5±0.2	28.0±0.2	25.9±0.2	26.7±0.4
		1.0·10 <sup>3</sup>	37.0±0.3	38.0±0.3	36.1±0.3	36.5±0.3
<b>G3z</b>	<b>Dye</b> -TAC CCA ACT GAA GCA GCA ACA GGG TA- <b>Quencher</b>	1.0·10 <sup>9</sup>	15.3±0.2	16.3±0.3	15.4±0.3	15.4±0.3
		1.0·10 <sup>6</sup>	25.7±0.3	26.4±0.3	25.6±0.3	25.8±0.3
		1.0·10 <sup>3</sup>	35.4±0.3	37.0±0.3	35.7±0.3	35.7±0.3
<b>G4z</b>	<b>Dye</b> -CTG AAC YTG TCG GCC ATC CTT TGG TT- <b>Quencher</b>	1.0·10 <sup>9</sup>	13.3±0.1	13.9±0.3	12.5±0.3	13.3±0.3
		1.0·10 <sup>6</sup>	23.6±0.2	24.6±0.3	22.6±0.3	23.6±0.3
		1.0·10 <sup>3</sup>	33.6±0.3	34.7±0.3	33.2±0.3	33.4±0.3
<b>P4z</b>	<b>Dye</b> -CAC TCT GAC TAC TAC CTT TAA ACA GAG CG- <b>Quencher</b>	1.0·10 <sup>8</sup>	22.7±0.1	23.7±0.1	22.3±0.2	23.4±0.2
		1.0·10 <sup>5</sup>	33.4±0.3	34.0±0.2	33.1±0.2	34.0±0.2
		1.0·10 <sup>2</sup>	-	-	41.0±0.5	-
<b>P8z</b>	<b>Dye</b> -GGA CTG CAG TCR TTG CTG TTG AAC- <b>Quencher</b>	1.0·10 <sup>9</sup>	14.1±0.2	14.8±0.3	13.5±0.3	14.2±0.4
		1.0·10 <sup>6</sup>	24.3±0.2	25.2±0.2	23.8±0.3	24.5±0.2
		1.0·10 <sup>3</sup>	35.2±0.3	35.7±0.3	35.0±0.3	36.1±0.3



**Fig. S7.** Fluorescence light-up (positive) and background (negative) in qPCR; ■ – Cy5-B2, ■ – sCy5-B2, ■ – Cy5-B3, ■ – sCy5-B3.