

Supporting Information

Modifications at the C-Terminus to Improve Py/Im Polyamide Activity in Cell Culture

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Compound	MALDI-ToF:	Purity	Compound	MALDI-ToF	Purity
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Number	Expected (M+H ⁺); Observed (M+H ⁺)	(HPLC, 310 nm)	Number	Expected (M+H): Observed (M+H):	(HPLC, 310 nm)
1	1274.49; 1274.51	99.3	23	1237.59; 1237.48	95.0
2	1246.50; 1246.81	99.3	24	1209.59; 1209.89	95.6
3	1231.49; 1231.77	99.7	25	1194.58; 1194.53	100.0
4	1175.43; 1175.61	98.7	26	1138.52; 1138.50	99.7
5	1191.42; 1191.38	97.2	27	1154.51; 1154.62	96.8
6	1394.52; 1394.46	98.0	28	1357.61; 1357.59	98.8
7	1379.50; 1379.72	99.3	29	1342.60; 1342.55	99.6
8	1323.44; 1323.52	99.9	30	1286.64; 1286.59	99.0
9	1323.44; 1323.44	95.6	31	1286.54; 1286.56	96.1
10	1297.44; 1297.48	95.9	32	1260.54; 1260.63	95.3
11	1319.42; 1319.52 ^a	97.2	33	1260.54; 1260.51	96.4
12	1238.58; 1238.51	96.5	34	1238.58; 1238.56	99.0
13	1210.59; 1210.76	98.1	35	1210.59; 1210.59	98.0
14	1195.58; 1195.57	98.7	36	1195.58; 1195.56	99.3
15	1139.51; 1139.49	98.8	37	1139.51; 1139.77	95.1
16	1155.51; 1155.83	98.4	38	1155.51; 1155.40	97.4
17	1358.60; 1358.66	97.8	39	1358.60; 1358.70	95.4
18	1343.59; 1343.76	99.9	40	1343.59; 1343.52	100.0
19	1287.53; 1287.56	99.4	41	1287.53; 1287.60	98.4
20	1287.53; 1287.73	97.6	42	1287.53; 1287.53	95.9
21	1261.53; 1261.87	97.3	43	1261.53; 1261.56	98.9
22	1261.53; 1261.51	98.8	44	1261.53; 1261.51	96.2

Table SI 1: MALDI-ToF and purity data for compounds **1-44**. MALDI-ToF data gives calculated exact mass for (M+H⁺) and observed mass for (M+H⁺) for polyamides **1-10** and **12-44**. ^a MALDI-ToF data for **11** corresponds to the (M+Na⁺) peak. Purity was measured by analytical HPLC at 310 nm; purity for oxime-linked compounds **9-11**, **20-22**, **31-33** and **42-44** included both E and Z isomers.

Compound	Tm (°C)	ΔTm (°C)	Compound	Tm (°C)	ΔTm (°C)
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Number			Number		
DNA Only	64.7 ± 0.2		DNA Only	60.8 ± 0.2	
1	77.1 ± 0.2	12.4	23	75.1 ± 0.3	14.3
2	79.5 ± 0.2	14.8	24	78.1 ± 0.2	17.3
3	77.4 ± 0.2	12.7	25	75.1 ± 0.2	14.3
4	78.2 ± 0.2	13.5	26	77.0 ± 0.2	16.2
5	75.3 ± 0.1	10.6	27	72.0 ± 0.3	11.2
6	76.4 ± 0.2	11.7	28	75.0 ± 0.3	14.2
7	73.7 ± 0.0	9.0	29	71.2 ± 0.2	10.4
8	74.2 ± 0.2	9.5	30	71.9 ± 0.2	11.1
9	76.1 ± 0.2	11.4	31	72.3 ± 0.2	11.5
10	74.1 ± 0.2	9.4	32	73.3 ± 0.1	12.5
11	75.2 ± 0.2	10.5	33	73.9 ± 0.3	13.1
12	66.3 ± 0.2	1.6	34	64.4 ± 0.2	3.6
13	67.8 ± 0.2	3.1	35	67.2 ± 0.1	6.4
14	66.8 ± 0.2	2.1	36	59.9 ± 0.2	-0.9
15	68.9 ± 0.0	4.1	37	66.1 ± 0.2	5.3
16	64.8 ± 0.2	0.1	38	62.8 ± 0.2	2.0
17	65.6 ± 0.2	0.9	39	64.8 ± 0.2	4.0
18	65.4 ± 0.2	0.7	40	63.4 ± 0.2	2.6
19	65.2 ± 0.2	0.5	41	62.3 ± 0.2	1.5
20	65.6 ± 0.3	0.8	42	63.4 ± 0.0	2.6
21	65.7 ± 0.2	0.9	43	64.8 ± 0.1	4.0
22	65.5 ± 0.2	0.8	44	64.2 ± 0.3	3.4

Table SI 2: Melting temperature and ΔT_m values for **1-44** on 14-mer oligonucleotides (oligonucleotide sequences provided in Materials and Methods). All values reported are averages of at least three melting temperature experiments (standard deviations indicated in parentheses.) $\Delta T_m = T_m(\text{DNA+polyamide}) - T_m(\text{DNA only})$.

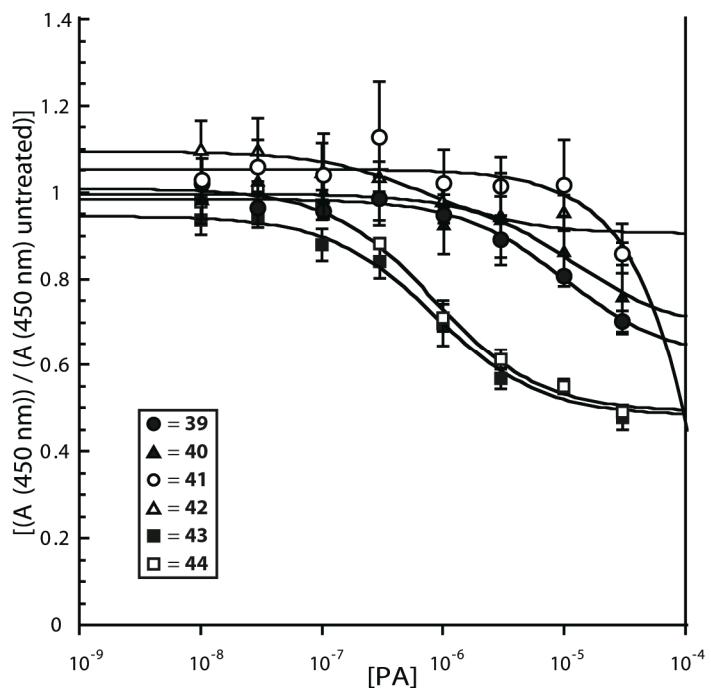


Figure SI 1: Cell viability data for **39-44** in LNCaP cells at 30, 10, 3, 1, 0.3, 0.1, 0.03 and 0.01 μM , performed according to the protocol provided in main text.

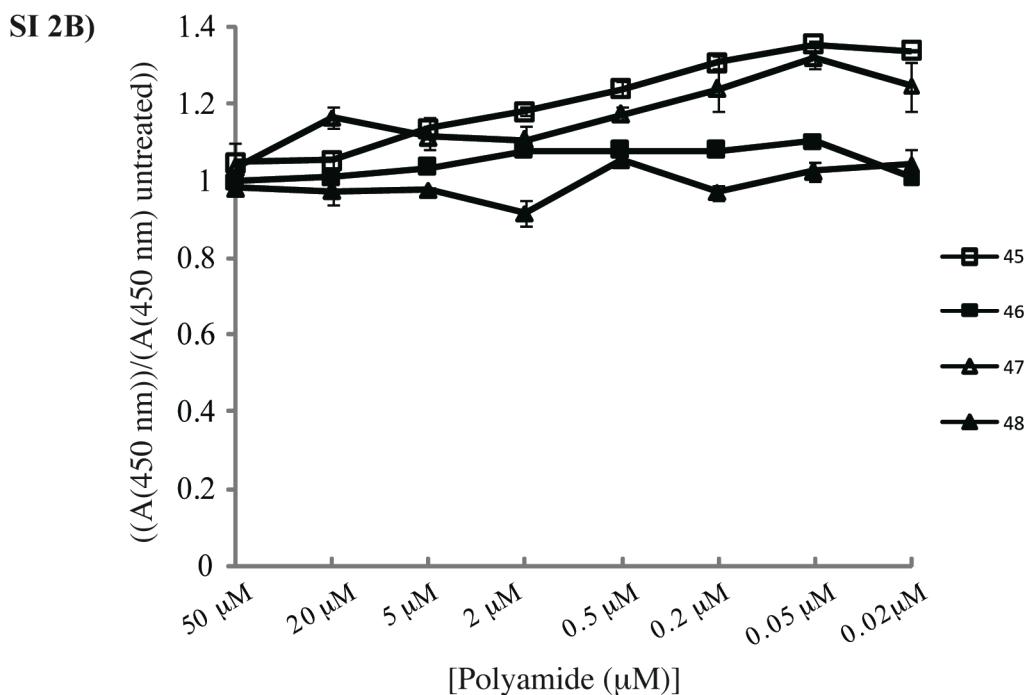
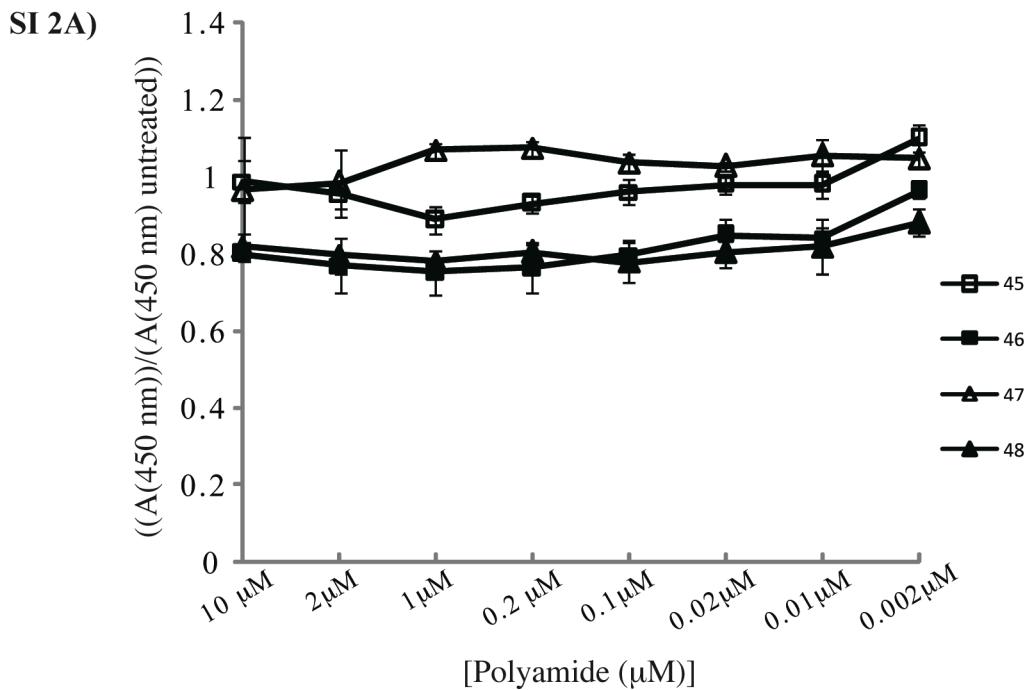


Figure SI 2: Cell viability data for **45-48** in U251 and LNCaP cells, performed according to the protocol provided in main text. A) Cell viability data for **45-48** in U251 cells at 10, 2, 1, 0.2, 0.1, 0.02, 0.01 and 0.002 μM . B) Cell viability data for **45-48** in LNCaP cells at 50, 20, 5, 2, 0.5, 0.2, 0.05 and 0.02 μM .