

Electronic Supplementary Information 4

Non-identical Electronic Characters of the Internucleotidic Phosphates in RNA Modulate the Chemical Reactivity of the Phosphodiester Bonds

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Content:

Figure S13. Panels (a1) – (a8), (b1) – (b10), (c1) – (c8), (d1) – (d12), (e1) – (e11), (f1) – (f9) and (g1) – (g8) show the Maldi Tof negative ion mode spectrum of the peaks separated by RP-Hplc and SMART™ RP-Hplc after 1h of alkali digestion for all heptameric ssRNAs. **p. S2 – S70**

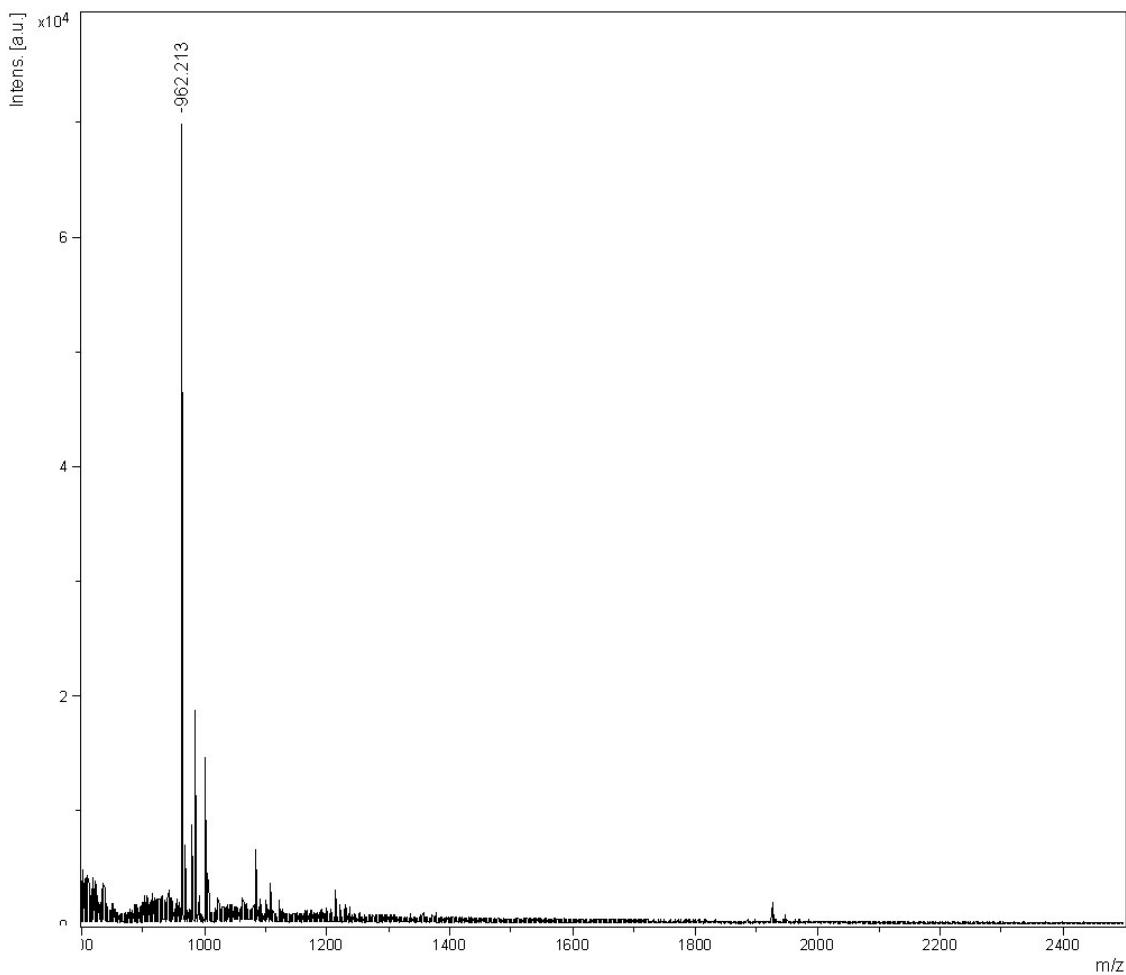
Figure S13

Figure S13(a1): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 21.52$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGAAC)-3'** (**5b**) after **1h** [for Hplc profile, see Fig **S12 (a2)**] showing the main peak at m/z **962.2** for **5'-CAA_{2'/3'-P}** [see Table S9(A)].

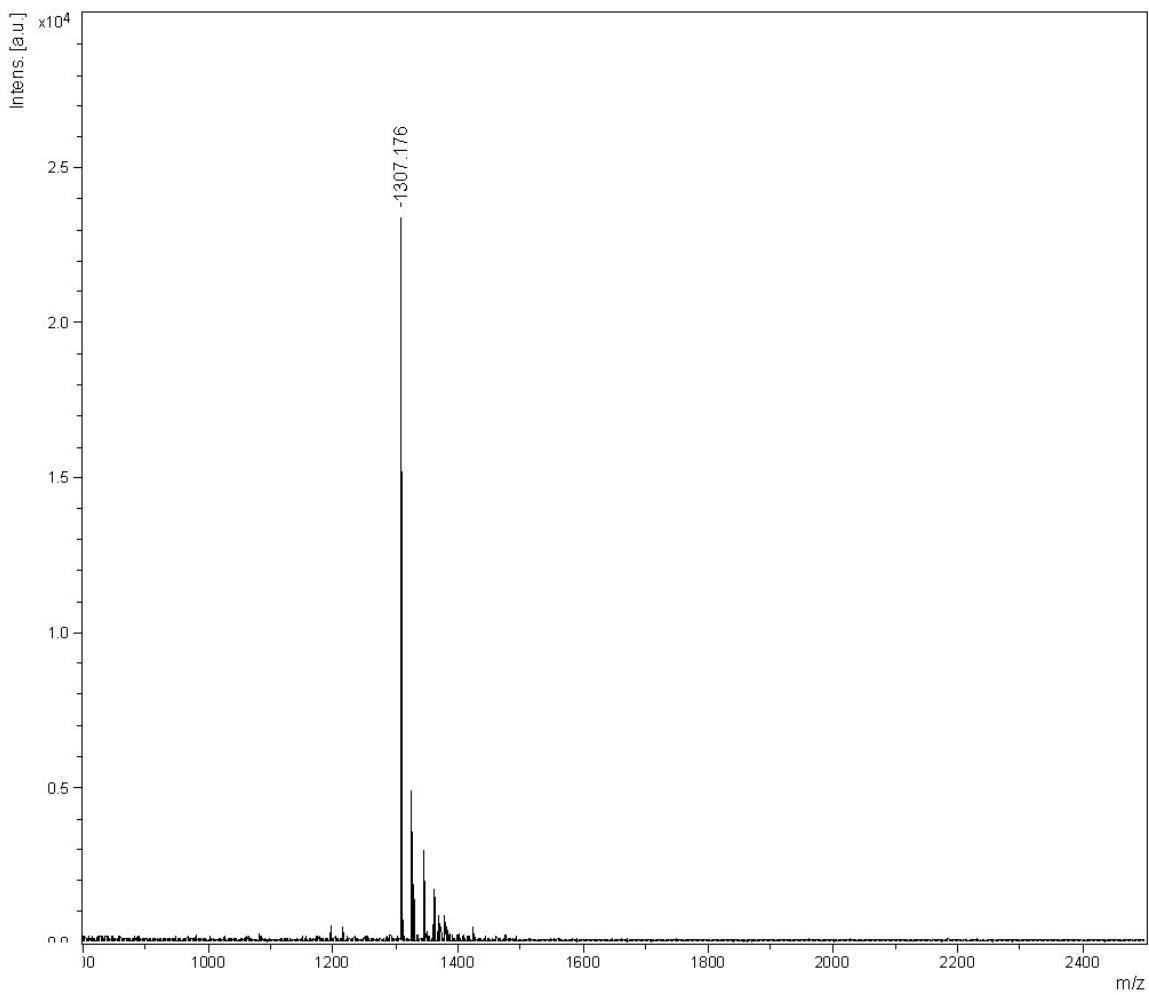


Figure S13(a2): Maldi Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 34.04$ min in the Fig S12(a2i). Corresponding RP-Hplc fraction is $R_T = 23.86$ min in the Fig S12(a2) [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGAAC)-3', 5b**]. The main peak at m/z **1307.3** is for **5'-CAAG_{2'},3'-cMP** [see Table S9(A)].

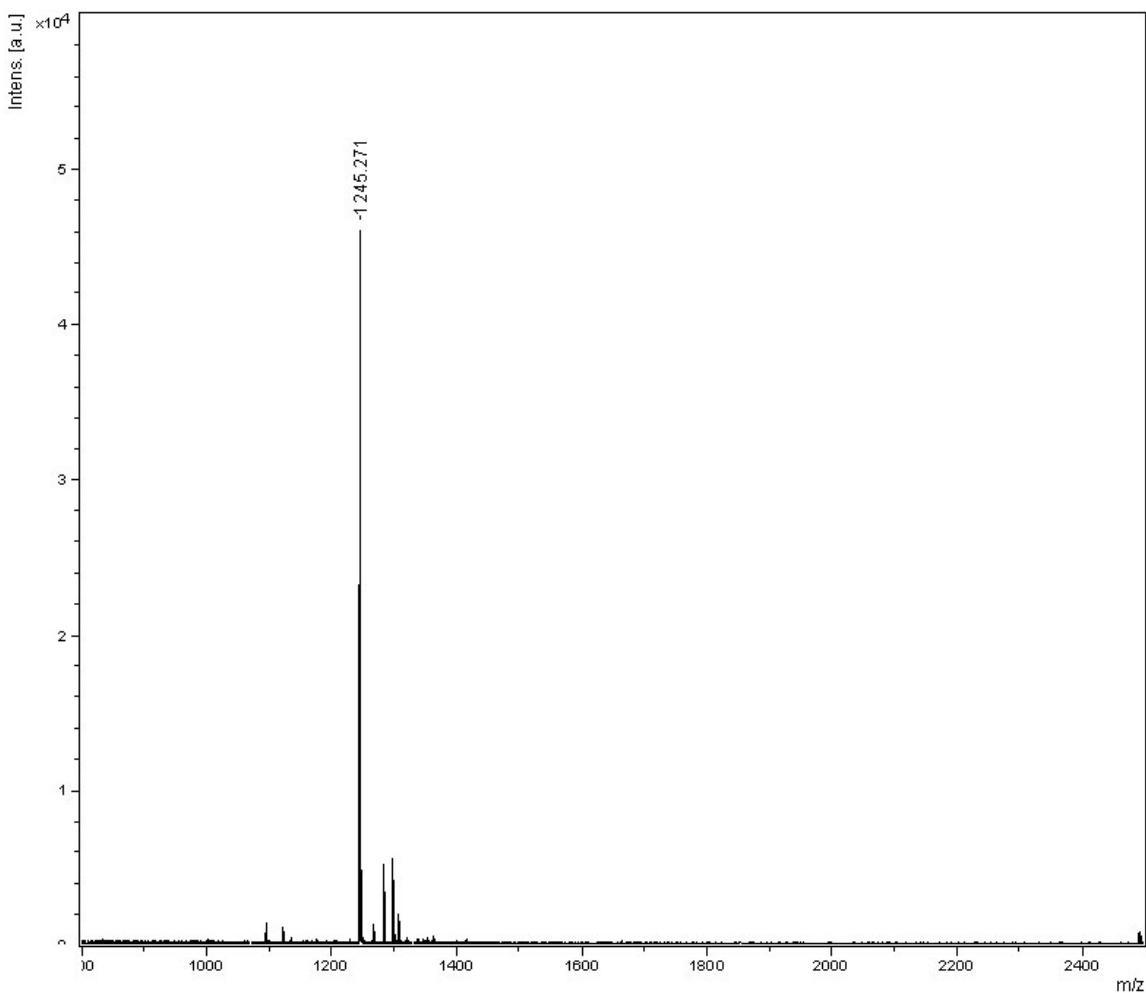


Figure S13(a3): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 34.77$ min in the **Fig S12(a2i)**. Corresponding RP-Hplc fraction is $R_T = 23.86$ min in the **Fig S12(a2)** [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGAAC)-3'**, **5b**]. The main peak at m/z **1245.3** is for **5'-GAAC-3'** [see Table S9(A)].

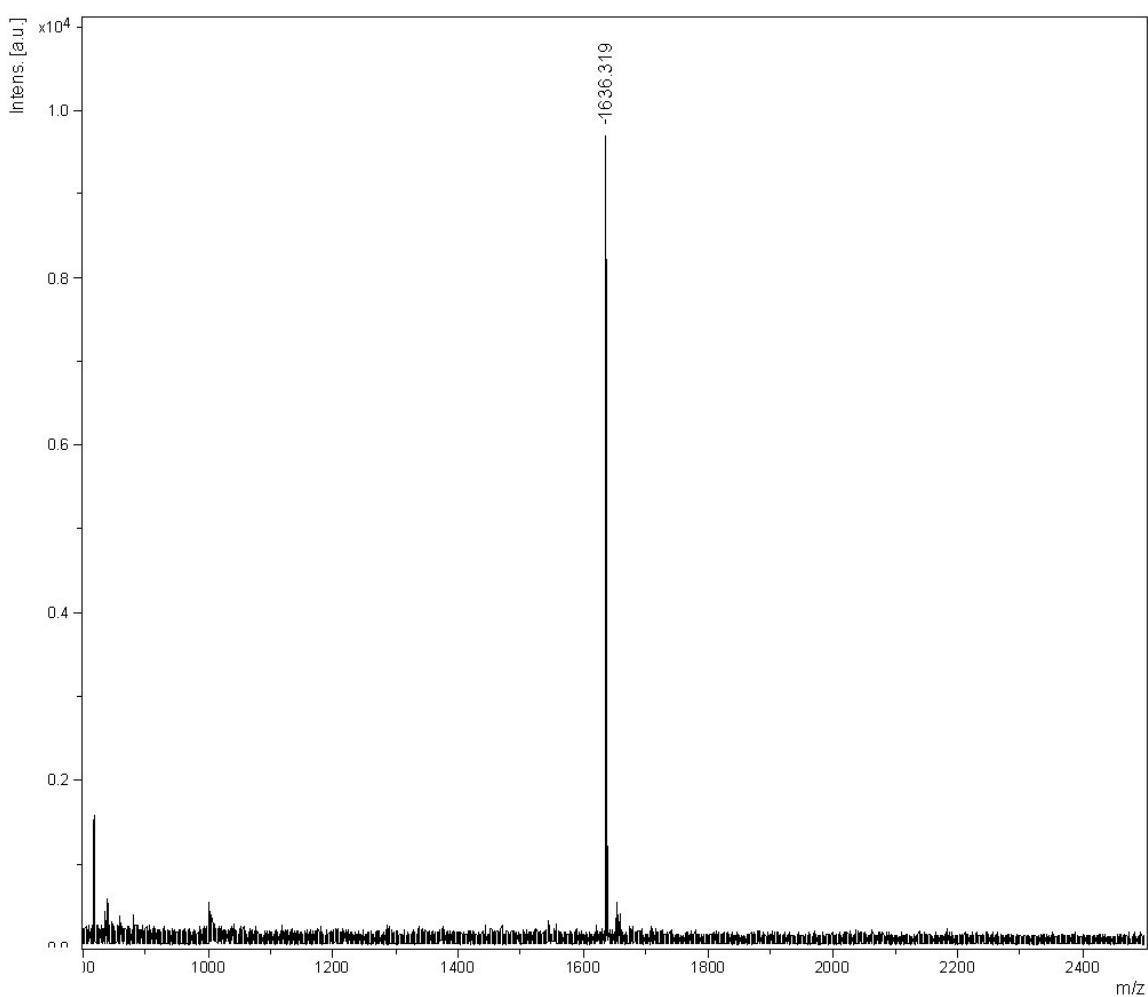


Figure S13(a4): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 27.12$ min in the Fig S12(a2j). Corresponding RP-Hplc fraction is $R_T = 25.44$ min in the Fig S12(a2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CAAGAAC)-3', 5b]. The main peak at m/z 1636.3 is for 5'-CAAGA_{2'},3'-cMP [see Table S9(A)].

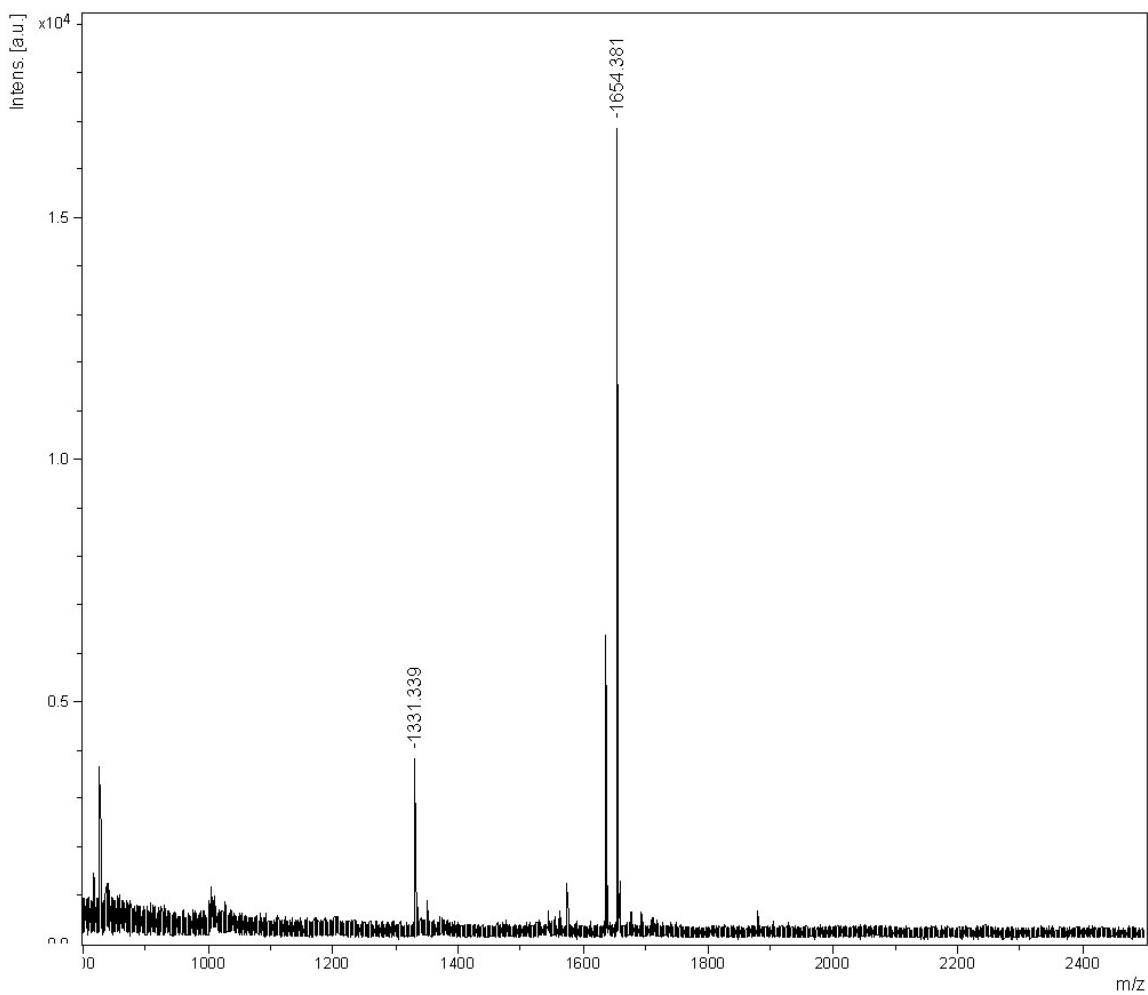


Figure S13(a5): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 27.41$ min in the Fig S12(a2j). Corresponding RP-Hplc fraction is $R_T = 25.44$ min in the Fig S12(a2) [after 1 h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CAAGAAC)-3', 5b]. The main peak at m/z 1654.4 is for 5'-CAAGA_{2'3'-P} and at m/z 1636.3 is for 5'-CAAGA_{2',3'-cMP} [see Table S9(A)].

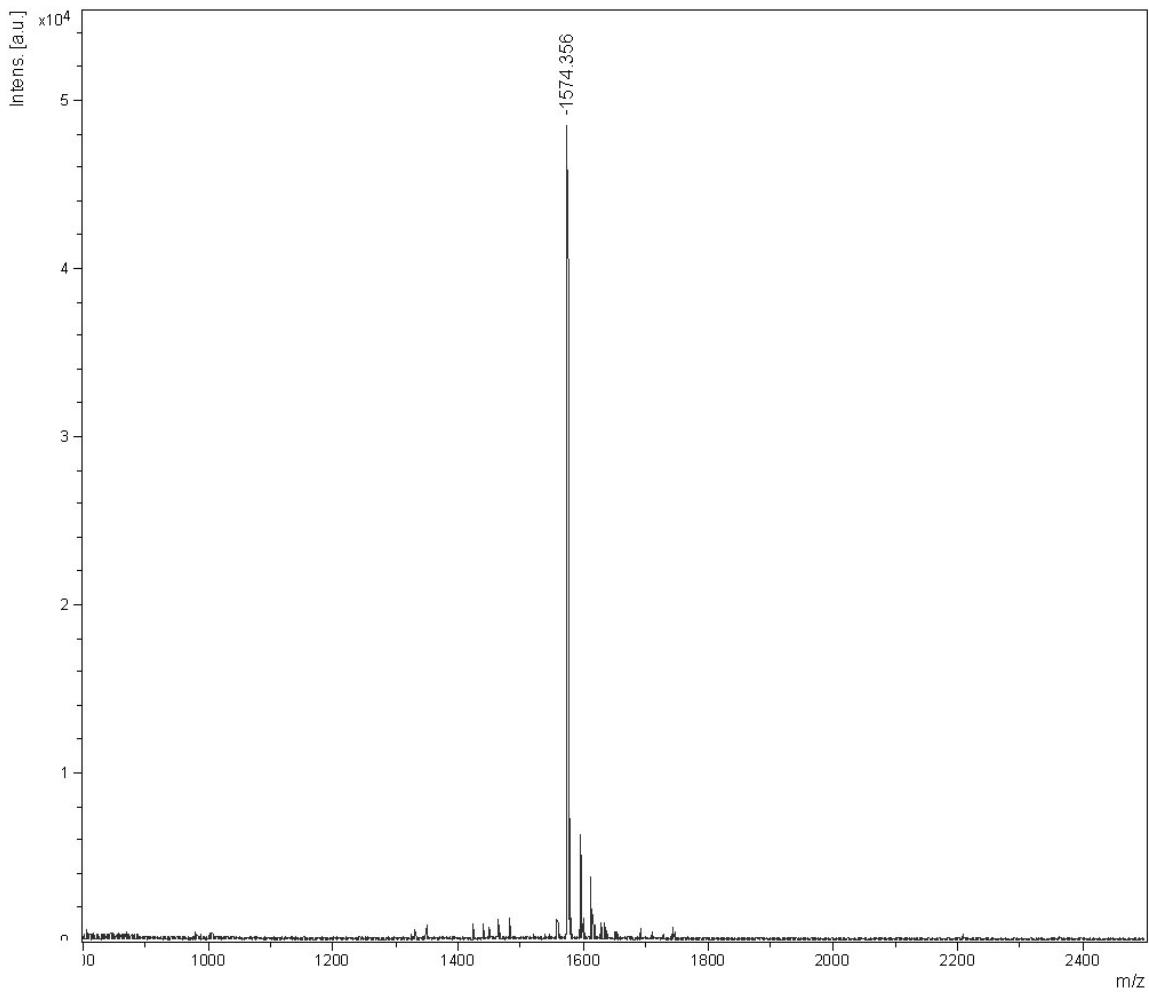


Figure S13(a6): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 27.93$ min in the **Fig S12(a2j)**. Corresponding RP-Hplc fraction is $R_T = 25.44$ min in the **Fig S12(a2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGAAC)-3'**, **5b**]. The main peak at m/z **1574.4** is for **5'-AGAAC-3'** [see Table S9(A)].

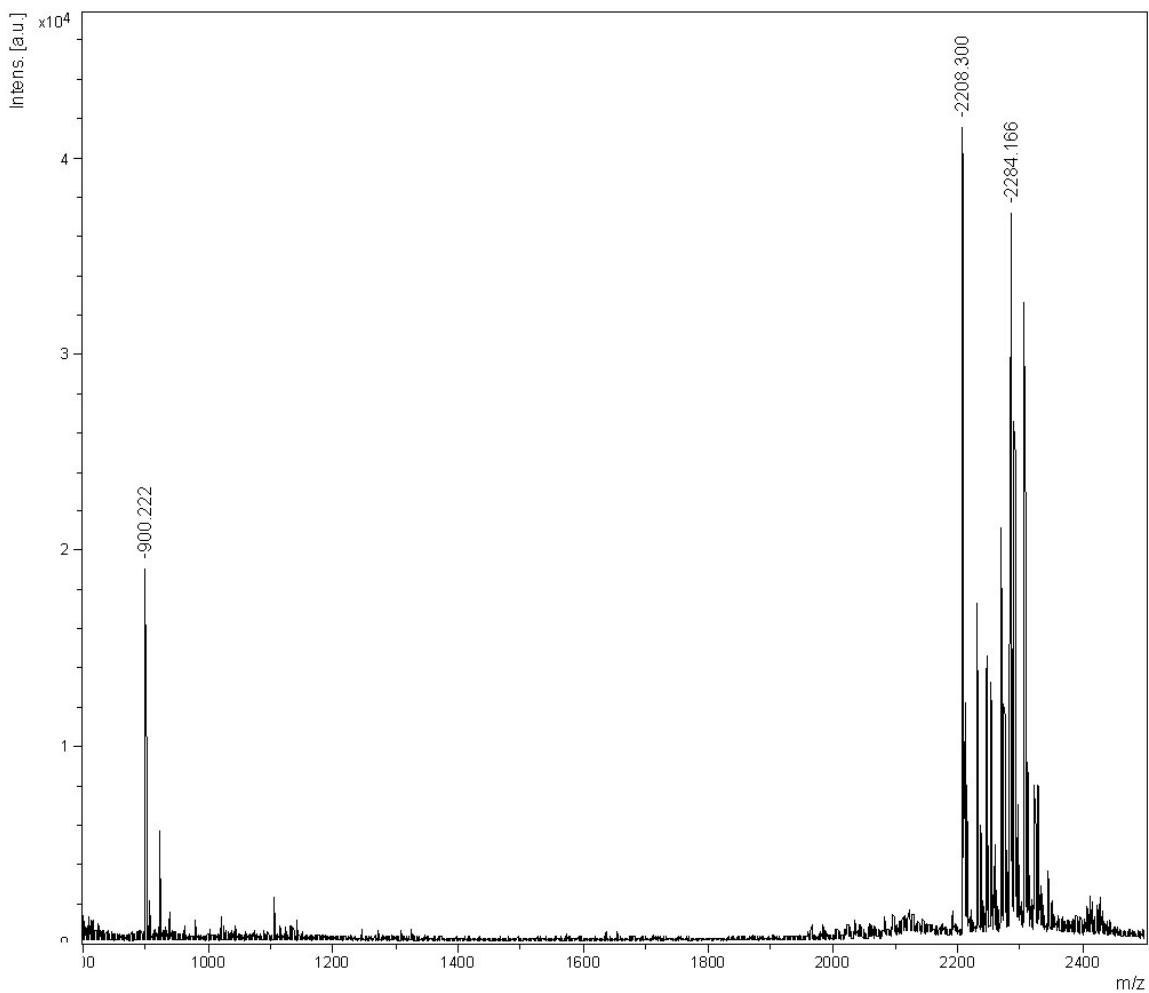


Figure S13(a7): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 26.21$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGAAC)-3'** (**5b**) after **1h** [for Hplc profile, see Fig **S12 (a2)**] showing the main peak at m/z **900.2** for 5'-AAC-3' (a fragmentation peak of the heptameric peak in the process of recording mass as confirmed by the variation of the laser power in the Maldi Tof), **2208.3** for the parent heptamer and **2284.2** for some non-nucleos(t)idic fragment [see Table S9(A)].

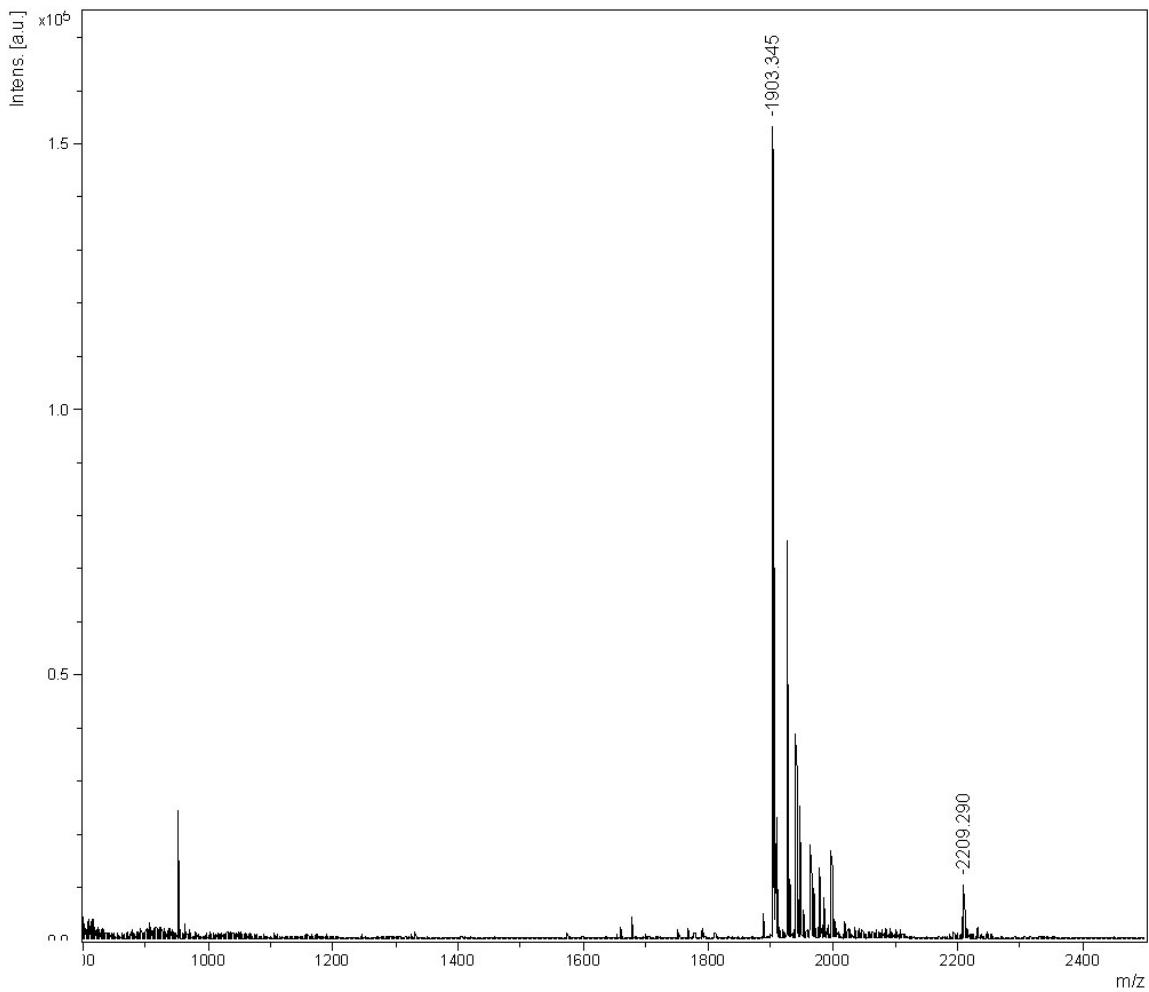


Figure S13(a8): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 27.10$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGAAC)-3'** (**5b**) after **1h** [for Hplc profile, see Fig **S12 (a2)**] showing the main peak at m/z **1903.3** for **5'-AAGAAC-3'** [see Table S9(A)].

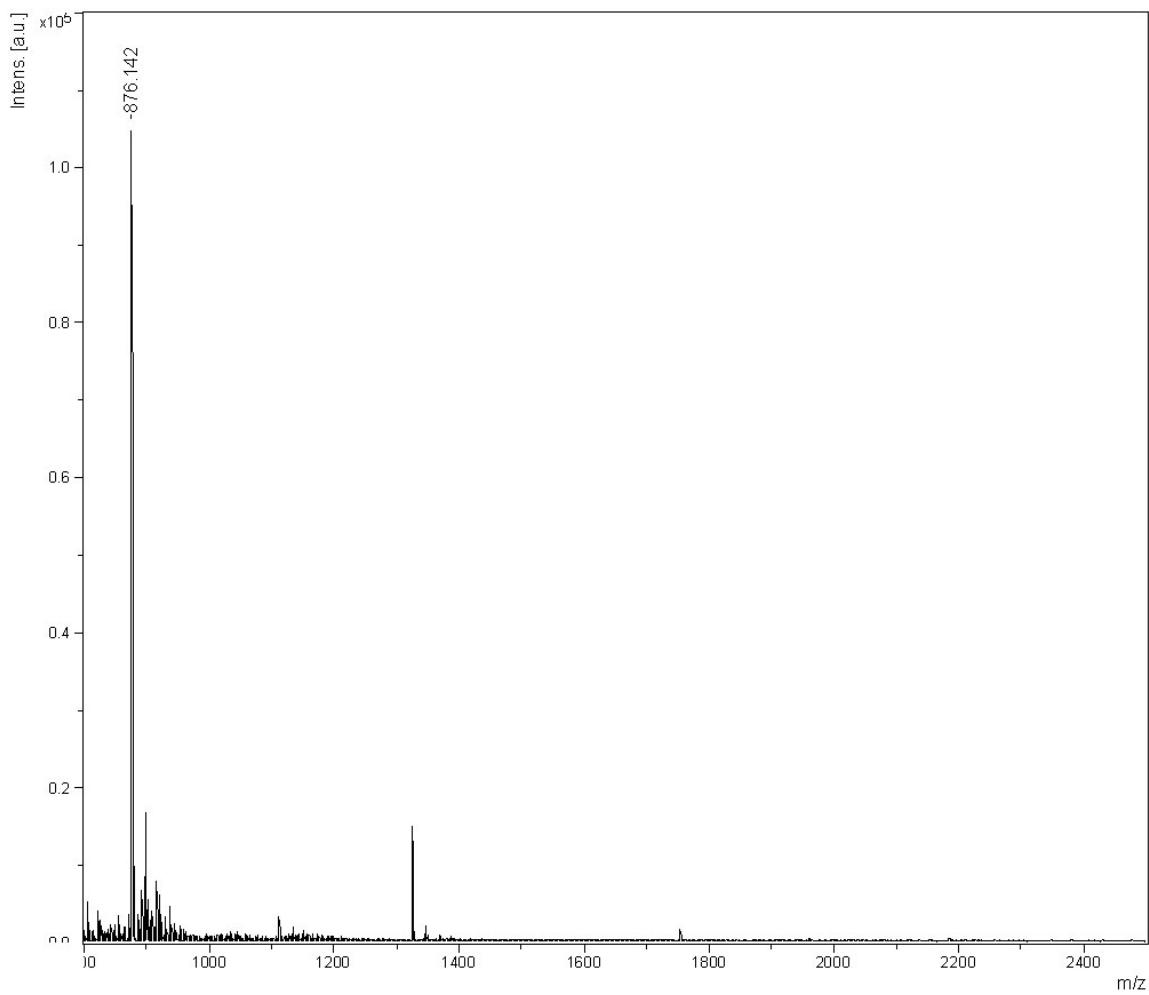


Figure S15(b1): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 19.87$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGCAC)-3' (6b)** after **1h** [for RP-Hplc profile, see Fig **S12 (b2)**] showing the main peak at m/z **876.1** for **5'-CAC-3'** [see Table S9(B)].

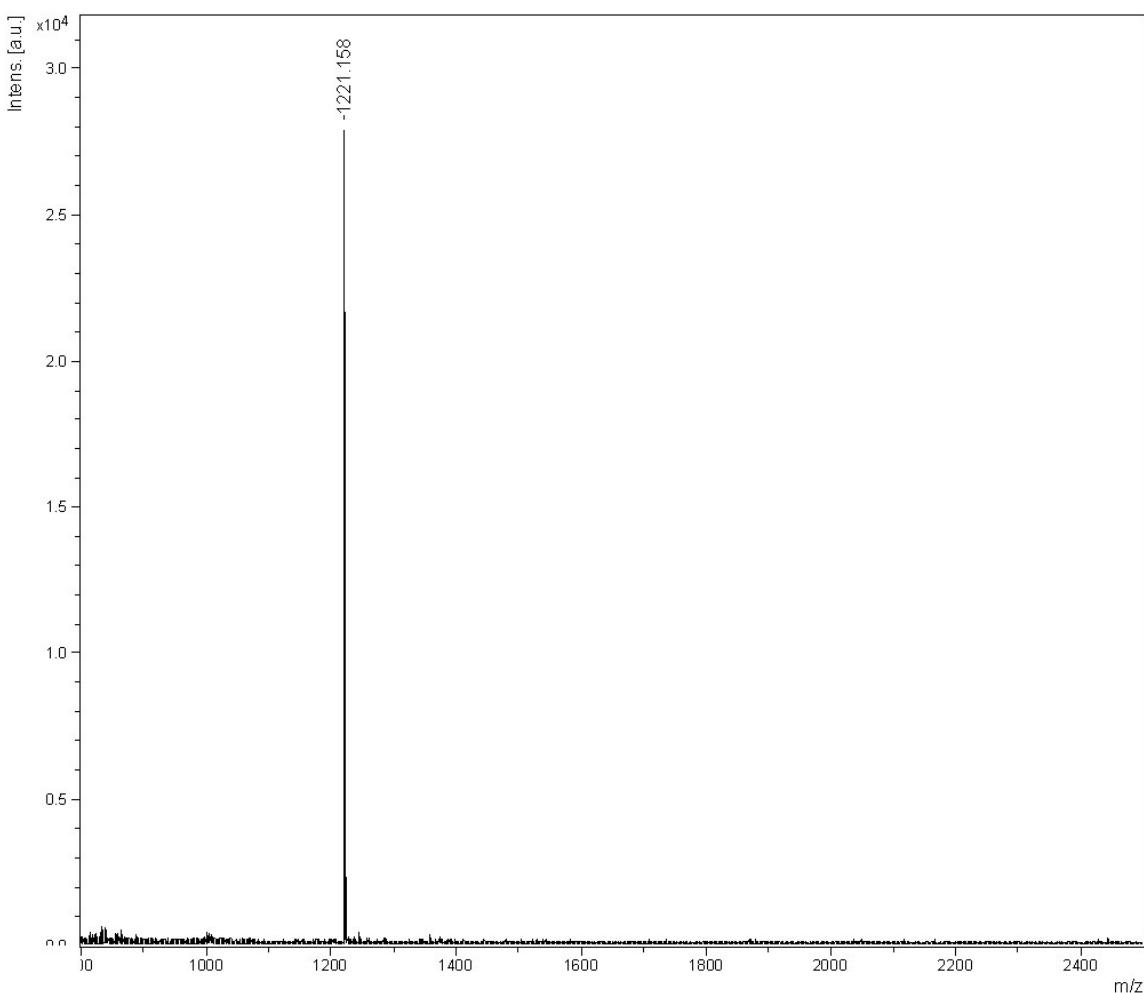


Figure S13(b2): Maldi-Tof mass-spectral analysis of the SMARTTM RP-Hplc fraction at $R_T = 20.68$ min in the **Fig S12(b2i)**. Corresponding RP-Hplc fraction is $R_T = 21.18$ min in the **Fig S12(b2)** [after **1h** alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGCAC)-3'**, **6b**]. The main peak at m/z **1221.3** is for **5'-GCAC-3'** [see Table S9(B)].

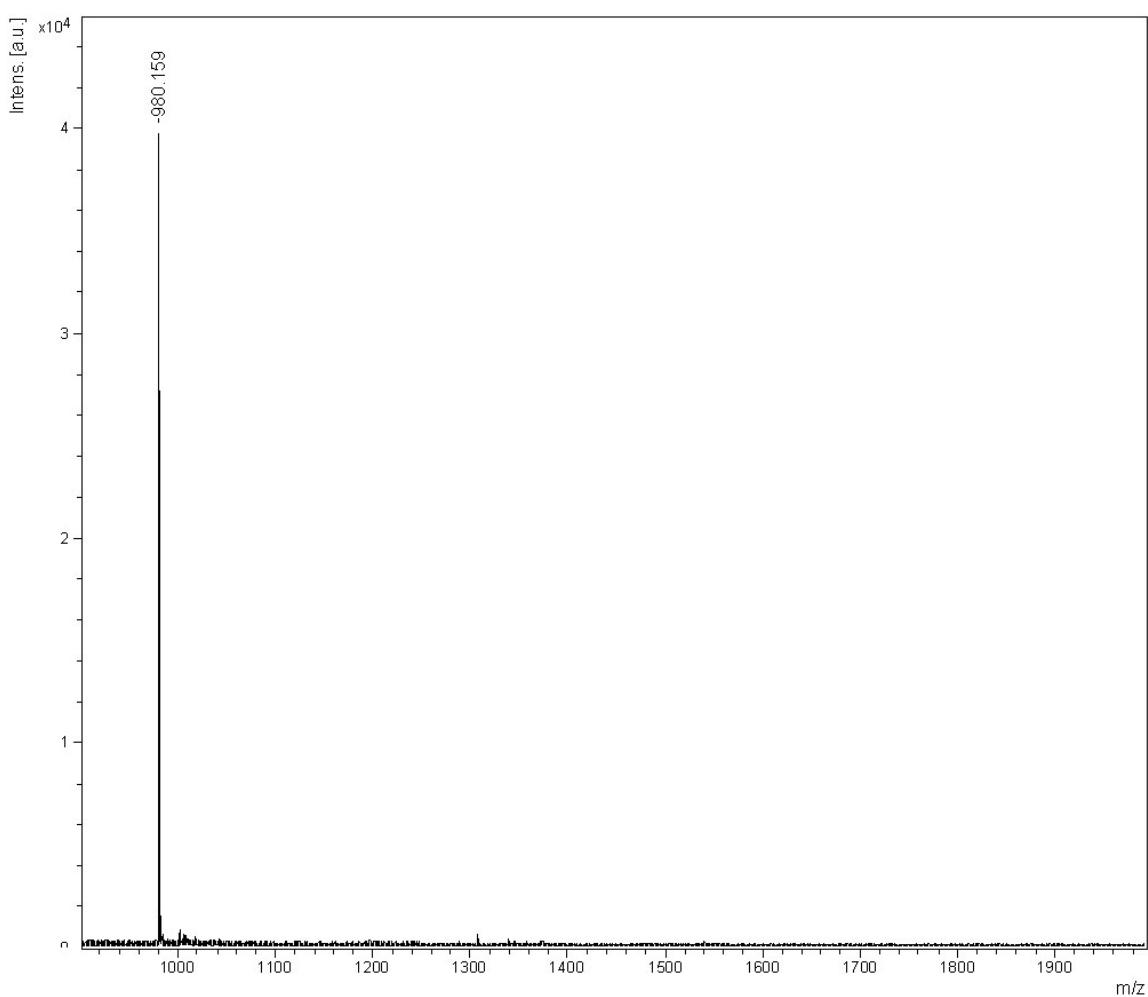


Figure S13(b3): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 22.16$ min and $R_T = 22.51$ min in the Fig S12(b2i). Corresponding RP-Hplc fraction is $R_T = 21.18$ min in the Fig S12(b2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CAAGCAC)-3', 6b]. The main peak at m/z 980.2 is for 5'-CAA_{2'3'-P} [see Table S9(B)].

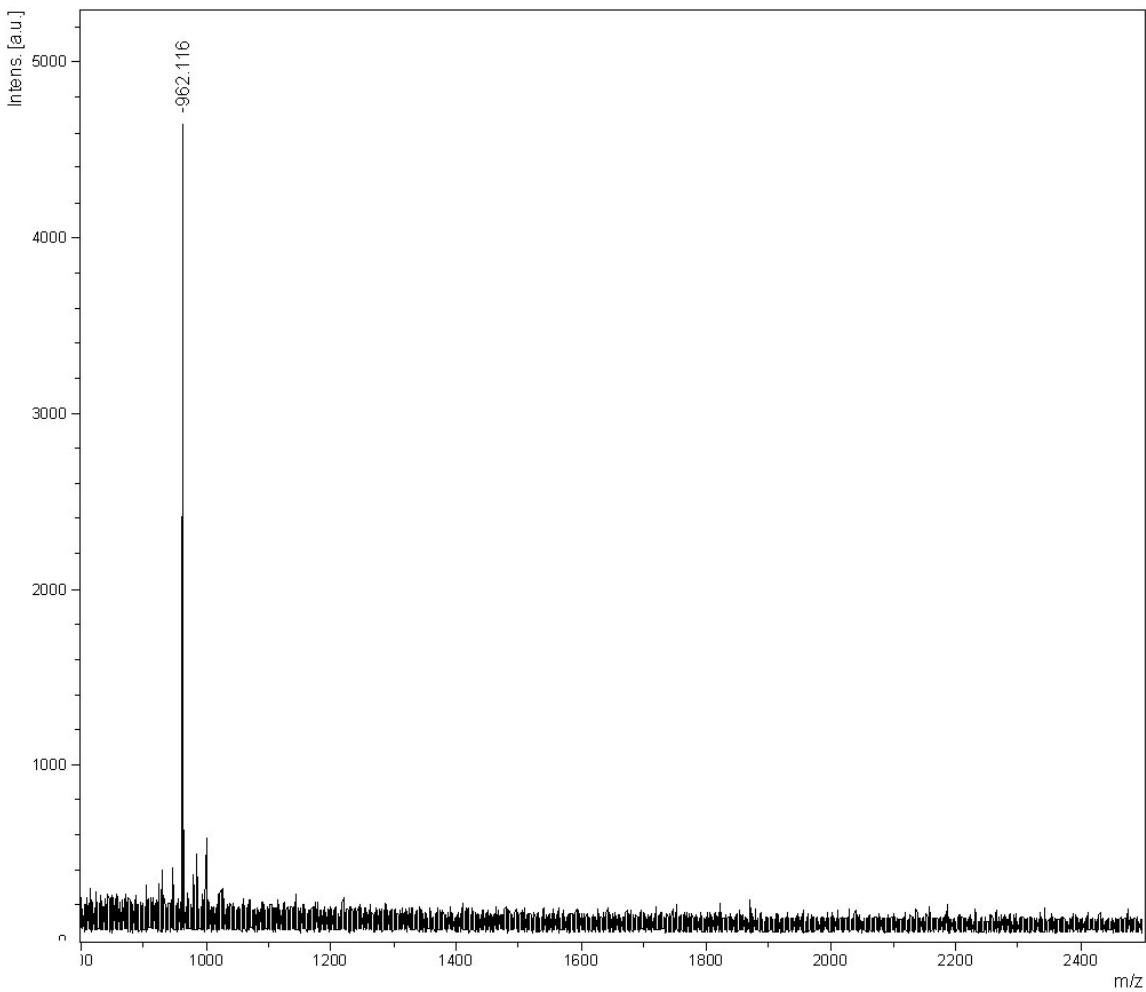


Figure S13(b4): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 21.77$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGCAC)-3' (6b)** after **1h** [for RP-Hplc profile, see Fig **S12 (b2)**] showing the main peak at m/z **962.1** for **5'-CAA₂, 3'-cMP** [see Table S9(B)].

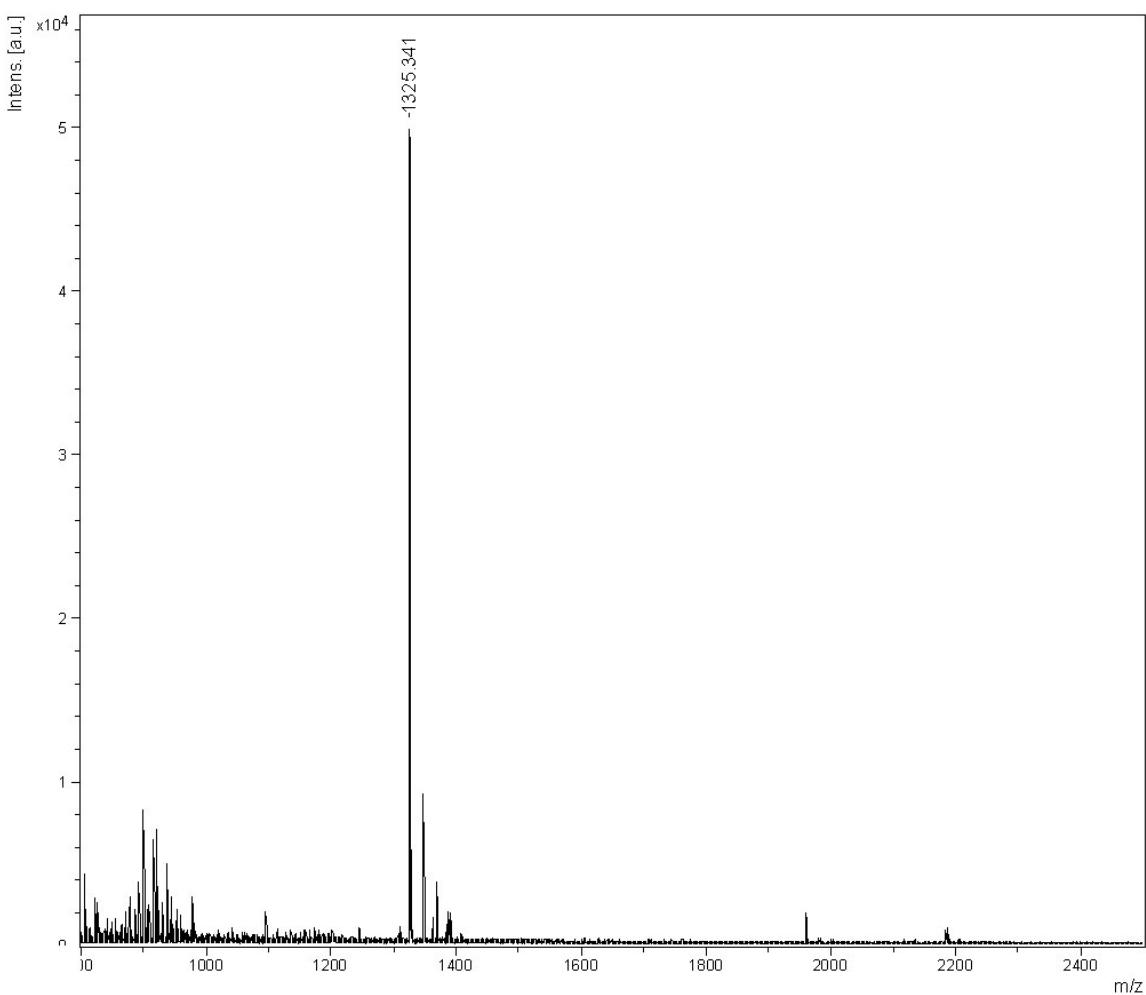


Figure S13(b5): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 22.76$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-rCAAGCAC-3' (6b)** after **1h** [for Hplc profile, see Fig **S12 (b2)**] showing the main peak at m/z **1325.3** for **5'-CAAG_{2'3'-P}** [see Table S9(B)].

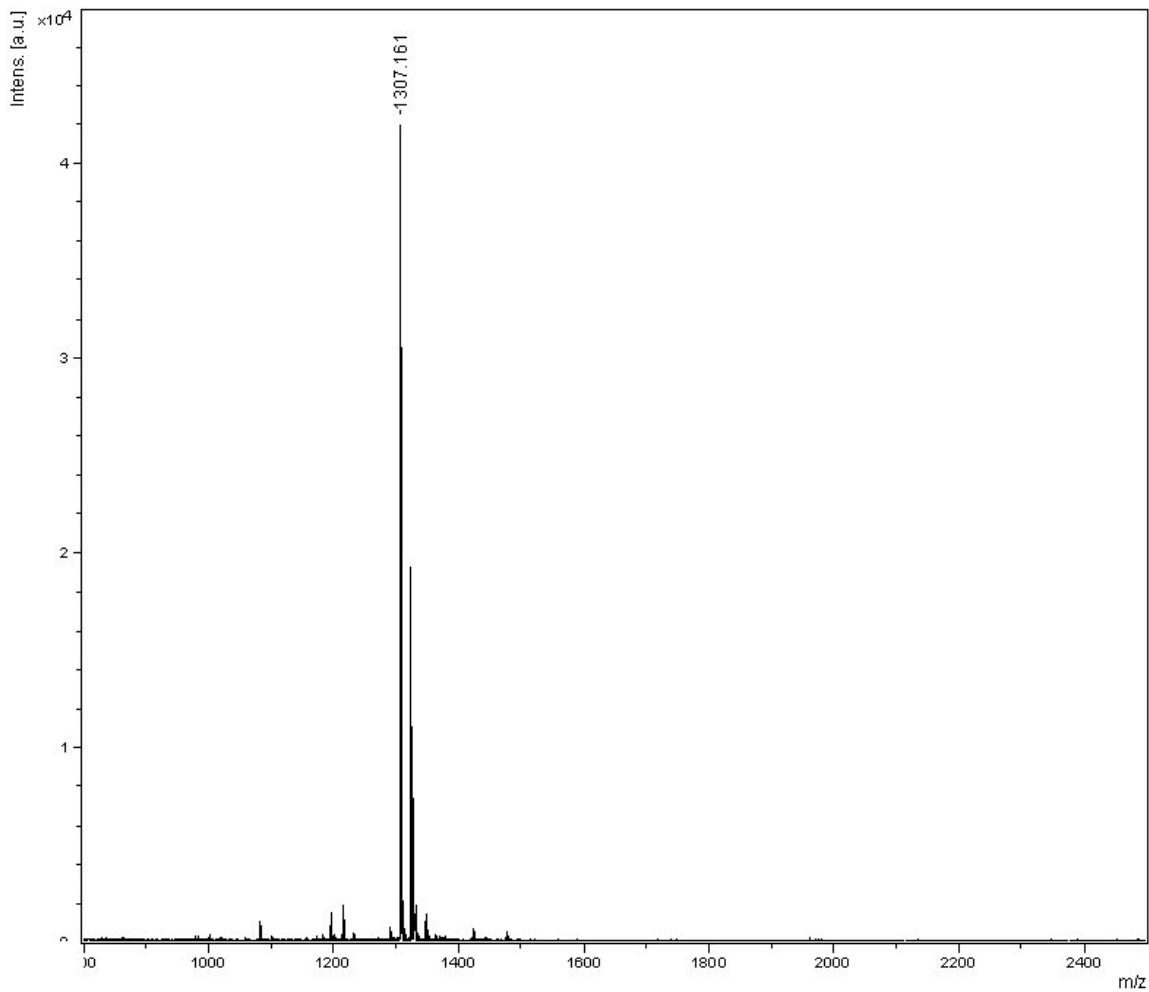


Figure S13(b6): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 29.33$ min in the **Fig S12(b2j)**. Corresponding RP-Hplc fraction is $R_T = 24.23$ min in the **Fig S12(b2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGCAC)-3'**, **6b**]. The main peak at m/z **1307.2** is for **5'-CAAG_{2'},3'-cMP** and **1325.2** is for **5'-CAAG_{2'/3'}-P** [see Table S9(B)].

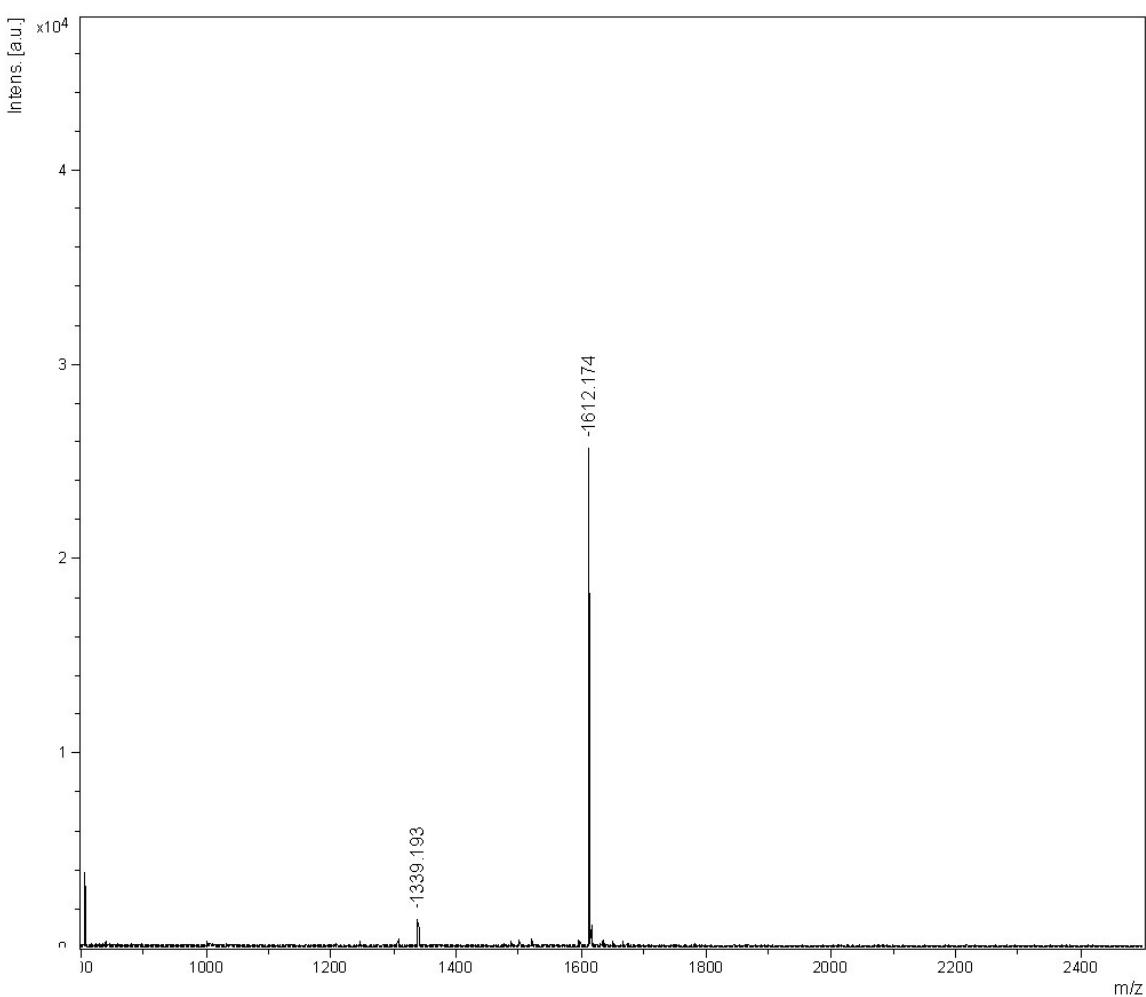


Figure S12(b7): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 30.12$ min in the Fig S12(b2j). Corresponding RP-Hplc fraction is $R_T = 24.23$ min in the Fig S12(b2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGCAC)-3'**, **6b**]. The main peak at m/z **1612.2** is for **5'-CAAGC_{2'},3'-cMP** [see Table S9(B)].

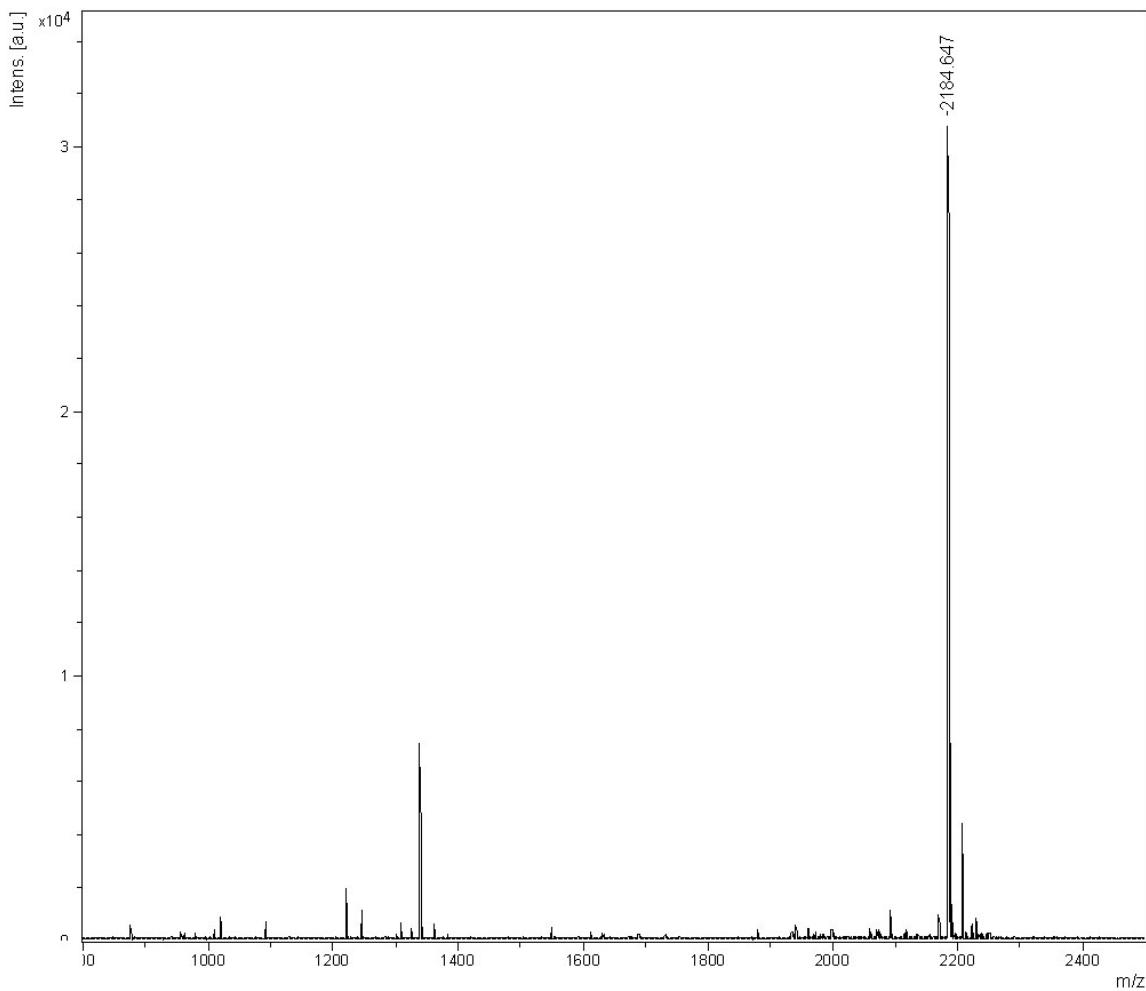


Figure S13(b8): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 25.27$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAGCAC)-3' (6b)** after **1h** [for RP-Hplc profile, see Fig S12 (b2)] showing the main peak at m/z **2184.7** for the parent heptamer [see Table S9(B)].

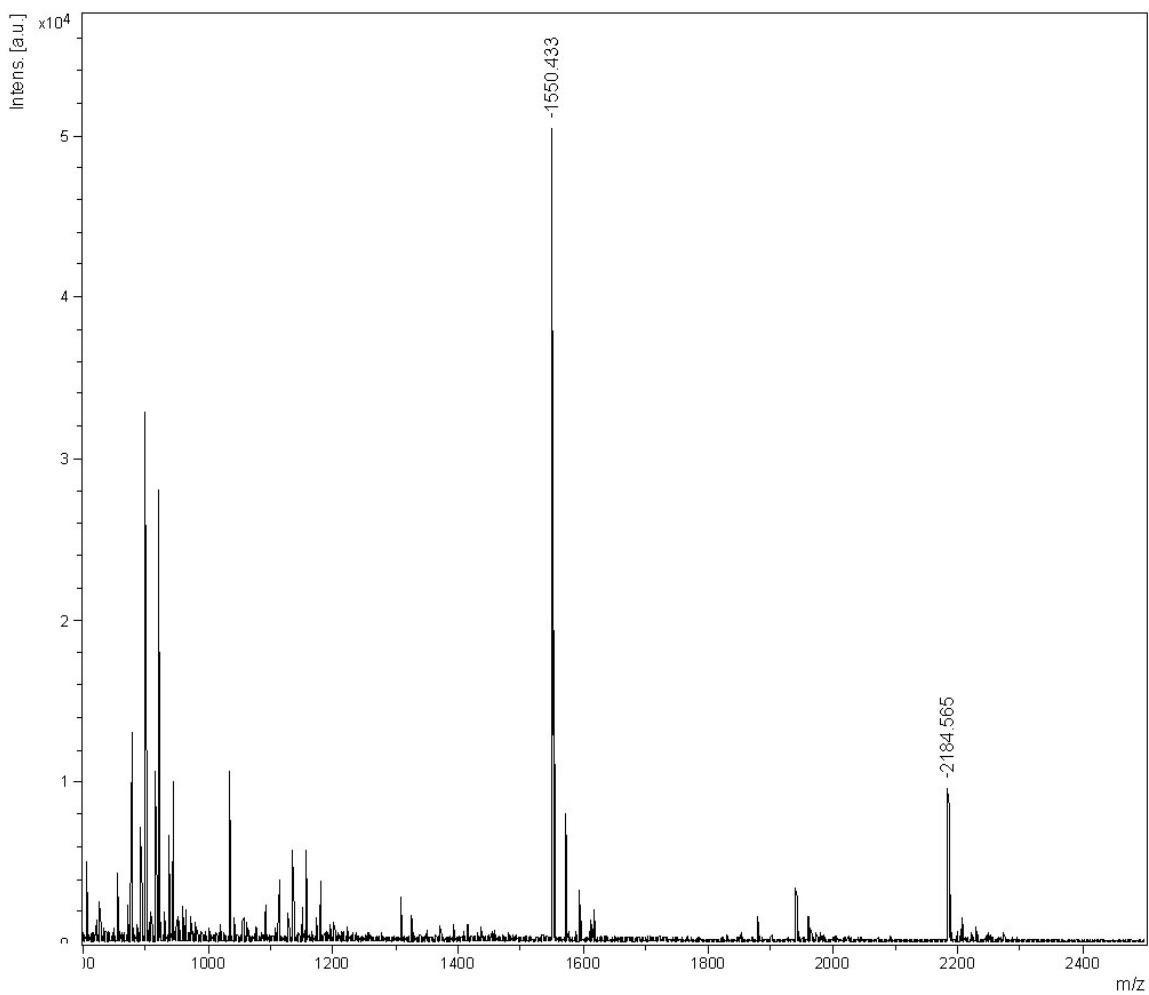


Figure S13(b9): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = \sim 25.5$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-rCAAGCAC-3' (6b)** after **1h** [for Hplc profile, see Fig S12 (b2)] showing the main peak at m/z **1550.4** for **5'-AGCAC-3'** and **2184.6** for the parent heptamer[see Table S9(B)].

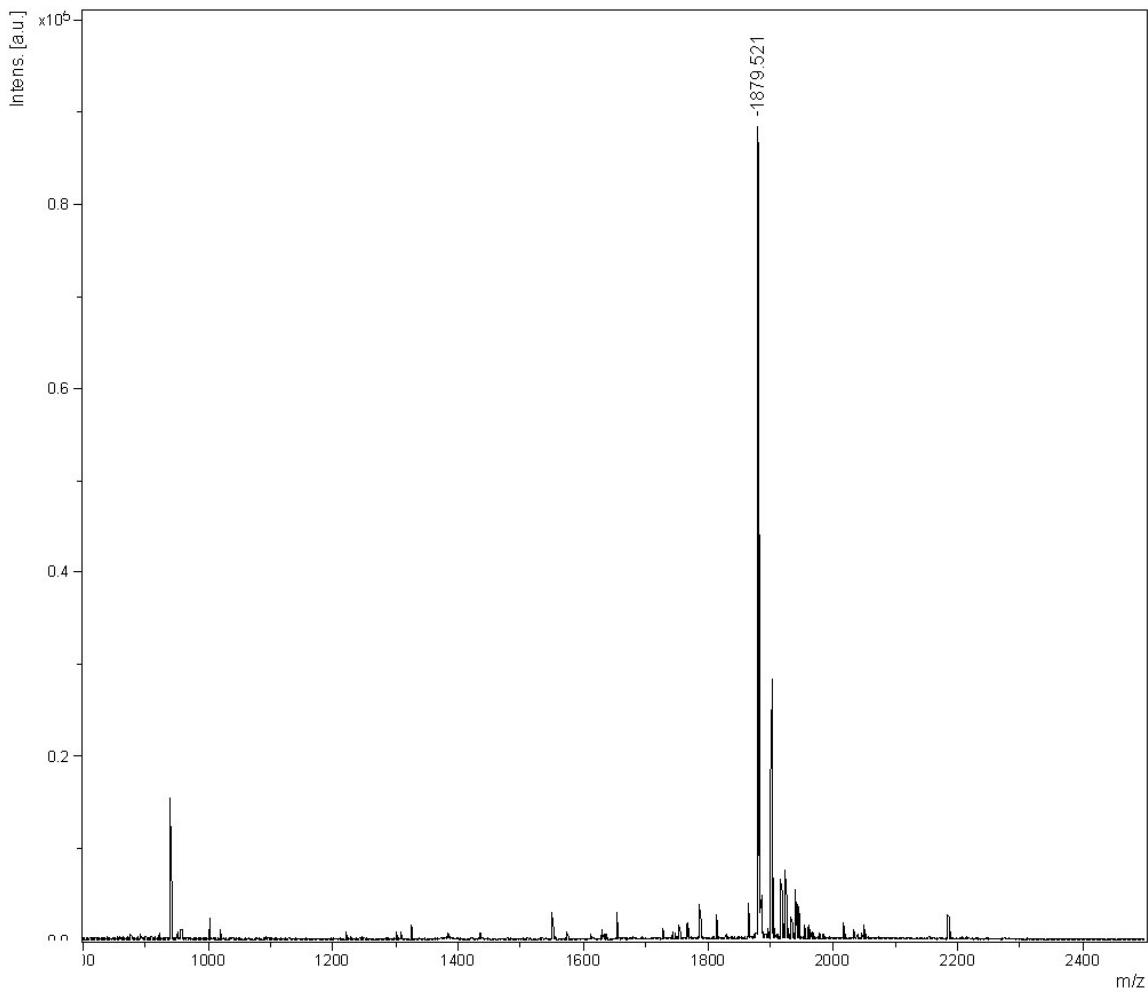


Figure S13(b10): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 27.11$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-rCAAGCAC-3'** (**6b**) after **1h** [for RP-Hplc profile, see Fig S12 (b2)] showing the main peak at m/z **1879.5** for **5'-AAGCAC-3'** [see Table S9(B)].

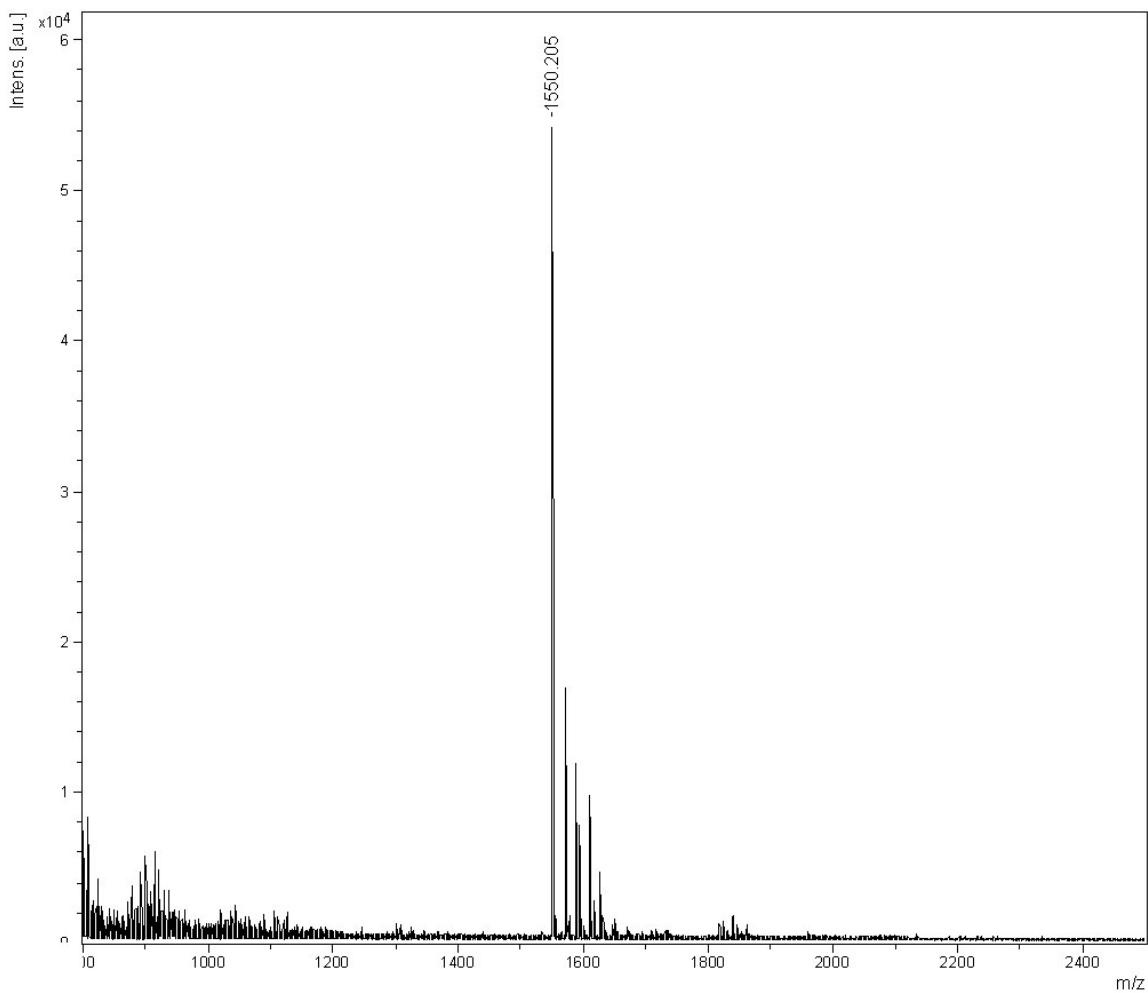


Figure S13(c1): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 23.18$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACGAAC)-3'** (**7b**) after **1h** [for RP-Hplc profile, see Fig S12 (c2)] showing the main peak at m/z **1550.2** for **5'-CGAAC-3'** and [see Table S9(C)].

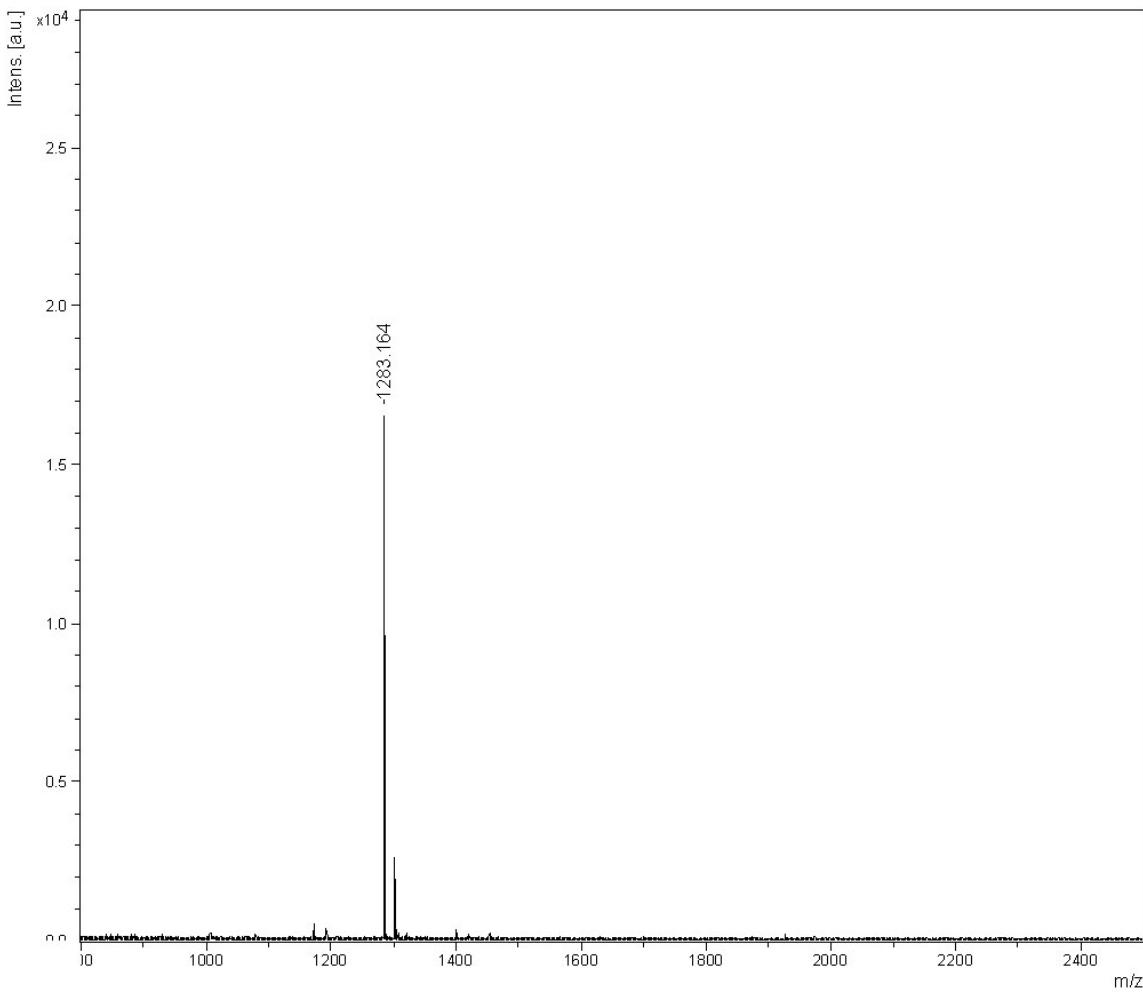


Figure S13(c2): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 30.27$ min in the Fig S13(c2i). Corresponding RP-Hplc fraction is $R_T = 24.97$ min in the Fig S12(c2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACGAAC)-3'**, **7b**]. The main peak at m/z 1283.2 is for **5'-CACG₂, 3'-cMP** [see Table S9(C)].

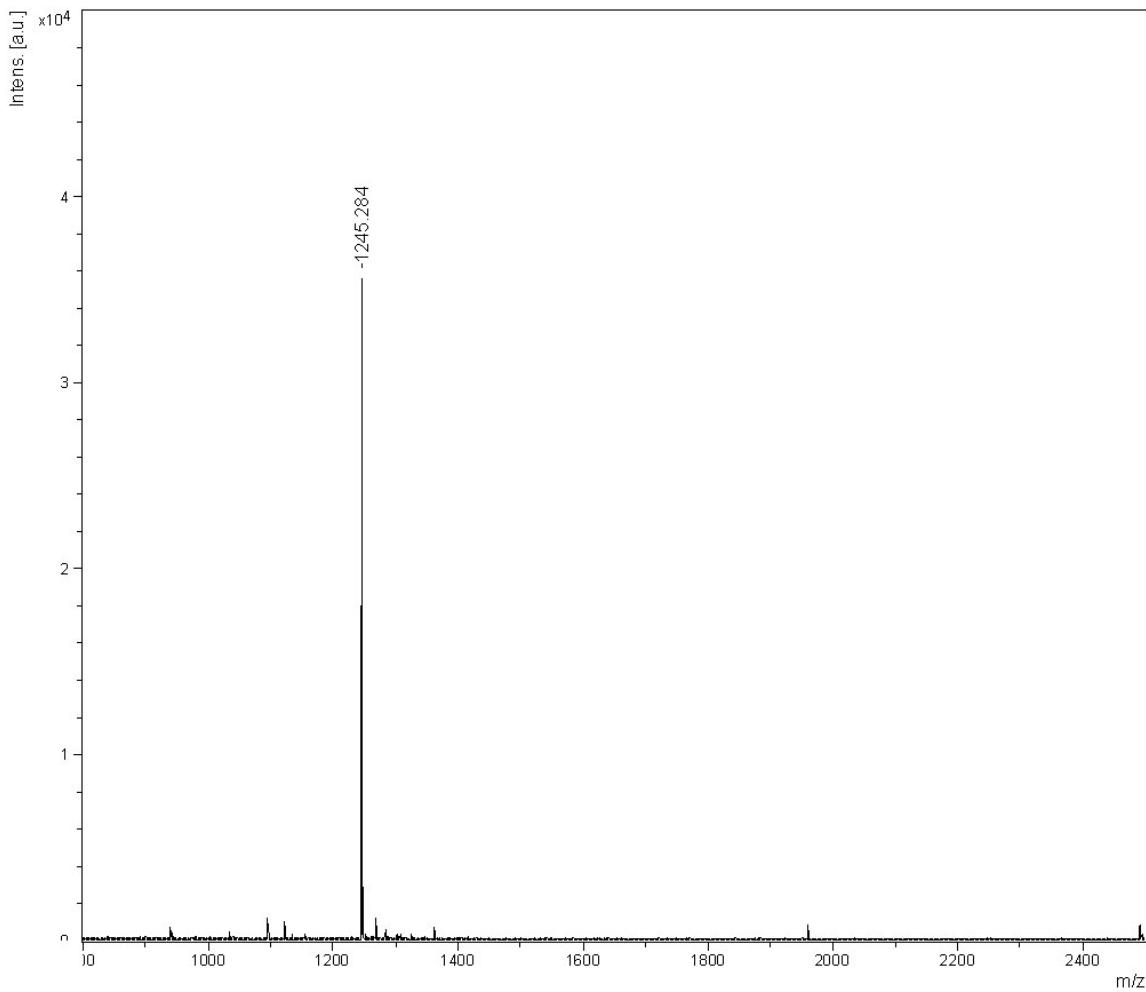


Figure S13(c3): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 31.48$ min in the Fig S12(c2i). Corresponding RP-Hplc fraction is $R_T = 24.97$ min in the Fig S12(c2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CACGAAC)-3', 7b]. The main peak at m/z 1245.3 is for 5'-GAAC-3' [see Table S9(C)].

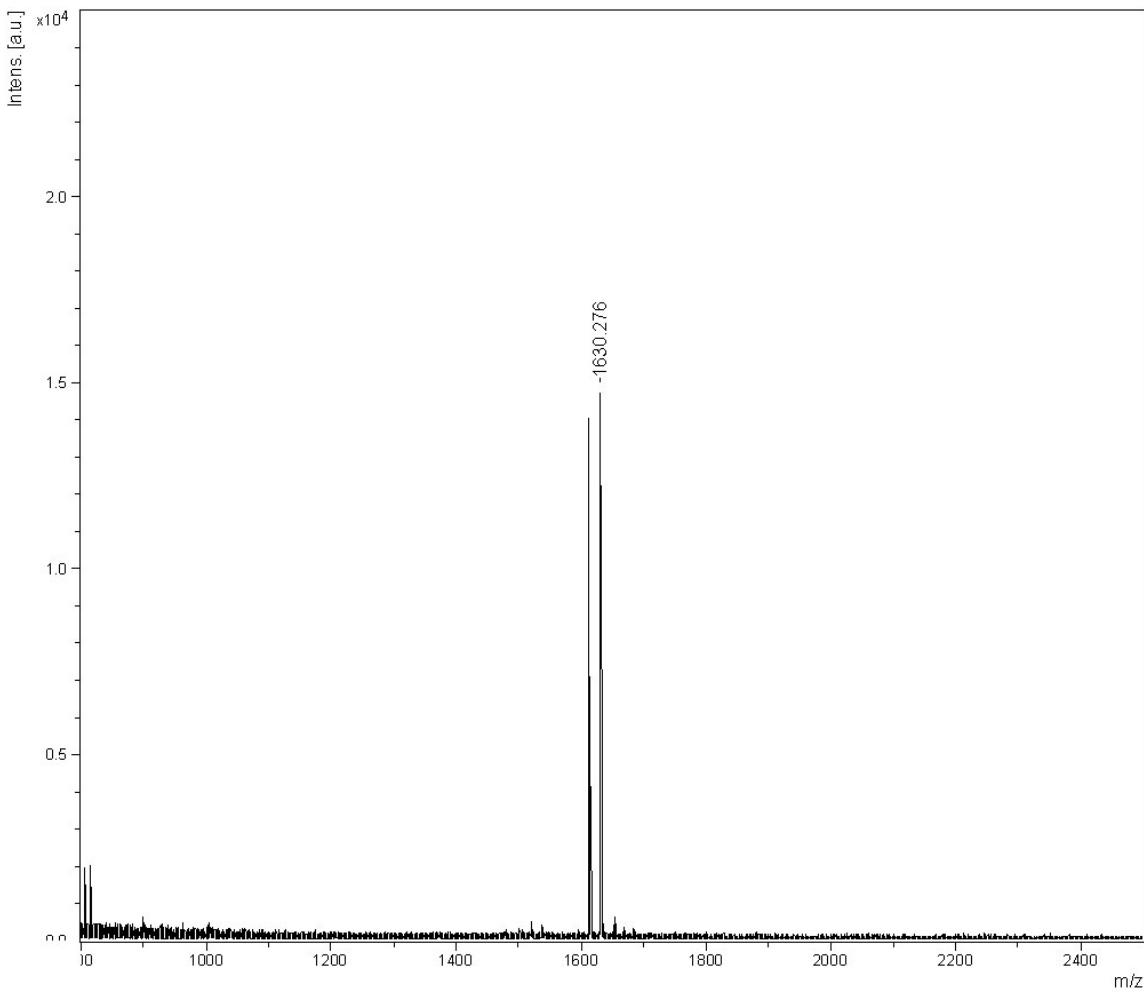


Figure S13(c4): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 30.63$ min in the Fig S12(c2j). Corresponding RP-Hplc fraction is $R_T = 26.50$ min in the Fig S12(c2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CACGAAC)-3', 7b]. The main peak at m/z 1612.3 is for 5'-CACGA_{2'},_{3'}-cMP and 1630.3 is for 5'-CACGA_{2'/3'}-P [see Table S9(C)].

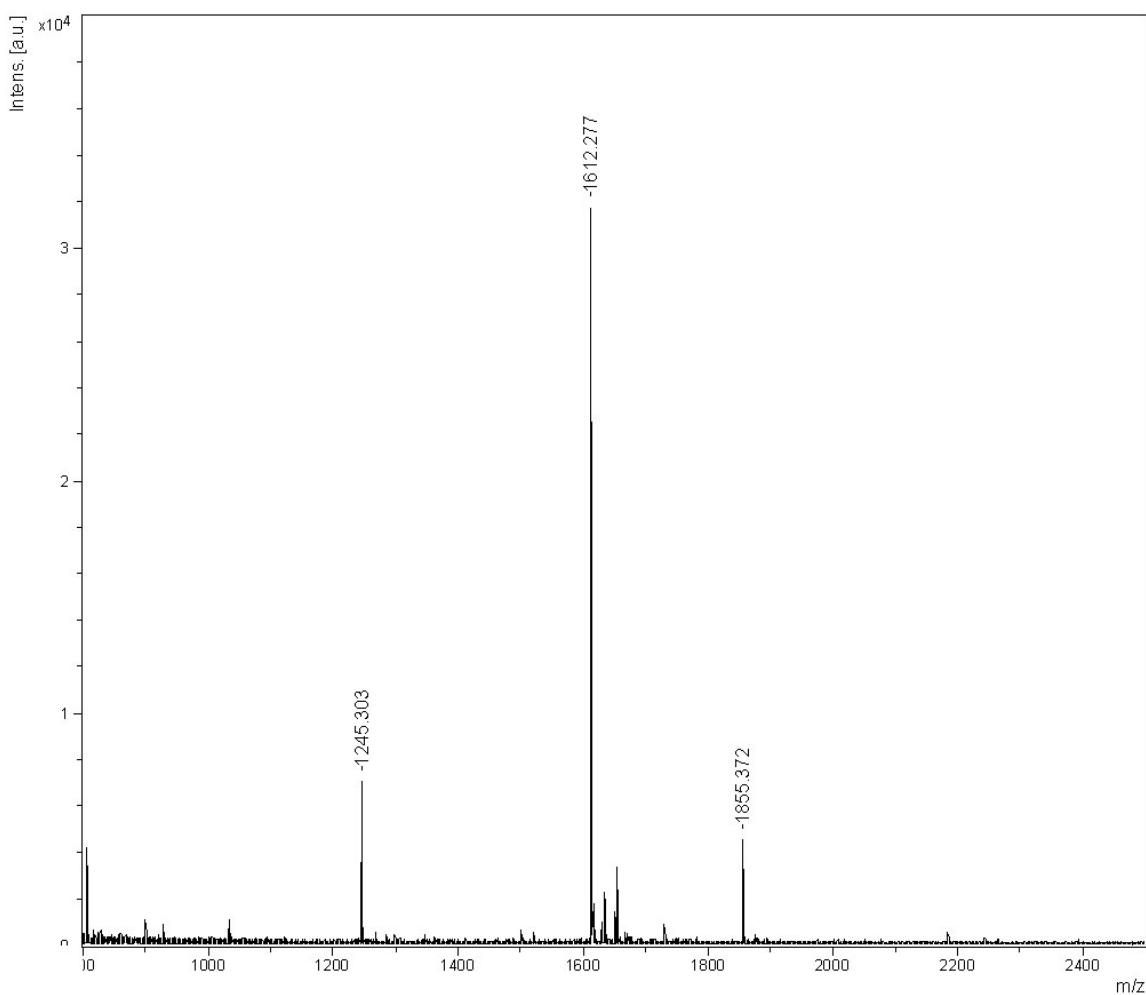


Figure S13(c5): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 30.97$ min in the Fig S12(c2j). Corresponding RP-Hplc fraction is $R_T = 26.50$ min in the Fig S12(c2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CACGAAC)-3', 7b]. The main peak at m/z 1612.3 is for 5'-CACGA_{2'},_{3'}-cMP [see Table S9(C)].

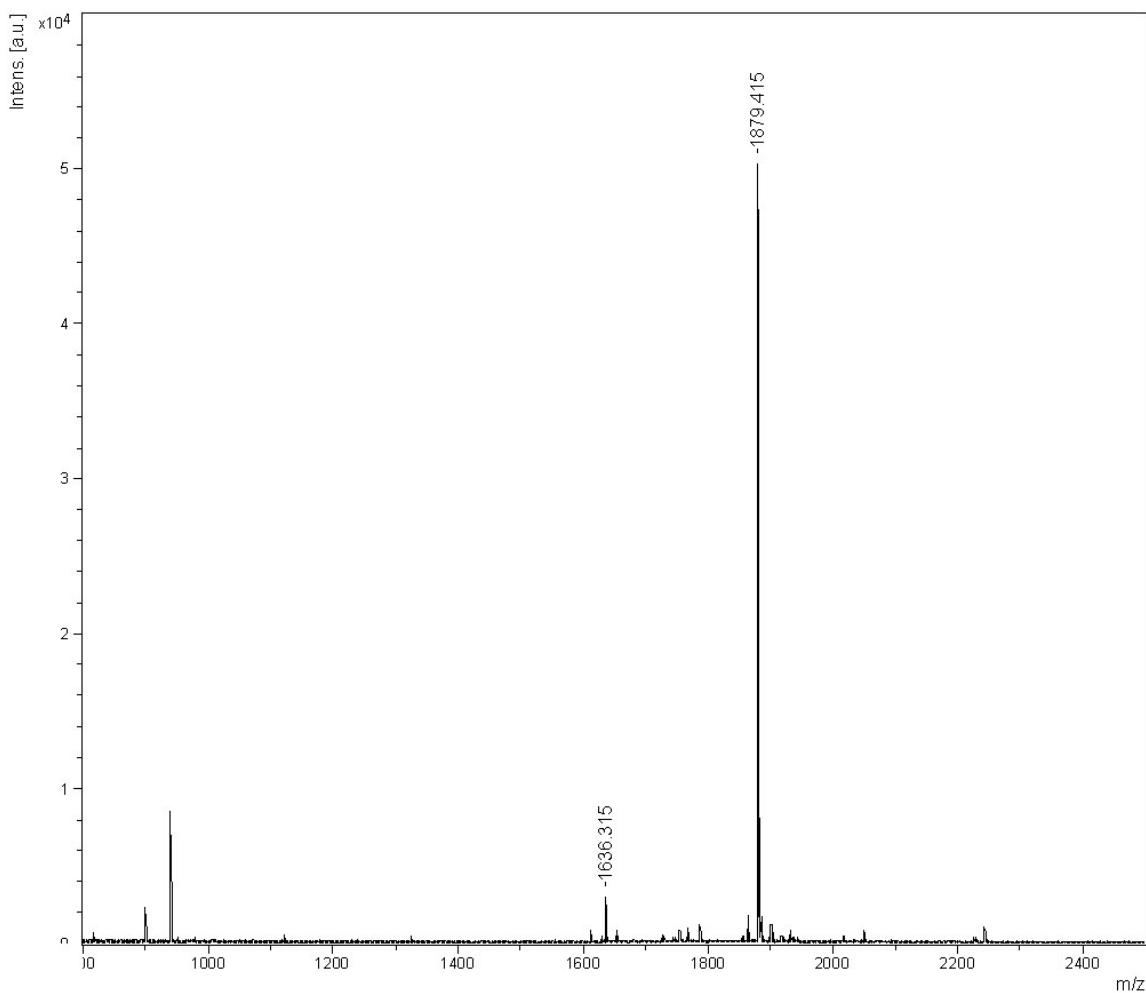


Figure S13(c6): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 31.49$ min in the Fig S12(c6). Corresponding RP-Hplc fraction is $R_T = 26.50$ min in the Fig S12(c2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CACGAAC)-3', 7b] the main peak at m/z 1879.4 is for 5'-ACGAAC-3' [see Table S9(C)].

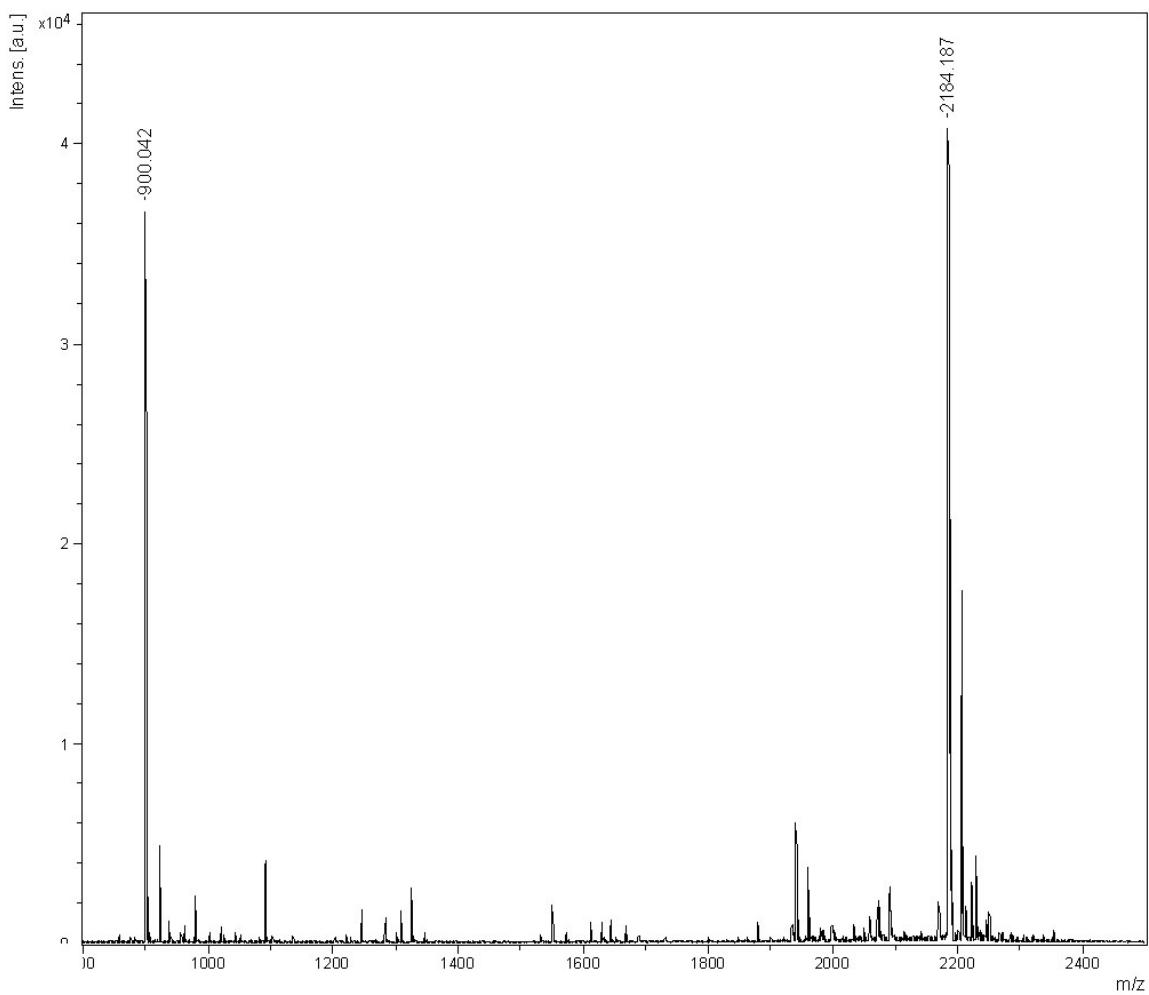


Figure S13(c7): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 27.36$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACGAAC)-3'** (**7b**) after **1h** [for Hplc profile, see Fig **S12 (c2)**] showing the main peak at m/z **900.0** for **5'-AAC-3'** (a fragmentation peak of the heptameric peak in the process of recording mass as confirmed by the variation of the laser power in the Maldi Tof) and **2184.2** for the parent heptamer [see Table S9(C)].

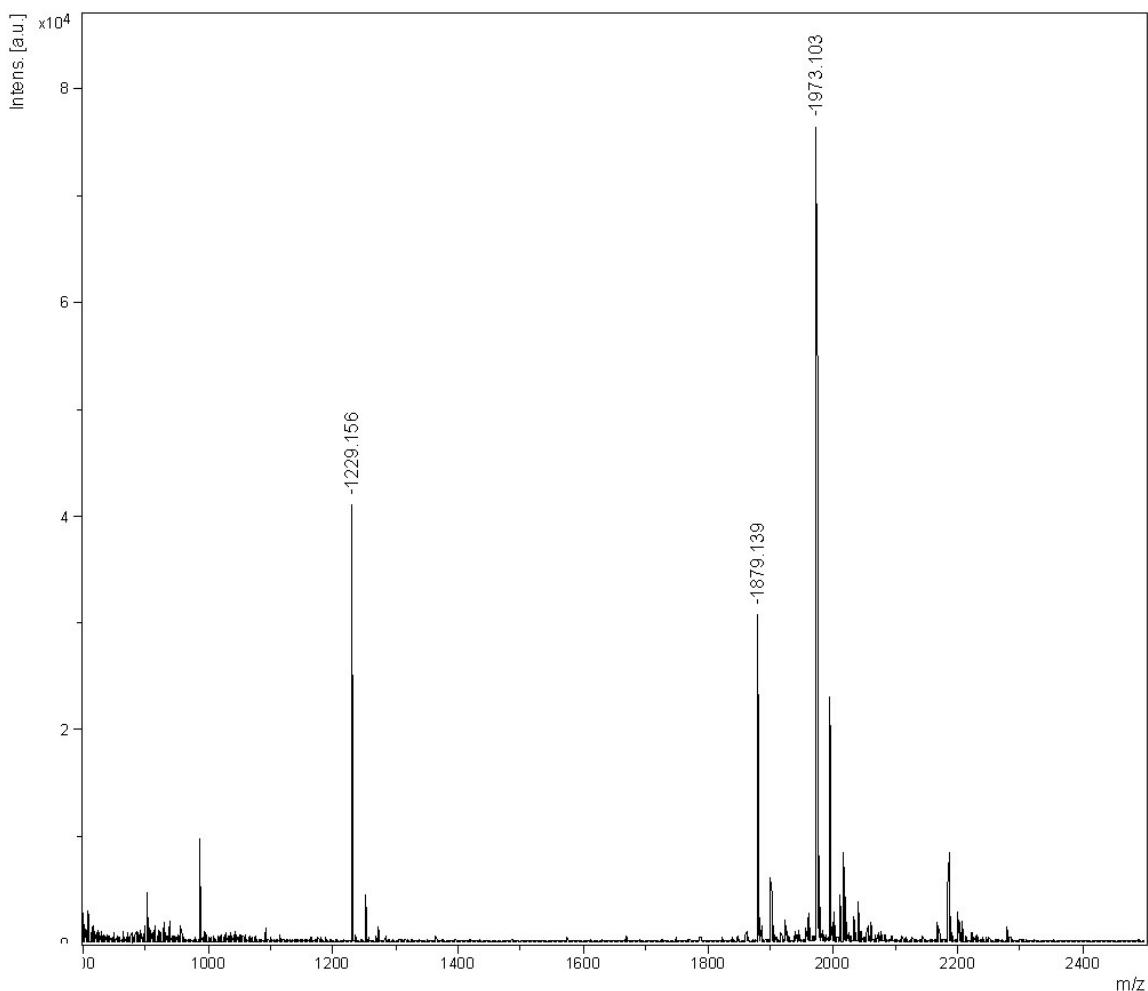


Figure S13(c8): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 28.06$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACGAAC)-3'** (**7b**) after **1h** [for Hplc profile, see Fig **S12 (c2)**] showing the main peak at m/z **1229.2** [non nucleos(t)idic fragment], **1879.1** for **5'-ACGAAC-3'** and **1973.1** [non nucleos(t)idic fragment] [see Table S9(C)].

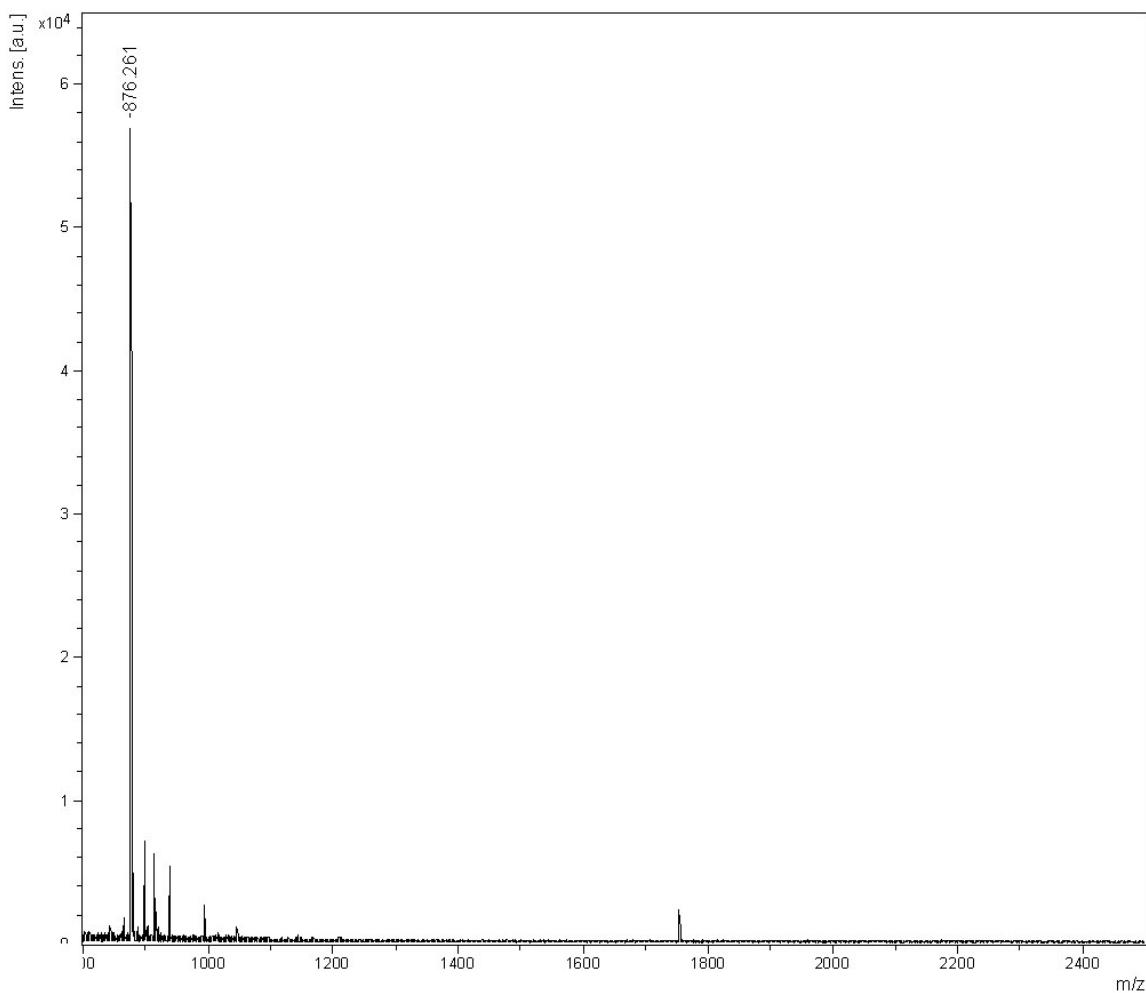


Figure S13(d1): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 18.10$ min in the **Fig S12(d2i)**. Corresponding RP-Hplc fraction is $R_T = 19.99$ min in the **Fig S12(d2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACGCAC)-3'**, **8b**]. The main peak at m/z **876.3** is for **5'-CAC-3'** [see Table S9(D)].

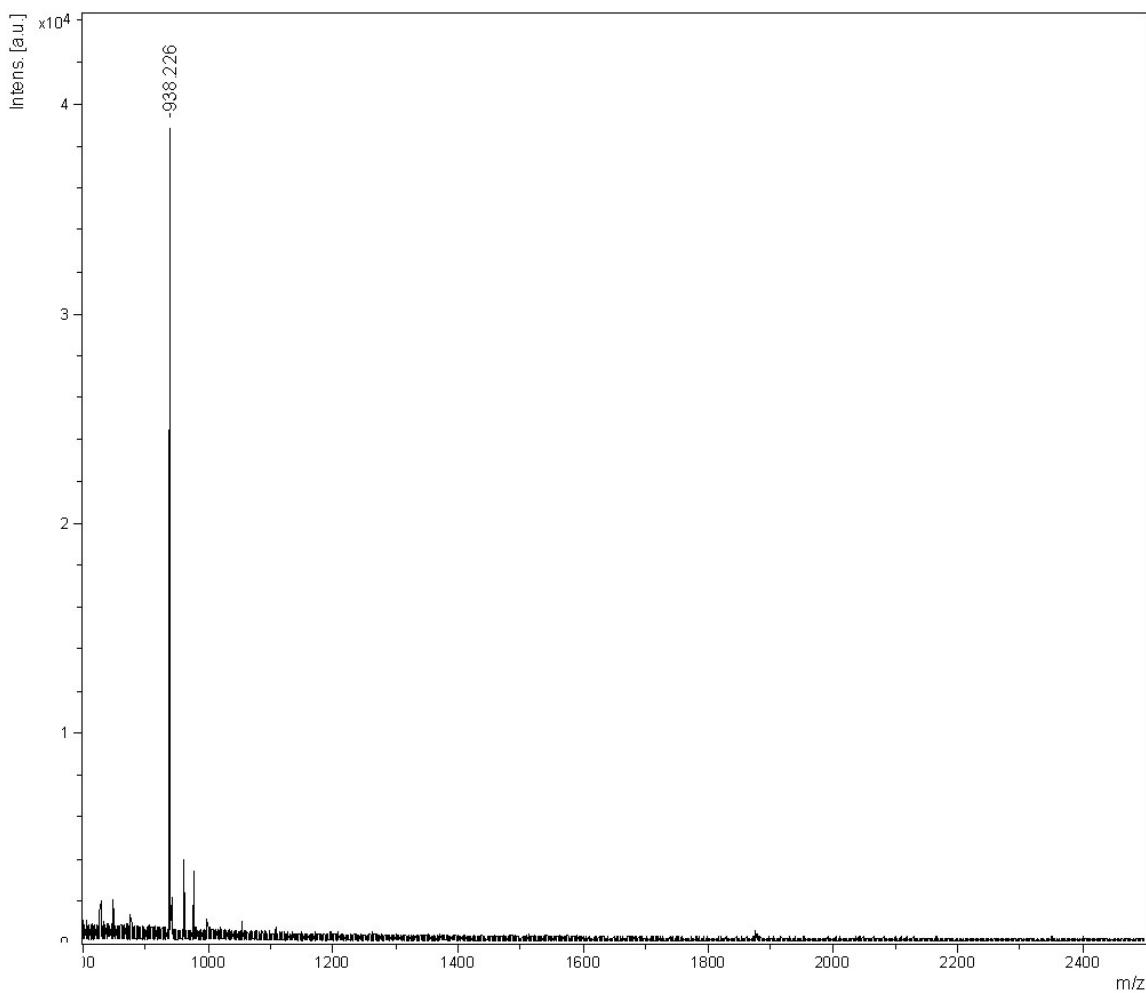


Figure S13(d2): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 18.55$ min in the **Fig S12(d2i)**. Corresponding RP-Hplc fraction is $R_T = 19.99$ min in the **Fig S12(d2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACGCAC)-3', 8b**]. The main peak at m/z **938.2** is for **5'-CAC_{2'}, 3'-cMP** [see Table S9(D)].

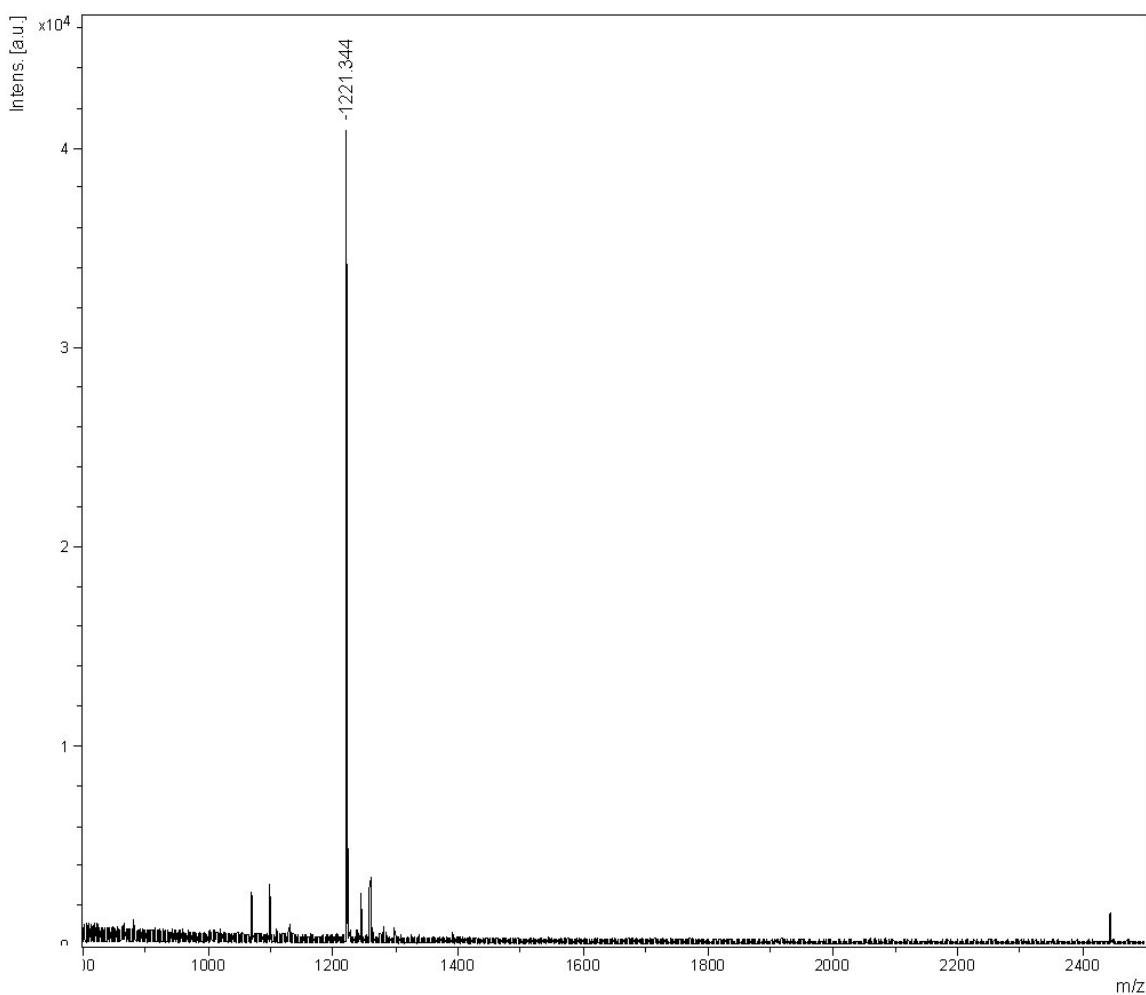


Figure S13(d3): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 20.58$ min in the **Fig S12(d2j)**. Corresponding RP-Hplc fraction is $R_T = 21.23$ min in the **Fig S12(d2)** [after **1 h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACGCAC)-3', 8b**]. The main peak at m/z **1221.3** is for **5'-GCAC-3'** [see Table S9(D)].

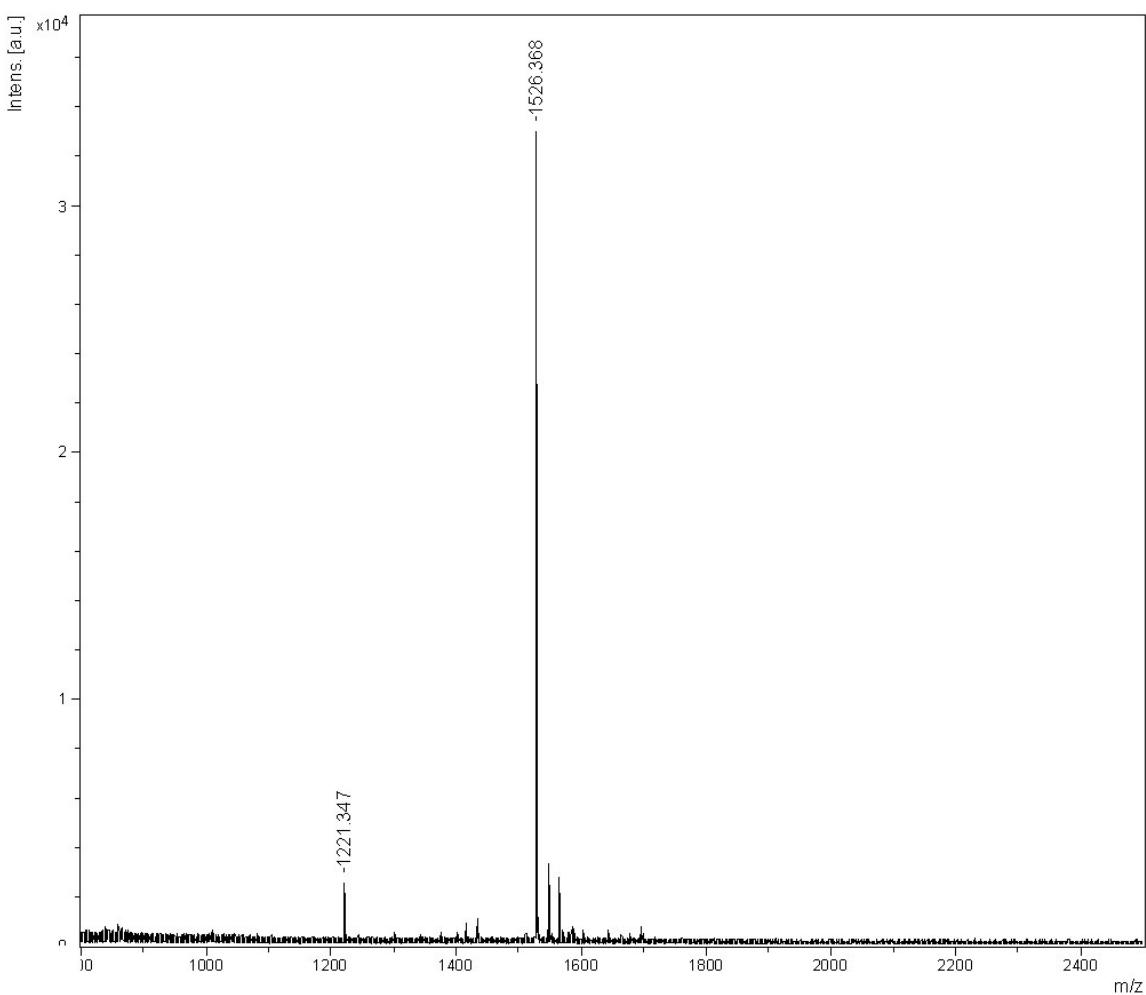


Figure S13(d4): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 21.15$ min in the Fig S12(d2j). Corresponding RP-Hplc fraction is $R_T = 21.23$ min in the Fig S12(d2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CACGCAC)-3', 8b]. The main peak at m/z 1526.4 is for 5'-CGCAC-3' [see Table S9(D)].

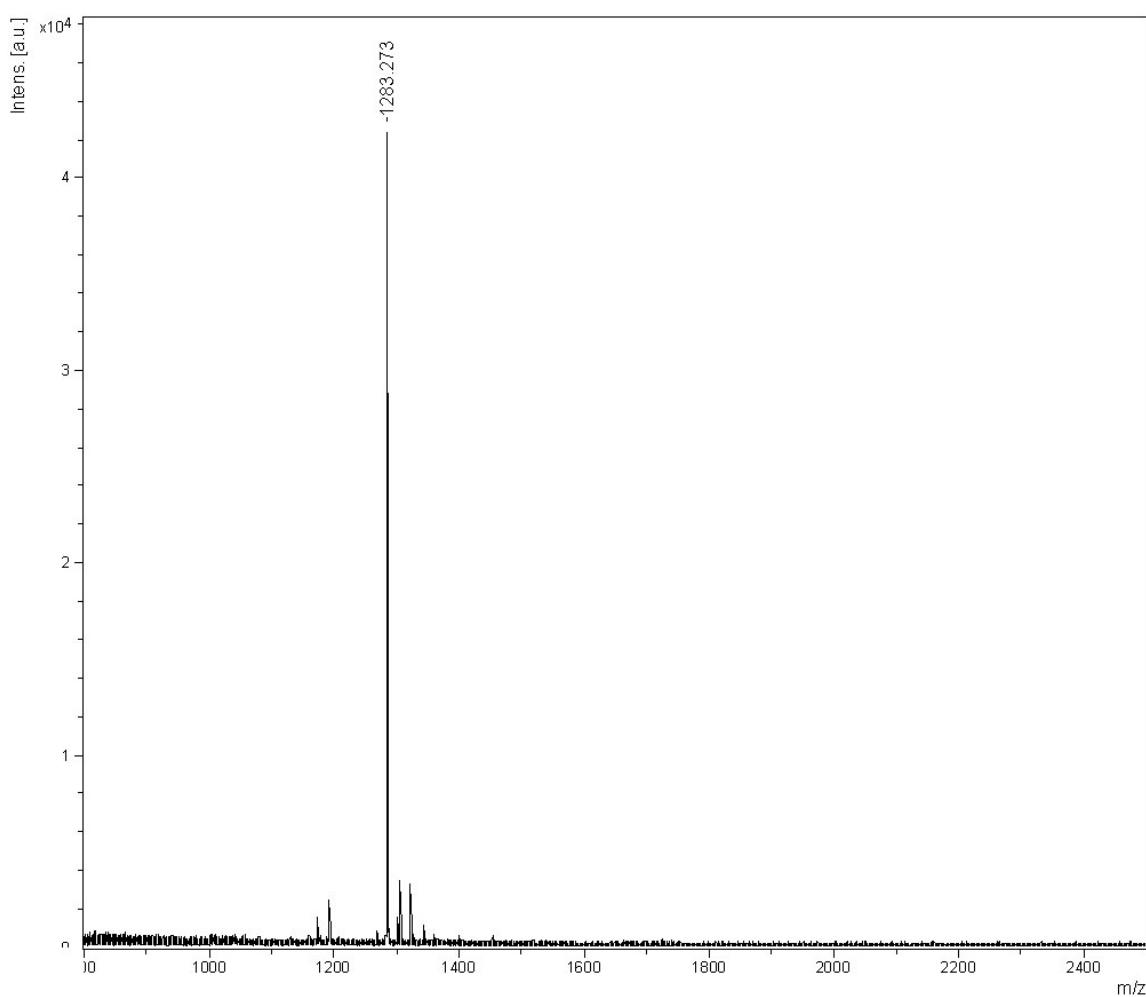


Figure S13(d5): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 21.79$ min in the Fig S12(d2k). Corresponding RP-Hplc fraction is $R_T = 24.14$ min in the Fig S12(d2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CACGCAC)-3', 8b]. The main peak at m/z 1283.3 is for 5'-CACG_{2',3'-cMP} [see Table S9(D)].

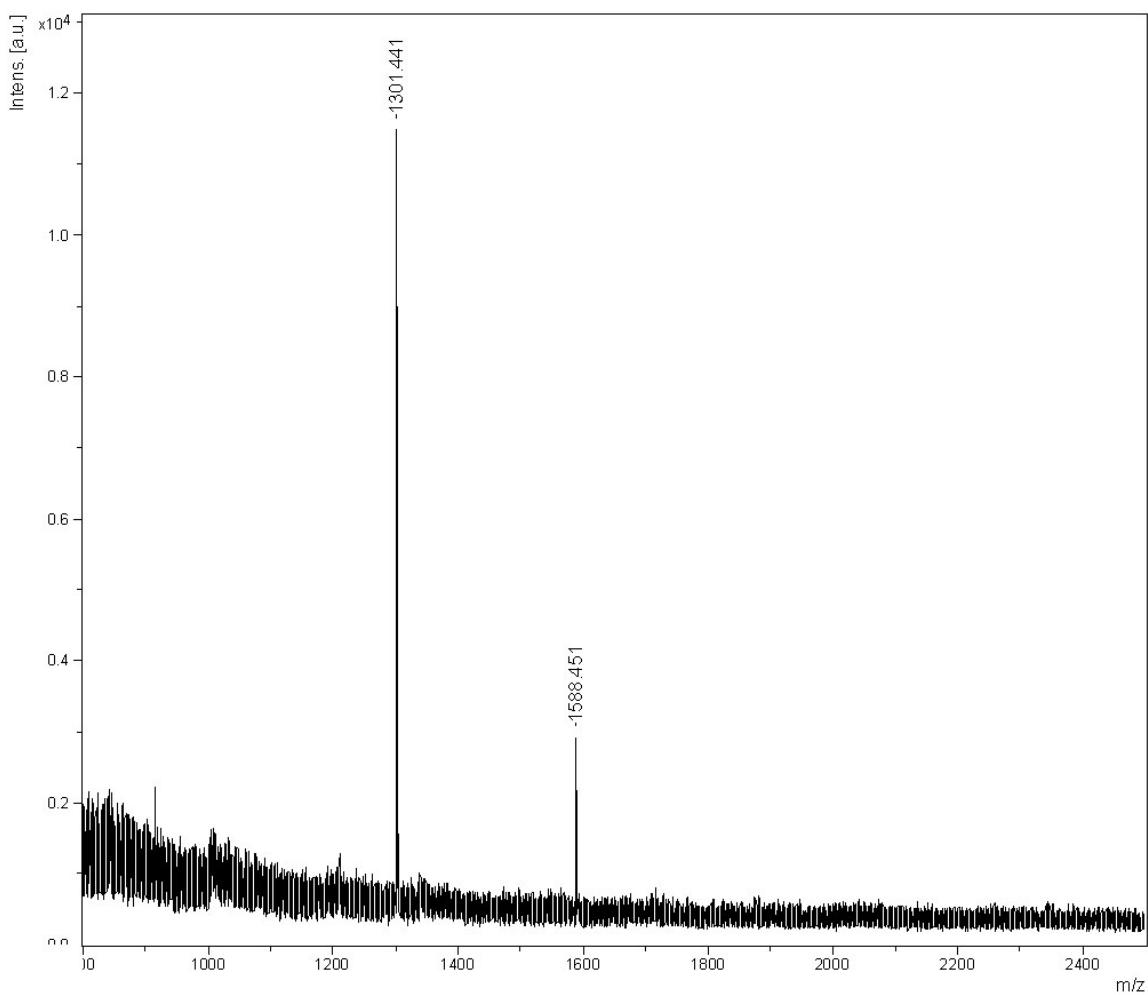


Figure S13(d6): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 22.72$ min in the **Fig S12(d2k)**. Corresponding RP-Hplc fraction is $R_T = 24.14$ min in the **Fig S12(d2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACGCAC)-3', 8b**]. The main peak at m/z **1301.4** for **5'-CACG_{2/3'-P}** and **1588.4** for **5'-CACGC_{2',3'-cMP}** [see Table S9(D)]

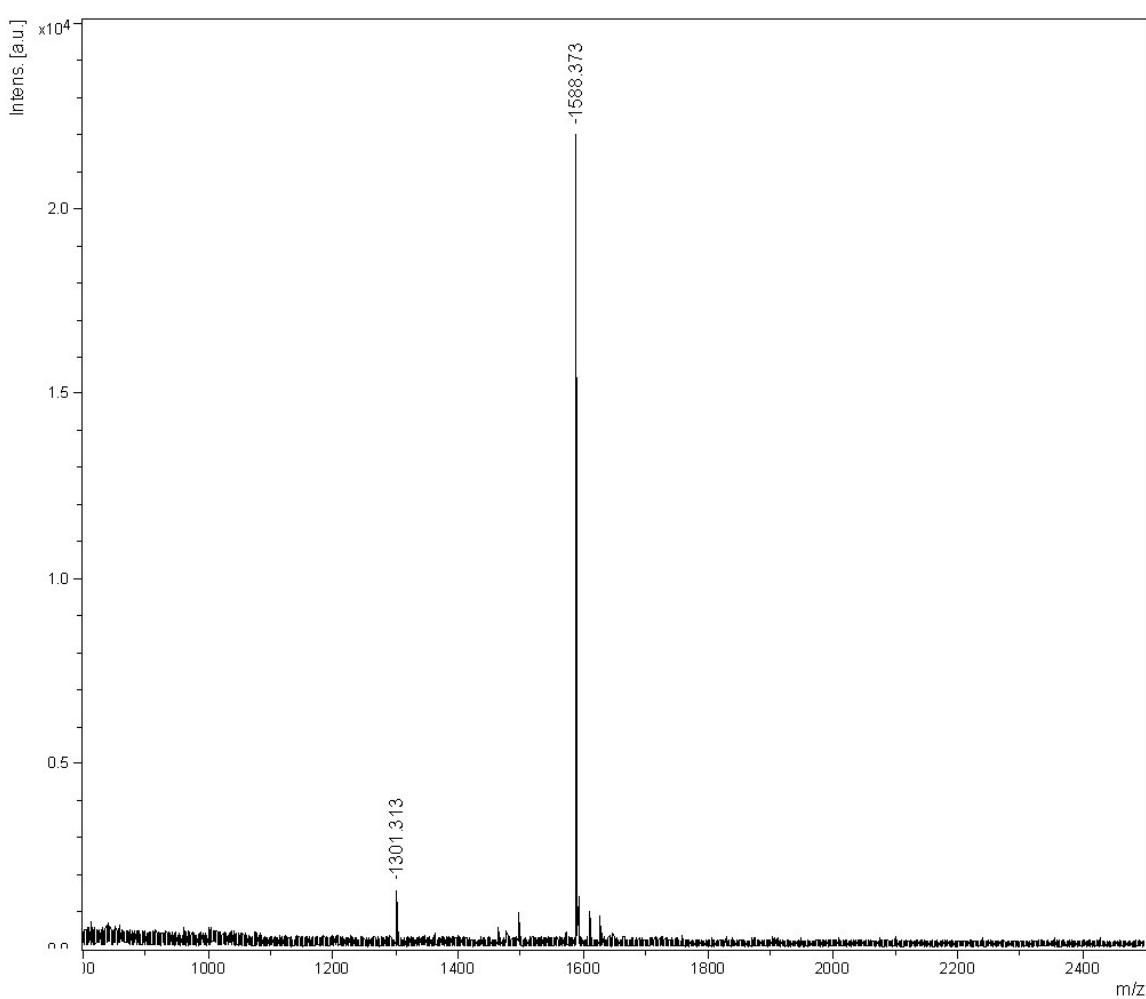


Figure S13(d7): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 23.04$ min in the Fig S12(d2k). Corresponding RP-Hplc fraction is $R_T = 24.14$ min in the Fig S12(d2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CACGCAC)-3', 8b]. The main peak at m/z 1588.4 for 5'-CACGC_{2'}, 3'-cMP [see Table S9(D)]

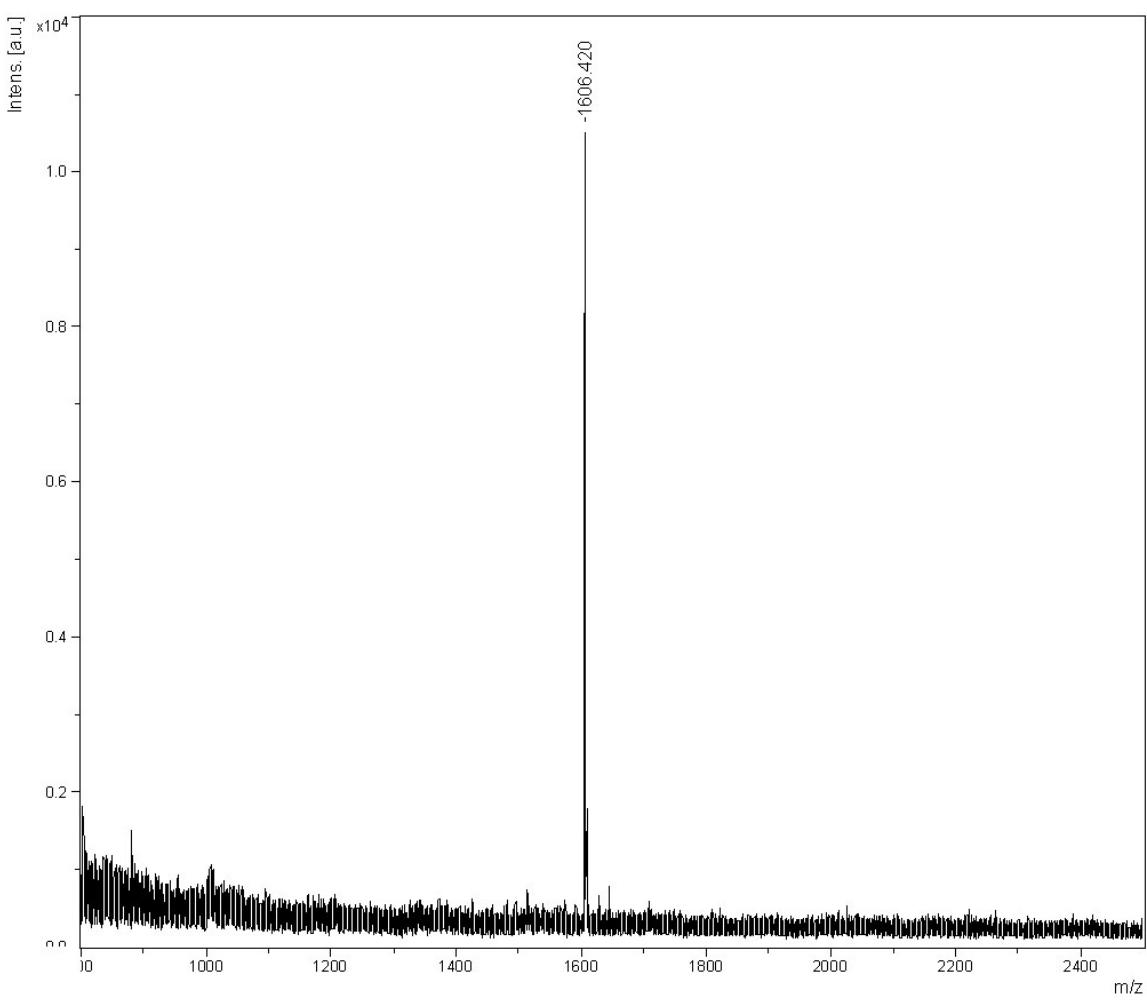


Figure S13(d8): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 24.16$ min in the Fig S12(d2k). Corresponding RP-Hplc fraction is $R_T = 24.14$ min in the Fig S12(d2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CACGCAC)-3', 8b]. The main peak at m/z 1606.4 for 5'-CACGC_{2'3'-P} [see Table S9(D)]

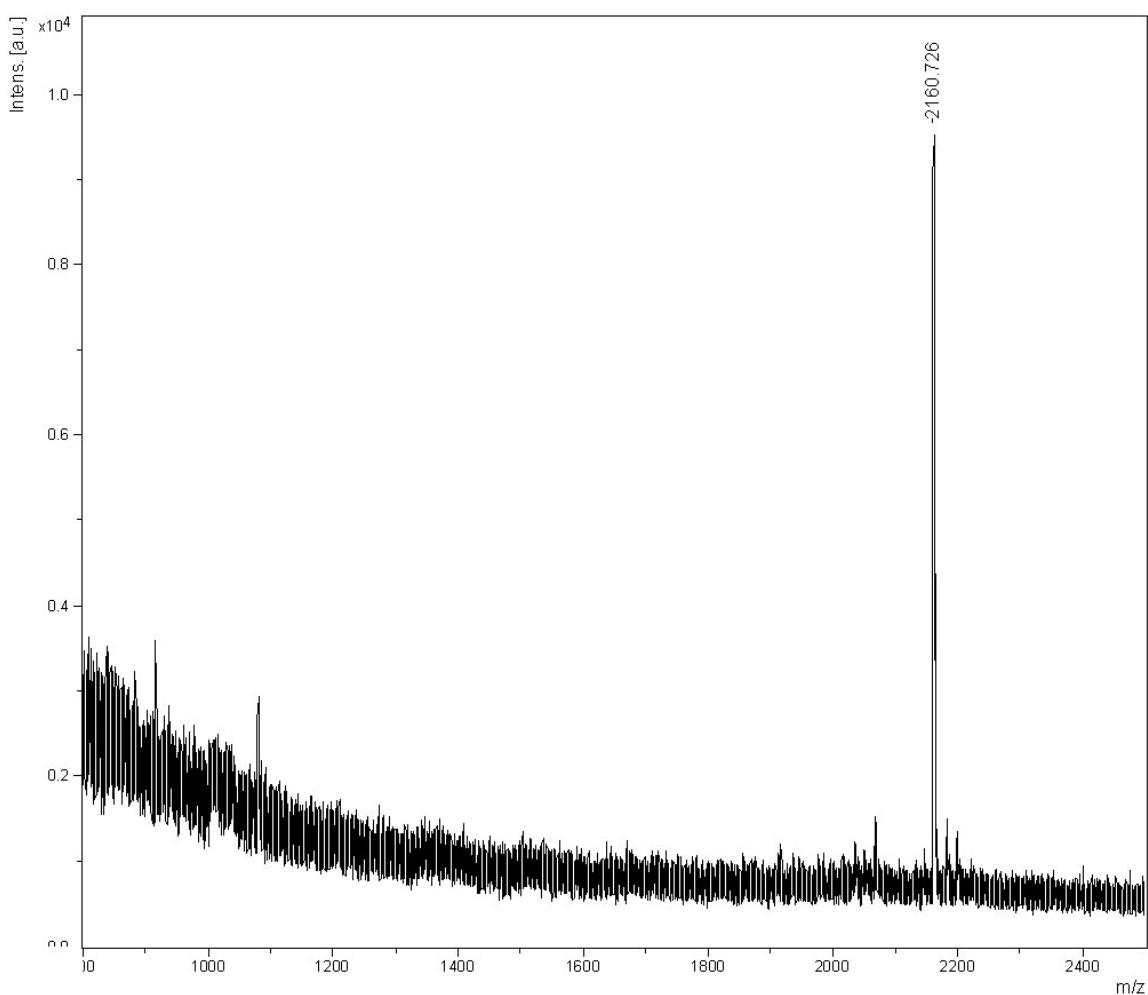


Figure S13(d9): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 26.20$ min in the **Fig S12(d2k)**. Corresponding RP-Hplc fraction is $R_T = 24.14$ min in the **Fig S12(d2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACGCAC)-3', 8b**]. The main peak at m/z **2160.7** is for the parent heptamer [see Table S9(D)].

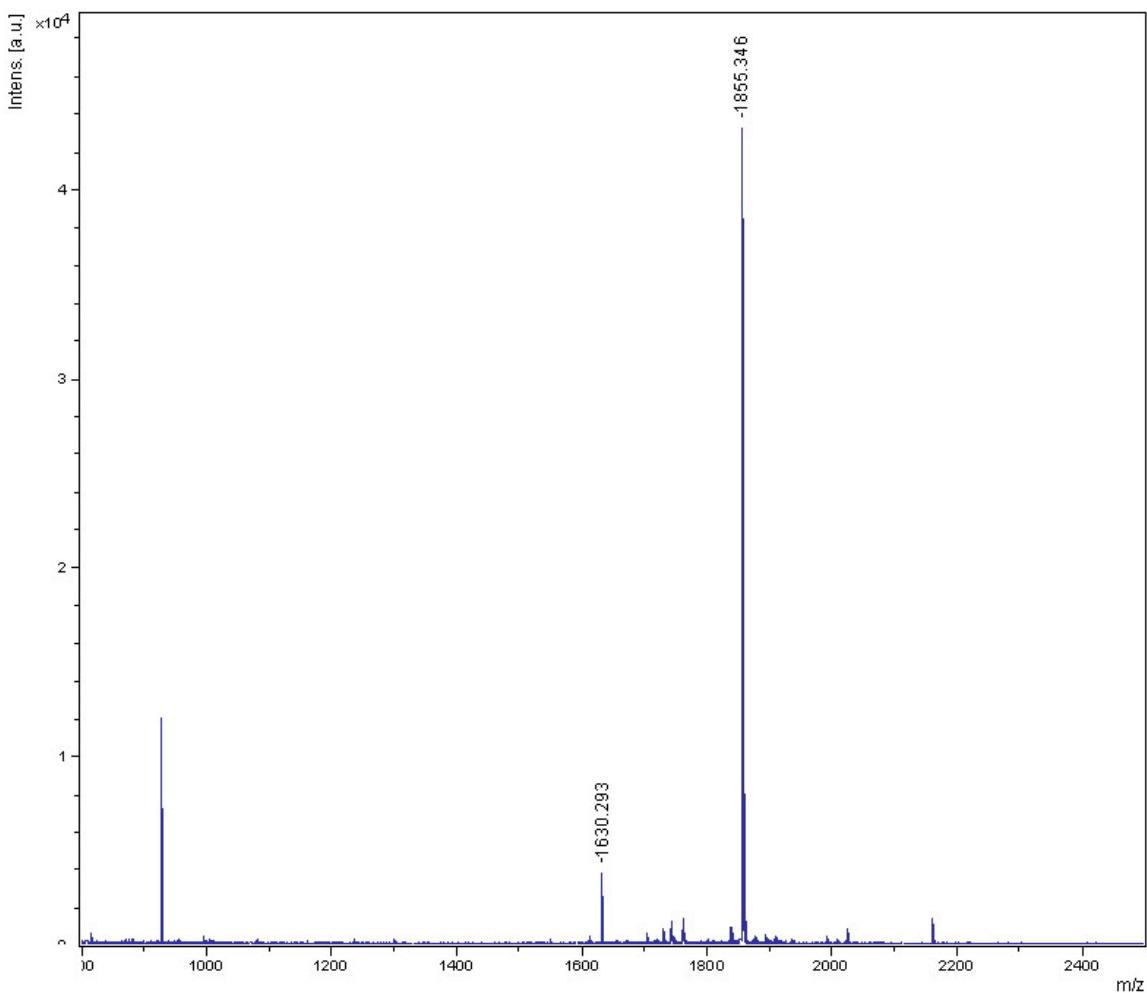


Figure S13(d10): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 14.11$ min in the **Fig S12(d2I)**. Corresponding RP-Hplc fraction is $R_T = 25.39$ min in the **Fig S12(d2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACGCAC)-3'**, **8b**]. The main peak at m/z **1855.4** for **5'-ACGCAC-3'** [see Table S9(D)]

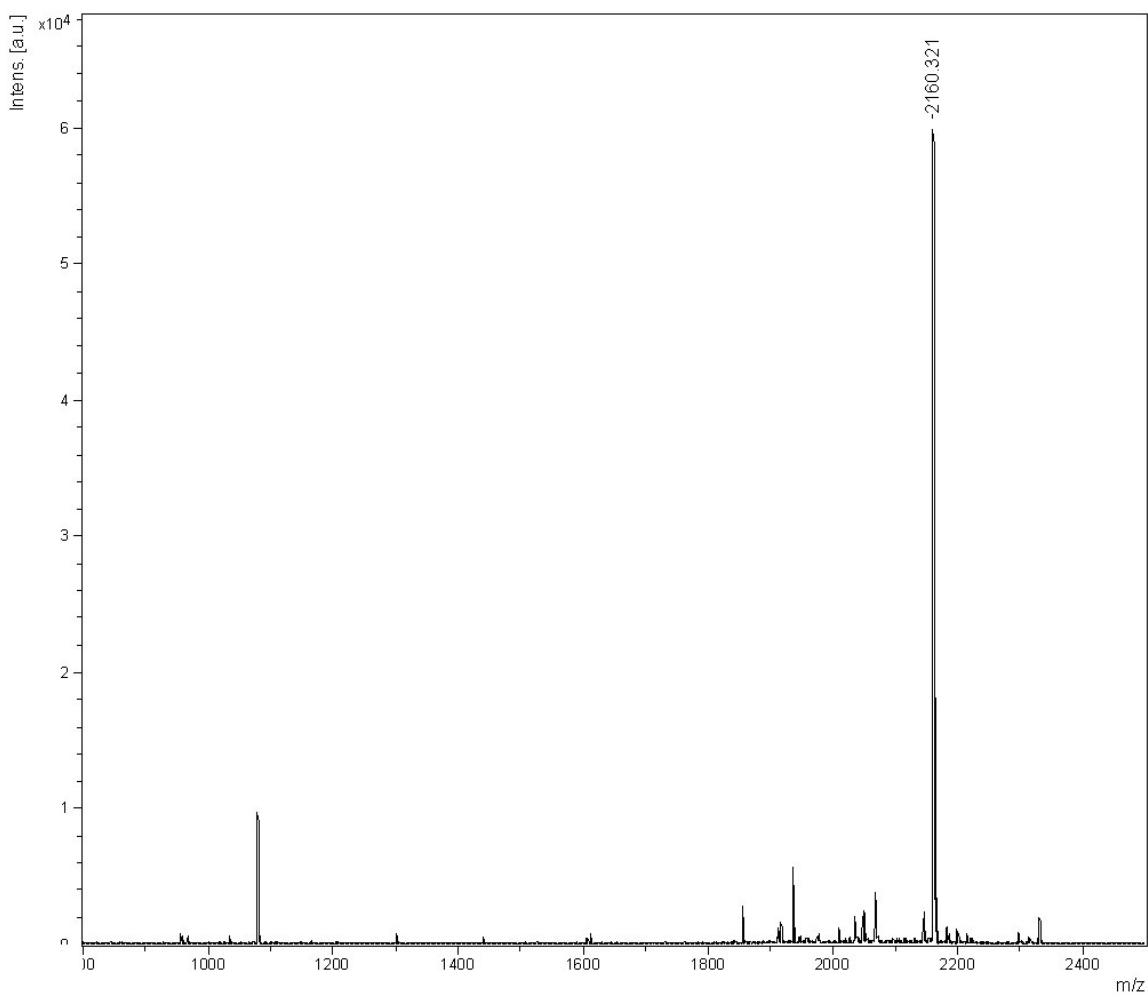


Figure S13(d11): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 14.75$ min in the **Fig S12(d2l)**. Corresponding RP-Hplc fraction is $R_T = 25.39$ min in the **Fig S12(d2)** [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACGCAC)-3', 8b**]. The main peak at m/z 2160.3 is for the parent heptamer [see Table S9(D)]

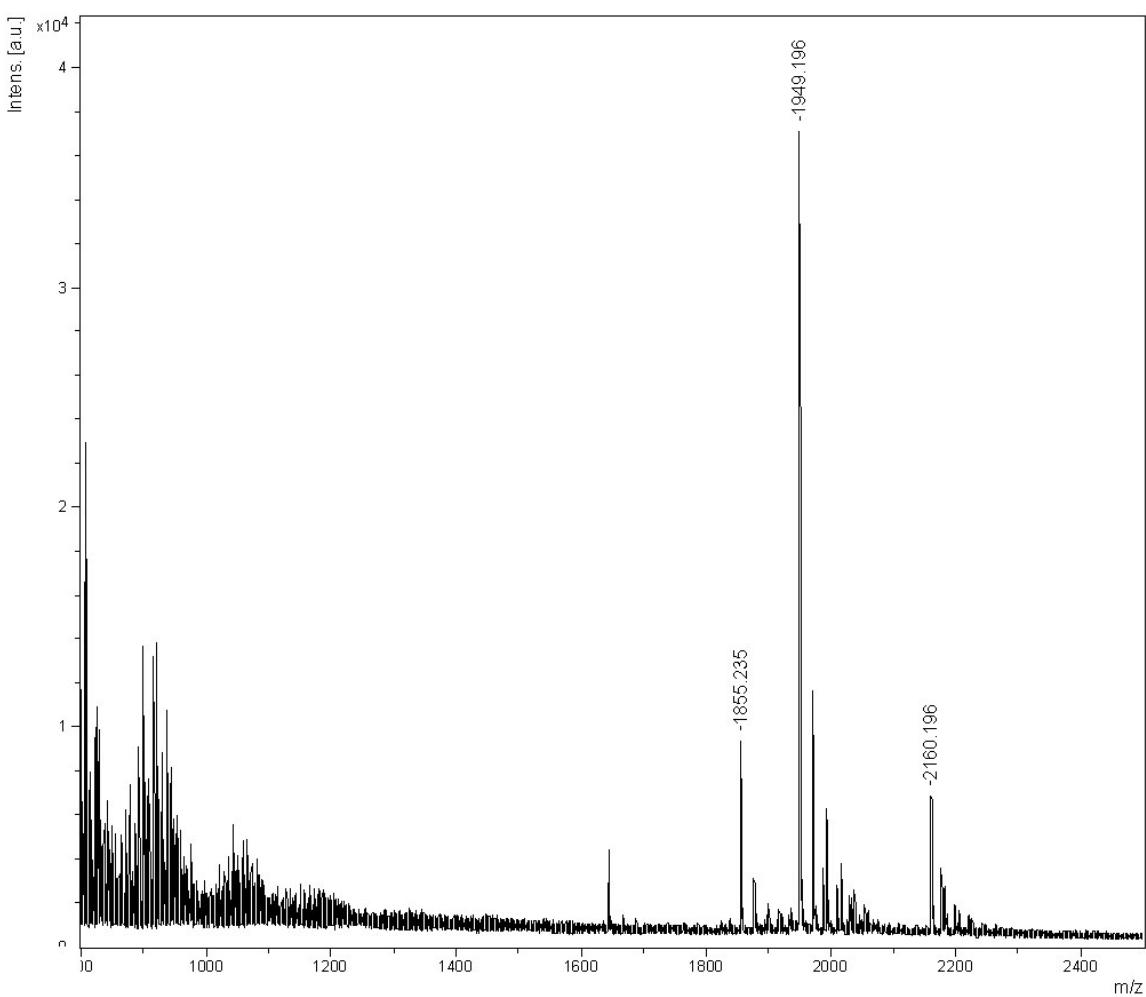


Figure S13(d12): Maldi-Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = \sim 25.70$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-rCACGCAC-3' (8b)** after **1h** [for Hplc profile, see Fig

S12 (d2)] showing the main peak at m/z **1855.2** for **5'-ACGCAC-3'**, **1949.2** for non nucleos(t)idic fragment and **2160.2** for the parent heptamer [see Table S9(D)].

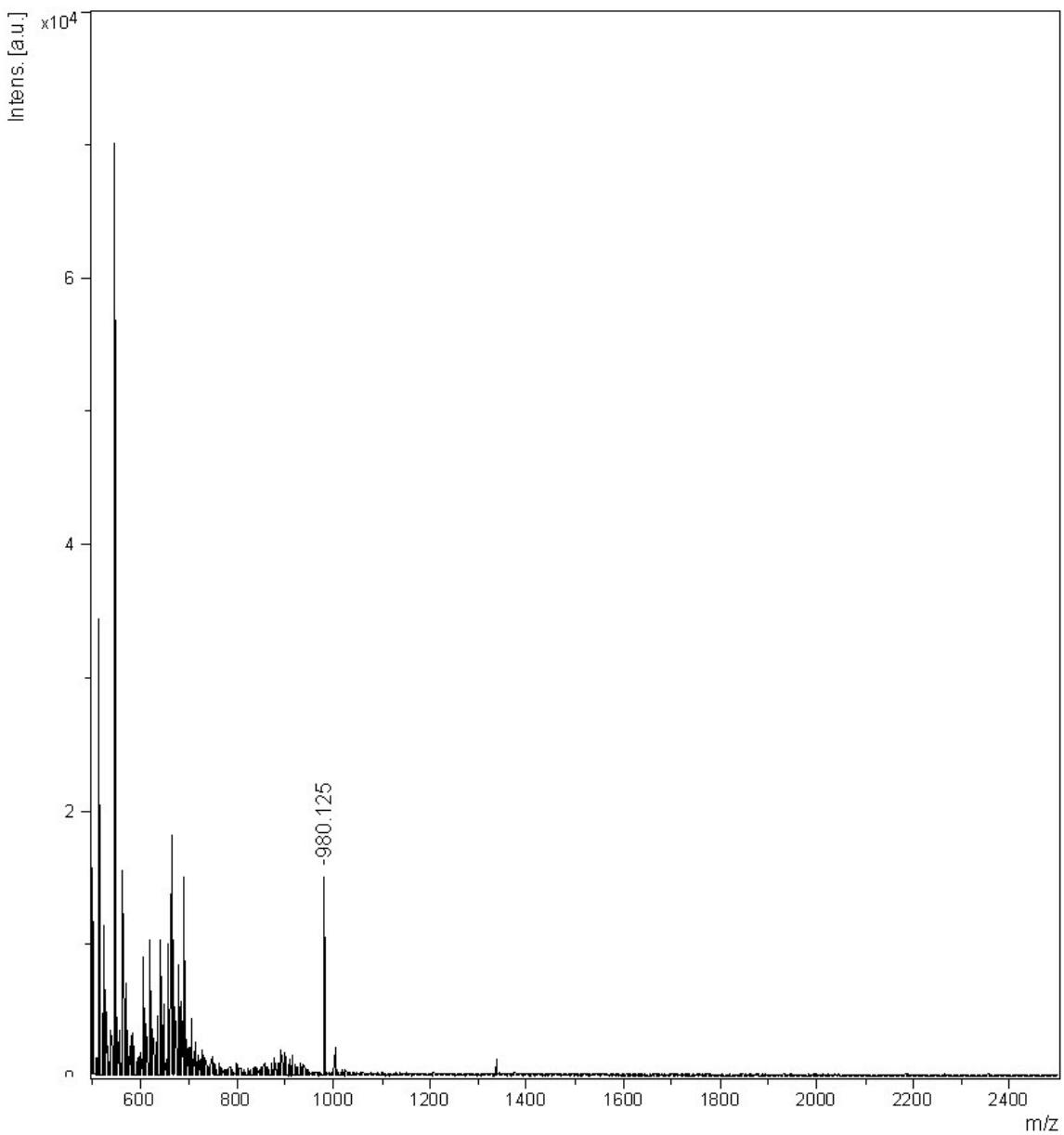


Figure S13(e1): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 20.39$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAG^{Me}AAC)-3'** (**5c**) after **1h** [for Hplc profile, see Fig S12 (e2)] showing the main peak at m/z **980.1** for **5'-CAA_{2'/3'-P}** [see Table S9(E)].

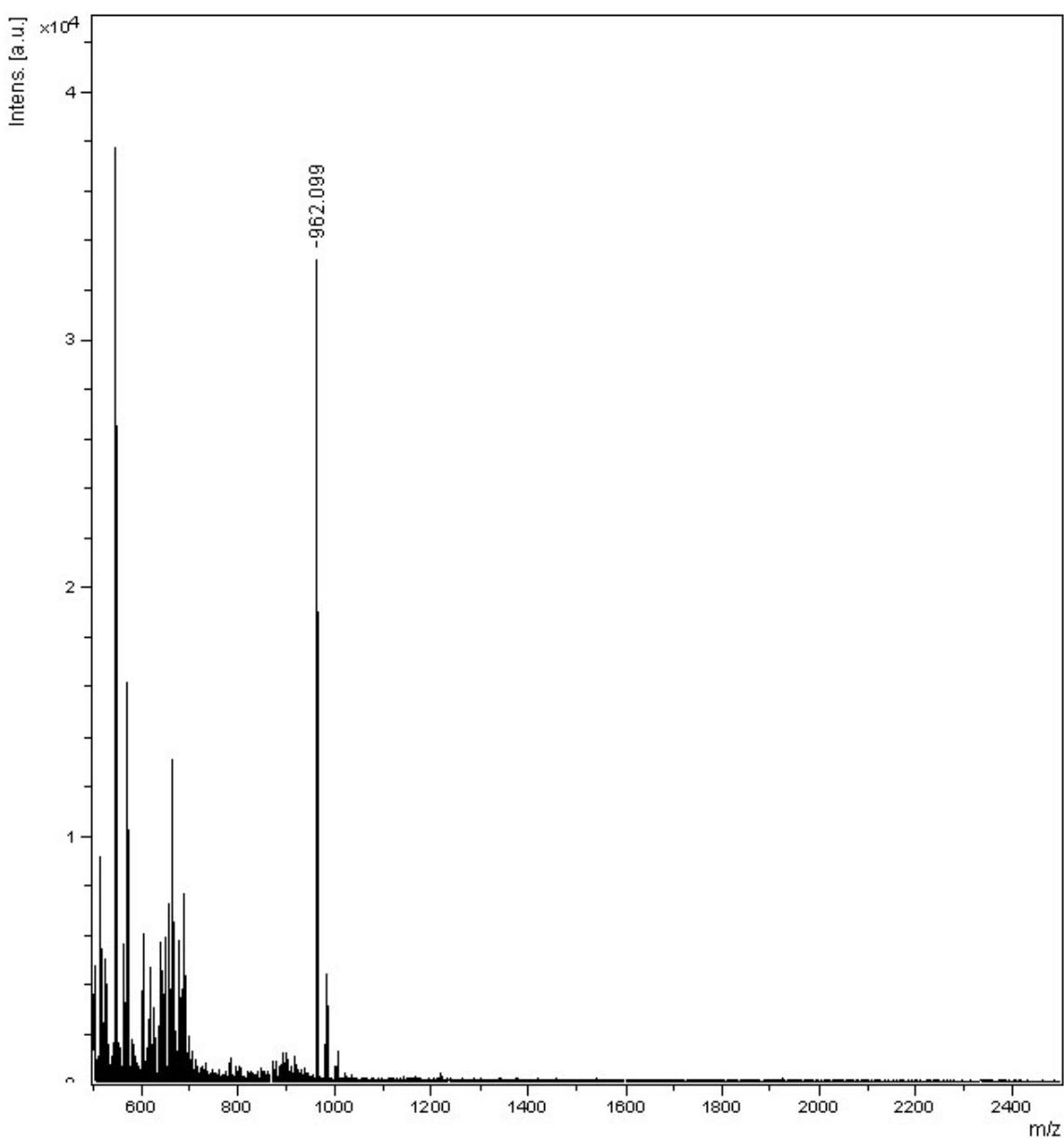


Figure S13(e2): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 21.36$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAG^{Me}AAC)-3'** (**5c**) after **1h** [for Hplc profile, see Fig **S12 (e2)**] showing the main peak at m/z **962.1** for **5'-CAA₂,3'-cMP** [see Table S9(E)].

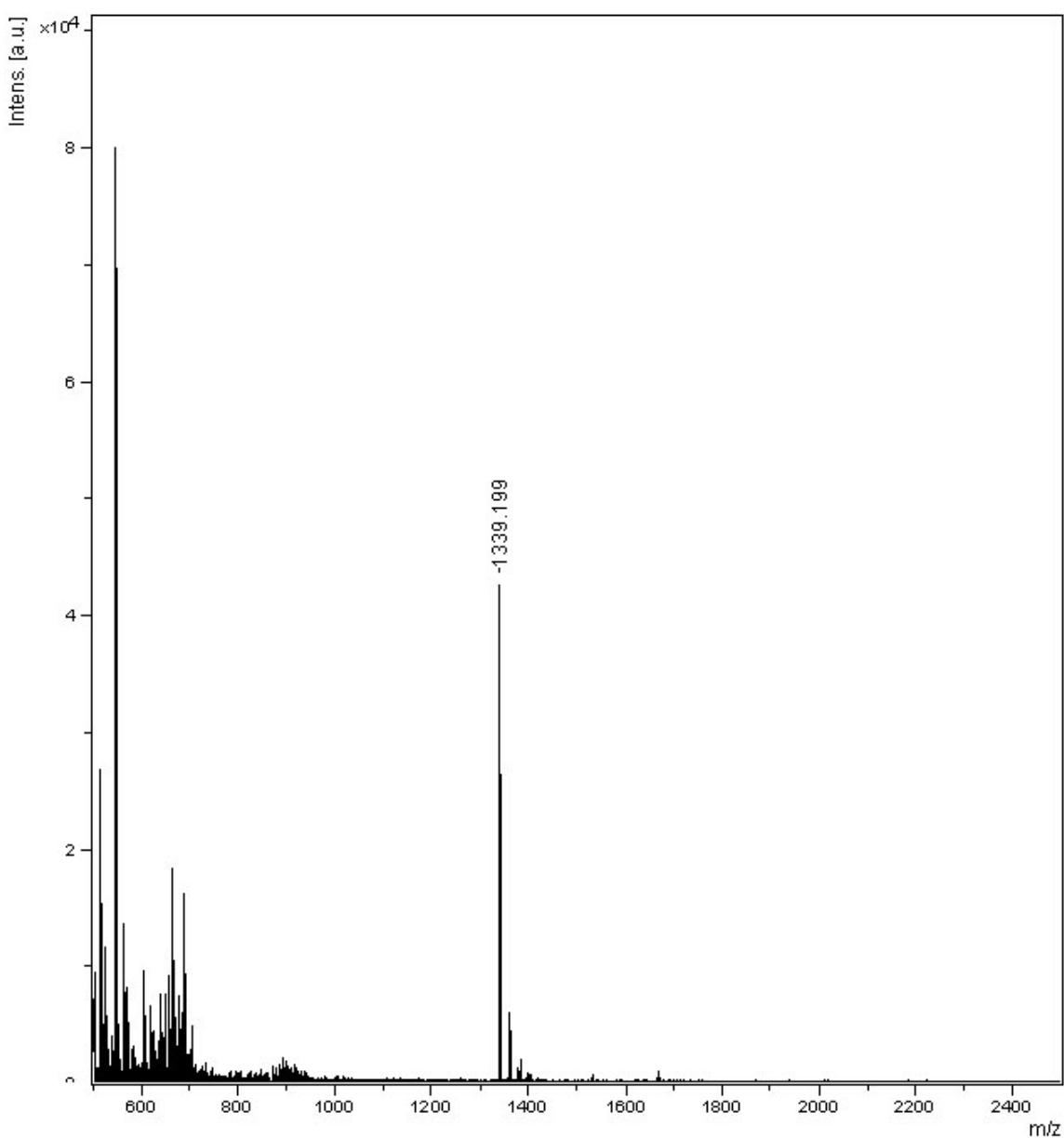


Figure S13(e3): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 22.89$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAG^{Me}AAC)-3' (5c)** after **1h** [for Hplc profile, see Fig S12 (e2)] showing the main peak at m/z **1339.2** for **5'-CAAG^{Me}_{2'/3'}-P** [see Table S9(E)].

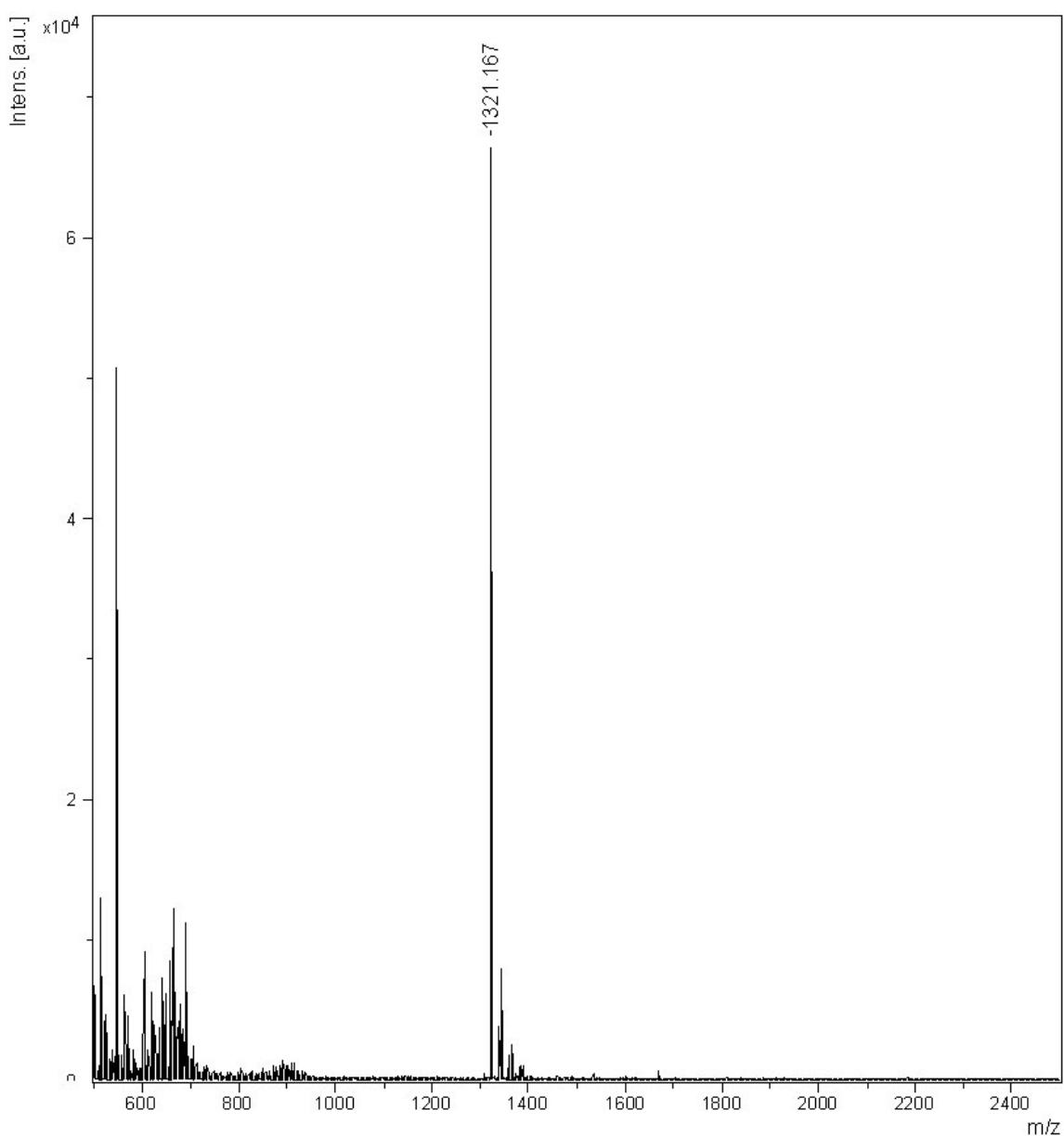


Figure S13(e4): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 23.37$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAG^{Me}AAC)-3' (5c)** after **1h** [for Hplc profile, see Fig **S12 (e2)**] showing the main peak at m/z **1321.2** for **5'-CAAG^{Me}_{2', 3'-cMP}** [see Table S9(E)].

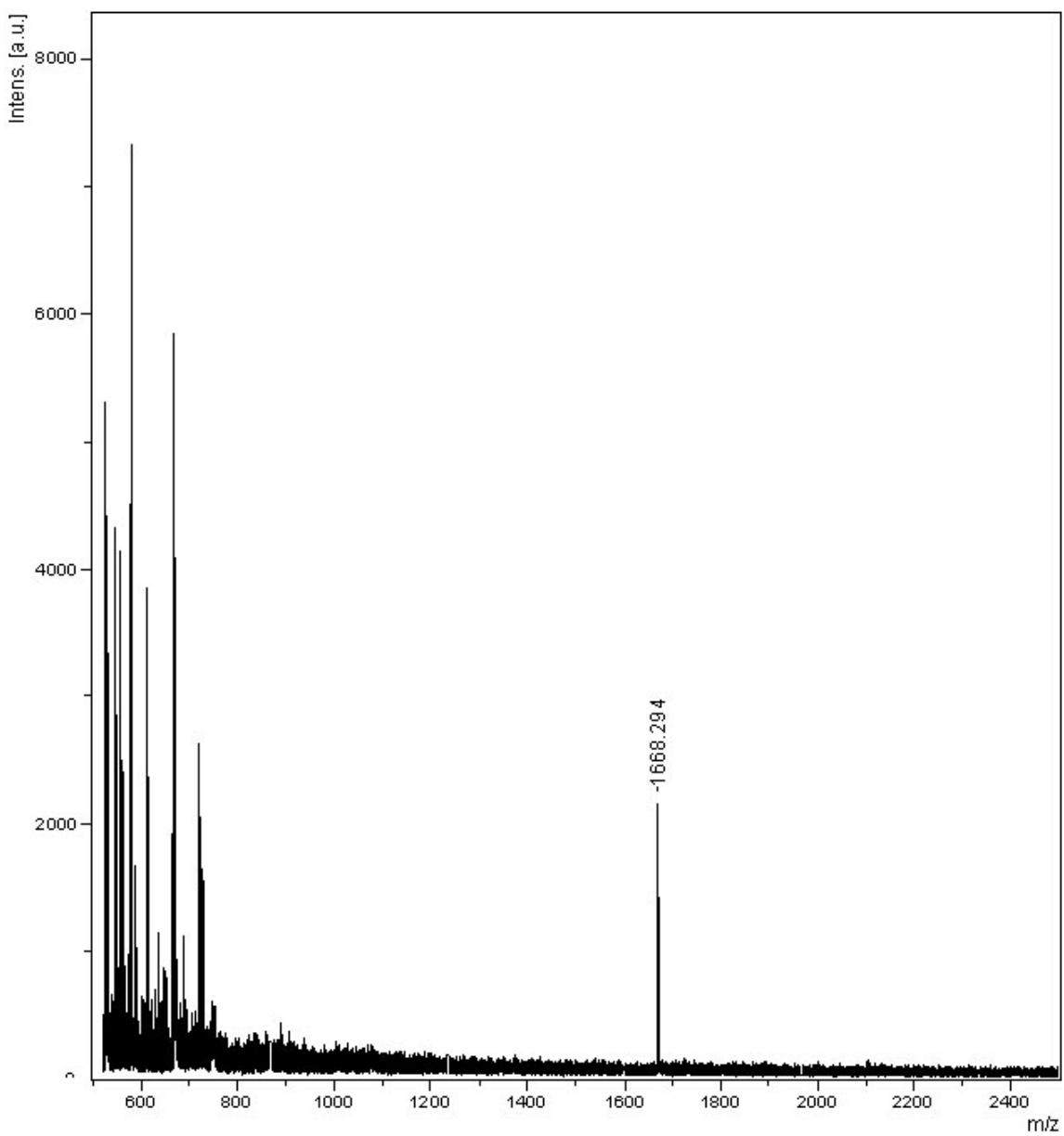


Figure S13(e5): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 29.41$ min in the Fig S12(e2i). Corresponding RP-Hplc fraction is $R_T = 25.50$ min in the Fig S12(e2) [after

1h of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAG^{Me}AAC)-3'**, **5c**] the main peak at m/z **1668.3** is for **5'-CAAG^{Me}A_{2'/3'}-P** [see Table S9(E)].

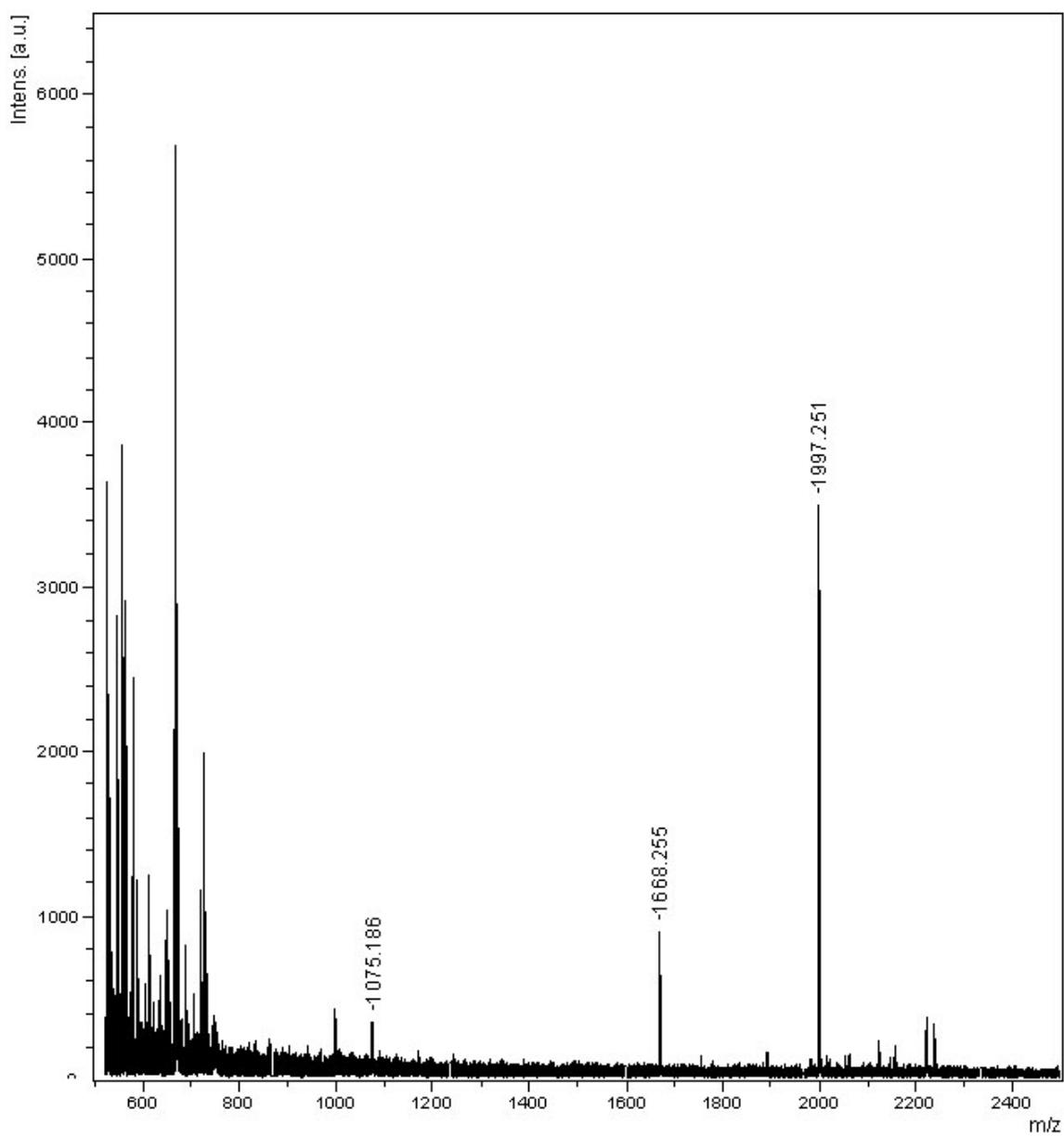


Figure S13(e6): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 30.08$ min in the Fig S12(e2i). Corresponding RP-Hplc fraction is $R_T = 25.50$ min in the Fig S12(e2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of 5'-r(CAAG^{Me}AAC)-3', 5c] the main peak at m/z 1997.3 is for 5'-CAAG^{Me}AA_{2'/3'-P} [see Table S9(E)].

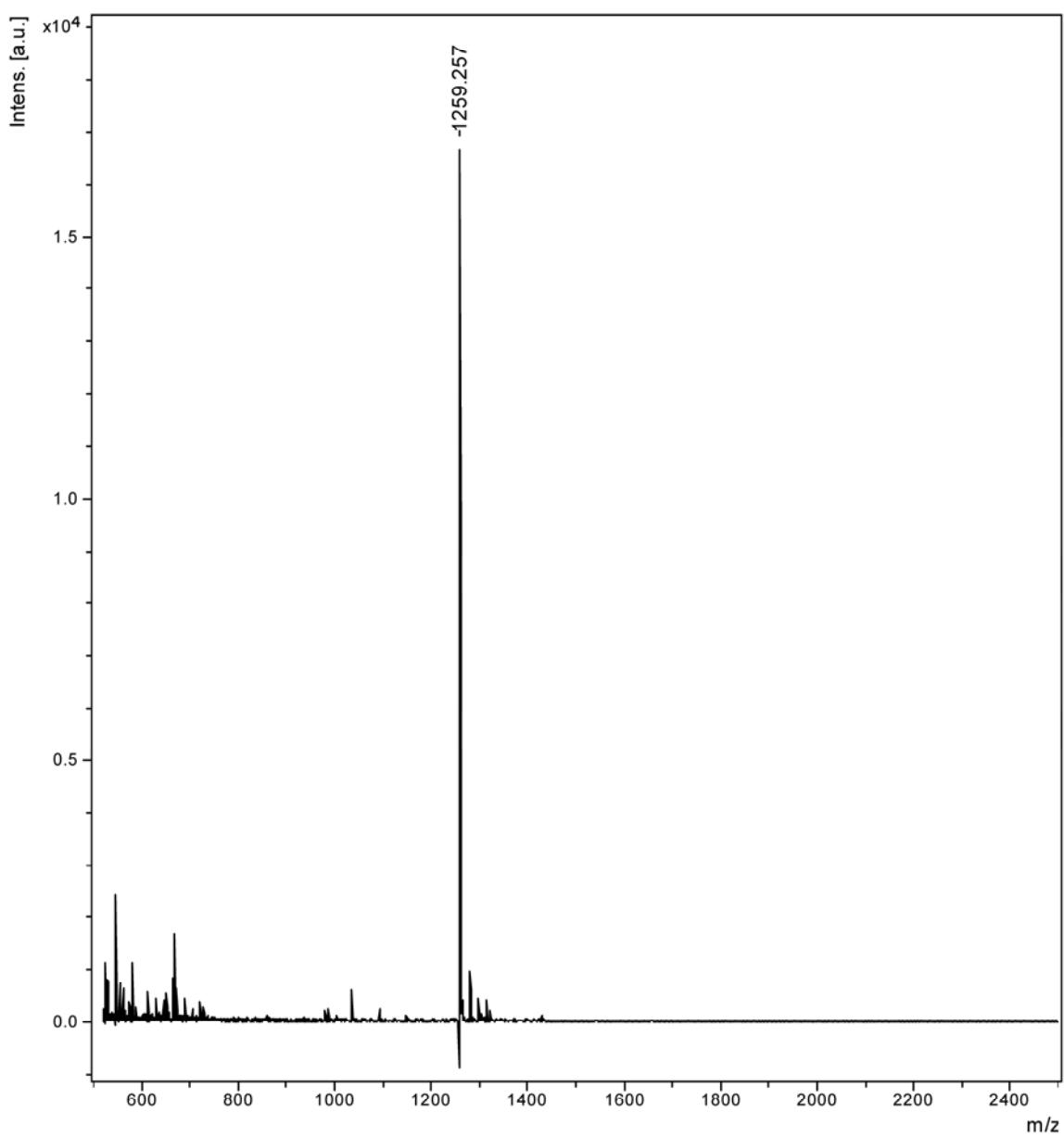


Figure S13(e7): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 27.71$ min in the **Fig S12(e2j)**. Corresponding RP-Hplc fraction is $R_T = 25.70$ min in the **Fig S12(e2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAG^{Me} AAC)-3', 5c**] the main peak at m/z **1259.3** is for **5'-CAAG^{Me} A_{2'}, 3'-cMP** [see Table S9(E)].

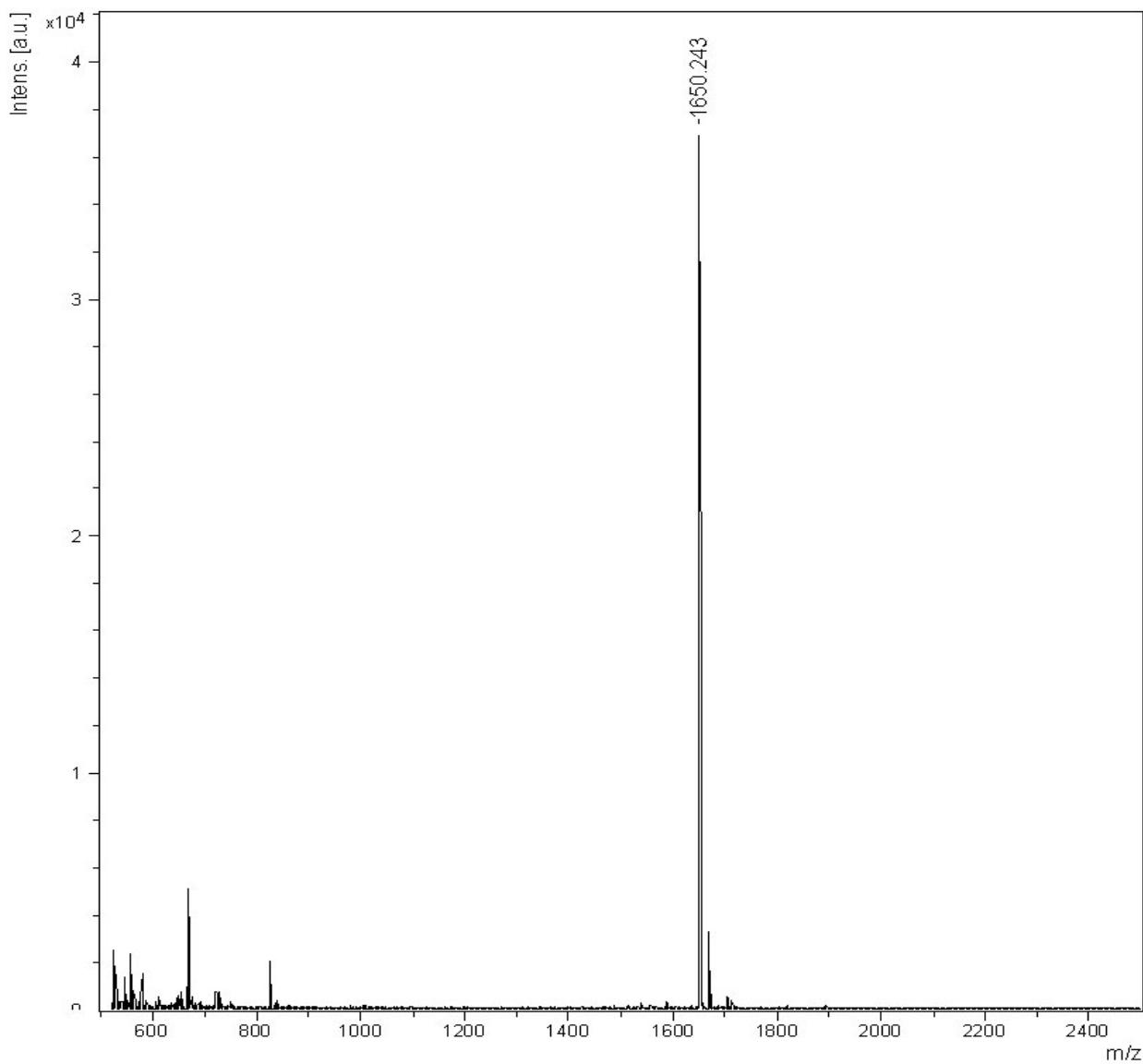


Figure S13(e8): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 29.24$ min in the Fig S12(e2j). Corresponding RP-Hplc fraction is $R_T = 25.70$ min in the Fig S12(e2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAG^{Me}AAC)-3'**, **5c**] the main peak at m/z **1650.2** is for **5'-G^{Me}AAC-3'** [see Table S9(E)].

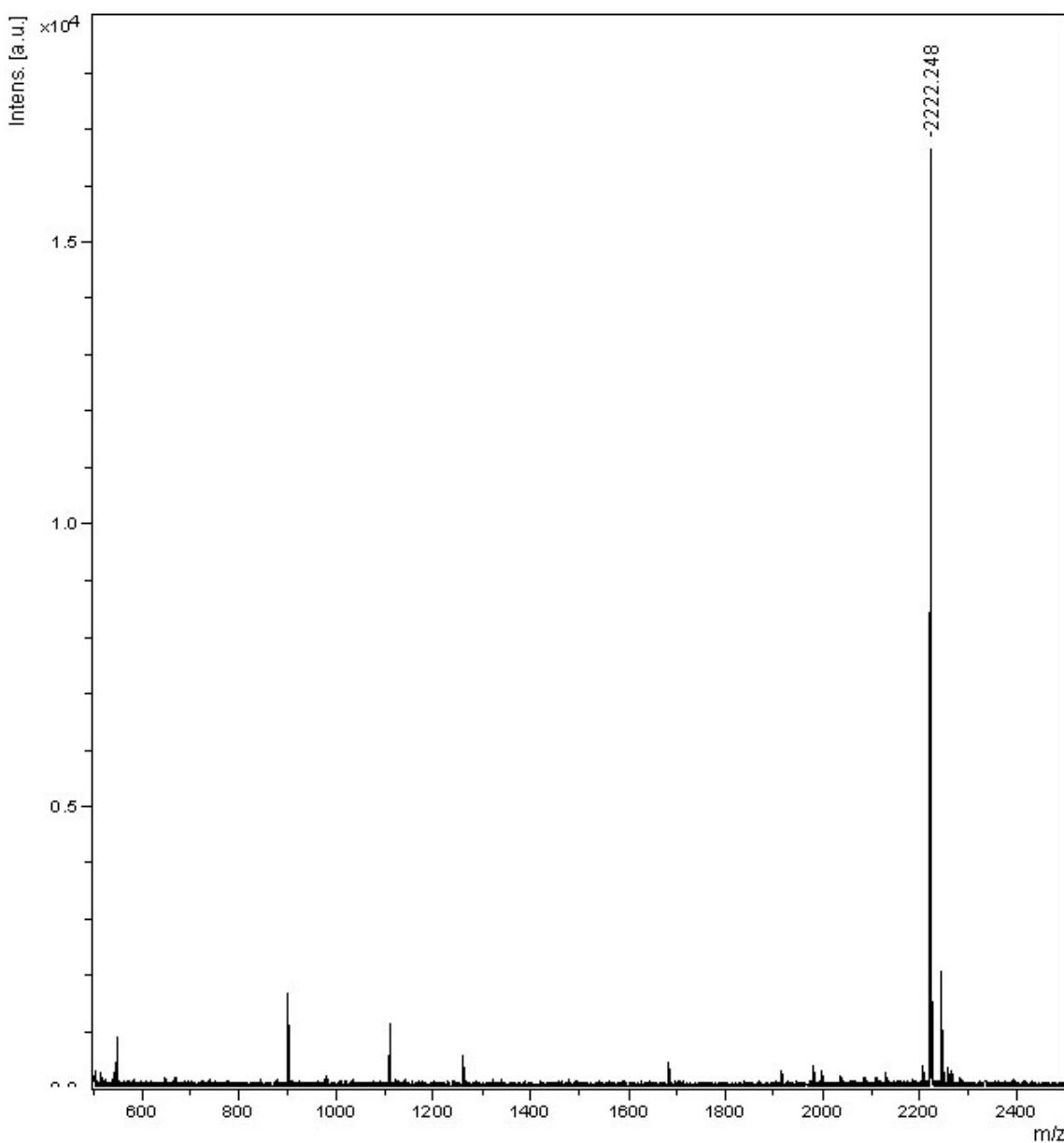


Figure S13(e9): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 26.16$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAG^{Me}AAC)-3'** (**5c**) after **1h** [for Hplc profile, see Fig **S12 (e2)**] showing the main peak at m/z **2222.2** for the parent heptameric peak [see Table S9(E)].

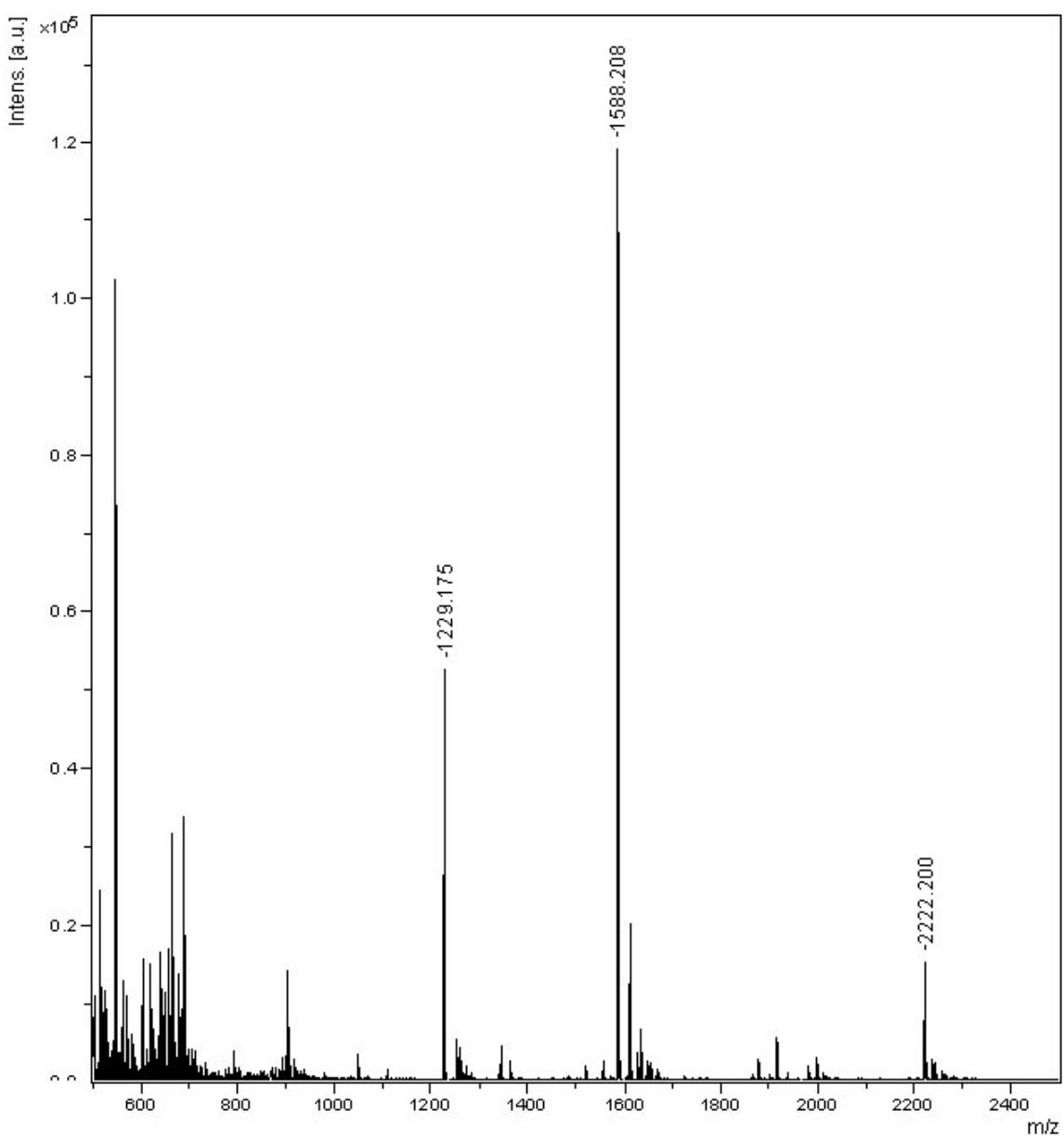


Figure S13(e10): Maldi ToF mass-spectral analysis of the RP-Hplc fraction at $R_T = 26.91$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAG^{Me}AAC)-3' (5c)** after **1h** [for Hplc profile, see Fig **S12 (e2)**] showing the main peak at m/z **1588.2** for **5'-AG^{Me}AAC-3'** along with some non-nucleos(t)idic peak at m/z **1229.2** [see Table S9(E)].

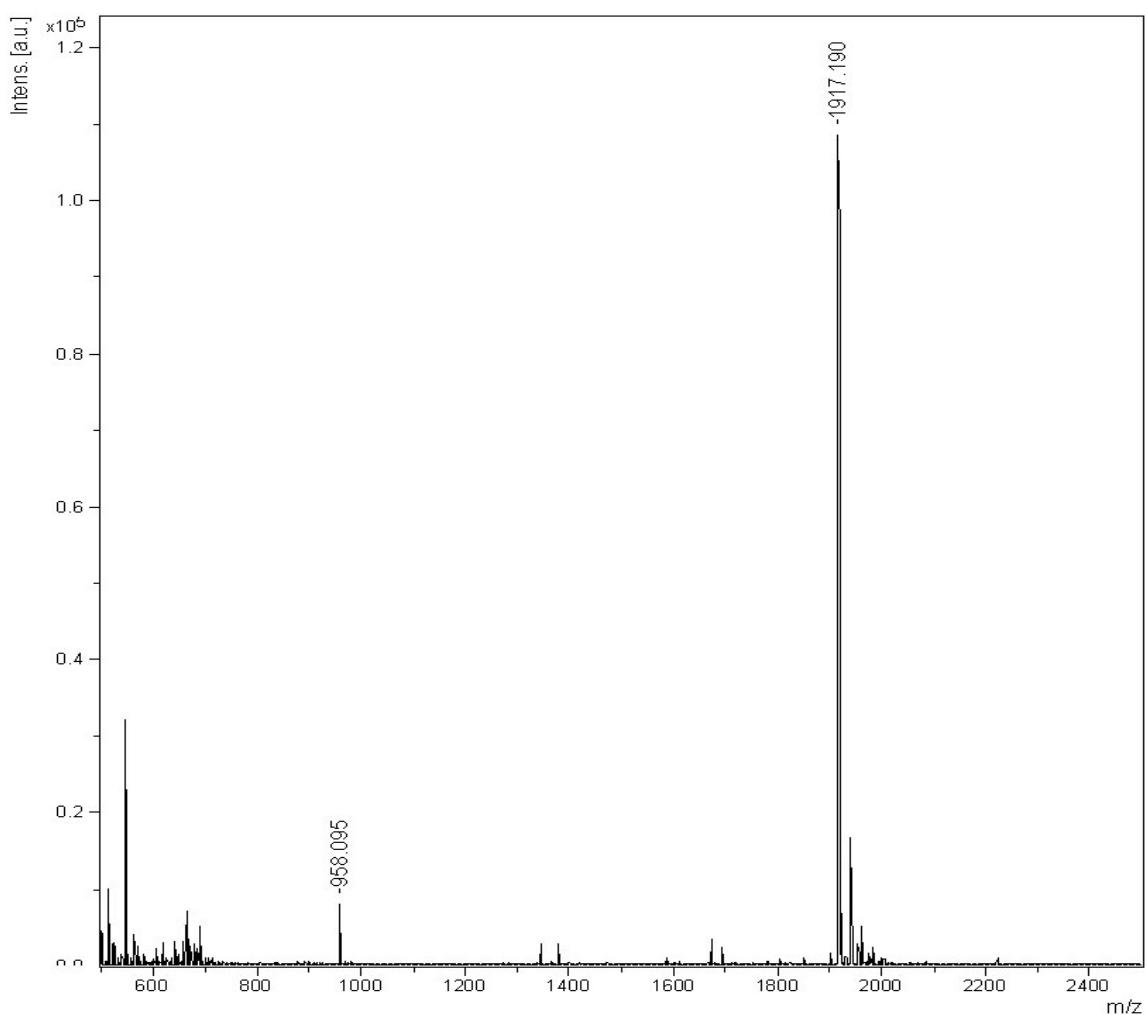


Figure S13(e11): Maldi ToF mass-spectral analysis of the RP-Hplc fraction at $R_T = 28.45$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CAAG^{Me}AAC)-3'** (**5c**) after **1h** [for Hplc profile, see Fig S12 (e2)] showing the main peak at m/z **1917.2** for **5'-AAG^{Me}AAC-3'** [see Table S9(E)].

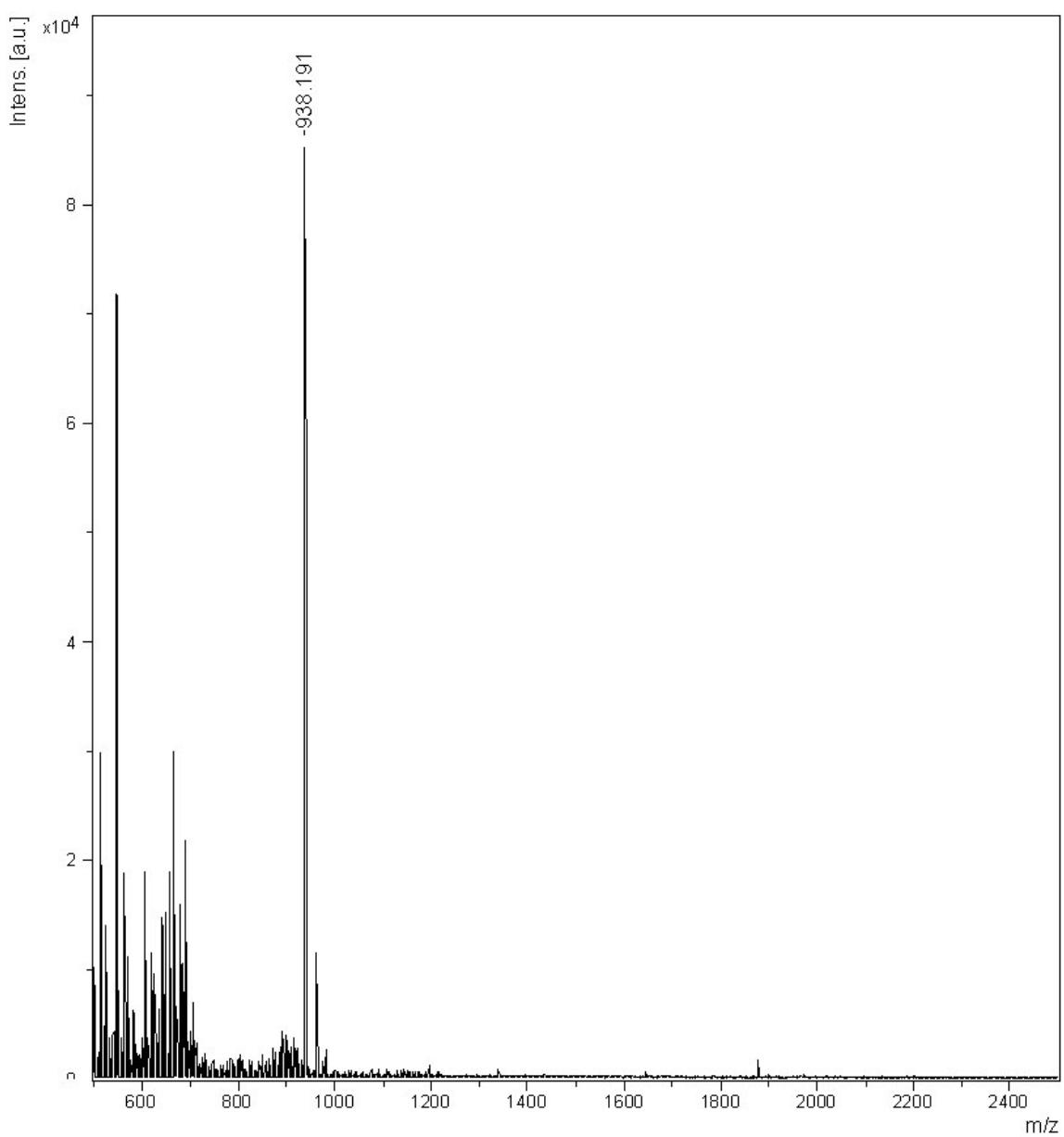


Figure S13(f1): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 19.68$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}AAC)-3'** (**7c**) after **1h** [for Hplc profile, see Fig **S12 (f2)**] showing the main peak at m/z **938.2** for **5'-CAC_{2'}, 3'-cMP** [see Table S9(F)].

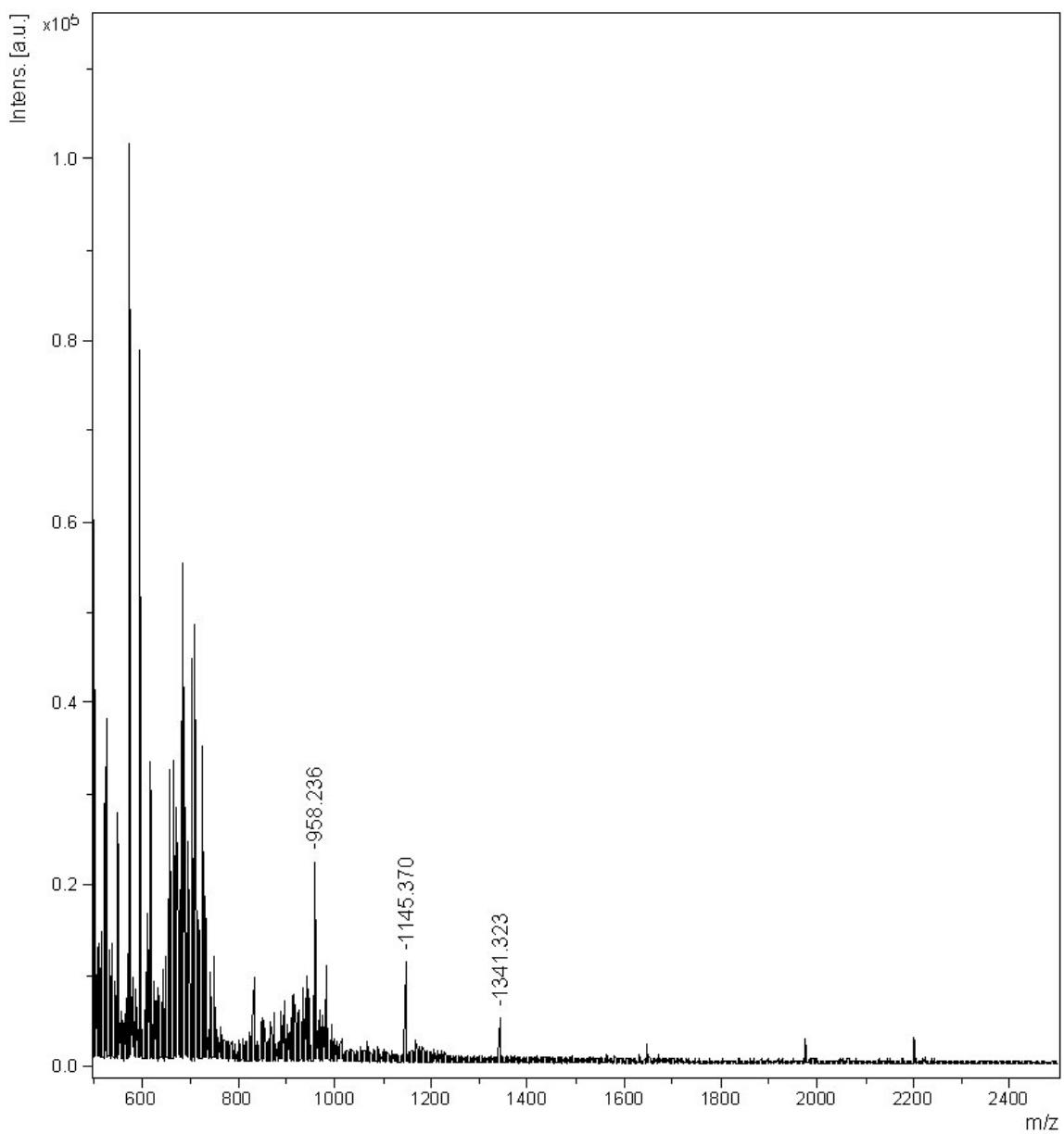


Figure S13(f2): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 21.81$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}AAC)-3'** (**7c**) after **1h** [for Hplc profile, see Fig S12 (f2)] showing the main peak at m/z **958.2** for **5'-CAC_{2'3'-P}** and other two peaks are non-nucleos(t)idic[see Table S9(F)].

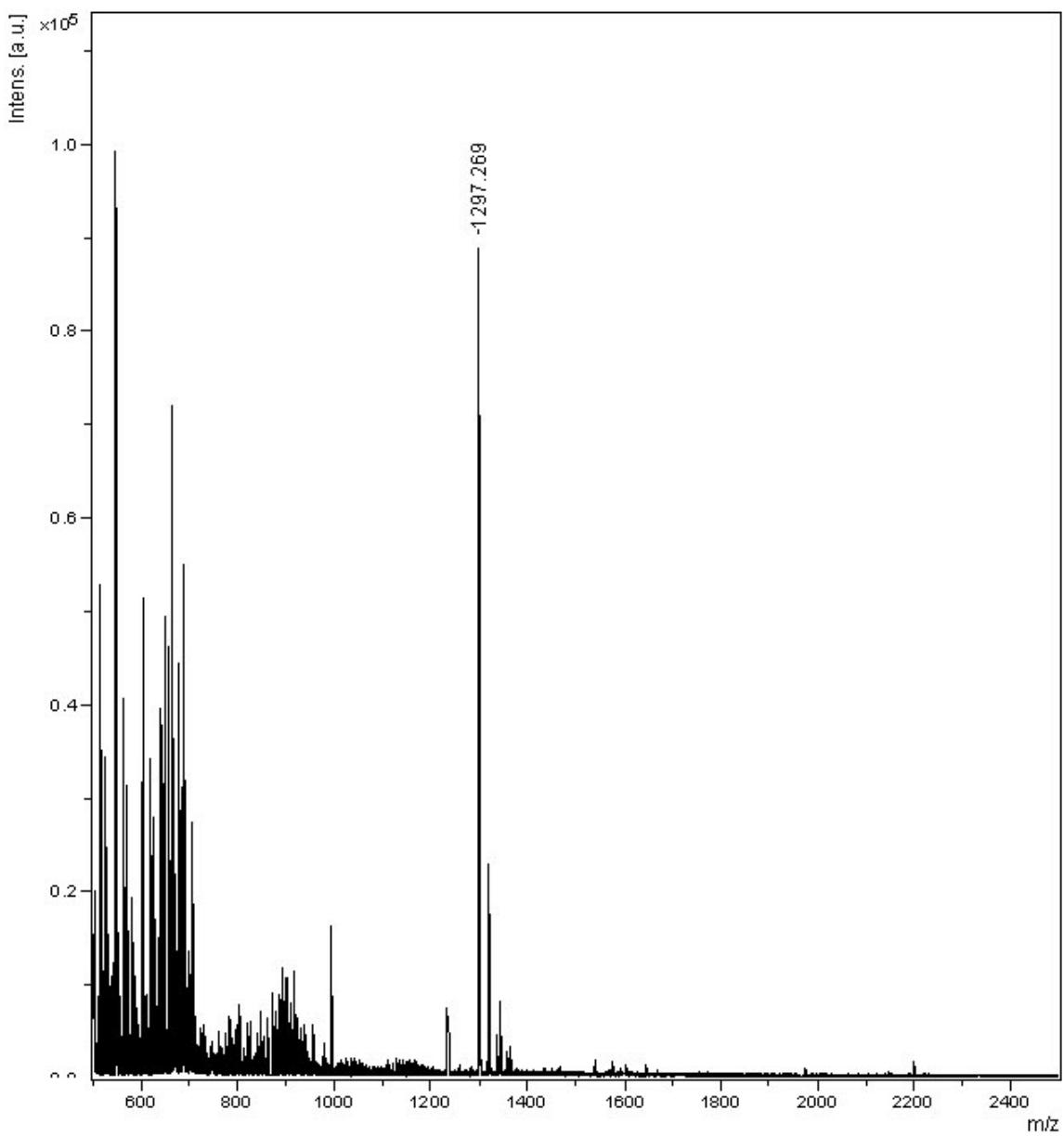


Figure S13(f3): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at **R_T = 22.89 min** for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}AAC)-3' (7c)** after **1h** [for Hplc profile, see Fig S12 (f2)] showing the main peak at m/z **1297.2** for **5'-CACG^{Me}_{2'},_{3'}-cMP** [see Table S9(F)].

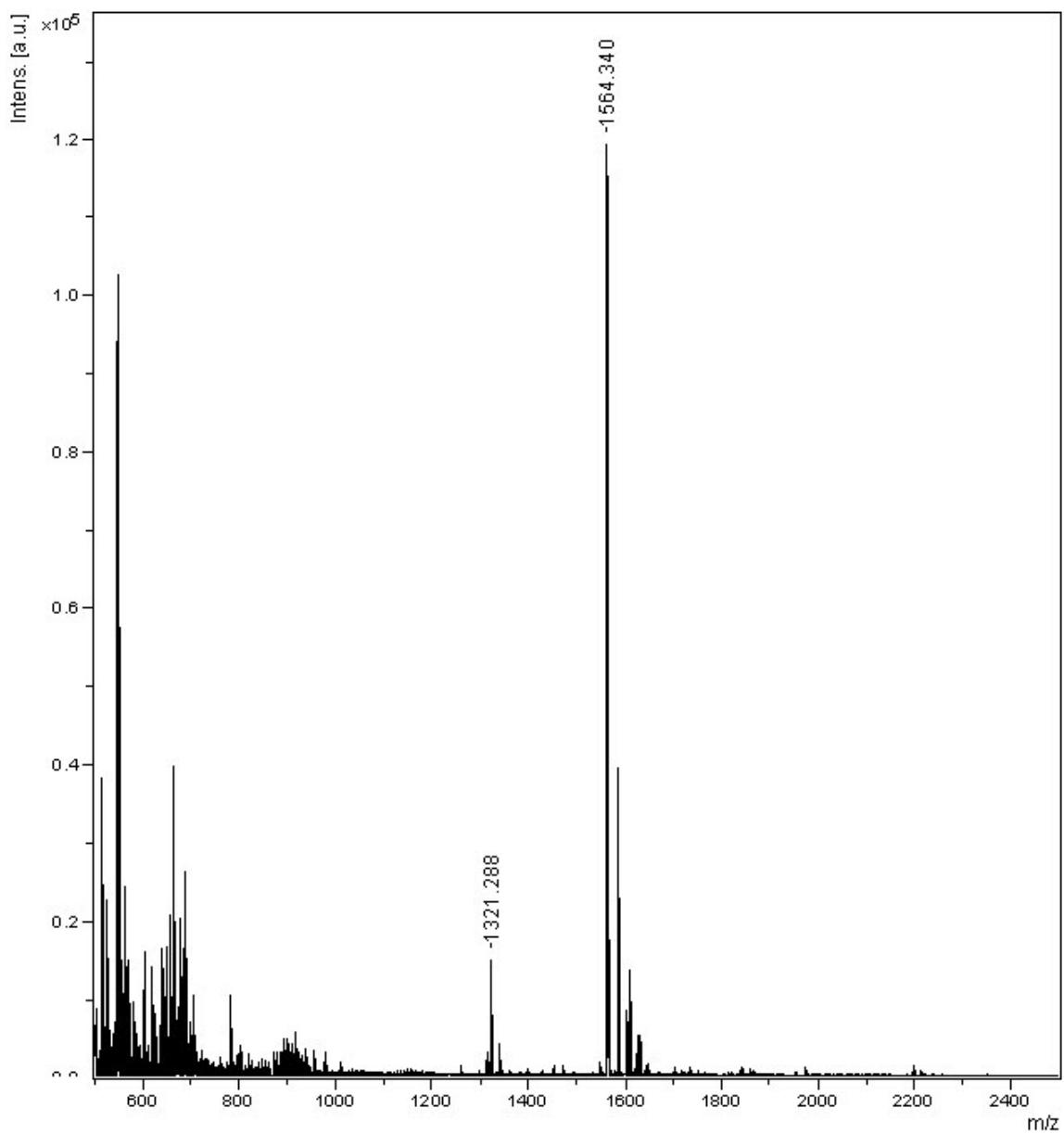


Figure S13(f4): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 23.85$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}AAC)-3'** (**7c**) after **1h** [for Hplc profile, see Fig S12 (f2)] showing the main peak at m/z **1564.3** for **5'-CG^{Me}AAC-3'** [see Table S9(F)].

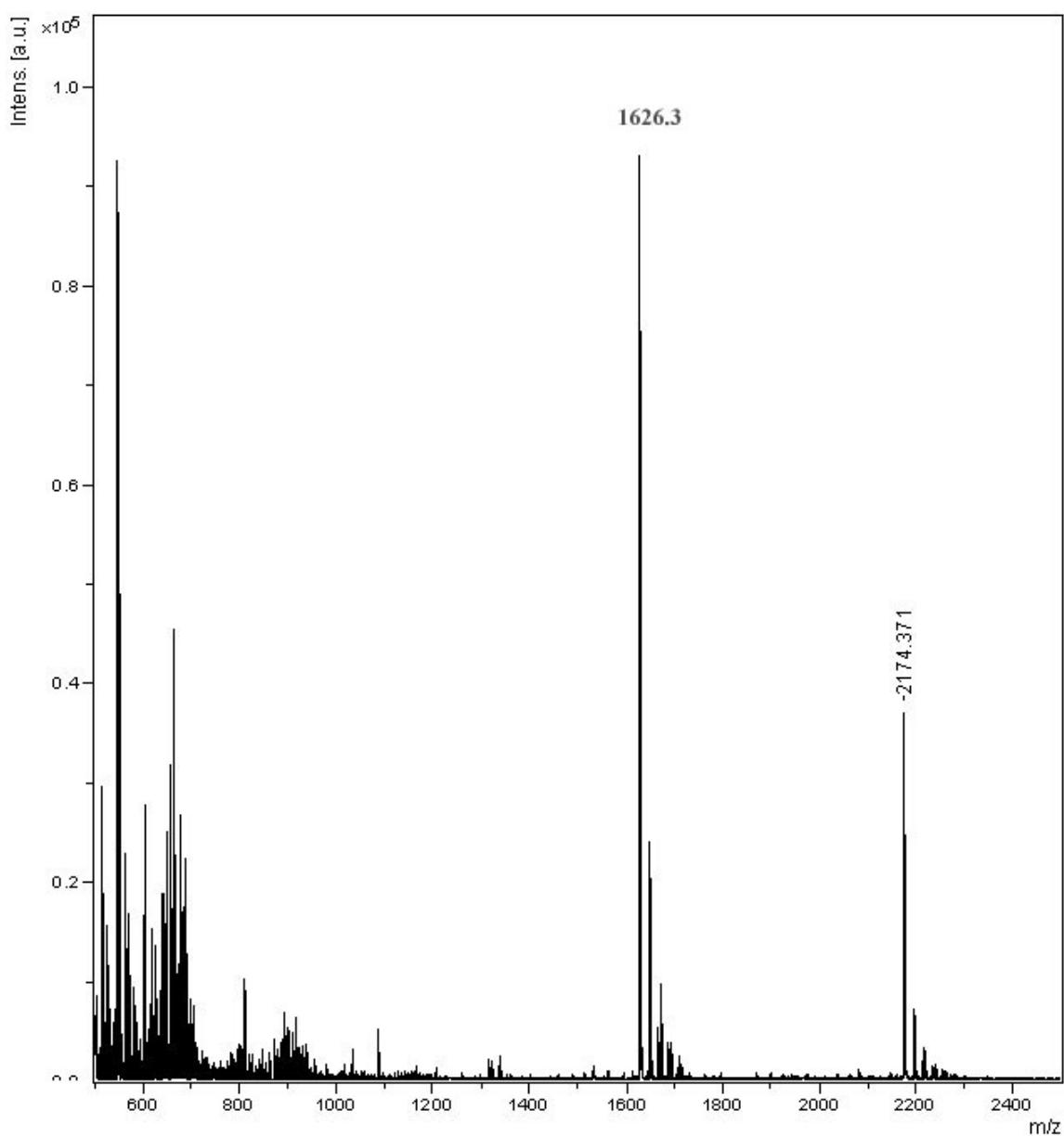


Figure S13(f5): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 24.77$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}AAC)-3' (7c)** after **1h** [for Hplc profile, see Fig S12 (f2)] showing the main peak at m/z **1626.3** for **5'-CACG^{Me}A-3'** along with some non-nucleos(t)idic peak at m/z **2174.3**. [see Table S9(F)].

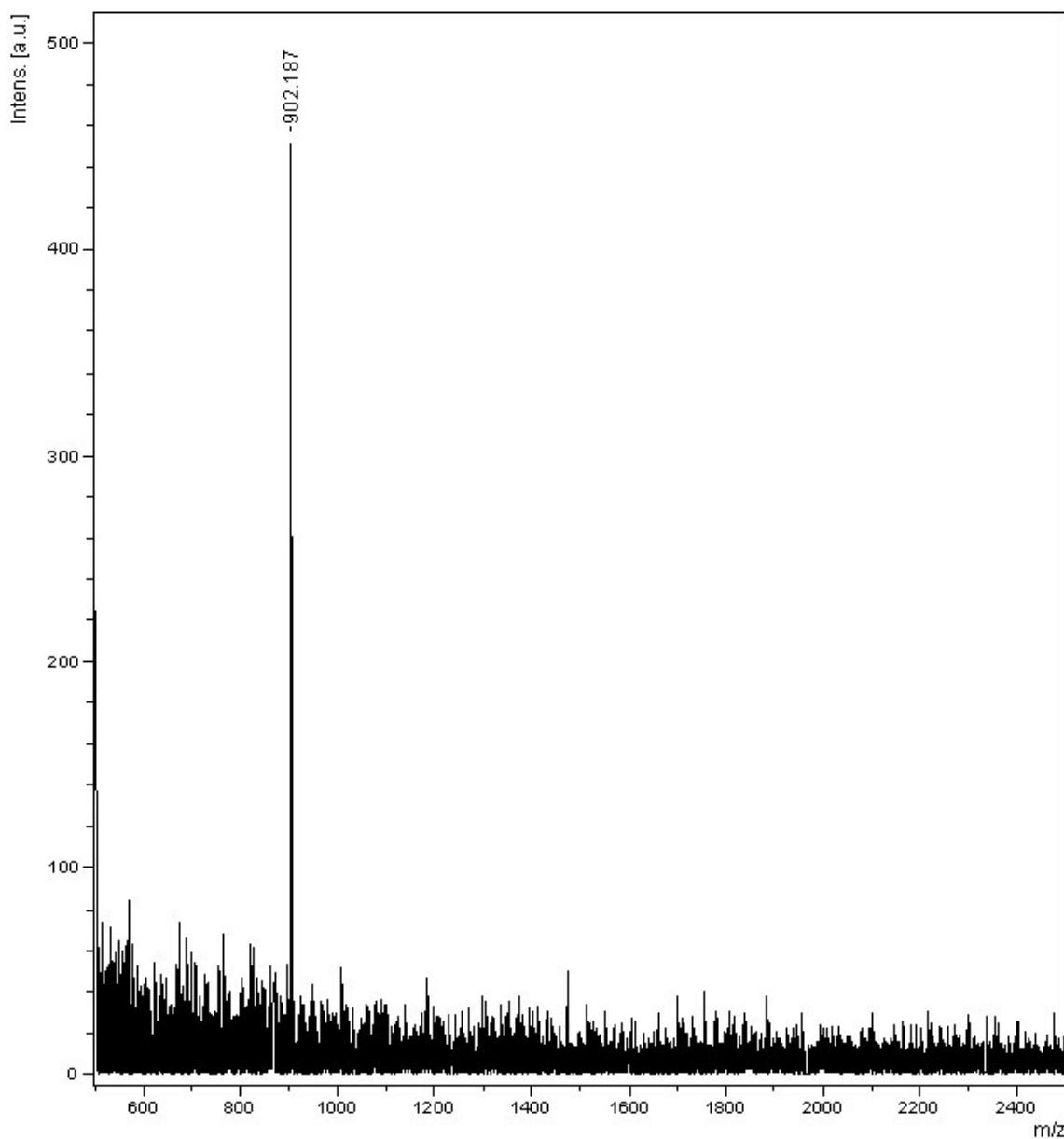


Figure S13(f6): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 25.15$ min and $R_T = 26.15$ in the **Fig S12(f2i)**. Corresponding mixed RP-Hplc fractions were at $R_T = 25.93$ min and $R_T = 26.49$ min in the **Fig S12(f2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}AAC)-3'**, **7c**] showing the main peak at m/z **902.3** for **5'-AAC-3'** [see Table S9(F)].

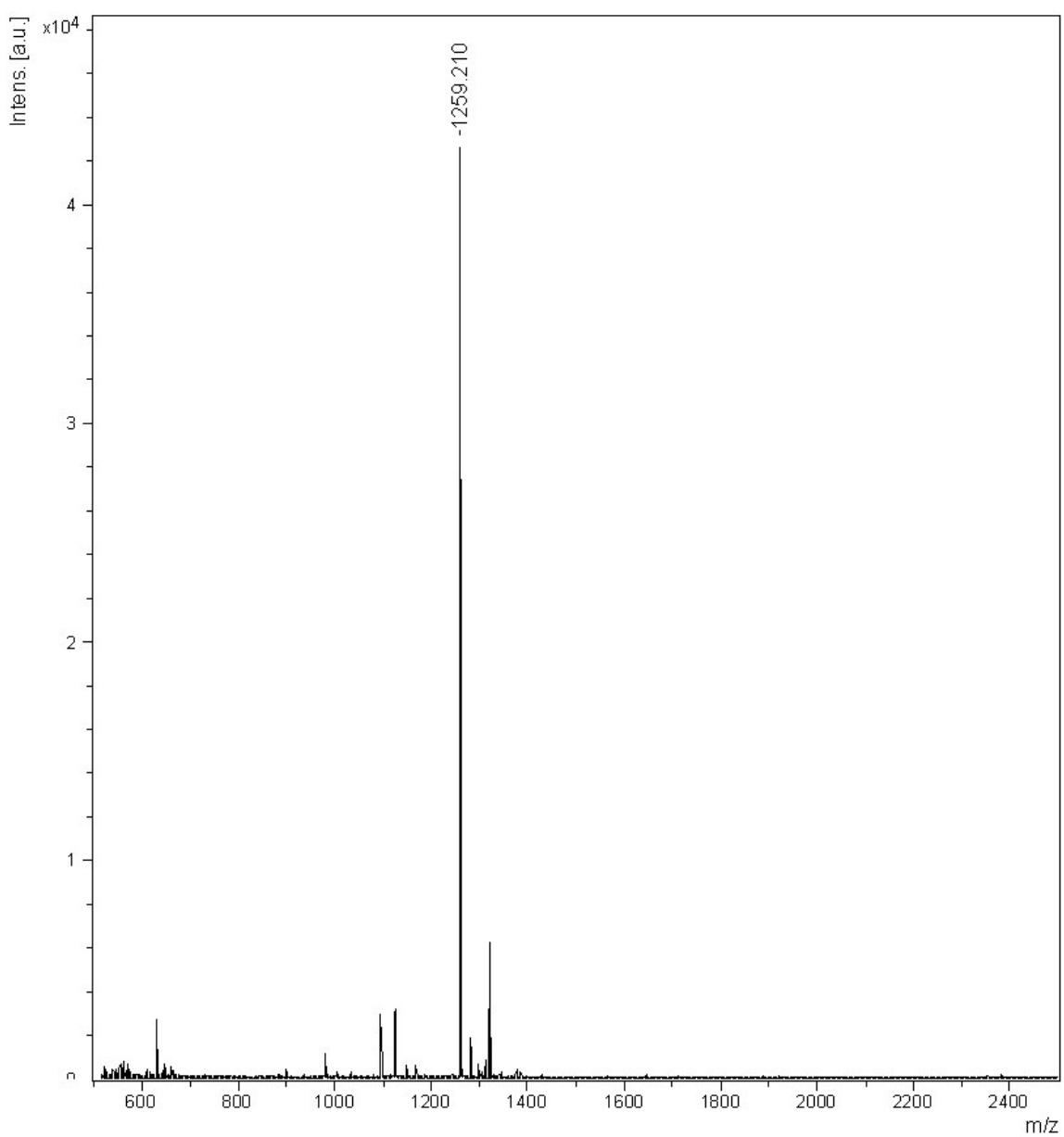


Figure S13(f7): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 27.71$ min in the Fig S12(f2i). Corresponding mixed RP-Hplc fractions were at $R_T = 25.93$ min and $R_T = 26.49$ min in the Fig S12(f2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}AAC)-3'**, **7c**] showing the main peak at m/z **1259.2** for **5'-G^{Me}AAC-3'** [see Table S9(F)].

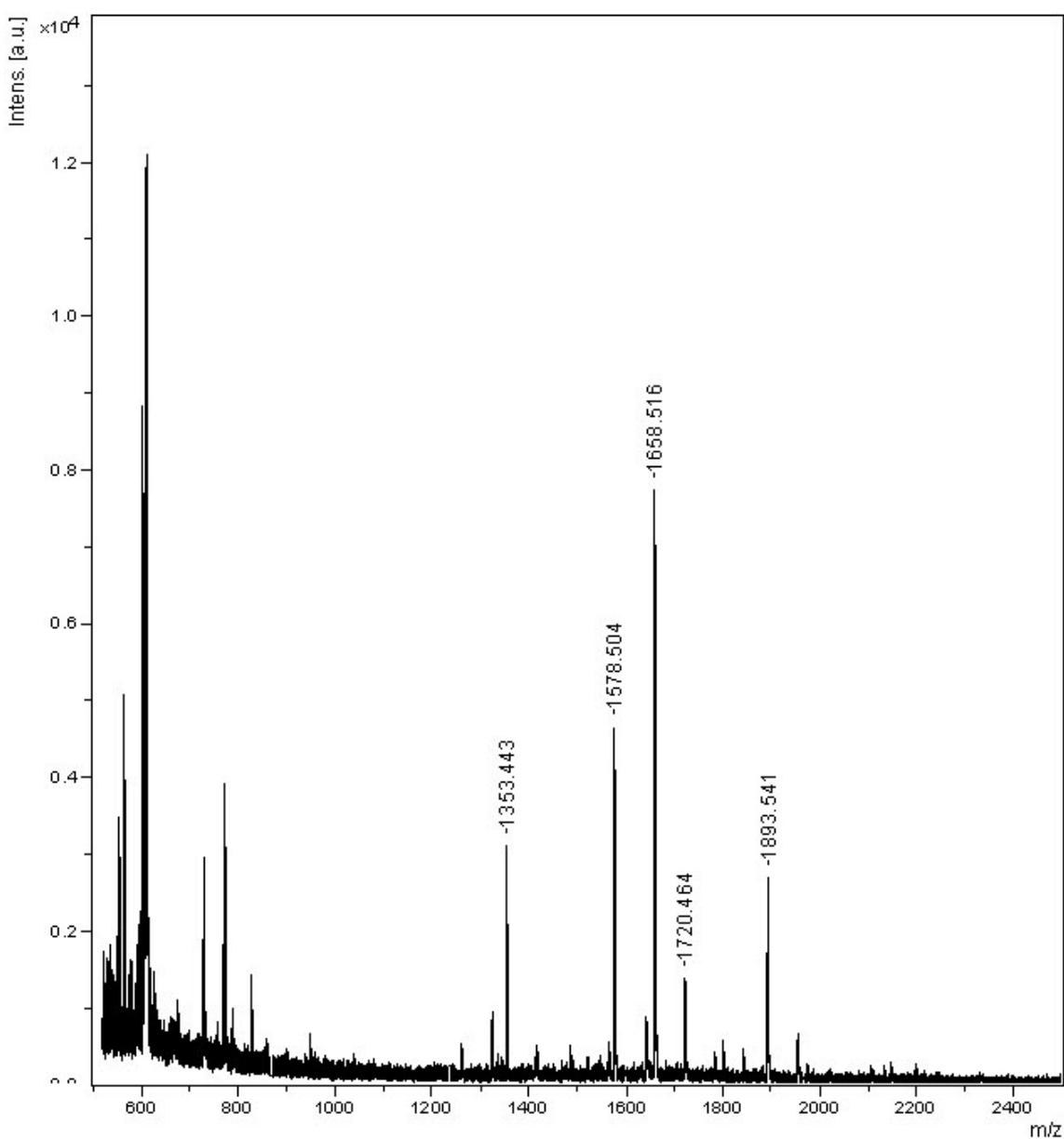


Figure S13(f8): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 29.00$ min in the Fig S12(f2i). Corresponding mixed RP-Hplc fractions were at $R_T = 25.93$ min and $R_T = 26.49$ min in the Fig S12(f2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}AAC)-3'**, **7c**] showing the peak at m/z 1894.5 for **5'-ACG^{Me}AAC-3'**. All other peaks are non-nucleos(t)idic.[see Table S9(F)].

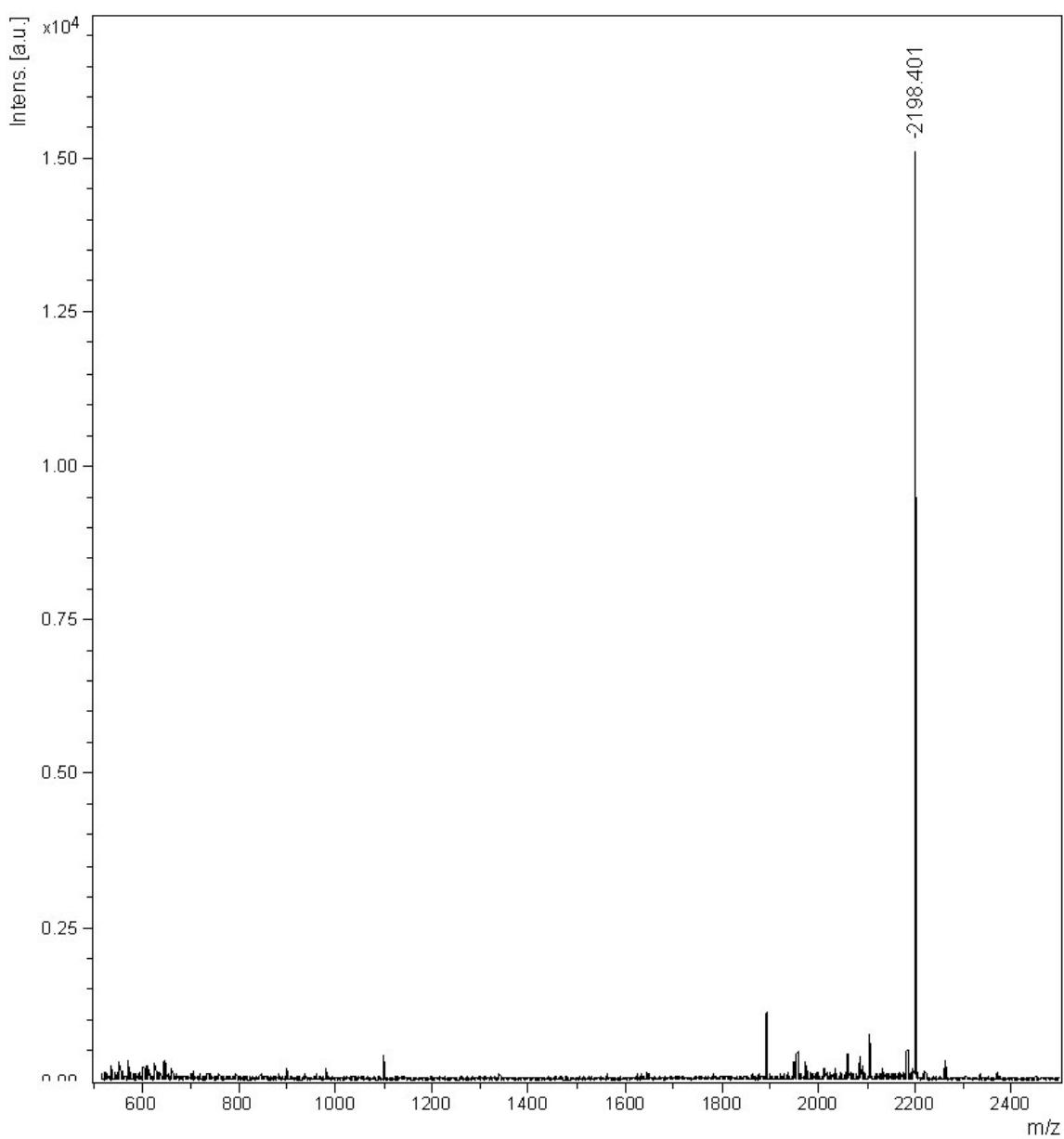


Figure S13(f9): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 29.91$ min in the Fig S12(f2i). Corresponding mixed RP-Hplc fractions were at $R_T = 25.93$ min and $R_T = 26.49$ min in the Fig S12(f2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}AAC)-3'**, **7c**] showing main the peak at m/z 2198.4 for **5'-CACG^{Me}AAC-3'**, the parent heptamer [see Table S9(F)].

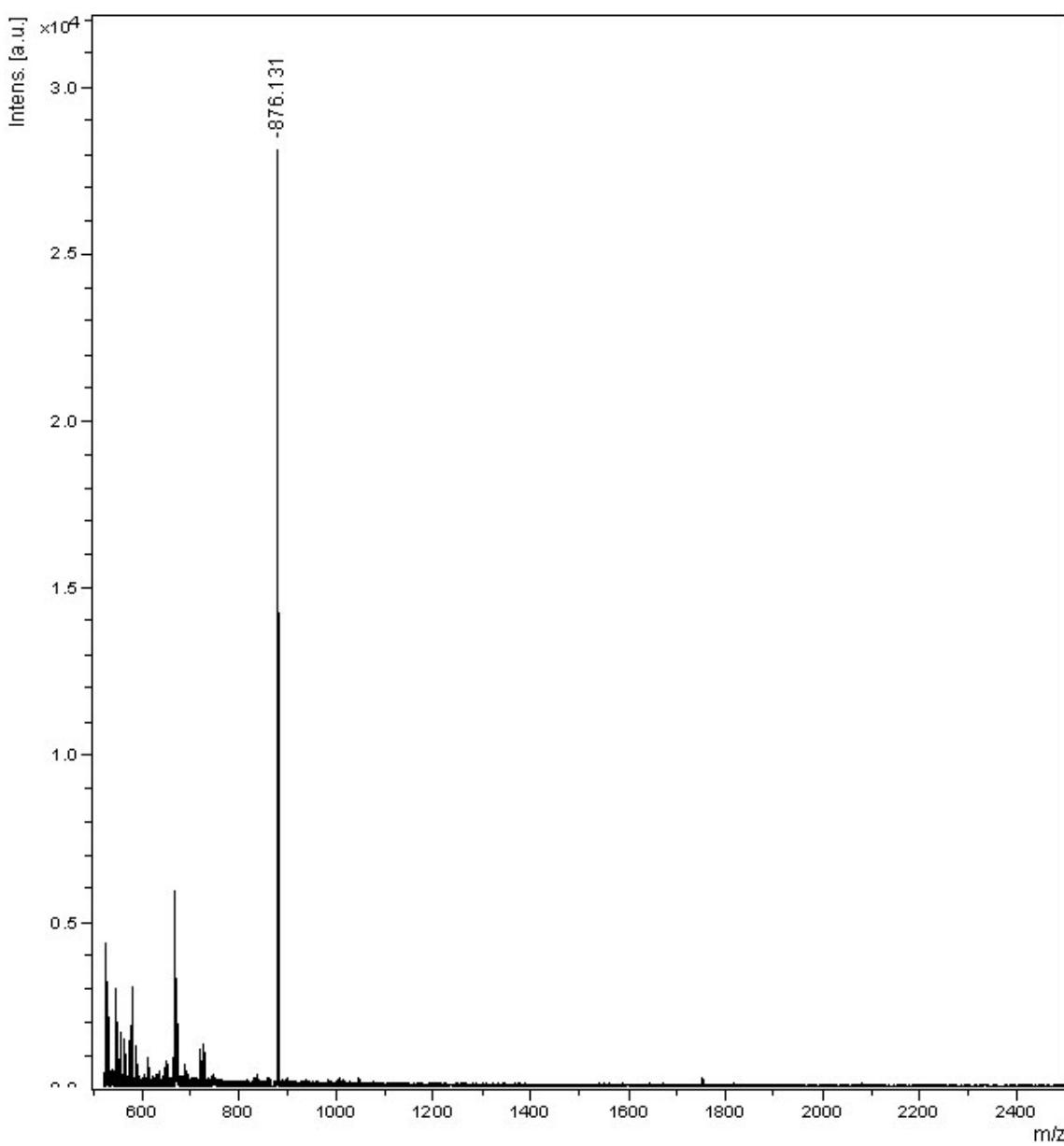


Figure S13(g1): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 20.09$ min in the Fig S12(g2i). Corresponding RP-Hplc fraction is $R_T = 19.68$ min in the Fig S12(g2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}CAC)-3'**, **8c**]. The main peak at m/z **876.1** for **5'-CAC-3'** [see Table S9(G)]

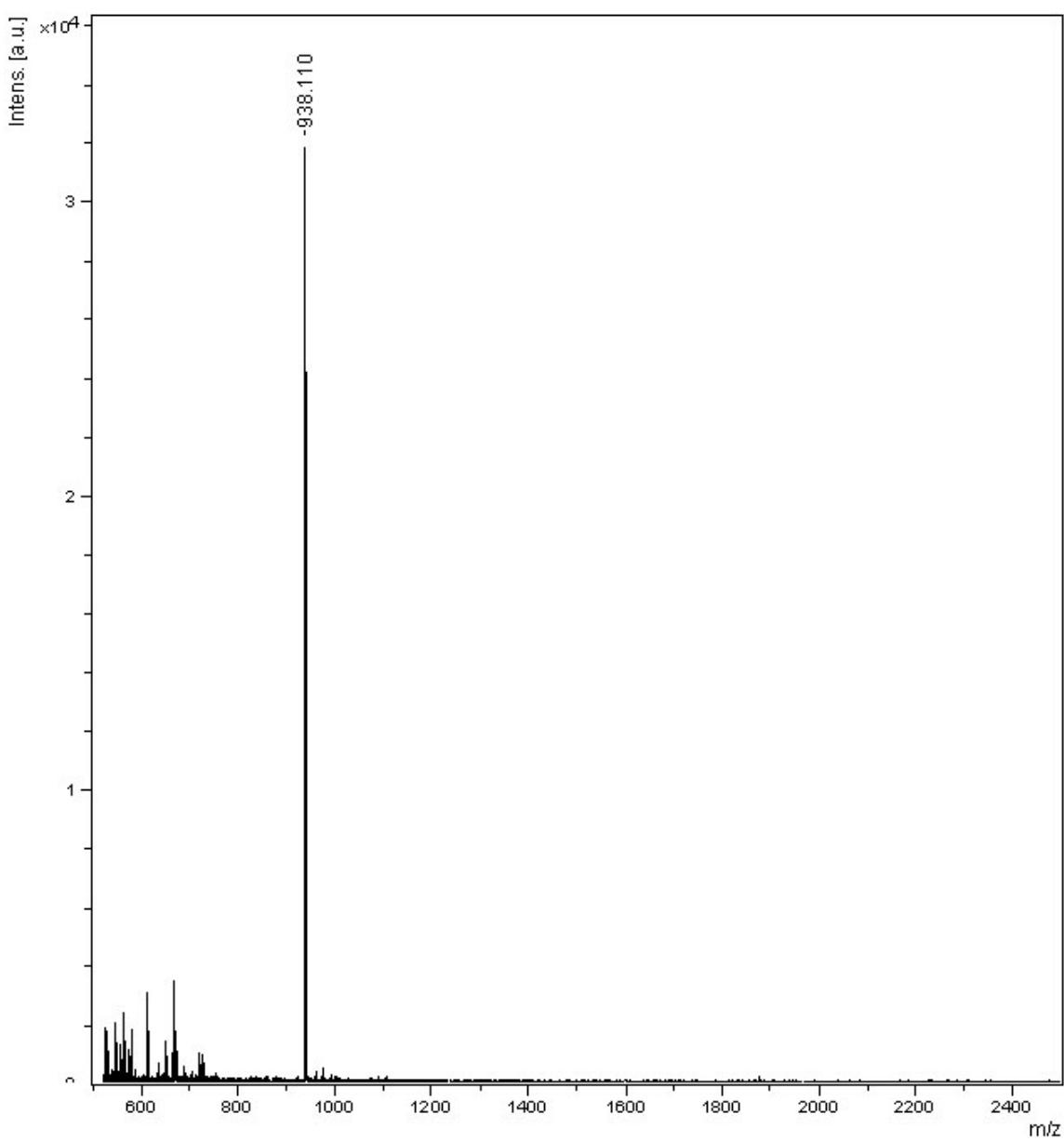


Figure S13(g2): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 20.80$ min in the Fig S12(g2i). Corresponding RP-Hplc fraction is $R_T = 19.68$ min in the Fig S12(g2) [after 1h of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}CAC)-3'**, **8c**]. The main peak at m/z **938.1** for **5'-CAC₂,3'-cMP** [see Table S9(G)].

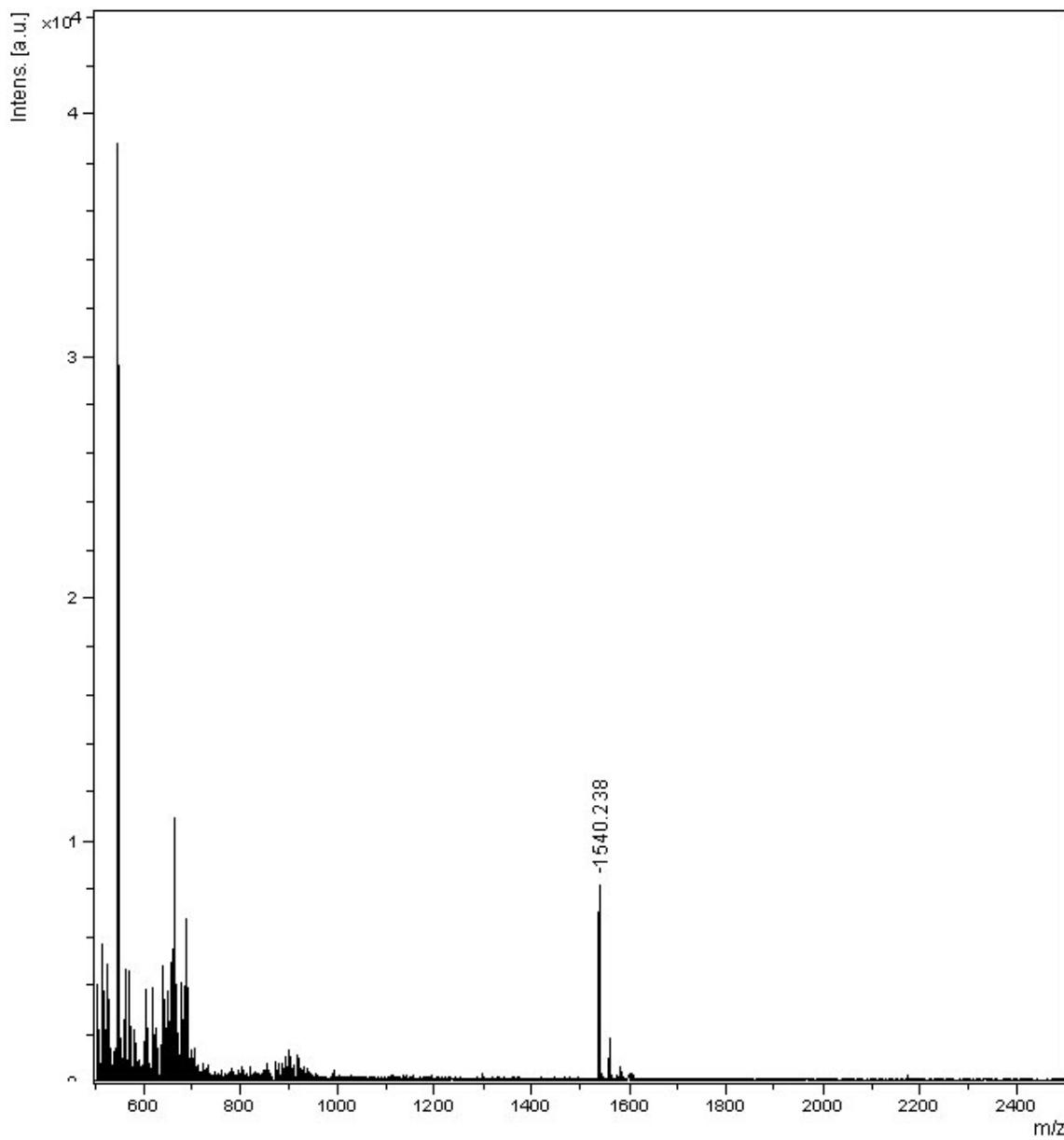


Figure S13(g3): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at **R_T = 22.55 min** for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}CAC)-3' (8c)** after **1h** [for Hplc profile, see Fig S12 (g2)] showing the main peak at m/z **1540.2** for **5'-CG^{Me}CAC-3'** [see Table S9(G)].

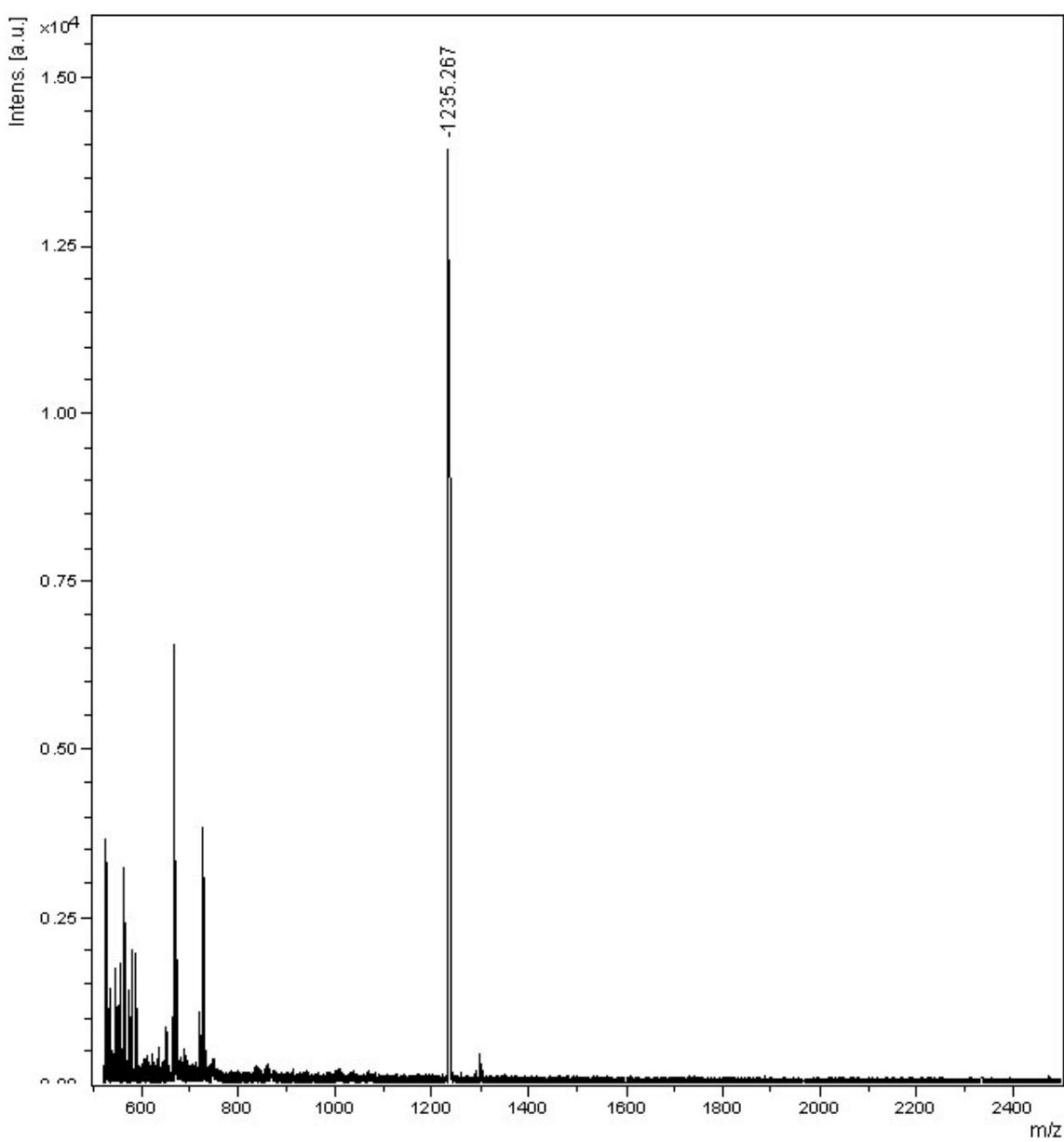


Figure S13(g4): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 26.73$ min in the **Fig S12(g2j)**. Corresponding RP-Hplc fraction is $R_T = 23.15$ min in the **Fig S12(g2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}CAC)-3'**, **8c**]. The main peak at m/z **1235.2** for **5'-CACG^{Me}2',3'-cMP** [see Table S9(G)].

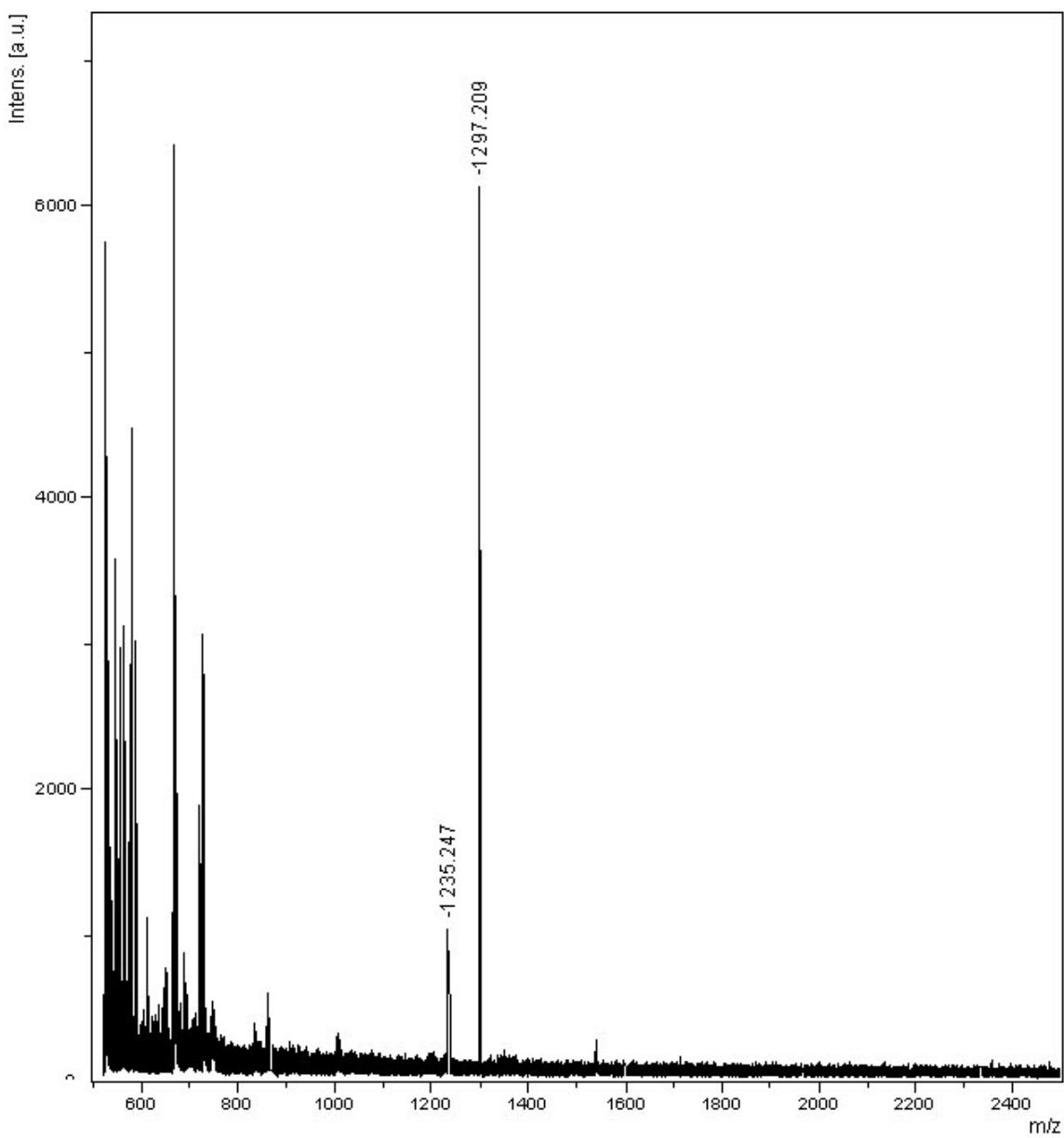


Figure S13(g5): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T \approx 27.00$ min in the **Fig S12(g2j)**. Corresponding RP-Hplc fraction is $R_T = 23.15$ min in the **Fig S12(g2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}CAC)-3'**, **8c**]. The main peak at m/z **1297.2** for **5'-G^{Me}CAC-3'** [see Table S9(G)].

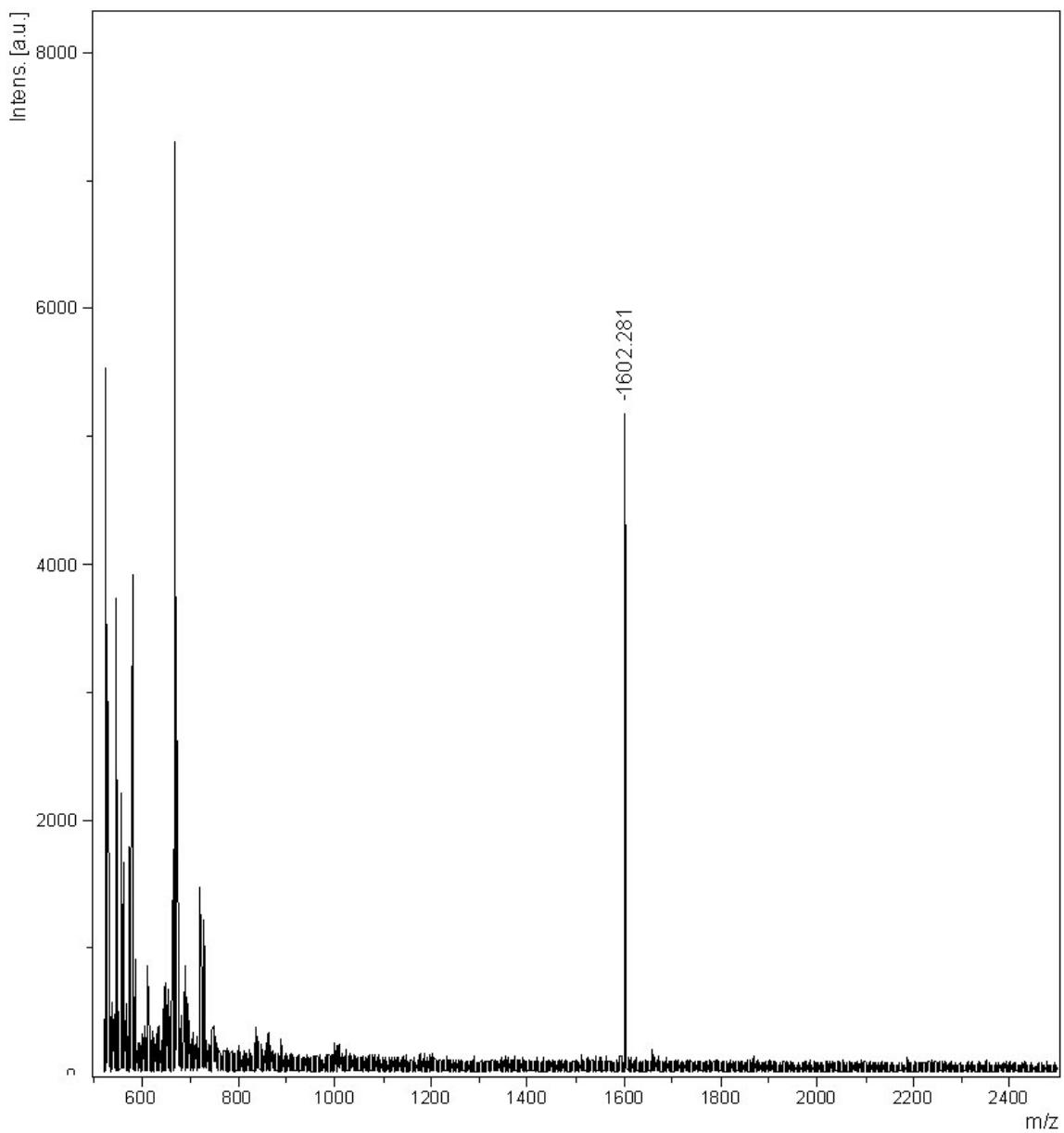


Figure S13(g6): Maldi-Tof mass-spectral analysis of the SMART™ RP-Hplc fraction at $R_T = 27.76$ min in the **Fig S12(g2j)**. Corresponding RP-Hplc fraction is $R_T = 23.15$ min in the **Fig S12(g2)** [after **1h** of alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}CAC)-3'**, **8c**]. The main peak at m/z **1602.2** for **5'-CACG^{Me}C_{2'}, 3'-cMP** [see Table S9(G)].

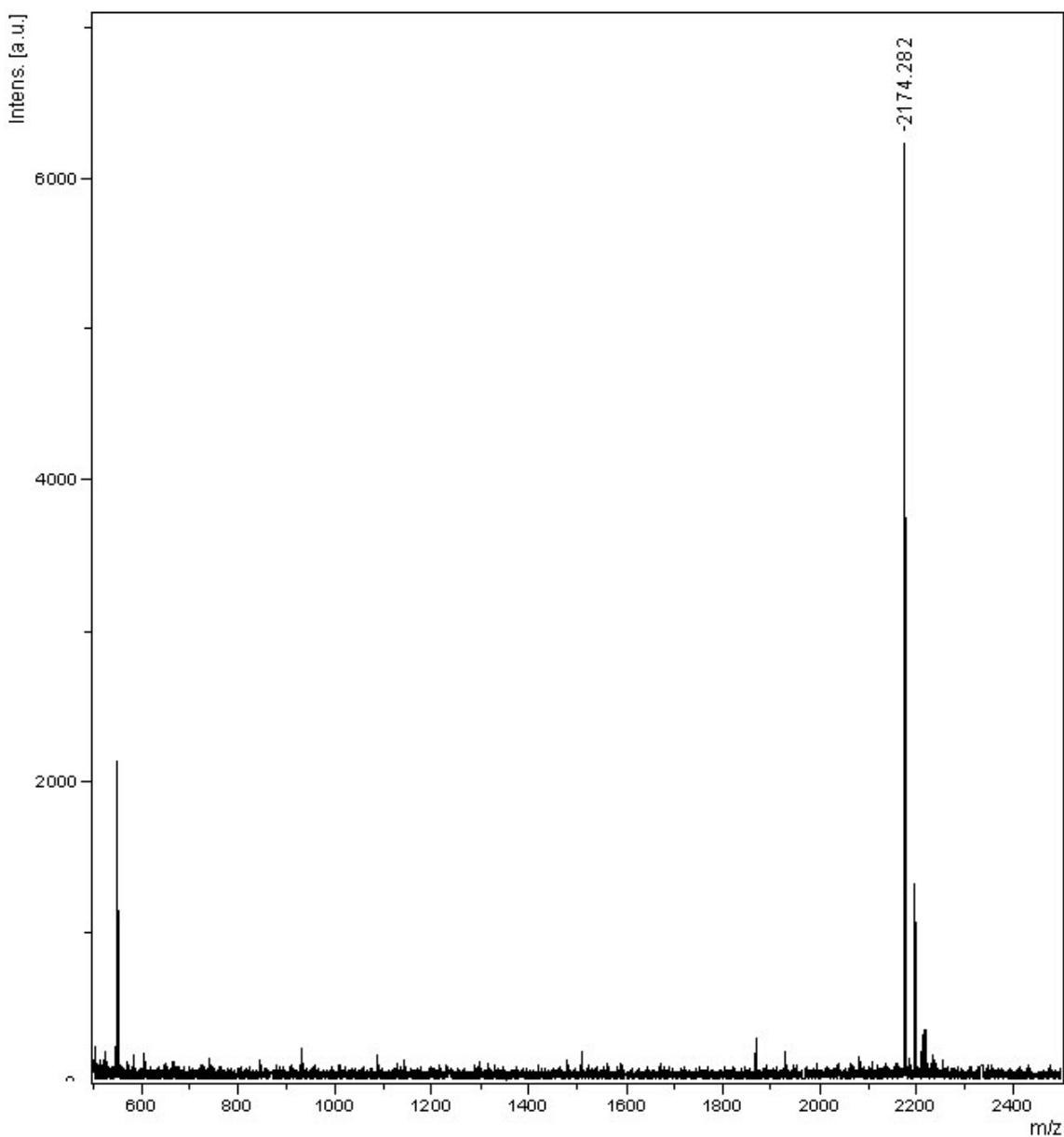


Figure S13(g7): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 24.87$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}CAC)-3' (8c)** after **1h** [for Hplc profile, see Fig **S12 (g2)**] showing the main peak at m/z **2174.2** for the parent heptamer [see Table S9(G)].

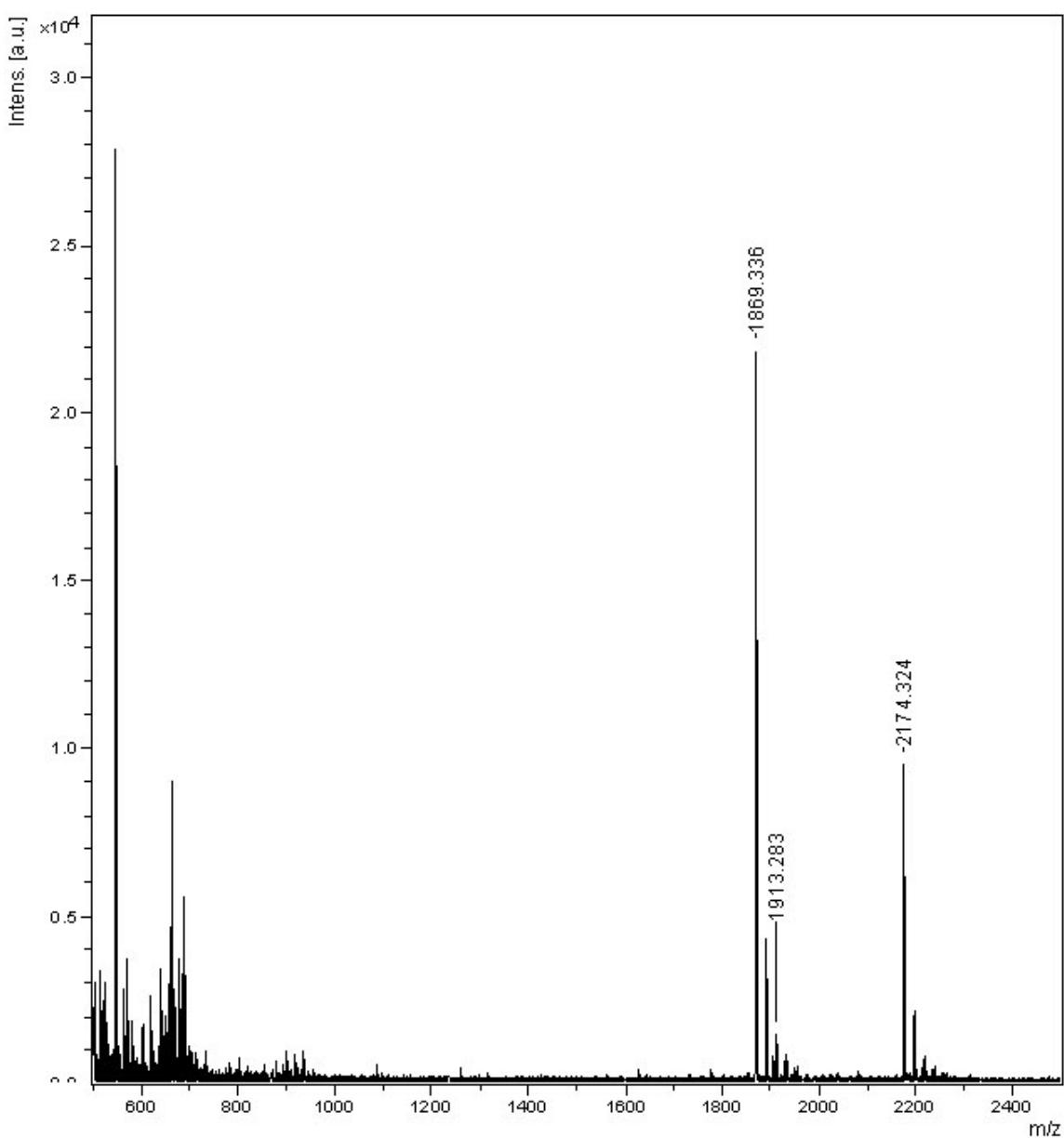


Figure S13(g8): Maldi Tof mass-spectral analysis of the RP-Hplc fraction at $R_T = 25.82$ min for alkaline hydrolysis (pH 12.5 / 20°C) of **5'-r(CACG^{Me}CAC)-3'** (**8c**) after **1h** [for Hplc profile, see Fig **S12 (g2)**] showing the main peak at m/z **1869.3** for **5'-ACG^{Me}CAC-3'** along with the parent heptameric peak[see Table S9(G)].