

Average Mass Differences Relative to the Full Length Product for Common Impurities

Possible problem	Mass difference (Dalton)		
PS backbone			96.05
PO backbone			79.98
PS-PO conversion			-16.07
Depurination of A base			-117.12
			-135.13
Depurination of G base			-133.11
			-151.13
Deamination			0.98
Reaction with methylamine			14.03
+Acetyl			42.04
+Cyanoethyl			53.06
+isobutyryl (G)			70.08
+Chloral (backbone)			147.39
+tert-butyl dimethylsilyl			114.26
+benzoyl			104.11
Modified C			80.08
+DMT			302.37
Depyrimidation			-52.03
Depyrimidation			-94.07
-dA (PO)	-313.21	-dA (PS)	-329.28
-dG (PO)	-329.21	-dG (PS)	-345.28
-dC (PO)	-289.18	-dC (PS)	-305.25
-5Me-dC (PO)	-303.21	-5Me-dC (PS)	-319.28
-dT (PO)	-304.19	-dT (PS)	-320.26
-dU (PO)	-290.17	-dU (PS)	-306.24
-rA (PO)	-329.21	-rA (PS)	-345.28
-rG (PO)	-345.21	-rG (PS)	-361.28
-rC (PO)	-305.18	-rC (PS)	-321.25
-5Me-rC (PO)	-319.21	-5Me-rC (PS)	-335.28
-rU (PO)	-306.17	-rU (PS)	-322.24
-rT (PO)	-320.19	-rT (PS)	-336.26
-mA (PO)	-343.23	-mA (PS)	-359.3
-mG (PO)	-359.23	-mG (PS)	-375.3
-mC (PO)	-319.21	-mC (PS)	-335.28

-5Me-mC (PO)	-333.24	-5Me-mC (PS)	-349.31
-mU (PO)	-320.19	-mU (PS)	-336.26
-mT (PO)	-334.22	-mT (PS)	-350.29
-fA (PO)	-331.2	-fA (PS)	-347.27
-fG (PO)	-347.2	-fG (PS)	-363.27
-fC (PO)	-307.17	-fC (PS)	-323.24
-5Me-fC (PO)	-321.2	-5Me-fC (PS)	-337.27
-fU (PO)	-308.16	-fU (PS)	-324.23
-fT (PO)	-322.18	-fT (PS)	-338.25
-LNA-A (PO)	-341.22	-LNA-A (PS)	-357.29
-LNA-G (PO)	-357.22	-LNA-G (PS)	-373.29
-LNA-C (PO)	-317.19	-LNA-C (PS)	-333.26
-LNA-U (PO)	-318.18	-LNA-U (PS)	-334.25
-5Me-LNA-C (PO)	-331.22	-5Me-LNA-C (PS)	-347.29
-LNA-T (PO)	-332.20	-LNA-T (PS)	-348.27
-dI (PO)	-314.19	-dI (PS)	-330.26
-rI (PO)	-330.19	-rI (PS)	-346.26
-mI (PO)	-344.22	-mI (PS)	-360.29
-cET-A (PO)	-355.24	-cET-A (PS)	-371.31
-cET-G (PO)	-371.26	-cET-G (PS)	-387.33
-cET-C (PO)	-331.22	-cET-C (PS)	-347.29
-5Me-cET-C (PO)	-345.25	-5Me-cET-C (PS)	-361.32
-cET-U (PO)	-332.2	-cET-U (PS)	-348.27
-UNA-A (PO)	-331.22	-UNA-A (PS)	-347.29
-UNA-G (PO)	-347.22	-UNA-G (PS)	-363.29
-UNA-C (PO)	-307.20	-UNA-C (PS)	-323.27
-UNA-U (PO)	-308.18	-UNA-U (PS)	-324.25
-GNA-A (PO)	-271.17	-GNA-A (PS)	-287.24
-GNA-G (PO)	-287.17	-GNA-G (PS)	-303.24
-GNA-C (PO)	-247.15	-GNA-C (PS)	-263.22
-GNA-T (PO)	-262.16	-GNA-T (PS)	-278.23
-TNA-A (PO)	-299.18	-TNA-A (PS)	-315.25
-TNA-G (PO)	-315.18	-TNA-G (PS)	-331.25
-TNA-C (PO)	-275.16	-TNA-C (PS)	-291.23
-TNA-T (PO)	-290.17	-TNA-T (PS)	-306.24
-MOE-A (PO)	-387.29	-MOE-A (PS)	-403.36
-MOE-G (PO)	-403.28	-MOE-G (PS)	-419.35

-5Me-MOE-C (PO)	-377.29	-5Me-MOE-C (PS)	-393.36
-MOE-T (PO)	-378.27	-MOE-T (PS)	-394.34
-Amino C6 (PO)	-179.16	-Amino C6 (PS)	-195.23
-Sp9 (PO)	-212.14	-Spacer 9 (PS)	-228.21
-Sp 18 (PO)	-344.30	-Sp 18 (PS)	-360.37
-SpC3 (PO)	-138.06	-SpC3 (PS)	-154.13
-SpC6 (PO)	-180.14	-SpC6 (PS)	-196.21
-SpC9 (PO)	-222.22	-SpC9 (PS)	-238.29
-SpC12 (PO)	-264.3	-SpC12 (PS)	-280.37
-VPmU (PO)	-396.18	-VPmU (PS)	-412.25
-VPmA (PO)	-419.23	-VPmA (PS)	-435.3
-VPmG (PO)	-435.22	-VPmG (PS)	-451.29
-VPmC (PO)	-395.2	-VPmC (PS)	-411.27