

Supplementary Materials: Effects of a Novel Thiadiazole Derivative with High Anticancer Activity on Cancer Cell Immunogenic Markers: Mismatch Repair System, PD-L1 Expression, and Tumor Mutation Burden

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Flow Cytometric Analysis Data

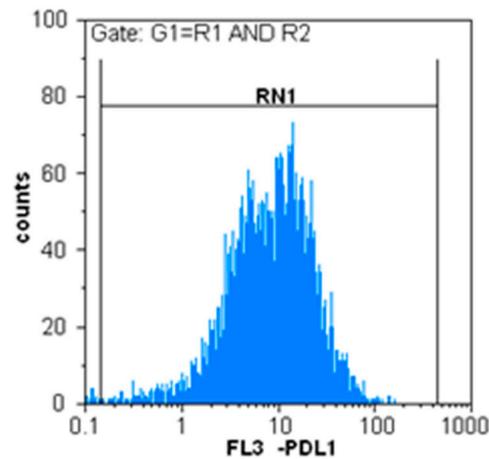


Figure S1. Flow cytometric analysis of PD-L1 expression on untreated DLD-1 cancer cells cultured for 48 h.

Table S1. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	6730	6730	-	65.61	2.37	65.80	12.13	47.06
RN1	G1	9784	3989	-	97.34	12.92	104.62	-	-
R2	<None>	4283	4283	-	41.75	0.52	488.28	0.26	10.38

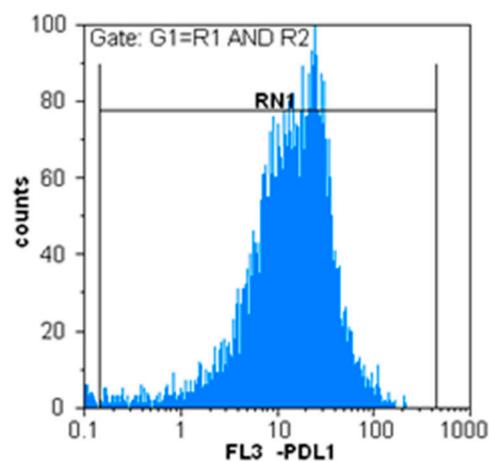


Figure S2. Flow cytometric analysis of PD-L1 expression on DLD-1 cancer cells treated with KA39 at IC₅₀ concentration (μ M) for 48 h.

Table S2. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	7320	7320	-	62.06	2.99	68.31	16.15	40.74
RN1	G1	11050	4991	-	93.69	20.14	97.27	-	-
R2	<None>	5472	5472	-	46.39	0.67	396.09	0.26	9.78

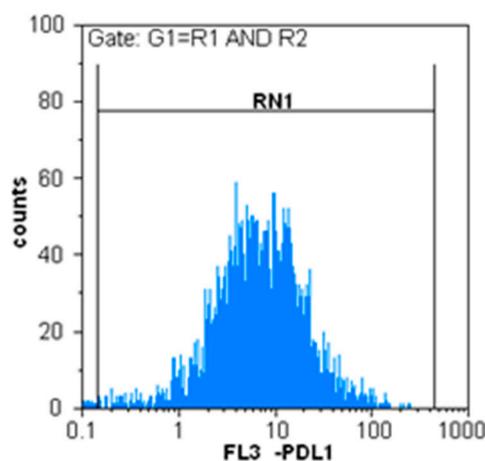


Figure S3. Flow cytometric analysis of PD-L1 expression on DLD-1 cancer cells treated with KA39 at IC₅₀ concentration (μ M) for 72 h.

Table S3. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	6798	6798	-	66.50	2.04	71.35	10.39	44.68
RN1	G1	9803	3383	-	97.05	12.30	144.61	-	-
R2	<None>	3628	3628	-	35.49	0.35	583.91	0.26	9.72

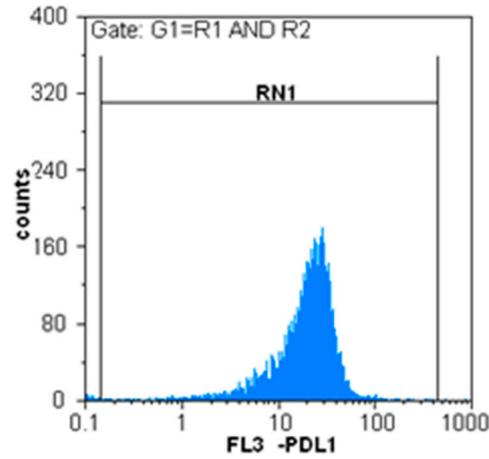


Figure S4. Flow cytometric analysis of PD-L1 expression on untreated DLD-1 cancer cells cultured for 72 h.

Table S4. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	11511	11511	-	55.75	3.21	64.50	18.92	42.29
RN1	G1	19656	5599	-	97.14	23.19	75.96	-	-
R2	<None>	6010	6010	-	29.11	2.14	267.75	0.28	10.09

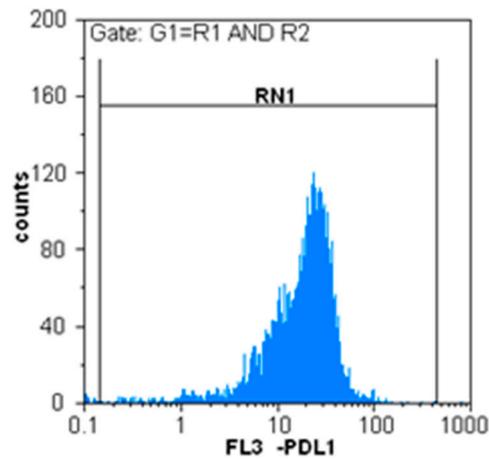


Figure S5. Flow cytometric analysis of PD-L1 expression on DLD-1 cancer cells treated with KA39 at TGI concentration (μ M) for 72 h.

Table S5. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	8954	8954	-	44.08	2.85	56.68	17.68	39.68
RN1	G1	19308	4172	-	97.20	22.27	77.37	-	-
R2	<None>	4711	4711	-	23.19	1.39	328.99	0.28	9.77

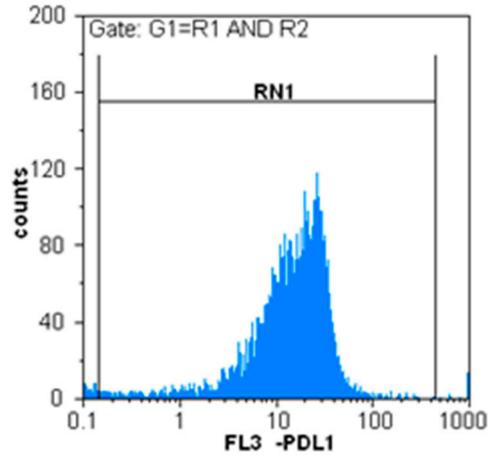


Figure S6. Flow cytometric analysis of PD-L1 expression on untreated HT-29 cancer cells cultured for 48 h.

Table S6. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	7454	7454	-	72.54	3.21	58.94	16.89	40.95
RN1	G1	9513	4681	-	91.03	19.56	104.78	-	-
R2	<None>	5277	5277	-	51.35	1.05	364.10	0.26	10.61

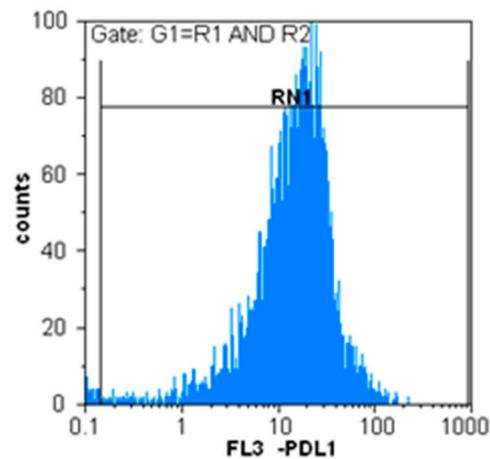


Figure S7. Flow cytometric analysis of PD-L1 expression on HT-29 cancer cells treated with KA39 at IC₅₀ concentration (μM) for 48 h.

Table S7. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	7237	7237	-	69.64	4.31	59.31	15.58	40.30
RN1	G1	9855	4524	-	93.88	20.13	92.52	-	-
R2	<None>	4908	4908	-	47.23	0.61	424.39	0.26	9.36

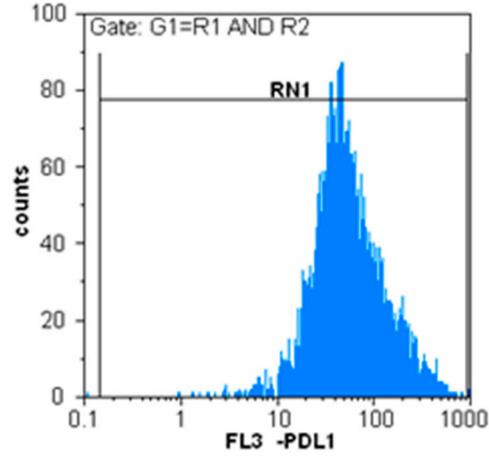


Figure S8. Flow cytometric analysis of PD-L1 expression on HT-29 cancer cells treated with KA39 at IC₅₀ concentration (μM) for 72 h.

Table S8. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	6746	6746	-	64.78	7.51	62.31	13.81	39.76
RN1	G1	10280	3698	-	99.76	80.82	108.17	-	-
R2	<None>	3752	3752	-	36.03	0.71	513.16	0.25	8.38

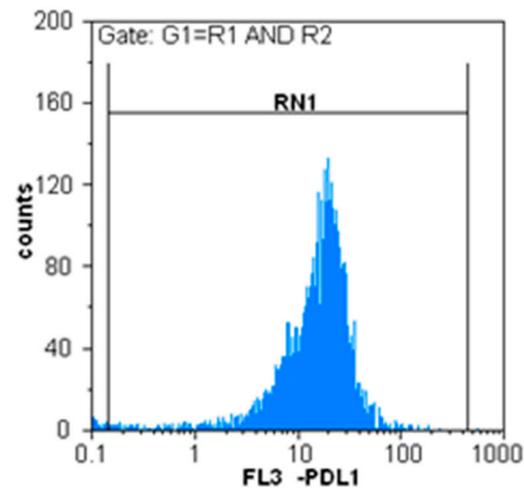


Figure S9. Flow cytometric analysis of PD-L1 expression on untreated HT-29 cancer cells cultured for 72 h.

Table S9. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	10905	10905	-	53.67	3.14	51.40	16.50	35.98
RN1	G1	19242	4057	-	95.28	19.40	75.77	-	-
R2	<None>	4516	4516	-	22.22	0.93	361.94	0.27	8.62

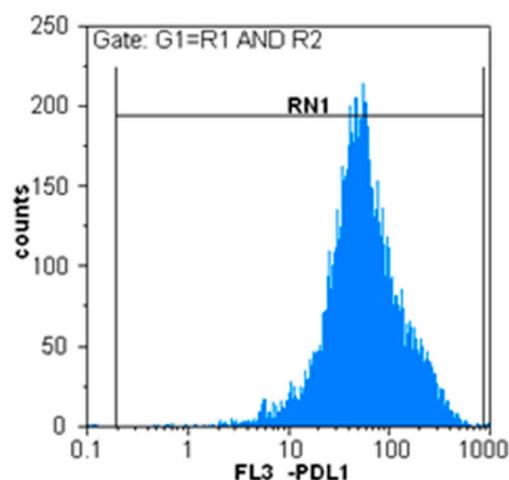


Figure S10. Flow cytometric analysis of PD-L1 expression on HT-29 cancer cells treated with KA39 at TGI concentration (μM) for 72 h.

Table S10. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	15069	15069	-	59.69	6.62	51.50	13.82	35.37
RN1	G1	24557	9083	-	99.67	78.56	95.78	-	-
R2	<None>	9409	9409	-	37.27	0.38	507.46	0.25	8.12

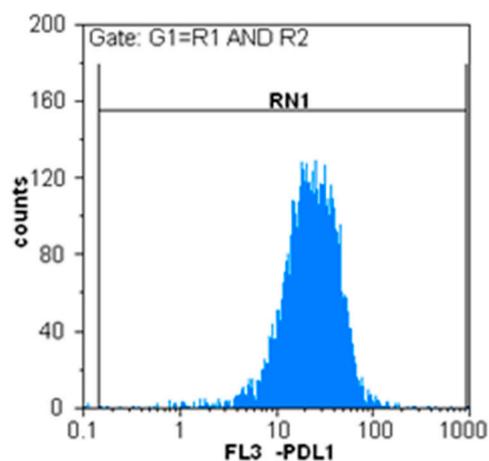


Figure S11. Flow cytometric analysis of PD-L1 expression on untreated DU-145 cancer cells cultured for 48 h.

Table S11. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	6999	6999	-	68.60	3.68	55.48	18.82	36.42
RN1	G1	9957	5387	-	99.57	27.99	79.41	-	-
R2	<None>	5591	5591	-	54.80	1.08	403.67	0.26	10.67

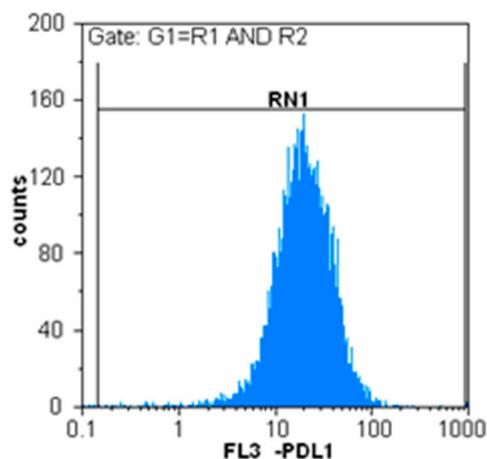


Figure S12. Flow cytometric analysis of PD-L1 expression on DU-145 cancer cells treated with KA39 at IC₅₀ concentration (µM) for 48 h.

Table S12. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	7513	7513	-	71.72	3.43	49.02	18.89	32.36
RN1	G1	10201	6007	-	99.29	24.48	77.58	-	-
R2	<None>	6175	6175	-	58.95	0.75	419.37	0.26	9.44

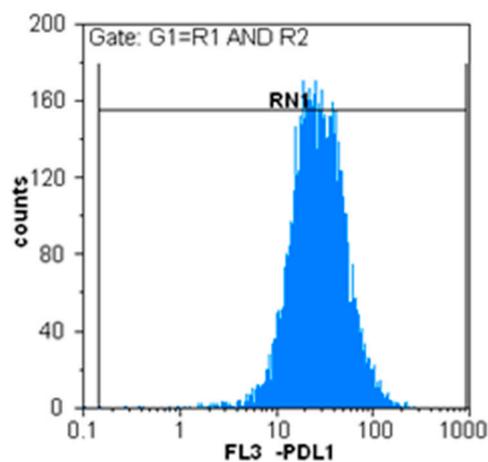


Figure S13. Flow cytometric analysis of PD-L1 expression on DU-145 cancer cells treated with KA39 at IC₅₀ concentration (µM) for 72 h.

Table S13. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	8257	8257	-	75.15	5.15	55.46	21.17	35.94
RN1	G1	10802	7069	-	99.80	33.55	69.66	-	-
R2	<None>	7199	7199	-	65.52	1.50	387.07	0.27	11.54

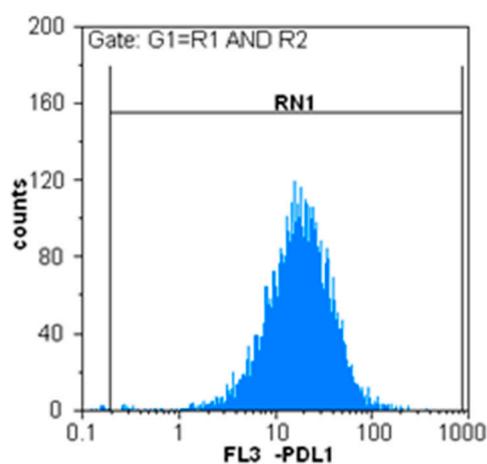


Figure S14. Flow cytometric analysis of PD-L1 expression on untreated DU-145 cancer cells cultured for 72 h.

Table S14. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	6907	6907	-	34.03	3.59	61.76	16.72	39.44
RN1	G1	19285	5088	-	99.10	23.46	86.03	-	-
R2	<None>	5726	5726	-	28.21	0.74	444.21	0.26	9.41

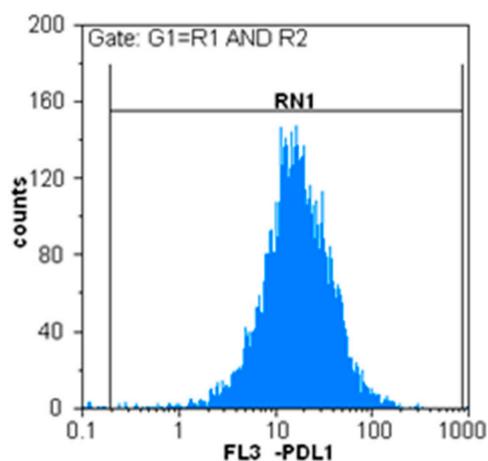


Figure S15. Flow cytometric analysis of PD-L1 expression on DU-145 cancer cells treated with KA39 at TGI concentration (μM) for 72 h.

Table S15. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	8572	8572	-	42.07	3.65	65.18	18.34	41.79
RN1	G1	19546	6736	-	99.44	23.02	94.70	-	-
R2	<None>	7311	7311	-	35.88	1.18	405.80	0.26	9.86

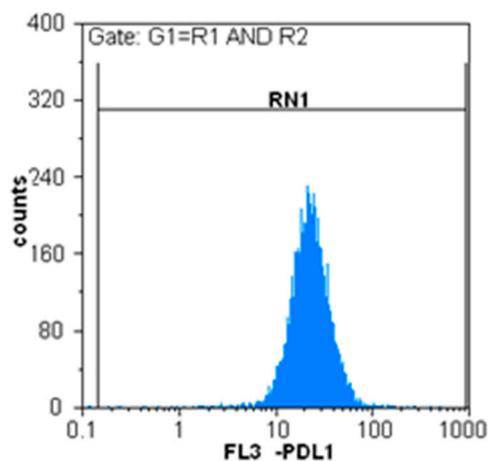


Figure S16. Flow cytometric analysis of PD-L1 expression on untreated PC-3 cancer cells cultured for 48 h.

Table S16. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	8410	8410	-	81.44	3.17	55.75	16.59	32.58
RN1	G1	10263	6134	-	99.89	25.91	64.58	-	-
R2	<None>	6212	6212	-	60.16	0.35	521.31	0.25	6.99

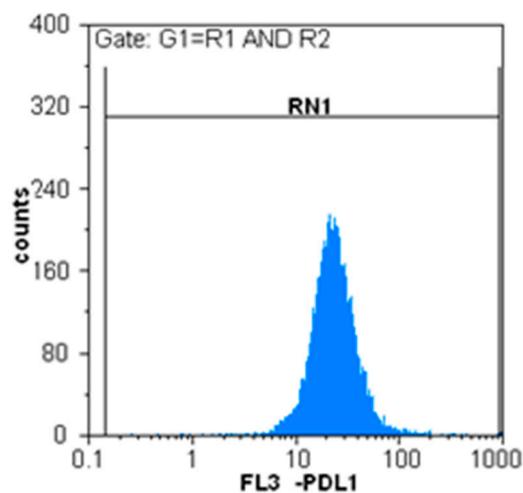


Figure S17. Flow cytometric analysis of PD-L1 expression on PC-3 cancer cells treated with KA39 at IC₅₀ concentration (μ M) for 48 h.

Table S17. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	8404	8404	-	80.95	4.19	61.38	17.63	34.04
RN1	G1	10257	6087	-	99.90	27.66	83.06	-	-
R2	<None>	6169	6169	-	59.42	0.42	444.45	0.25	7.22

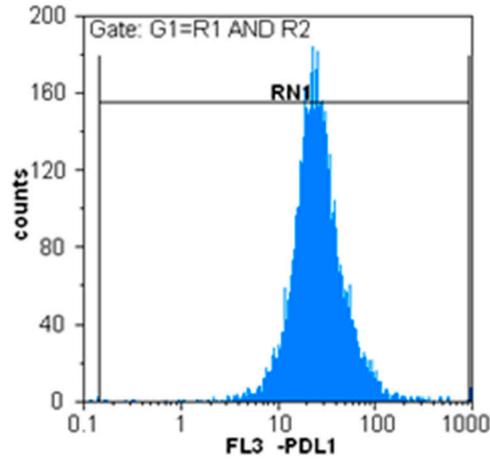


Figure S18. Flow cytometric analysis of PD-L1 expression on PC-3 cancer cells treated with KA39 at IC₅₀ concentration (µM) for 72 h.

Table S18. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	8381	8381	-	80.02	6.37	53.97	16.00	36.76
RN1	G1	10305	5663	-	99.68	32.98	108.90	-	-
R2	<None>	5751	5751	-	54.91	0.34	475.01	0.25	7.03

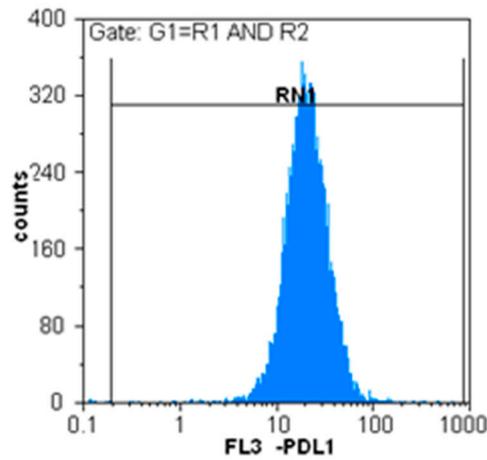


Figure S19. Flow cytometric analysis of PD-L1 expression on untreated PC-3 cancer cells cultured for 72 h.

Table S19. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	14475	14475	-	61.79	3.23	47.03	17.16	27.35
RN1	G1	22965	10842	-	99.83	23.93	69.66	-	-
R2	<None>	11321	11321	-	48.33	0.33	460.49	0.25	6.47

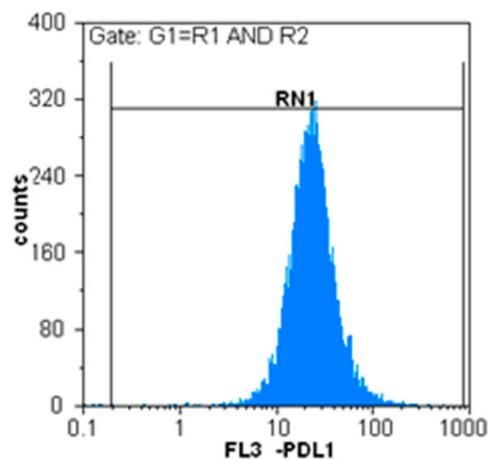


Figure S20. Flow cytometric analysis of PD-L1 expression on PC-3 cancer cells treated with KA39 at TGI concentration (μM) for 72 h.

Table S20. Region analysis – calculations as resulted from the CyFlow® Partec Report software.

Region	Gate	Ungated	Count	Count/mL	%Gated	Mean-x	CV-x%	Mean-y	CV-y%
R1	<None>	12827	12827	-	61.02	5.46	51.65	17.35	33.49
RN1	G1	20598	9903	-	99.84	28.21	74.87	-	-
R2	<None>	10297	10297	-	48.98	0.36	469.83	0.26	7.40

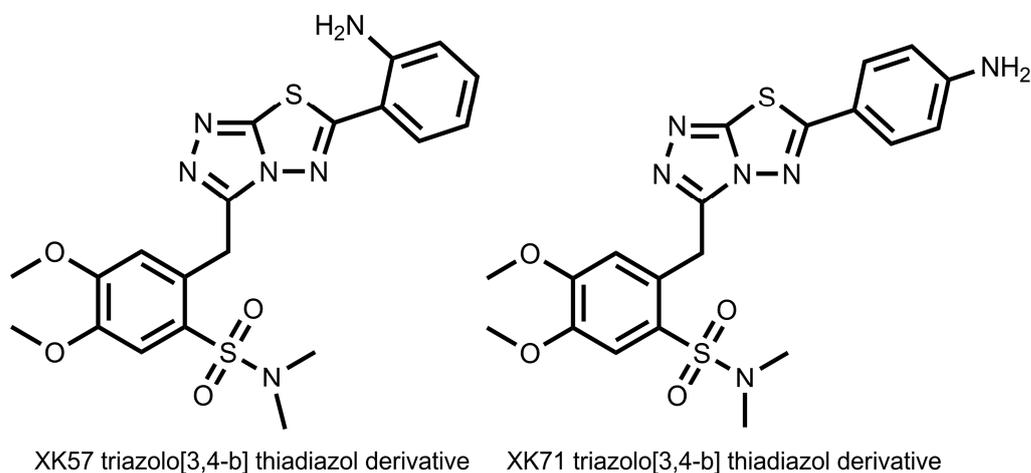


Figure S21. Chemical structures of XK 57: 2-((6-(2-aminophenyl)-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol-3-yl)methyl)-4,5-dimethoxy-*N,N*-dimethylbenzenesulfonamide and XK 71: 2-((6-(4-aminophenyl)-[1,2,4]triazolo[3,4-b][1,3,4]thiadiazol-3-yl)methyl)-4,5-dimethoxy-*N,N*-dimethylbenzenesulfonamide.

Table S21. Cytostatic (GI₅₀, TGI) and cytotoxic effects (IC₅₀) induced by XK57 and XK71 on the tested human cancer cell lines, DLD-1, HT-29, LS174T, DU-145 and PC-3.

Cancer Cell Lines	XK57 GI ₅₀	XK57 TGI	XK57 IC ₅₀	XK71 GI ₅₀	XK71 TGI	XK71 IC ₅₀
DLD-1	90 μM	>100 μM	>100 μM	>100 μM	>100 μM	>100 μM
HT-29	>100 μM	>100 μM	>100 μM	>100 μM	>100 μM	>100 μM
LS174T	>100 μM	>100 μM	>100 μM	>100 μM	>100 μM	>100 μM
DU-145	>100 μM	>100 μM	>100 μM	>100 μM	>100 μM	>100 μM
PC-3	>100 μM	>100 μM	>100 μM	>100 μM	>100 μM	>100 μM

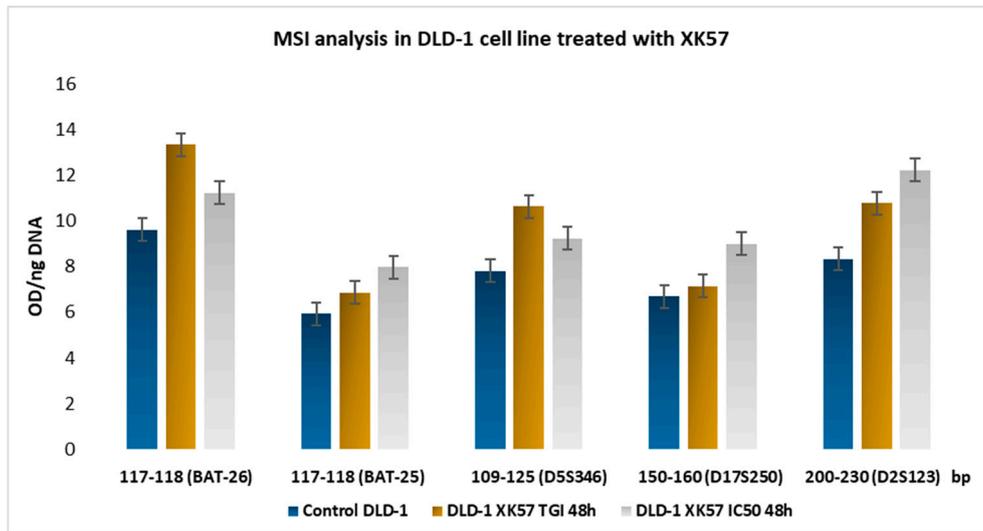
Table S22. The absolute values of PD-L1 expression as defined by flow cytometric analysis in untreated (control) and treated cells with XK57 derivative at IC₅₀ concentration (μM) for 48 and 72 h, as well as at TGI concentration (μM) for 72 h. The units of measurements are expressed as mean of immuno-fluorescence intensity counted per cell.

Cancer cell lines	PD-L1 Expression				
	Control 48 h	XK57 IC ₅₀ 48 h	XK57 IC ₅₀ 72 h	Control 72 h	XK57 TGI 72 h
DLD-1	16.89 ± 0.84	19.88 ± 0.99	15.77 ± 0.94	25.09 ± 1.25	26.54 ± 1.59
HT-29	23.34 ± 1.4	24.79 ± 1.98	26.34 ± 1.31	21.23 ± 2.12	23.19 ± 1.62
DU-145	28.89 ± 2.02	27.19 ± 2.17	32.09 ± 2.24	24.56 ± 1.96	25.18 ± 1.25
PC-3	27.14 ± 2.44	28.44 ± 1.42	29.08 ± 2.03	27.88 ± 1.39	29.67 ± 2.07

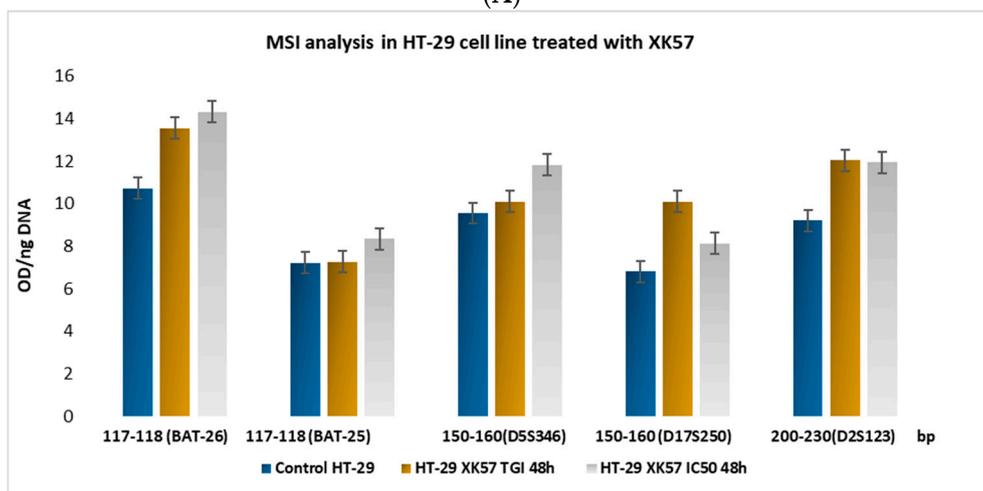
Table S23. The absolute values of PD-L1 expression as defined by flow cytometric analysis in untreated (control) and treated cells with XK71 derivative at IC₅₀ concentration (μM) for 48 and 72 h, as well as at TGI concentration (μM) for 72 h. The units of measurements are expressed as mean of immuno-fluorescence intensity counted per cell.

Cancer cell lines	PD-L1 expression				
	Control 48 h	XK71 IC ₅₀ 48 h	XK71 IC ₅₀ 72 h	Control 72 h	XK71 TGI 72 h
DLD-1	14.09 ± 0.7	17.45 ± 1.04	13.43 ± 0.67	22.81 ± 1.14	21.19 ± 1.05
HT-29	22.78 ± 1.36	21.31 ± 1.06	27.32 ± 1.63	20.59 ± 1.23	24.64 ± 1.47
DU-145	25.35 ± 1.26	27.53 ± 1.65	28.77 ± 2.01	26.89 ± 2.15	25.04 ± 1.75
PC-3	28.19 ± 1.97	30.98 ± 2.78	26.19 ± 1.83	25.12 ± 1.75	27.23 ± 2.17

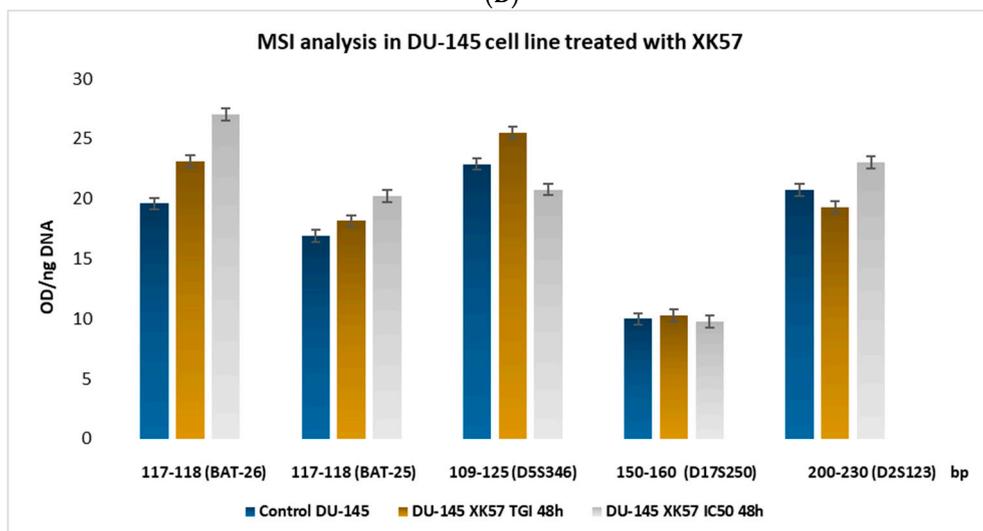
MSI analysis in DLD-1, HT-29, DU-145, PC-3 and LS174T cancer cells treated with XK71 and XK57 derivatives



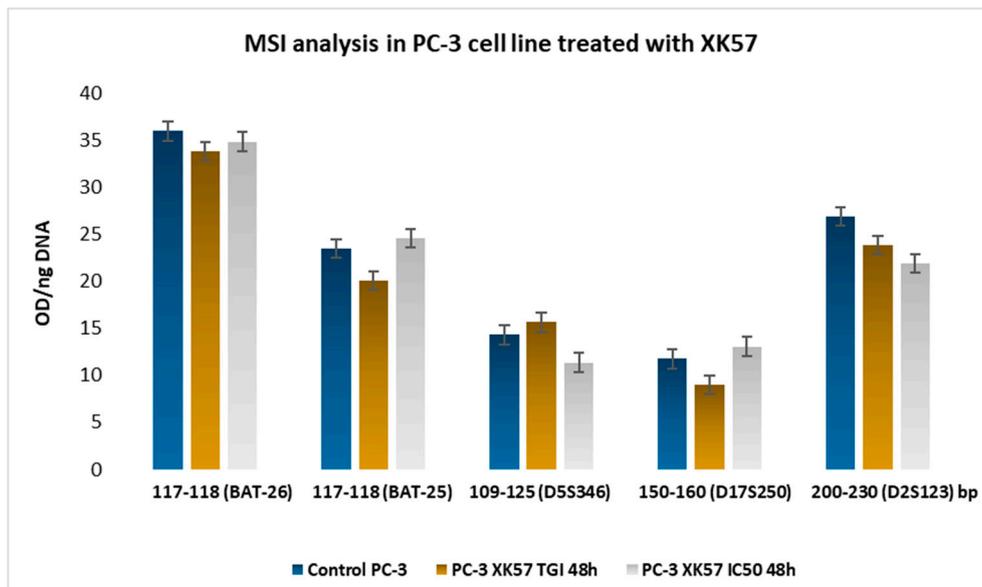
(A)



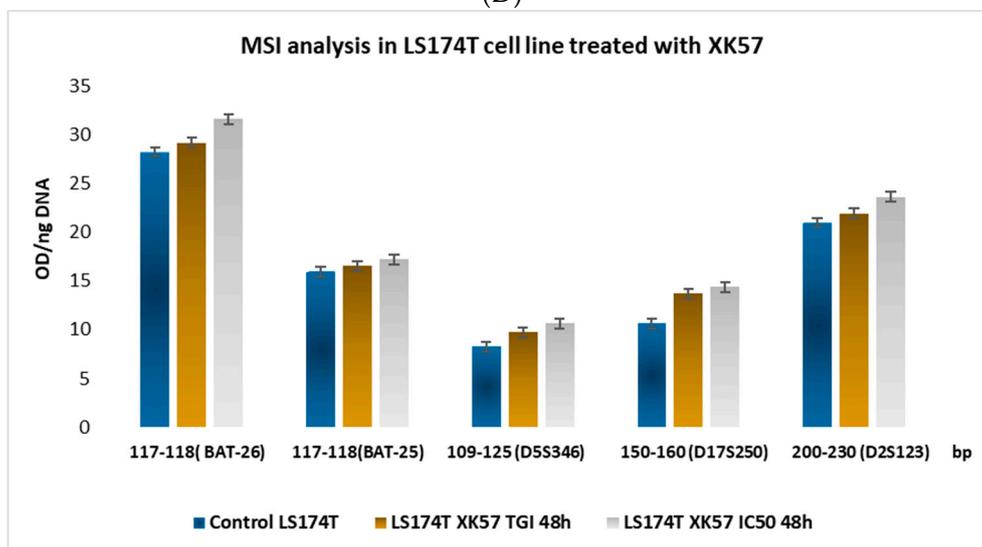
(B)



(C)

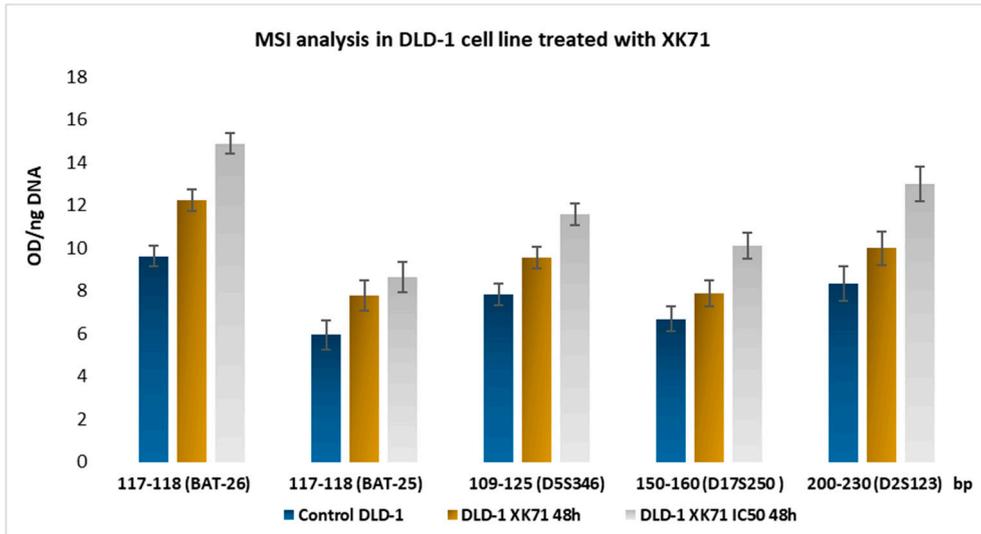


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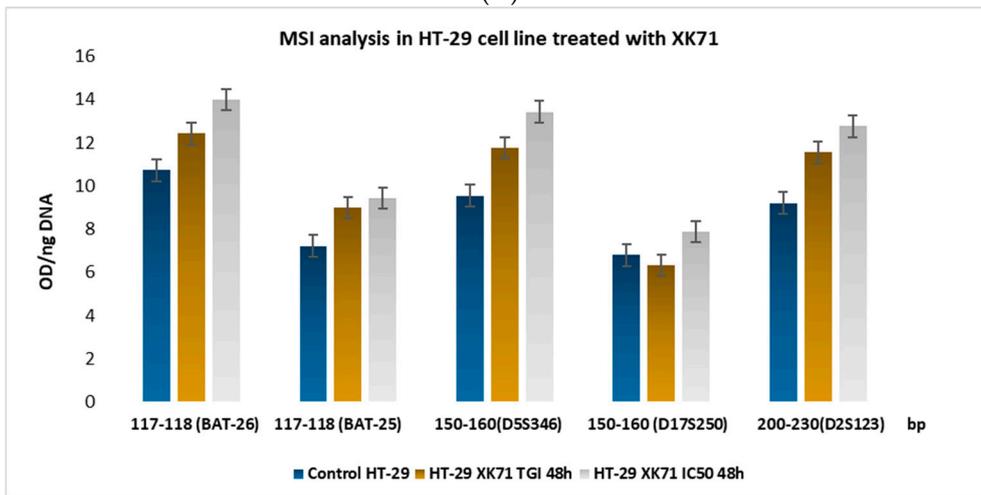


(E)

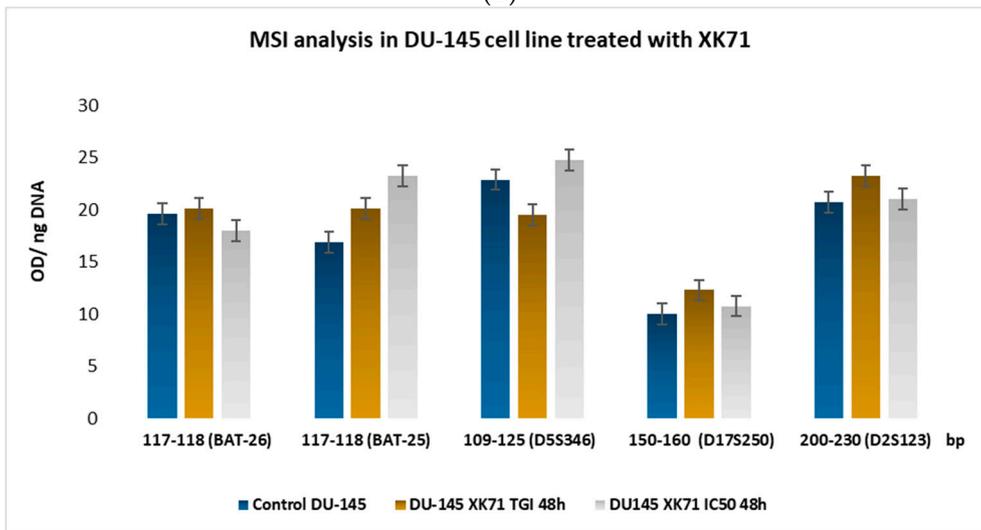
Figure S22. MSI alterations (mean \pm SEM) induced by XK57 triazolo[3,4-b]thiadiazole derivative (per ng DNA) in five human cancer cell lines. (A), (B), (C), (D) and (E) demonstrate the MSI alterations in DLD-1, HT-29, DU-145, PC-3, and LS174T cancer cells, treated with XK57 at the TGI concentration and IC₅₀ (μ M) for 48h, respectively. Quantitative MSI fragments analysis conducted in Bethesda panel (BAT-26, BAT-25, D5S346, D17S250 and D2S123). Student t-test; $p > 0.01$; no statistical significance level.



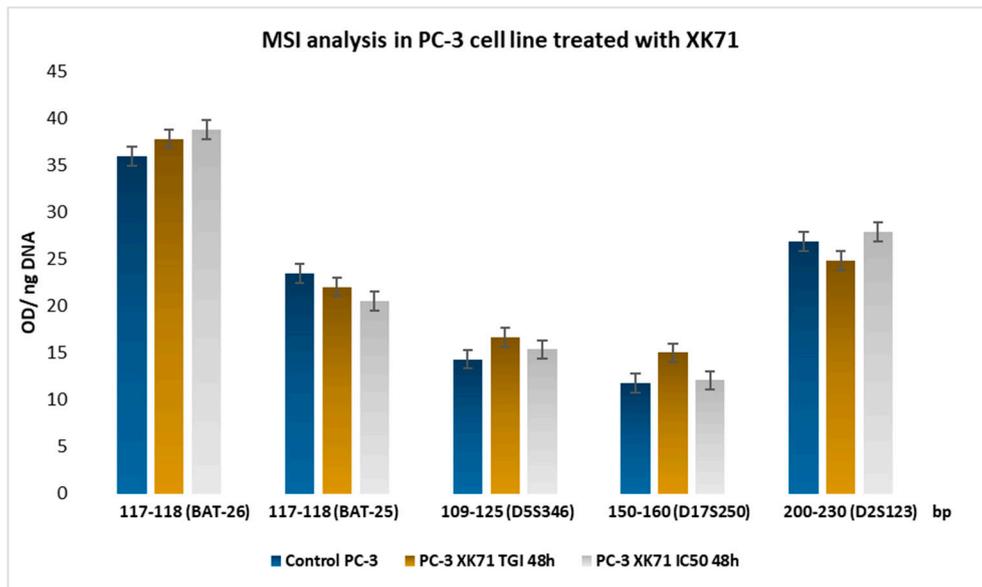
(A)



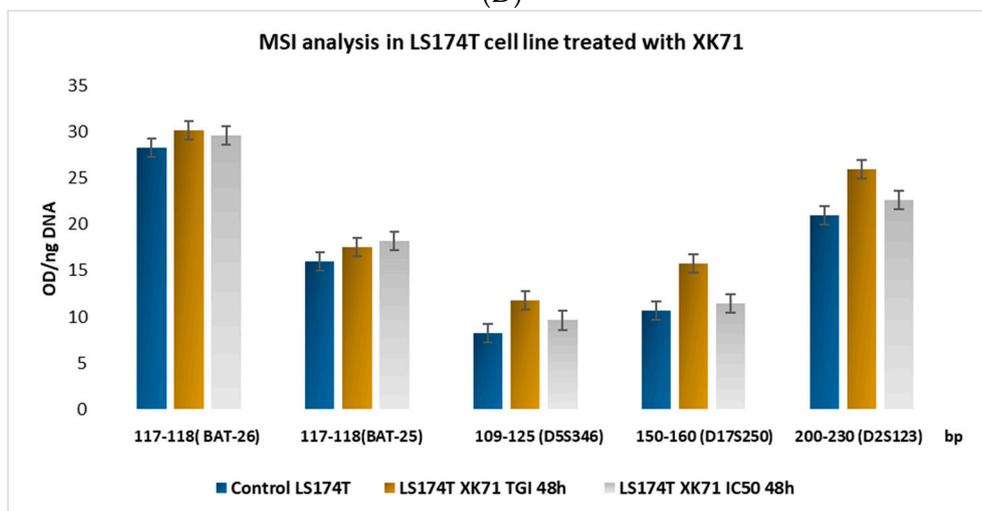
(B)



(C)



(D)



(E)

Figure S23. MSI alterations (mean \pm SEM) induced by XK71 triazolo[3,4-*b*]thiadiazole derivative (per ng DNA) in five human cancer cell lines. (A), (B), (C), (D) and (E) show the MSI alterations in DLD-1, HT-29, DU-145, PC-3, and LS174T cancer cells, treated with XK71 at the TGI concentration and IC₅₀ (μ M) for 48h, respectively. Quantitative MSI fragments analysis conducted in Bethesda panel (BAT-26, BAT-25, D5S346, D17S250 and D2S123). Student t-test; $p > 0.01$; no statistical significance level.