

Supplementary Data

CNS Cancer		
Drug	Indication	Mechanism of Action
Temozolomide	glioblastoma concomitantly with radiotherapy and refractory anaplastic astrocytoma	DNA and RNA Alkylation
Lomustine	primary and metastatic brain tumors	DNA and RNA Alkylation
Carmustine	brain tumors glioblastoma, brainstem glioma, medulloblastoma, astrocytoma, ependymoma, and metastatic brain tumors	DNA and RNA alkylation; Enzymatic Carbamoylation
Everolimus	subependymal giant cell astrocytoma (SEGA) that requires therapeutic intervention	mTOR inhibitor; HIF inhibitor

Prostate Cancer		
Drug	Indication	Mechanism of Action
Rucaparib	metastatic Castration-Resistant Prostate Cancer (mCRPC)	PARP Inhibitor
Enzalutamide	mCRPC and metastatic Castration-Sensitive Prostate Cancer (mCSPC)	Androgen Receptor Inhibitor
Abiraterone Acetate	mCRPC and mCSPC	CYP17 Inhibitor
Cabazitaxel	mCRPC	Antimicrotubular Antineoplastic Agent
Docetaxel	mCRPC	Antimicrotubular Antineoplastic Agent
Leuprolide Acetate	palliative treatment of advanced prostate cancer	GnRH Agonist
Mitoxantrone	advanced prostate cancer not responding to hormone treatment	Topoisomerase Inhibitor
Olaparib	mCRPC	PARP Inhibitor

Non-Small Cell Lung Cancer		
Drug	Indication	Mechanism of Action
Alectinib	ALK-positive metastatic NSCLC	ALK and RET Inhibitor
Pemetrexed	locally advanced or metastatic, non-squamous NSCLC	Antimetabolite
Carboplatin	chemotherapeutic treatment for NSCLC	Alkylating Agent
Crizotinib	ALK or ROS1-positive metastatic NSCLC	ALK and ROS1 Inhibitor
Docetaxel	locally advanced or metastatic NSCLC after failure of prior platinum-based chemotherapy	Taxoid
Doxorubicin	chemotherapeutic treatment for NSCLC	Anthracycline Antibiotic
Erlotinib	EGFR gene mutated metastatic NSCLC	EGFR Tyrosine Kinase Inhibitor
Everolimus	unresectable, locally advanced, or metastatic NSCLC	mTOR Kinase Inhibitor
Gefitinib	EGFR gene mutated metastatic NSCLC; in combination with Everolimus	EGFR Tyrosine Kinase Inhibitor
Lorlatinib	ALK-positive metastatic NSCLC	ALK and ROS1 Inhibitor
Mechlorethamine	NSCLC	Alkylating Agent
Trametinib	in combination with dabrafenib for BRAF V600E mutated metastatic NSCLC	MEK Kinase Inhibitor
Methotrexate	NSCLC	Folate Analog Metabolic Inhibitor
Paclitaxel	chemotherapeutic treatment for NSCLC	Taxane; Plant Alkaloid
Vinorelbine	alone or in combination with Cisplatin for locally advanced or metastatic NSCLC	Plant Alkaloid

Table S1 The indication and mechanism of FDA-approved anticancer drugs for (A)CNS cancer, (B) prostate cancer (C) NSCLC. Literature review of indication, mechanism of FDA-approved CNS, prostate, NSCLC cancers.

A

	Cell Lines	NMI	Temozolomide	Lomustine	Carmustine	Everolimus	Max	Min	Max-Min
GI50 (μM) Min-max normalization	SF-268	1.820	100.000	20.900	31.600	0.269	100.000	0.269	99.731
		0.016	1.000	0.207	0.314	0.000			
		e.g. Min-max normalization of GI50 of NMI: $(1.820 - 0.269) / (99.731) = 0.016$							
GI50 (μM) Min-max normalization	SF-295	1.910	100.000	91.200	39.800	0.010	100.000	0.010	99.990
		0.019	1.000	0.912	0.398	0.000			
		(GI50 - Min) / (Max - Min)							
GI50 (μM) Min-max normalization	SF-539	1.200	100.000	16.600	31.600	0.010	100.000	0.010	99.990
		0.012	1.000	0.166	0.316	0.000			
		(GI50 - Min) / (Max - Min)							
GI50 (μM) Min-max normalization	SNB-19	1.120	100.000	100.000	63.100	0.257	100.000	0.257	99.743
		0.009	1.000	1.000	0.630	0.000			
		(GI50 - Min) / (Max - Min)							
GI50 (μM) Min-max normalization	SNB-75	1.290	63.000	23.400	50.100	0.010	63.000	0.010	62.990
		0.020	1.000	0.371	0.795	0.000			
		(GI50 - Min) / (Max - Min)							
GI50 (μM) Min-max normalization	U251	0.740	100.000	15.800	31.600	0.011	100.000	0.011	99.989
		0.007	1.000	0.158	0.316	0.000			
		(GI50 - Min) / (Max - Min)							
TGI (μM) Min-max normalization	SF-268	6.760	100.000	57.500	100.000	20.900	100.000	6.760	93.240
		0.000	1.000	0.544	1.000	0.152			
		(TGI - Min) / (Max - Min)							
TGI (μM) Min-max normalization	SF-295	4.070	100.000	100.000	251.000	17.300	251.000	4.070	246.930
		0.000	0.388	0.388	1.000	0.054			
		(TGI - Min) / (Max - Min)							
TGI (μM) Min-max normalization	SF-539	2.690	100.000	33.900	100.000	17.300	100.000	2.690	97.310
		0.000	1.000	0.321	1.000	0.150			
		(TGI - Min) / (Max - Min)							
TGI (μM) Min-max normalization	SNB-19	3.630	100.000	100.000	251.000	19.500	251.000	3.630	247.370
		0.000	0.390	0.390	1.000	0.064			
		(TGI - Min) / (Max - Min)							
TGI (μM) Min-max normalization	SNB-75	3.890	100.000	64.600	126.000	14.100	126.000	3.890	122.110
		0.000	0.787	0.497	1.000	0.084			
		(TGI - Min) / (Max - Min)							
TGI (μM) Min-max normalization	U251	2.090	100.000	51.300	63.100	18.200	100.000	2.090	97.910
		0.000	1.000	0.503	0.623	0.165			
		(TGI - Min) / (Max - Min)							
LC50 (μM) Min-max normalization	SF-268	100.000	100.000	100.000	251.000	53.700	251.000	53.700	197.300
		0.235	0.235	0.235	1.000	0.000			
		(LC50 - Min) / (Max - Min)							
LC50 (μM) Min-max normalization	SF-295	8.710	100.000	100.000	251.000	63.100	251.000	8.710	242.290
		0.000	0.377	0.377	1.000	0.224			
		(LC50 - Min) / (Max - Min)							
LC50 (μM) Min-max normalization	SF-539	6.030	100.000	70.800	251.000	41.700	251.000	6.030	244.970
		0.000	0.384	0.264	1.000	0.146			
		(LC50 - Min) / (Max - Min)							
LC50 (μM) Min-max normalization	SNB-19	28.200	100.000	100.000	251.000	44.700	251.000	28.200	222.800
		0.000	0.322	0.322	1.000	0.074			
		(LC50 - Min) / (Max - Min)							
LC50 (μM) Min-max normalization	SNB-75	13.800	100.000	100.000	251.000	38.900	251.000	13.800	237.200
		0.000	0.363	0.363	1.000	0.106			
		(LC50 - Min) / (Max - Min)							
LC50 (μM) Min-max normalization	U251	5.010	100.000	100.000	126.000	42.700	126.000	5.010	120.990
		0.000	0.785	0.785	1.000	0.312			
		(LC50 - Min) / (Max - Min)							
Cumulative Score	SF-268	0.250	2.235	0.986	2.314	0.152			
		0.019	1.765	1.677	2.398	0.278			
		0.012	2.384	0.751	2.316	0.296			
		0.009	1.712	1.712	2.630	0.138			
		0.020	2.150	1.232	2.795	0.189			
		0.007	2.785	1.446	1.939	0.476			
e.g. Cumulative score of min-max normalization of GI50, TGI and LC50 of NMI: $0.016 + 0.000 + 0.235 = 0.250$									

B

	Cell Lines	NMI	Rucaparib	Enzalutamide	Abriraterone Acetate	Cabazitaxel	Doxetaxel	Leuprolide Acetate	Mitoxantrone	Olaparib	Max	Min	Max-Min
GI50 (μM) Min-max normalization	PC-3	1.700	23.400	20.000	3.980	0.010	0.010	100.000	0.141	51.288	100.000	0.010	99.990
		0.017	0.234	0.200	0.040	0.000	0.000	1.000	0.001	0.513			(GI50 - Min) / (Max - Min)
		e.g. Min-max normalization of GI50 of NMI: $(1.700 - 0.010) / (99.990 - 0.017) = 0.017$											
GI50 (μM) Min-max normalization	DU-145	3.020	18.200	31.800	20.000	0.010	0.010	100.000	0.018	100.000	100.000	0.010	99.990
		0.030	0.182	0.316	0.200	0.000	0.000	1.000	0.000	1.000			(GI50 - Min) / (Max - Min)
		e.g. Min-max normalization of GI50 of NMI: $(3.020 - 0.030) / (99.990 - 0.030) = 0.030$											
TGI (μM) Min-max normalization	PC-3	3.980	100.000	100.000	100.000	79.400	100.000	100.000	1.445	100.000	100.000	1.445	98.555
		0.026	1.000	1.000	1.000	0.791	1.000	1.000	0.000	1.000			(TGI - Min) / (Max - Min)
		e.g. Min-max normalization of TGI of NMI: $(3.980 - 0.026) / (98.555 - 0.026) = 0.026$											
TGI (μM) Min-max normalization	DU-145	11.000	36.300	100.000	100.000	0.010	0.010	77.625	100.000	0.468	100.000	0.010	99.990
		0.110	0.363	1.000	1.000	0.000	0.000	0.776	1.000	0.005	1.000		(TGI - Min) / (Max - Min)
		e.g. Min-max normalization of TGI of NMI: $(11.000 - 0.110) / (99.990 - 0.110) = 0.110$											
LC50 (μM) Min-max normalization	PC-3	9.550	100.000	100.000	100.000	100.000	100.000	100.000	10.471	100.000	100.000	9.550	90.450
		0.000	1.000	1.000	1.000	1.000	1.000	1.000	0.010	1.000			(LC50 - Min) / (Max - Min)
		e.g. Min-max normalization of LC50 of NMI: $(9.550 - 0.000) / (90.450 - 0.000) = 0.099$											
LC50 (μM) Min-max normalization	DU-145	34.700	72.400	100.000	100.000	100.000	100.000	100.000	11.482	100.000	100.000	11.482	88.518
		0.262	0.688	1.000	1.000	1.000	1.000	1.000	0.000	1.000			(LC50 - Min) / (Max - Min)
		e.g. Min-max normalization of LC50 of NMI: $(34.700 - 0.262) / (88.518 - 0.262) = 0.262$											
Cumulative Score	PC-3	0.043	2.234	2.200	2.040	1.791	2.000	3.000	0.011	2.513			
		0.402	1.233	2.318	2.200	1.000	1.776	3.000	0.005	3.000			GI50+TGI+LC50 Min-max normalization

(Continued)

(Continued)

C

	Cell Line #	N MI	Brklnb	Trm + Brkb	Ale + Brkb	Pmt + Brkb	Carboxyln	Crotnb	Oxvite	Oxvite	Everbmls	G + Brkb	Lotslnb	Mchbrms	Mchbrms	Pctbrms	Mntrbrms	Max.	MI	Max-MI	
	A545/A/TCC	3.220 0.032	9.120 0.091	0.030 0.000	0.013 0.006	1.350 0.590	0.590 1.000	1.170 0.012	0.010 0.000	0.063 0.001	0.010 0.000	7.590 0.076	25.119 0.251	2.818 0.028	0.010 0.000	0.011 0.000	100.000 0.000	0.019 0.000	0.010 0.000	99.990 99.990	
		e.g. MI=MI/(MI+MI) = 0.220/(0.220+0.106)=0.632																			
G50 (µM)		2.420 0.024	0.191 0.002	0.013 0.000	1.550 0.015	1.000 1.000	5.130 0.051	3.720 0.037	0.420 0.004	0.010 0.000	0.045 0.000	16.218 0.162	6.026 0.060	10.000 0.100	0.977 0.010	0.010 0.000	100.000 0.000	0.010 0.000	0.010 0.000	99.990 99.990	
G50 (µM)	EKVX	2.100 0.021	15.100 0.151	0.251 0.002	0.850 0.008	0.720 0.007	100.000 1.000	1.910 0.010	0.010 0.000	0.068 0.001	0.010 0.000	10.200 0.102	22.909 0.229	5.248 0.052	0.016 0.000	0.031 0.000	100.000 0.000	0.015 0.000	0.010 0.000	99.990 99.990	
G50 (µM)	HOP-42	1.570 0.016	4.210 0.014	0.010 0.000	0.390 0.004	100.000 1.000	0.560 0.050	9.120 0.091	0.100 0.001	0.020 0.000	0.100 0.000	7.410 0.074	2.512 0.025	0.312 0.026	0.743 0.079	0.525 0.070	0.047 0.005	100.000 0.000	0.010 0.000	0.010 0.000	99.990 99.990
G50 (µM)	HOP-92	1.570 0.016	4.210 0.014	0.010 0.000	0.390 0.004	100.000 1.000	0.560 0.050	9.120 0.091	0.100 0.001	0.020 0.000	0.100 0.000	7.410 0.074	2.512 0.025	0.312 0.026	0.743 0.079	0.525 0.070	0.047 0.005	100.000 0.000	0.010 0.000	0.010 0.000	99.990 99.990
G50 (µM)	NCI-H226	1.120 0.017	42.000 0.427	4.370 0.043	1.740 0.017	100.000 1.000	1.820 0.018	0.360 0.003	0.050 0.000	0.350 0.003	0.150 0.000	40.738 0.407	5.318 0.083	2.344 0.023	1.047 0.010	0.030 0.000	100.000 0.000	0.010 0.000	0.010 0.000	99.970 99.990	
G50 (µM)	NCI-H226	1.120 0.017	42.000 0.427	4.370 0.043	1.740 0.017	100.000 1.000	1.820 0.018	0.360 0.003	0.050 0.000	0.350 0.003	0.150 0.000	40.738 0.407	5.318 0.083	2.344 0.023	1.047 0.010	0.030 0.000	100.000 0.000	0.010 0.000	0.010 0.000	99.970 99.990	
G50 (µM)	NCI-H22	1.940 0.019	26.200 0.282	2.820 0.008	100.000 1.000	2.950 0.021	0.010 0.000	0.150 0.001	0.320 0.003	0.170 0.000	0.084 0.000	59.884 0.589	1.318 0.013	0.087 0.001	0.012 0.000	0.019 0.000	100.000 0.000	0.010 0.000	0.010 0.000	99.990 99.990	
G50 (µM)	NCI-H22 M	3.010 0.021	0.998 0.001	0.012 0.000	2.340 0.023	100.000 1.000	5.230 0.052	0.010 0.000	0.040 0.005	0.220 0.005	0.085 0.002	33.113 0.331	16.849 0.331	0.427 0.021	0.009 0.000	0.010 0.000	100.000 0.000	0.020 0.000	0.010 0.000	99.990 99.990	
G50 (µM)	NCI-H459	1.770 0.018	5.890 0.059	3.160 0.015	1.480 0.003	100.000 1.000	0.890 0.000	0.010 0.000	0.017 0.000	0.150 0.000	6.610 0.060	28.184 0.282	0.129 0.026	0.006 0.001	0.010 0.000	100.000 0.000	0.010 0.000	0.010 0.000	99.995 99.995		
G50 (µM)	NCI-H22	1.160 0.016	9.955 3.800	1.170 0.110	18.200 18.200	100.000 100.000	1.510 1.510	0.010 0.010	0.029 0.029	0.160 0.160	6.760 5.744	0.871 0.871	1.445 1.445	0.004 0.004	0.023 0.023	100.000 0.000	0.010 0.000	0.010 0.000	99.996 99.996		
TG1 (µM)		13.200 0.129	100.000 1.000	43.700 0.435	3.990 0.028	100.000 1.000	11.200 0.010	81.300 0.010	0.290 0.000	22.400 0.000	21.950 0.000	100.000 100.000	32.359 32.359	10.000 10.000	70.795 0.000	33.884 0.000	100.000 0.000	0.250 0.020	99.710 94.500		
TG1 (µM)	A545/A/TCC	12.300 0.077	100.000 1.000	7.000 0.691	6.920 0.015	100.000 1.000	18.200 0.014	0.500 0.000	5.500 0.000	21.400 0.000	17.400 0.000	100.000 100.000	26.303 26.303	10.000 10.000	85.114 0.000	18.621 0.000	100.000 0.000	5.500 5.500	94.500 94.500		
TG1 (µM)	EKVX	12.300 0.077	100.000 1.000	7.000 0.691	6.920 0.015	100.000 1.000	18.200 0.014	0.500 0.000	5.500 0.000	21.400 0.000	17.400 0.000	100.000 100.000	26.303 26.303	10.000 10.000	85.114 0.000	18.621 0.000	100.000 0.000	5.500 5.500	94.500 94.500		
TG1 (µM)	HOP-42	5.120 0.038	56.200 0.563	5.100 0.006	1.000 1.000	1.000 0.000	0.050 0.000	0.018 0.000	0.208 0.000	0.342 0.000	1.000 0.000	0.020 0.000	0.020 0.000	0.020 0.000	0.036 0.036	0.230 0.230	100.000 0.000	0.010 0.000	0.010 0.000	99.990 99.990	
TG1 (µM)	HOP-92	5.120 0.038	56.200 0.563	5.100 0.006	1.000 1.000	1.000 0.000	0.050 0.000	0.018 0.000	0.208 0.000	0.342 0.000	1.000 0.000	0.020 0.000	0.020 0.000	0.020 0.000	0.036 0.036	0.230 0.230	100.000 0.000	0.010 0.000	0.010 0.000	99.990 99.990	
TG1 (µM)	NCI-H22	4.110 0.039	47.900 1.000	4.370 0.000	1.000 1.000	1.000 0.000	7.240 0.000	6.300 0.000	0.250 0.000	23.400 0.000	33.100 0.000	100.000 100.000	39.811 39.811	10.000 10.000	19.955 0.000	36.308 0.000	100.000 0.000	0.250 0.020	99.750 93.050		
TG1 (µM)	NCI-H22 M	5.090 0.042	19.500 1.034	38.900 0.046	1.000 0.000	100.000 1.000	16.500 0.014	7.600 0.014	0.010 0.000	0.630 0.000	0.000 0.000	0.232 0.022	1.000 0.000	0.397 0.000	0.008 0.000	0.189 0.000	0.361 0.000	100.000 0.000	7.244 6.122	41.887 37.722	100.000 99.990
TG1 (µM)	NCI-H22	5.090 0.042	19.500 1.034	38.900 0.046	1.000 0.000	100.000 1.000	16.500 0.014	7.600 0.014	0.010 0.000	0.630 0.000	0.000 0.000	0.232 0.022	1.000 0.000	0.397 0.000	0.008 0.000	0.189 0.000	0.361 0.000	100.000 0.000	7.244 6.122	41.887 37.722	100.000 99.990
TG1 (µM)	NCI-H459	5.010 0.046	91.200 0.912	93.300 0.933	4.070 0.036	100.000 1.000	5.630 0.000	40.700 0.000	1.250 0.000	19.500 0.000	20.000 0.000	100.000 100.000	27.544 27.544	10.000 10.000	40.738 0.000	0.447 0.000	100.000 0.000	0.447 0.000	0.447 0.000	99.553 99.553	
TG1 (µM)	NCI-H22	5.010 0.046	91.200 0.912	93.300 0.933	4.070 0.036	100.000 1.000	5.630 0.000	40.700 0.000	1.250 0.000	19.500 0.000	20.000 0.000	100.000 100.000	27.544 27.544	10.000 10.000	40.738 0.000	0.447 0.000	100.000 0.000	0.447 0.000	0.447 0.000	99.553 99.553	
TG1 (µM)	NCI-H22 M	5.010 0.046	91.200 0.912	93.300 0.933	4.070 0.036	100.000 1.000	5.630 0.000	40.700 0.000	1.250 0.000	19.500 0.000	20.000 0.000	100.000 100.000	27.544 27.544	10.000 10.000	40.738 0.000	0.447 0.000	100.000 0.000	0.447 0.000	0.447 0.000	99.553 99.553	
LC50 (µM)		39.900 0.033	75.900 2.240	91.200 0.024	10.000 0.000	100.000 1.000	46.800 0.000	10.000 0.000	0.374 0.000	1.021 0.000	0.421 0.000	0.834 0.000	0.459 0.000	0.813 0.000	0.000 0.000	1.000 1.000	0.590 0.590	0.590 0.590	0.590 0.590		
LC50 (µM)	HOP-42	3.328 0.017	7.372 1.000	5.002 0.000	1.000 1.000	100.000 1.000	4.048 0.000	0.239 0.000	0.928 0.000	0.654 0.000	0.519 0.000	0.723 0.000	1.000 1.000	0.000 0.000	0.038 0.000	0.841 0.000	0.841 0.000	0.841 0.000	0.841 0.000	0.841 0.000	
LC50 (µM)	HOP-92	3.328 0.017	7.372 1.000	5.002 0.000	1.000 1.000	100.000 1.000	4.048 0.000	0.239 0.000	0.928 0.000	0.654 0.000	0.519 0.000	0.723 0.000	1.000 1.000	0.000 0.000	0.038 0.000	0.841 0.000	0.841 0.000	0.841 0.000	0.841 0.000	0.841 0.000	
LC50 (µM)	NCI-H22	4.110 0.012	17.000 1.000	1.000 0.000	1.000 1.000	100.000 1.000	27.500 0.000	10.000 0.000	0.100 0.000	51.300 0.000	50.100 0.000	47.500 0.000	100.000 100.000	22.809 22.809	10.000 10.000	93.325 0.000	35.481 0.000	100.000 0.000	0.250 0.020	99.750 93.050	
LC50 (µM)	NCI-H22 M	4.110 0.012	17.000 1.000	1.000 0.000	1.000 1.000	100.000 1.000	27.500 0.000	10.000 0.000	0.100 0.000	51.300 0.000	50.100 0.000	47.500 0.000	100.000 100.000	22.809 22.809	10.000 10.000	93.325 0.000	35.481 0.000	100.000 0.000	0.250 0.020	99.750 93.050	
LC50 (µM)	NCI-H459	4.110 0.012	17.000 1.000	1.000 0.000	1.000 1.000	100.000 1.000	27.500 0.000	10.000 0.000	0.100 0.000	51.300 0.000	50.100 0.000	47.500 0.000	100.000 100.000	22.809 22.809	10.000 10.000	93.325 0.000	35.481 0.000	100.000 0.000	0.250 0.020	99.750 93.050	
LC50 (µM)	NCI-H22	9.660 0.010	28.800 1.000	100.000 1.000	12.600 0.010	85.100 0.000	2.820 0.000	8.100 0.000	0.000 0.000	47.900 0.000	81.100 0.000	100.000 100.000	10.000 10.000	6.026 0.000	10.000 0.000	74.131 0.000	10.195 0.000	100.000 0.000	2.820 2.820	97.190 97.190	
LC50 (µM)	HOP-42	9.660 0.010	28.800 1.000	100.000 1.000	12.600 0.010	85.100 0.000	2.820 0.000	8.100 0.000	0.000 0.000	47.900 0.000	81.100 0.000	100.000									

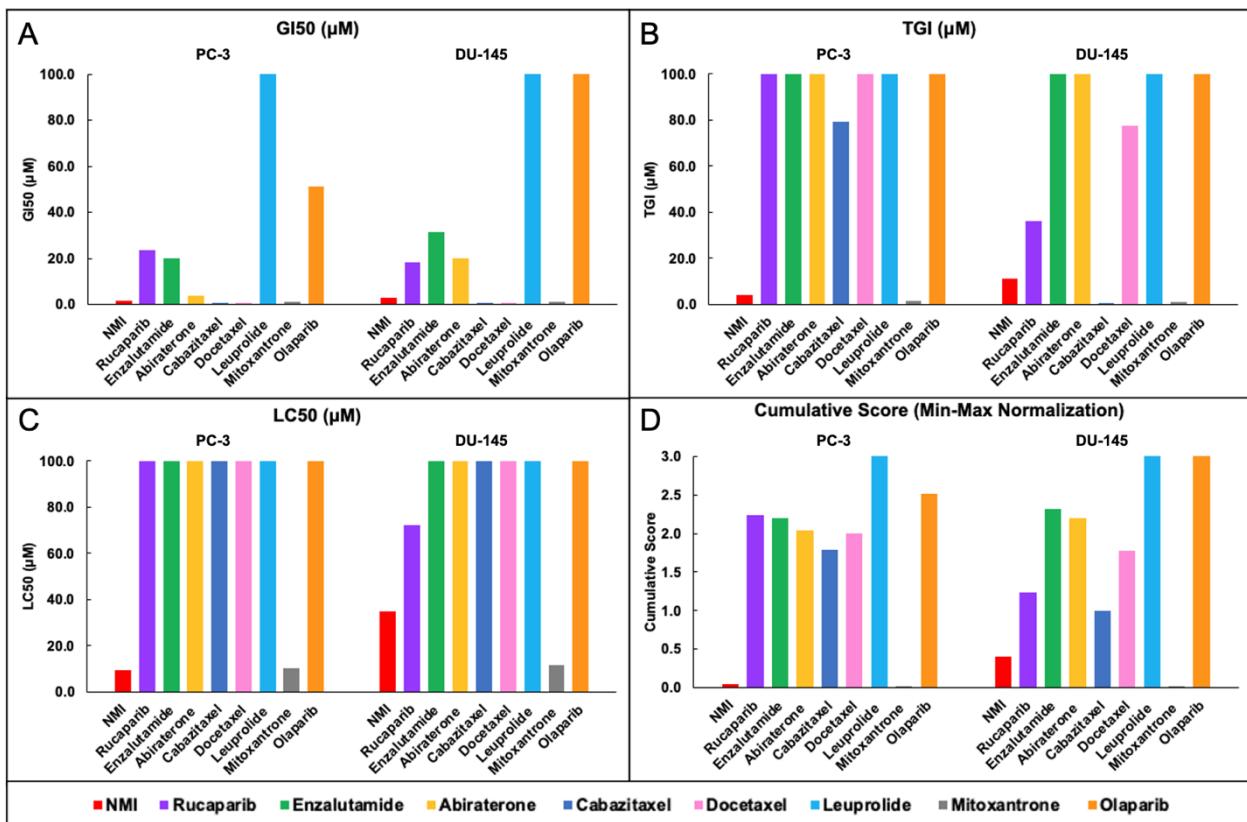
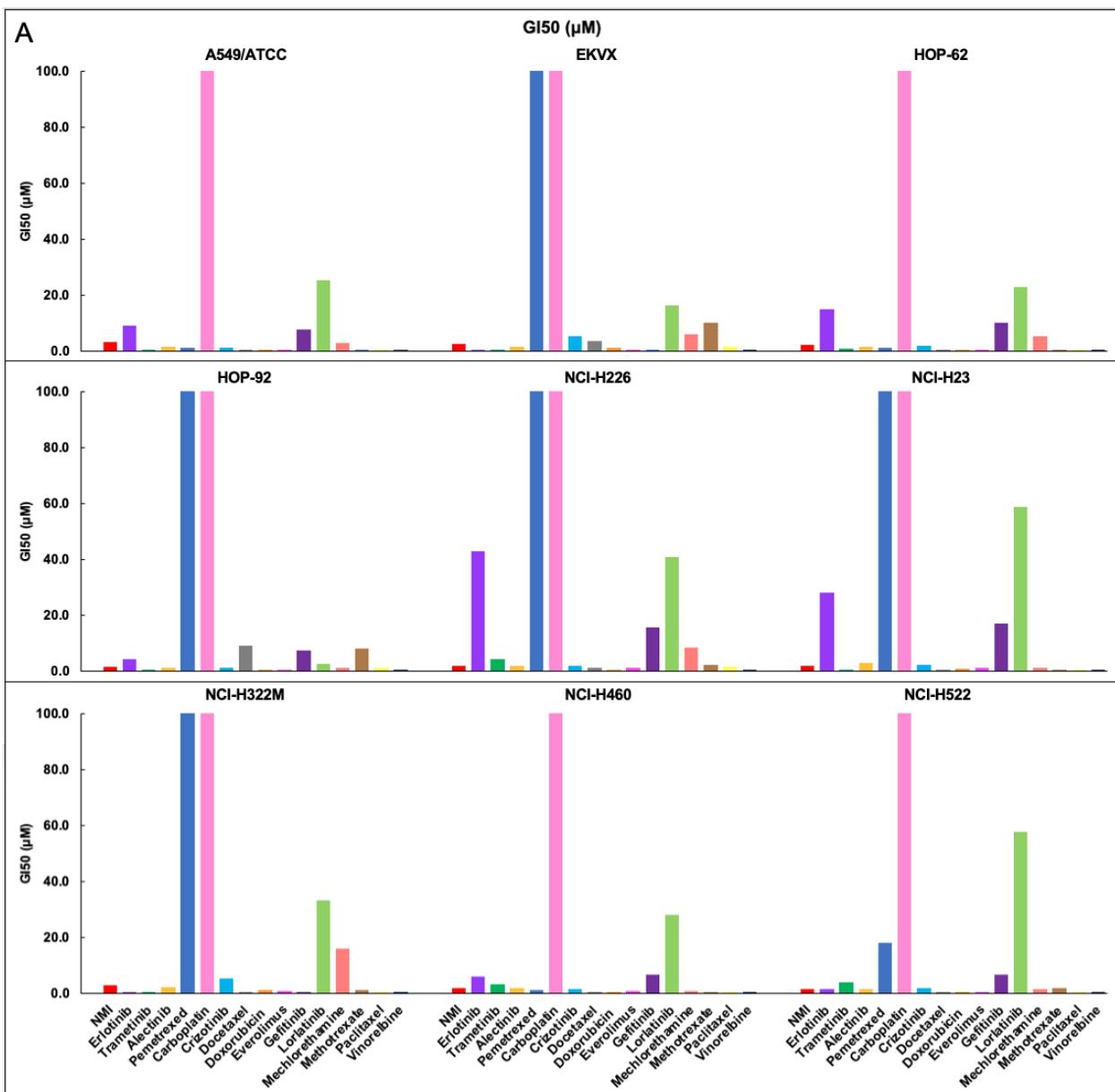
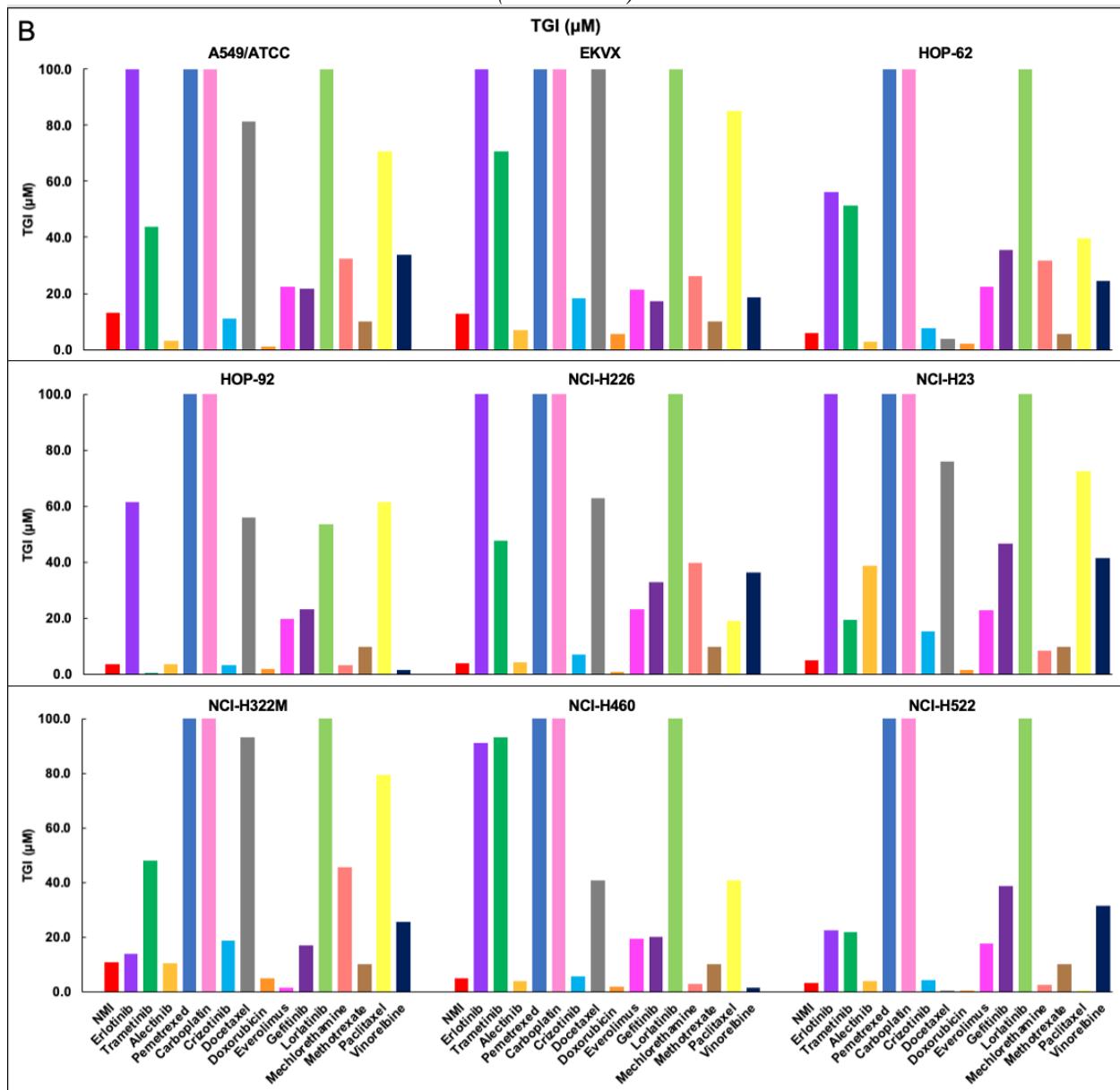


Figure S1 The comparison of (A)GI₅₀, (B)TGI, (C) LC₅₀, (D)cumulative score of NMI to FDA-approved prostate cancer drugs. NMI shows higher potency to prostate cancer cell lines than most FDA-approved prostate cancer drugs. Y-axis indicates concentration of GI₅₀, TGI, LC₅₀ (0 μM to 100 μM), and cumulative score (0.0 to 3.0). Drugs are color-coded: NMI (Red); Rucaparib (Purple); Enzalutamide (Green); Abiraterone (Yellow); Cabazitaxel (Blue); Docetaxel (Pink); Leuprolide (Light Blue); Mitoxantrone (Grey); Olaparib (Orange).



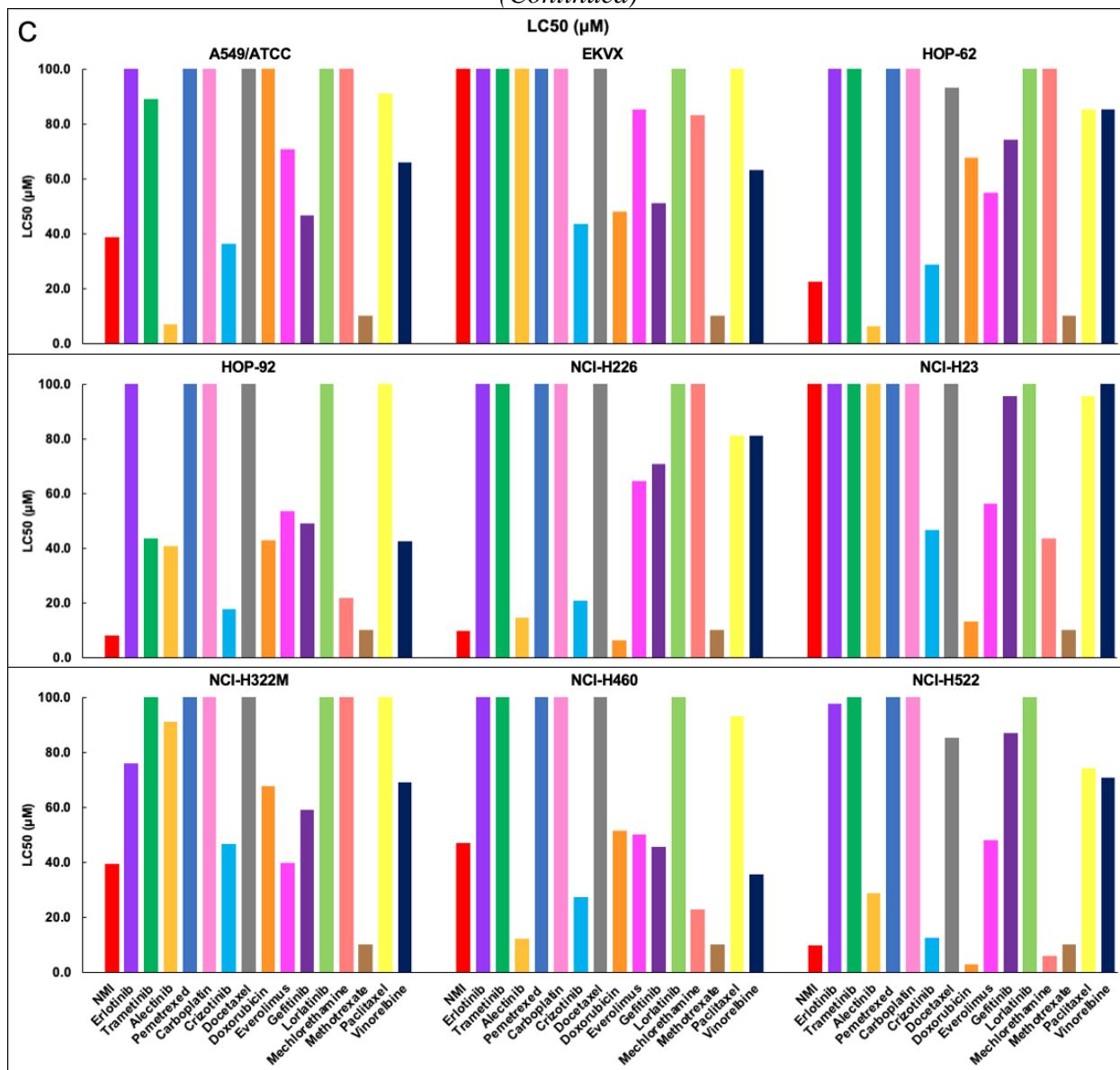
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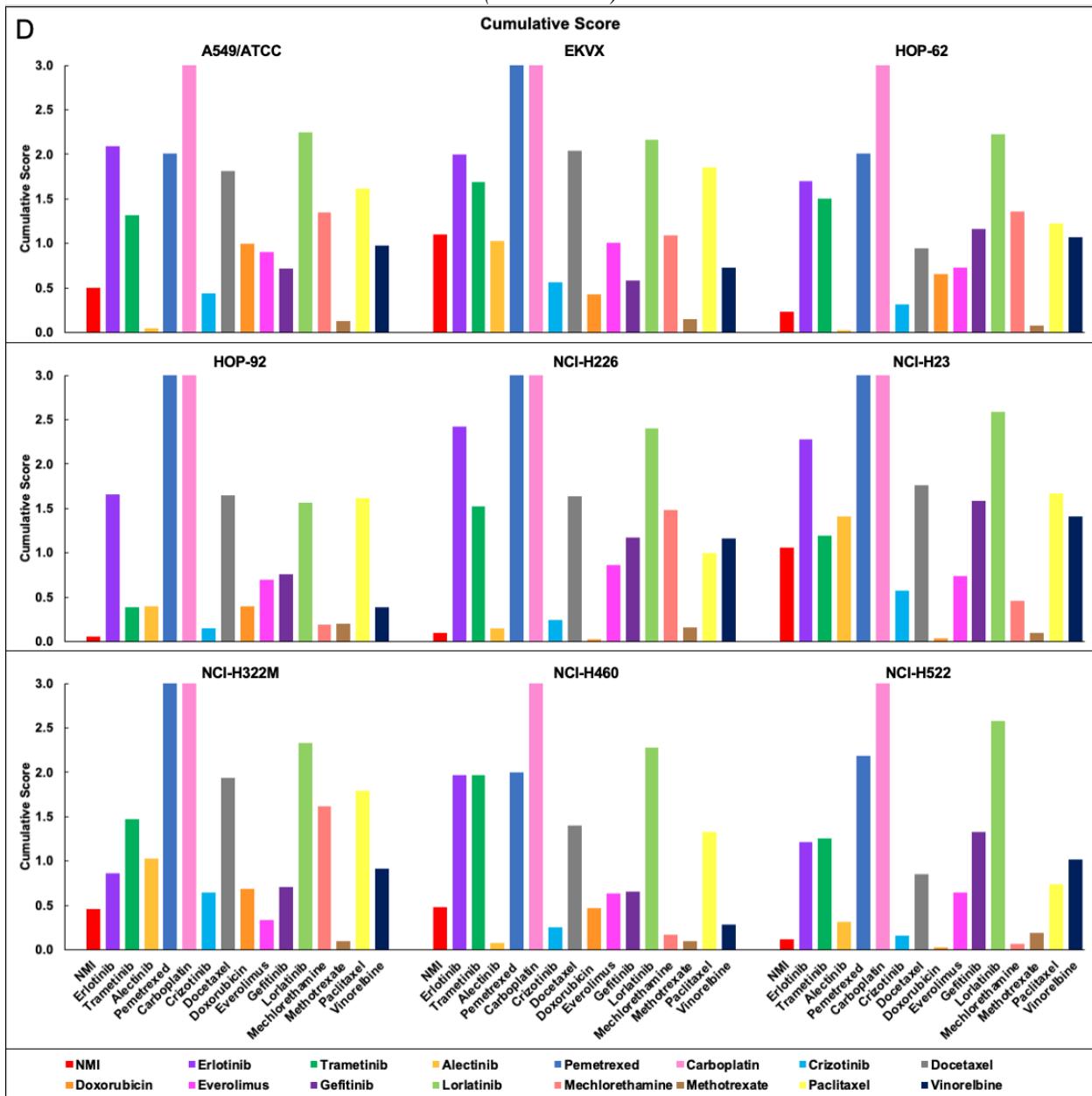
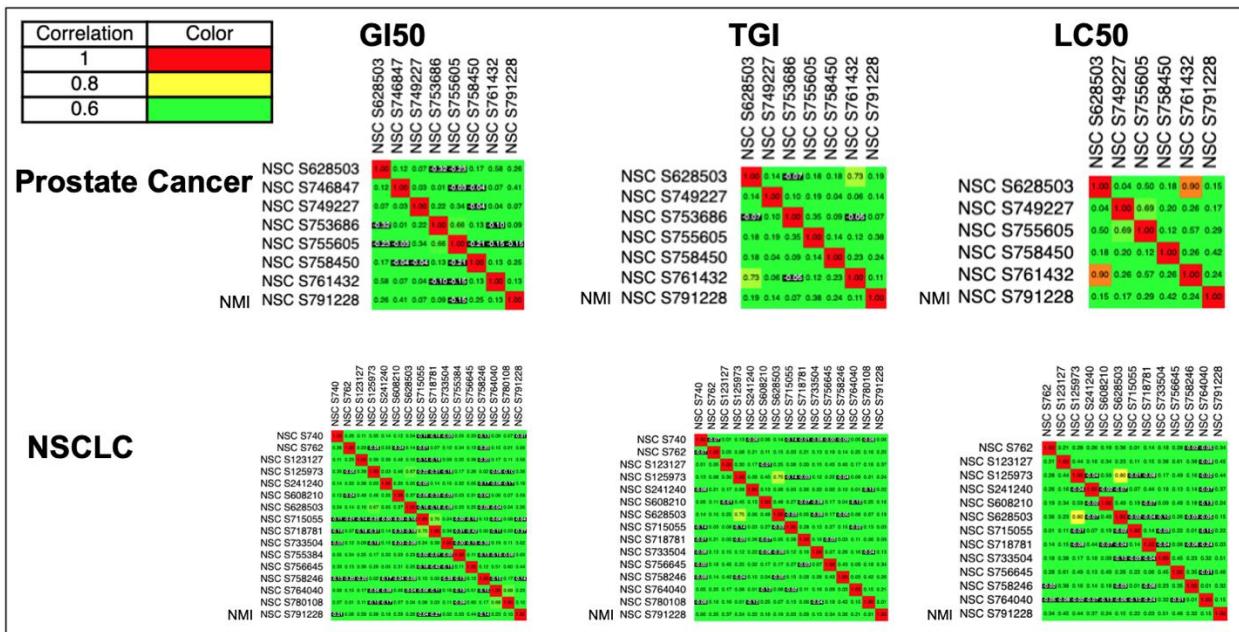


Figure S2 The comparison of (A) GI₅₀, (B) TGI, (C) LC₅₀, (D) cumulative score of NMI to FDA-approved NSCLC drugs. NMI shows higher potency to NSCLC cell lines than most FDA-approved NSCLC cancer drugs. Y-axis indicates concentration of GI₅₀, TGI, LC₅₀ (0 μM to 100 μM), and cumulative score (0.0 to 3.0). Drugs are color-coded: NMI (■); Erlotinib (□); Trametinib (■); Alectinib (□); Pemetrexed (■); Carboplatin (■); Crizotinib (□); Docetaxel (■); Doxorubicin (□); Everolimus (■); Gefitinib (□); Lorlatinib (■); Mechlorethamine (■); Methotrexate (■); Paclitaxel (□); Vinorelbine (■).

Table S3. The COMPARE results between NMI and FDA-approved CNS cancer, prostate cancer, and NSCLC drugs. All PCC values lower than 0.8, NMI has unique mechanisms than FDA-approved CNS, prostate, and NSCLC cancer drugs. Table of COMPARE results of Figure 5 and Figure S3. Exact values generated by COMPARE algorithm are provided.

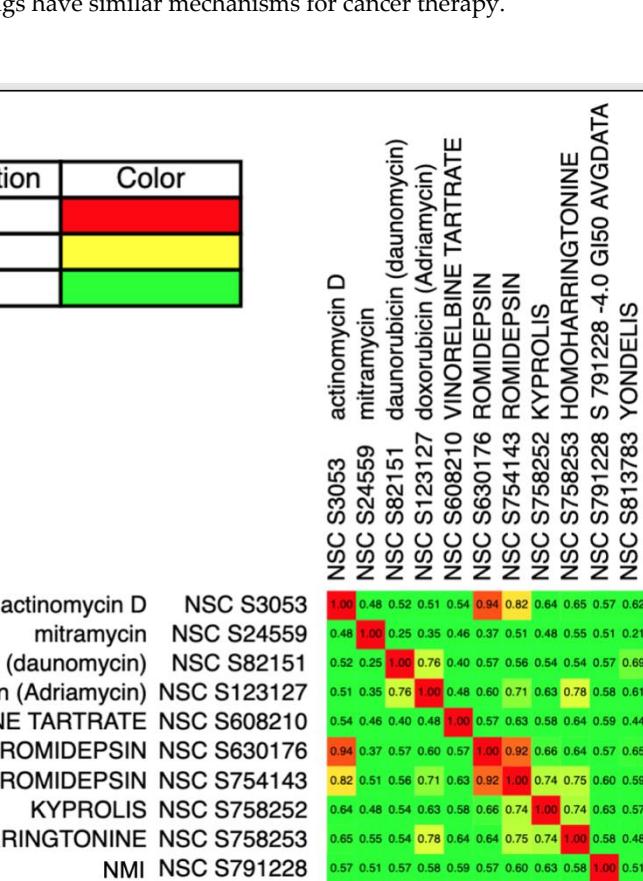
CNS Cancer				
Drug	NSC	PCC-GI50	PCC-TGI	PCC-LC50
Temozolomide	362856	0.00	-0.08	No data (StdDev = 0)
Lomustine	79037	0.48	0.04	0.23
Carmustine	409962	0.31	-0.04	0.27
Everolimus	733504	0.02	0.13	0.51
Prostate Cancer				
Drug	NSC	PCC-GI50	PCC-TGI	PCC-LC50
Rucaparib	756644	No data	No data	No data
Enzalutamide	755605	-0.15	0.38	0.29
Abiraterone Acetate	749227	0.07	0.14	0.17
Cabazitaxel	761432	0.13	0.11	0.24
Docetaxel	628503	0.26	0.19	0.15
Leuprolide Acetate	746847	0.41	No data (StdDev = 0)	No data (StdDev = 0)
Mitoxantrone	758450	0.25	0.24	0.42
Olaparib	753686	0.09	0.07	No data (StdDev = 0)
Non-Small Cell Lung Cancer				
Drug	NSC	PCC-GI50	PCC-TGI	PCC-LC50
Erlotinib	718781	-0.27	0.09	0.03
Trametinib	758246	-0.14	0.26	0.32
Alectinib	764040	0.23	0.15	0.21
Pemetrexed	755384	0.03	No data (StdDev = 0)	No data (StdDev = 0)
Carboplatin	241240	No data (StdDev = 0)	No data (StdDev = 0)	No data (StdDev = 0)
Crizotinib	756645	0.44	0.46	0.34
Docetaxel	628503	0.26	0.19	0.15
Doxorubicin	123127	0.58	0.37	0.45
Everolimus	733504	0.02	0.13	0.51
Gefitinib	715055	-0.04	0.03	0.22
Lorlatinib	780108	0.1	0.01	No data (StdDev = 0)
Mechlorethamine	762	0.06	0.25	0.34
Methotrexate	740	-0.01	0.06	No data (StdDev = 0)
Paclitaxel	125973	0.38	0.24	0.44
Vinorelbine	608210	0.59	0.16	0.24



A

Correlation

Correlation	Color
1	Red
0.8	Yellow
0.6	Green



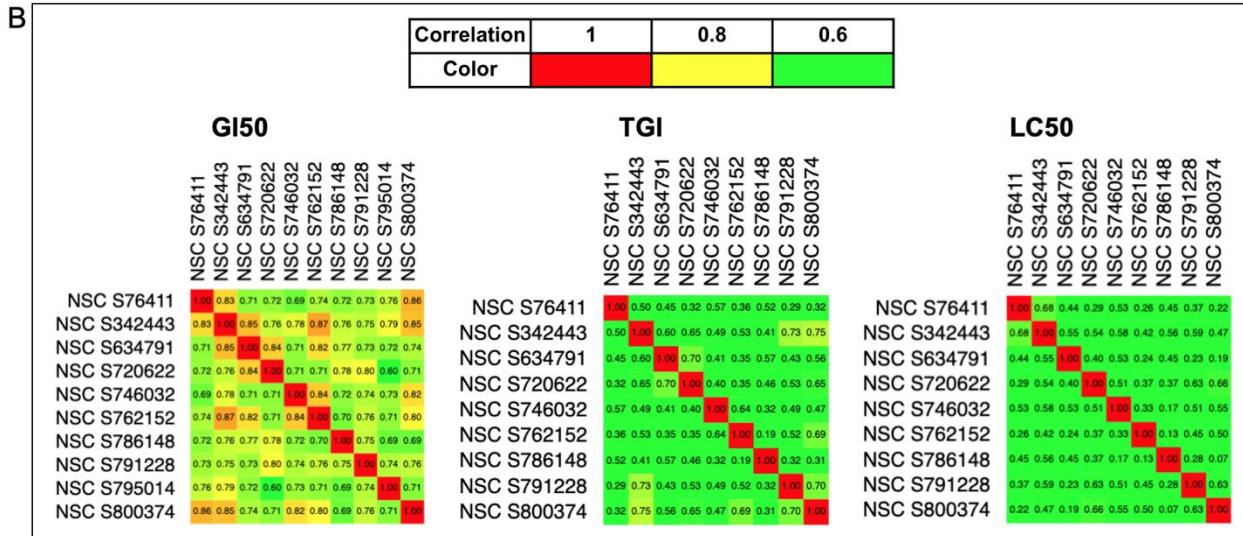


Figure S4. COMPARE plots of NMI (NSC 791228) to (A)marketed, (B)unmarked drugs in all cancers (top 10) PCC were shown. The result showed NMI has unique mechanism compared to all marketed anti-cancer drugs ; NMI might have similar mechanism with some unmarketed drugs. The Pairwise Pearson Correlation Coefficient (PCC) was determined by searching PUBLIC COMPARE Web Site Navigation for all marketed or unmarketed drugs in all different cancers in NCI60 database.

Table S4. Table of the COMPARE results of Figure S4A, with the drug names and their mechanisms.

NSC No.	Drug Name	PCC-GI50	Mechanism of Action
758252	Carfilzomib (Kyprolis)	0.63	Selective Proteasome Inhibitor
754143	Romidepsin	0.60	Histone Deacetylase Inhibitor
608210	Vinorelbine Tartrate	0.59	Antimicrotubular Antineoplastic Agent
123127	Doxorubicin	0.58	Topoisomerase Inhibitor
758253	Homoharringtonine	0.58	Protein Synthesis Inhibitor
3053	Actinomycin D	0.57	Transcription Inhibitor
82151	Daunorubicin	0.57	Topoisomerase Inhibitor
630176	Romidepsin	0.57	Histone Deacetylase Inhibitor
24559	Mitramycin	0.51	RNA Synthesis Inhibitor
813783	Trabectedin (Yondelis)	0.51	Alkylating Drug
791228	NMI	1.00	MAO A Inhibition; HIF1 α Stabilization