

**Identification of Polo-like kinase 1 interaction inhibitors  
using a novel cell-based assay**

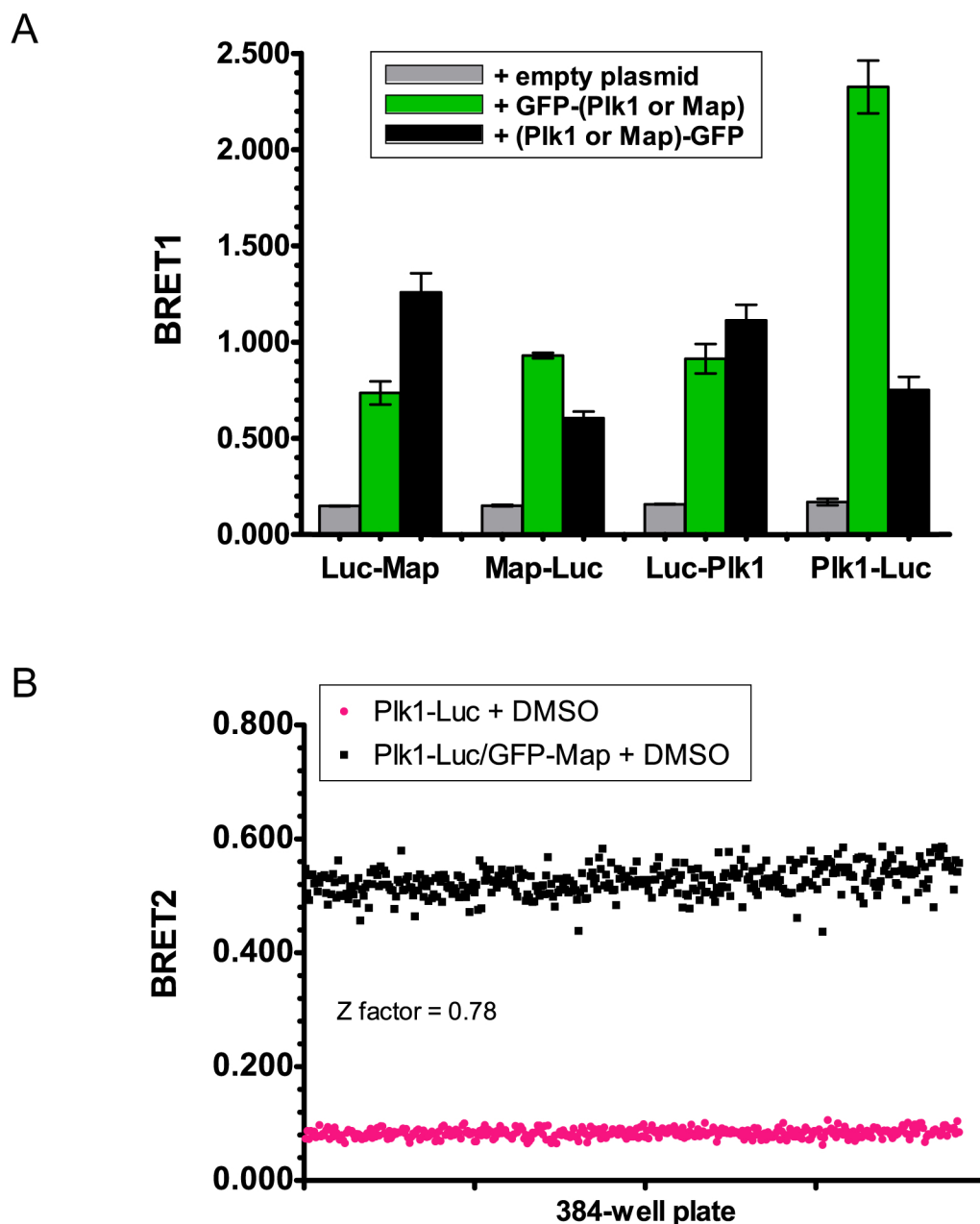
Karine Normandin<sup>1</sup>, Jean-François Lavallée<sup>1</sup>, Marie Futter<sup>1</sup>, Alexandre Beaudrait<sup>1</sup>, Jean Duchaine<sup>1</sup>, Sébastien Guiral<sup>1</sup>, Anne Marinier<sup>1,2</sup> & Vincent Archambault<sup>1,3\*</sup>

<sup>1</sup>Institute for Research in Immunology and Cancer, Université de Montréal;

<sup>2</sup>Département de chimie, Université de Montréal; <sup>3</sup>Département de biochimie et médecine moléculaire, Université de Montréal.

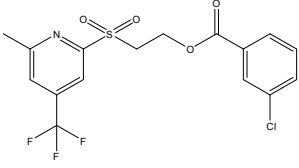
\*Corresponding author: [vincent.archambault.1@umontreal.ca](mailto:vincent.archambault.1@umontreal.ca)

**SUPPLEMENTARY FIGURES AND TABLES**

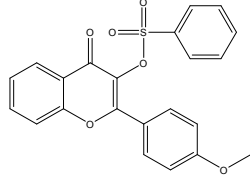


**Figure S1. Development of the BRET assay used in the screen.** A. BRET results for all combinations of Plk1 and Map<sub>254-416</sub> tagged with RLucII and GFP10 on their N-terminus or C-terminus. Experiment was performed in triplicate. Error bars: standard deviation. The Plk1-Luc/GFP-Map combination was selected because it produced the highest BRET dynamic window. Clonal stable cell lines co-expressing these proteins were then generated. B. Statistical validation of the BRET assay. One 384-well plate containing stable HEK293T Plk1-Luc/GFP-Map cells and one 384-well plate containing stable HEK293T Plk1-Luc cells (negative control) were treated for 2 hours with 0.5% DMSO (vehicle) before reading the BRET2. A Z-factor of 0.78 was obtained.

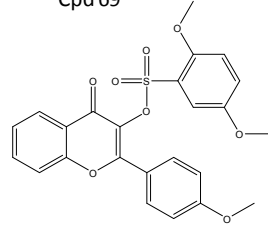
Cpd 16



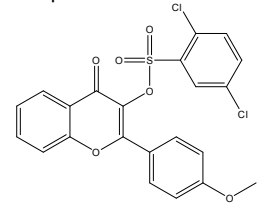
Cpd 46



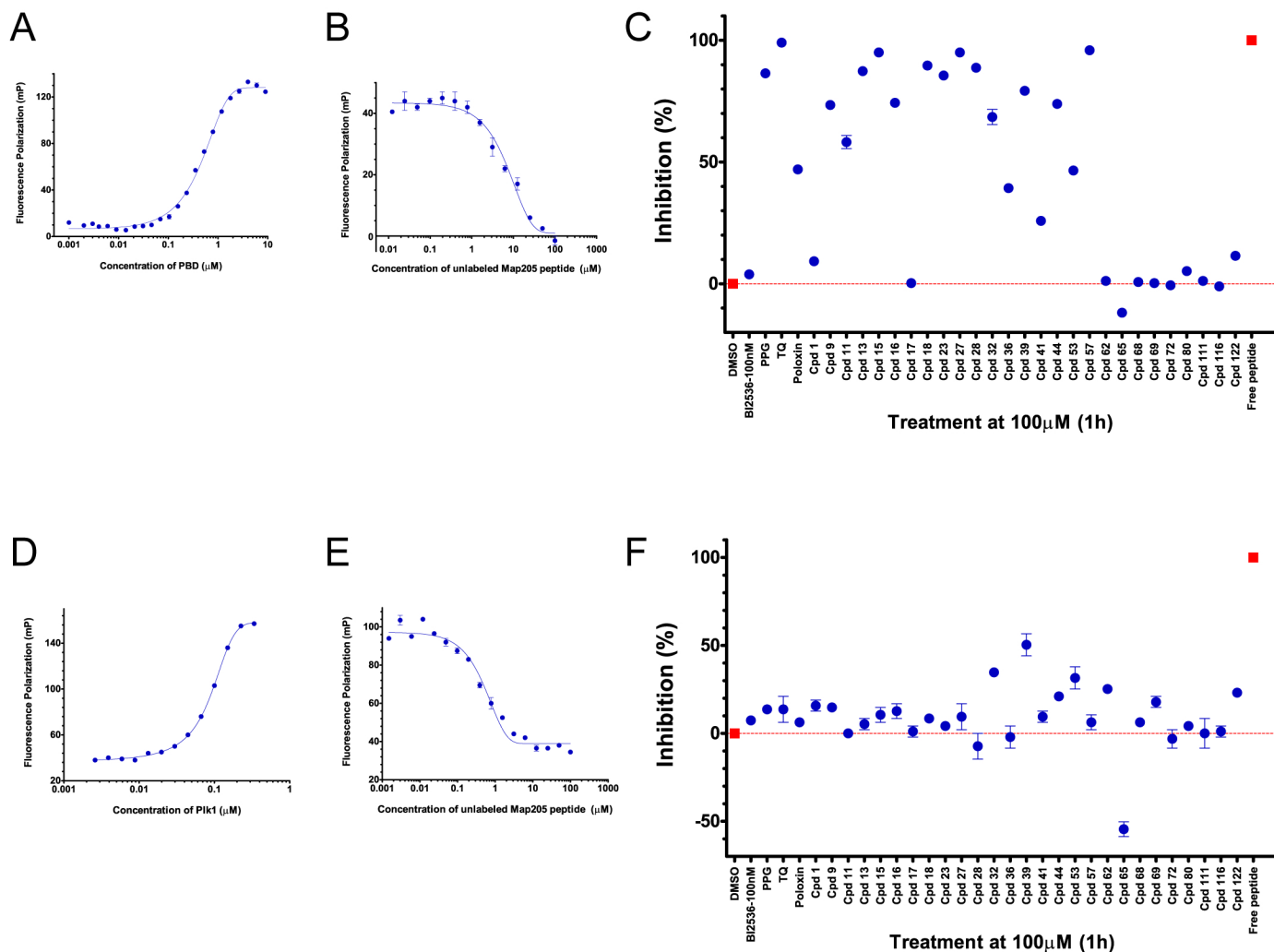
Cpd 69



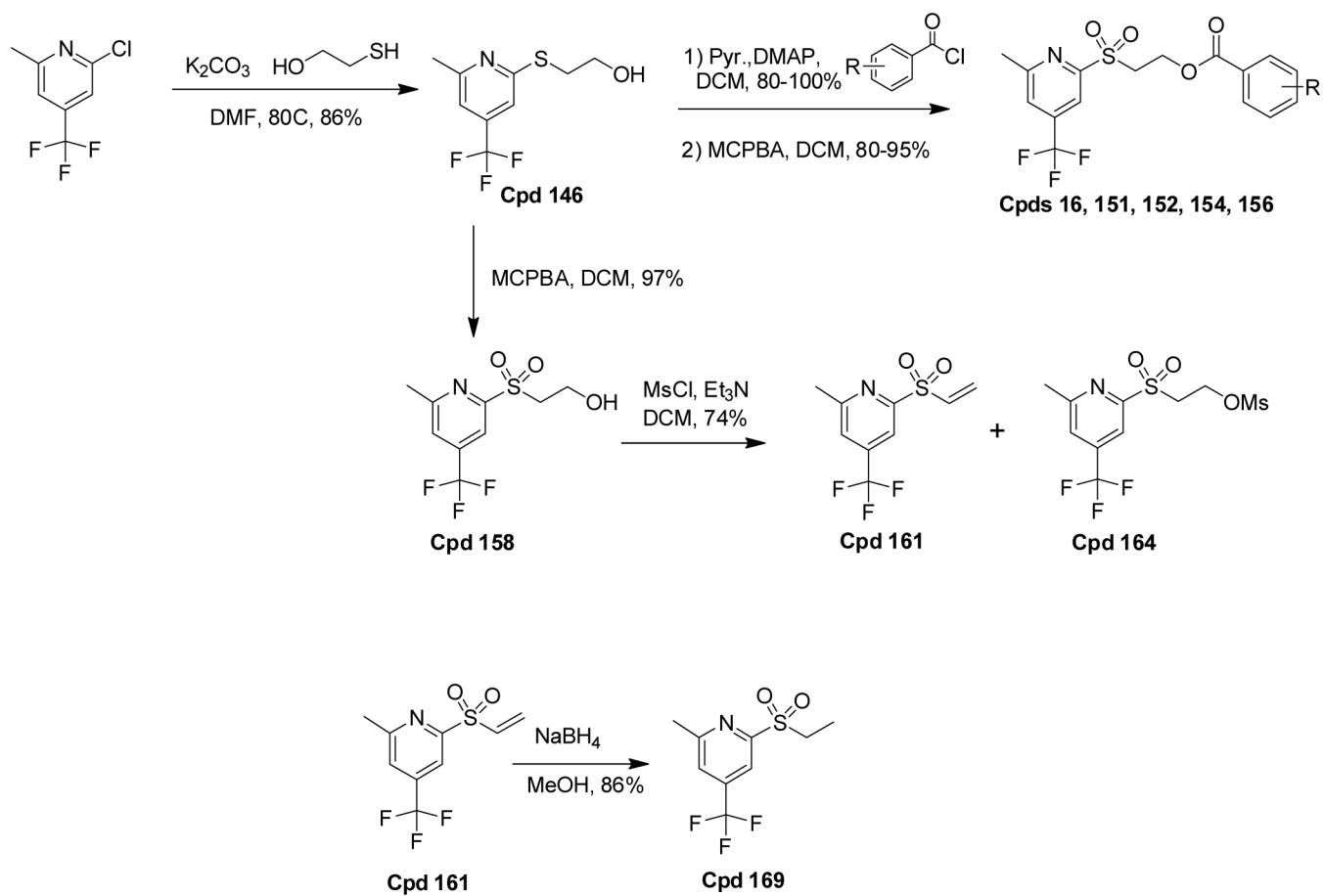
Cpd 122



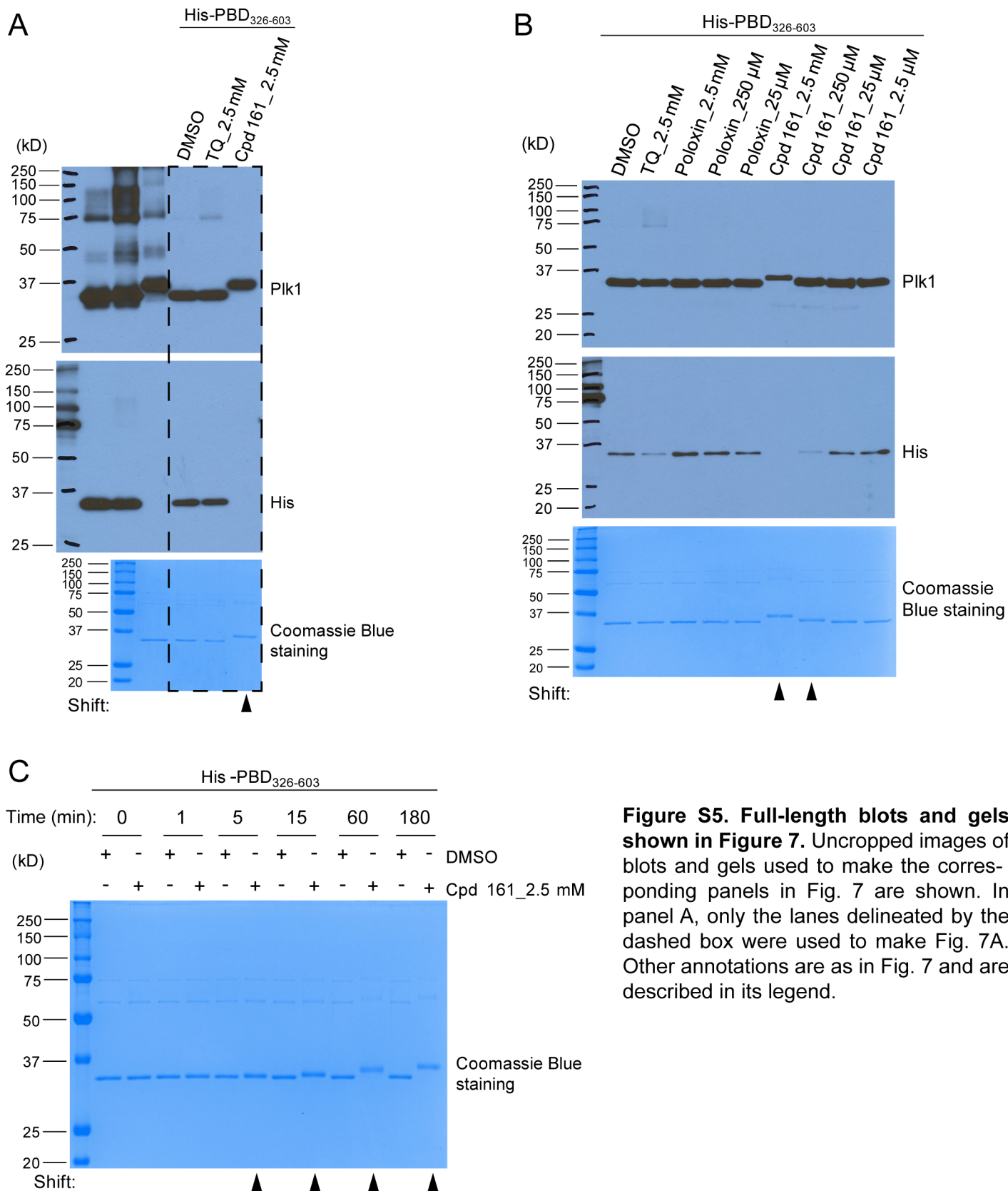
**Figure S2. Chemical structures of the BRET hits also active in the mitotic index assay.**



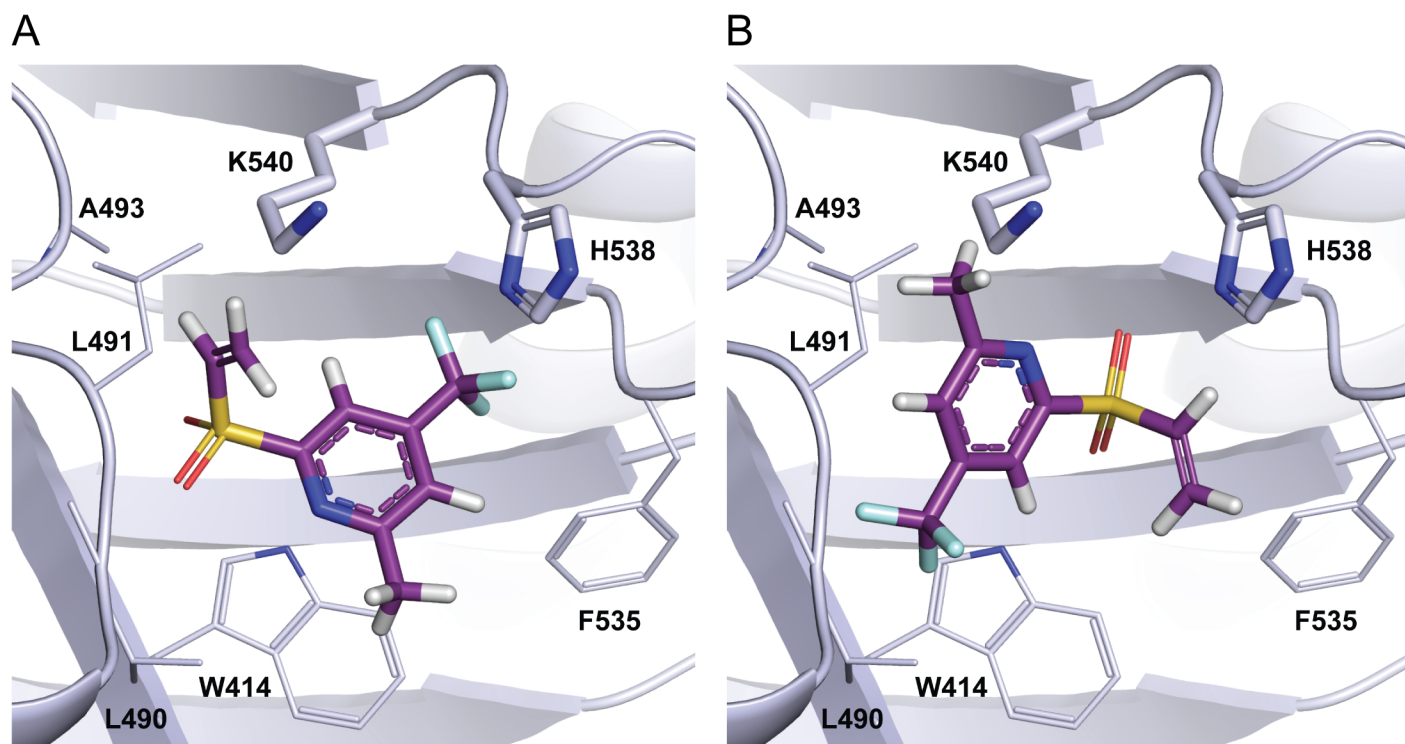
**Figure S3. Compounds identified interfere with the interaction between a Map peptide and the PBD alone (A-C) or whole Plk1 (D-F) in a Fluorescence Polarization assay.** A and D. Binding of the labeled FITC-Map peptide (10 nM) with increasing concentrations of His-PBD<sub>326-603</sub> (A) or with whole Plk1 (D). B and E. Competitive displacement of the FITC-Map peptide by the unlabeled Map peptide, added at increasing concentrations, with the Plk1 His-PBD<sub>326-603</sub> (used at EC<sub>65</sub> = 700 nM) (B) or with the whole Plk1 (used at EC<sub>65</sub> = 125 nM) (E). C and F. Targeted screen of selected compounds in the FP assay with the Map peptide. His-PBD<sub>326-603</sub> (1800 nM) (C) and whole Plk1 (130 nM) (F) were treated for 1 hour with different compounds at 100 μM before addition of the FITC-Map peptide. Values are means of duplicates from the same experiment (± range between values).



**Figure S4. Synthesis schemes for SAR around Cpd 16.** Other compounds were obtained commercially.



**Figure S5. Full-length blots and gels shown in Figure 7.** Uncropped images of blots and gels used to make the corresponding panels in Fig. 7 are shown. In panel A, only the lanes delineated by the dashed box were used to make Fig. 7A. Other annotations are as in Fig. 7 and are described in its legend.



**Figure S6. Putative binding modes of Cpd 161 on the PBD.** Molecular docking predicts that the preferred binding site of Cpd 161 on the PBD (PDB entry 1UMW) is its phosphopeptide binding pocket (where W414, H538 and K540 are crucial residues). Two alternative poses of Cpd 161, with non-significant difference in their respective estimated docking score, are shown (A and B). PBD is shown as cartoon and sticks, carbon and nitrogen atoms are in grey and blue color, respectively. Cpd 161 is shown as sticks, colored as follows: purple: carbon atoms, blue: nitrogen, yellow: sulfur; red: oxygen, light blue: fluorine, white: hydrogen.

**Table S1. Raw data from the LUMIER assay (Figure 3).** Luminescence on beads are presented for each of the 4 plates (one per tab). DMSO was included as an internal control on each plate (first and last column, in gray) and all compounds were tested in duplicate (side by side) on a same plate. Due to the time needed for the readings between columns in this assay, we found that counts systematically increased from left to right in each plate, as the enzymatic reaction progressed. To circumvent this problem, we analyzed the data in the following way. The mean luminescence value for each compound was divided by the maximum mean value of each column of compounds (in orange, normalized to 1). Resulting values are expressed as Relative Luminescence Units (RLU). The data should be viewed as rough indications of activity in co-IP.

PLATE 1

BEADS

Column #1	Column #2	Column #3	Column #4	Column #5	Column #6	Column #7	Column #8	Column #9	Column #10
DMSO: 390,066	Cpd 1: 15,076	9,654	Cpd 9: 74,244	87,986	Cpd 17: 886,164	1,022,272	Cpd 25: 1,428,771	1,306,371	DMSO: 2,585,917
DMSO: 453,836	Cpd 2: 159,976	152,582	Cpd 10: 1,132,468	1,141,840	Cpd 18: 48,113	66,908	Cpd 26: 1,841,391	1,130,286	DMSO: 2,910,820
DMSO: 423,655	Cpd 3: 669,840	695,778	Cpd 11: 374,254	328,854	Cpd 19: 18,046	28,452	Cpd 27: 42,504	39,770	DMSO: 2,967,920
DMSO: 359,412	Cpd 4: 459,296	589,011	Cpd 12: 641,210	771,760	Cpd 20: 1,032,000	1,384,114	Cpd 28: 470,276	374,830	DMSO: 2,979,922
DMSO: 401,916	Cpd 5: 723,890	358,615	Cpd 13: 38,822	51,866	Cpd 21: 728,406	1,095,957	Cpd 29: 2,579,044	2,456,177	DMSO: 2,864,320
DMSO: 425,204	Cpd 6: 221,644	208,600	Cpd 14: 1,055,362	1,184,868	Cpd 22: 1,037,198	1,434,797	Cpd 30: 1,964,228	1,946,733	DMSO: 2,482,486
DMSO: 398,126	Cpd 7: 422,540	537,002	Cpd 15: 18,060	19,824	Cpd 23: 75,154	249,262	Cpd 31: 1,563,040	639,044	DMSO: 2,315,537
DMSO: 57,336	Cpd 8: 114,916	163,133	Cpd 16: 21,904	28,648	Cpd 24: 315,073	464,828	Cpd 32: 26,500	26,938	DMSO: 217,502

Average-Beads	
DMSO- Column #1	407,459
DMSO- Column #10	2,729,560

Relative Luminescence Unit	
DMSO	0.907
Cpd 1	0.018
Cpd 2	0.229
Cpd 3	1.000
Cpd 4	0.768
Cpd 5	0.793
Cpd 6	0.315
Cpd 7	0.703
Cpd 8	0.204
Cpd 9	0.071
Cpd 10	1.000
Cpd 11	0.309
Cpd 12	0.621
Cpd 13	0.040
Cpd 14	0.985
Cpd 15	0.017
Cpd 16	0.022
Cpd 17	0.772
Cpd 18	0.047
Cpd 19	0.019
Cpd 20	0.977
Cpd 21	0.738
Cpd 22	1.000
Cpd 23	0.131
Cpd 24	0.315
Cpd 25	0.543
Cpd 26	0.590
Cpd 27	0.016
Cpd 28	0.168
Cpd 29	1.000
Cpd 30	0.777
Cpd 31	0.437
Cpd 32	0.011

DMSO	Column #1	Column #10	Average
	0.859	0.868	
	1.000	0.977	
	0.933	0.996	
	0.792	1.000	
	0.886	0.961	
	0.937	0.833	
	0.877	0.777	0.907



BEADS

	Column #1	Column #2	Column #3	Column #4	Column #5	Column #6	Column #7	Column #8	Column #9	Column #10
DMSO:	467,370	Cpd 33: 681,071	421,712	Cpd 41: 44,182	81,310	Cpd 49: 1,167,230	1,294,606	Cpd 57: 40,968	73,000	DMSO: 1,701,762
DMSO:	582,677	Cpd 34: 428,644	568,092	Cpd 42: 863,290	1,410,060	Cpd 50: 1,045,308	927,518	Cpd 58: 3,263,326	3,402,838	DMSO: 1,696,280
DMSO:	521,296	Cpd 35: 979,956	1,207,571	Cpd 43: 1,101,437	1,094,104	Cpd 51: 939,792	902,486	Cpd 59: 2,047,466	2,427,149	DMSO: 2,104,498
DMSO:	568,680	Cpd 36: 76,196	88,128	Cpd 44: 804,820	998,832	Cpd 52: 837,456	1,133,424	Cpd 60: 1,817,867	2,478,250	DMSO: 2,287,766
DMSO:	553,748	Cpd 37: 665,926	1,047,022	Cpd 45: 1,340,842	1,309,842	Cpd 53: 556,422	586,142	Cpd 61: 1,789,324	2,256,664	DMSO: 2,730,653
DMSO:	467,094	Cpd 38: 722,513	959,044	Cpd 46: 584,340	619,608	Cpd 54: 746,228	1,283,113	Cpd 62: 1,444,410	2,073,600	DMSO: 2,541,782
DMSO:	535,678	Cpd 39: 66,264	109,924	Cpd 47: 463,475	812,822	Cpd 55: 587,900	1,095,266	Cpd 63: 1,340,537	2,218,073	DMSO: 2,516,589
	258,198	Cpd 40: 227,542	285,071	Cpd 48: 2,760,910	2,724,993	Cpd 56: 608,977	913,864	Cpd 64: 858,840	1,249,824	990,406

Average-Beads

DMSO- Column #1 528,078

DMSO- Column #10 2,225,619

Relative Luminescence Unit

	Relative Luminescence Unit
DMSO	0.861
Cpd 33	0.504
Cpd 34	0.456
Cpd 35	1.000
Cpd 36	0.075
Cpd 37	0.783
Cpd 38	0.769
Cpd 39	0.081
Cpd 40	0.234
Cpd 41	0.023
Cpd 42	0.414
Cpd 43	0.400
Cpd 44	0.329
Cpd 45	0.483
Cpd 46	0.219
Cpd 47	0.233
Cpd 48	1.000
Cpd 49	1.000
Cpd 50	0.801
Cpd 51	0.748
Cpd 52	0.801
Cpd 53	0.464
Cpd 54	0.824
Cpd 55	0.684
Cpd 56	0.619
Cpd 57	0.017
Cpd 58	1.000
Cpd 59	0.671
Cpd 60	0.644
Cpd 61	0.607
Cpd 62	0.528
Cpd 63	0.534
Cpd 64	0.316

DMSO	Column #1	Column #10	Average
	0.802	0.623	
	1.000	0.621	
	0.895	0.771	
	0.976	0.838	
	0.950	1.000	
	0.802	0.931	
	0.919	0.922	0.861

BEADS

	Column #1	Column #2	Column #3	Column #4	Column #5	Column #6	Column #7	Column #8	Column #9	Column #10	Column #11
DMSO:	149,068	Cpd 65: 4,502	6,462	Cpd 73: 449,800	511,712	Cpd 81: 227,038	247,708	Cpd 89: 325,044	362,996	Cpd 97: 528,542	681,268
DMSO:	140,140	Cpd 66: 175,566	226,524	Cpd 74: 405,492	355,371	Cpd 82: 311,754	318,770	Cpd 90: 536,006	498,564	Cpd 98: 492,874	816,915
DMSO:	129,470	Cpd 67: 290,576	360,997	Cpd 75: 424,156	532,396	Cpd 83: 424,392	454,828	Cpd 91: 593,357	451,615	Cpd 99: 410,970	289,453
DMSO:	130,420	Cpd 68: 109,900	138,380	Cpd 76: 444,633	594,914	Cpd 84: 356,900	434,891	Cpd 92: 600,246	595,235	Cpd 100: 689,748	497,264
DMSO:	116,602	Cpd 69: 232,744	331,226	Cpd 77: 477,280	570,393	Cpd 85: 454,990	421,466	Cpd 93: 565,666	439,402	DMSO: 629,410	609,232
DMSO:	128,070	Cpd 70: 265,246	317,708	Cpd 78: 287,174	461,071	Cpd 86: 299,360	479,828	Cpd 94: 521,318	506,278	DMSO: 613,608	387,714
DMSO:	115,826	Cpd 71: 165,275	238,710	Cpd 79: 295,998	343,468	Cpd 87: 266,608	435,000	Cpd 95: 407,448	414,768	DMSO: 413,108	334,016
	12,484	Cpd 72: 64,794	178,690	Cpd 80: 121,111	185,792	Cpd 88: 170,771	302,793	Cpd 96: 146,777	357,380	DMSO: 321,470	67,770

Average-Beads

DMSO- Column #1 129,942

DMSO- Columns #10-11 472,651

Relative Luminescence Unit

DMSO	Relative Luminescence Unit
DMSO	0.816
Cpd 65	5.482
Cpd 66	201,045
Cpd 67	325,787
Cpd 68	124,140
Cpd 69	281,985
Cpd 70	291,477
Cpd 71	201,993
Cpd 72	121,742
Cpd 73	480,756
Cpd 74	380,432
Cpd 75	478,276
Cpd 76	519,774
Cpd 77	523,837
Cpd 78	374,123
Cpd 79	319,733
Cpd 80	153,452
Cpd 81	237,373
Cpd 82	315,262
Cpd 83	439,610
Cpd 84	395,896
Cpd 85	438,228
Cpd 86	389,594
Cpd 87	350,804
Cpd 88	236,782
Cpd 89	344,020
Cpd 90	517,285
Cpd 91	522,486
Cpd 92	597,741
Cpd 93	502,534
Cpd 94	513,798
Cpd 95	411,108
Cpd 96	252,079
Cpd 97	604,905
Cpd 98	654,895
Cpd 99	350,212
Cpd 100	593,506

DMSO	Column #1	Columns #10-11	Average
	1.000	1.000	
	0.940	0.975	
	0.869	0.656	
	0.875	0.511	
	0.782	1.000	
	0.859	0.636	
	0.777	0.548	0.816

BEADS

	Column #1	Column #2	Column #3	Column #4	Column #5	Column #6	Column #7	Column #8	Column #9	Column #12
DMSO:	250,062	Cpd 101: 377,313	353,900	Cpd 109: 275,224	381,837	Cpd 117: 370,854	531,853	TQ: 221,826	304,556	DMSO: 1,595,057
DMSO:	259,951	Cpd 102: 412,144	595,808	Cpd 110: 555,674	535,226	Cpd 118: 757,471	832,352	Poloxin: 1,325,160	1,298,916	DMSO: 1,442,344
DMSO:	228,044	Cpd 103: 400,971	520,906	Cpd 111: 96,962	94,360	Cpd 119: 801,555	773,018			DMSO: 1,414,942
DMSO:	286,524	Cpd 104: 425,618	630,168	Cpd 112: 702,052	676,557	Cpd 120: 888,117	828,624			DMSO: 1,095,800
DMSO:	185,230	Cpd 105: 415,974	547,728	Cpd 113: 690,682	745,816	Cpd 121: 783,868	856,250	Cpd 122: 909,984	1,127,164	DMSO: 1,122,604
DMSO:	207,044	Cpd 106: 316,228	486,338	Cpd 114: 557,726	625,664					DMSO: 1,067,646
DMSO:	216,178	Cpd 107: 246,040	392,276	Cpd 115: 364,440	560,460	BI 2536: 47,611	56,678			DMSO: 877,613
DMSO:	107,904	Cpd 108: 175,324	197,258	Cpd 116: 248,798	364,698	PPG: 290,044	312,040			DMSO: 359,786

Average-Beads

DMSO- Column #1 233,290

DMSO- Column #12 1,230,858

Relative Luminescence Unit

	Relative Luminescence Unit
DMSO	0.793
Cpd 101	0.693
Cpd 102	0.955
Cpd 103	0.873
Cpd 104	1.000
Cpd 105	0.913
Cpd 106	0.760
Cpd 107	0.605
Cpd 108	0.353
Cpd 109	0.457
Cpd 110	0.759
Cpd 111	0.133
Cpd 112	0.960
Cpd 113	1.000
Cpd 114	0.824
Cpd 115	0.644
Cpd 116	0.427
Cpd 117	0.526
Cpd 118	0.926
Cpd 119	0.917
Cpd 120	1.000
Cpd 121	0.955
BI 2536	0.061
PPG	0.351
TQ	0.201
Poloxin	1.000
Cpd 122	0.776

DMSO	Column #1	Column #12	Average
	0.873	1.000	
	0.907	0.904	
	0.796	0.887	
	1.000	0.687	
	0.646	0.704	
	0.723	0.669	
	0.754	0.550	0.793

Table S2. BRET, mitotic index and co-IP results compilation for the 122 BRET hits.

Cpd #	ID	Batch	Supplier	Supplier ID	Primary screen Plk1-Map (stable)		Primary screen CRAF-BRAF (transient)		Confirmation Plk1-Map (stable)		Confirmation Plk1-Map (transient)		Confirmation CRAF-BRAF (transient)		Mitotic index		Co-IP
					% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	Mitotic index (%)
1	UM0107177	01	MAYBRIDGE	AW 00482	50.82	79.13	21.44	80.08	66.71	81.02	63.61	69.40	-24.74	72.14	1.59	668	0.018
2	UM0107221	01	MAYBRIDGE	AW 00592	55.07	85.96	31.45	75.12	35.46	90.44	-34.13	73.50	0.35	69.42	5.64	1121	0.229
3	UM0107266	01	MAYBRIDGE	AW 01093	53.30	85.61	61.48	88.22	30.16	93.58	-12.13	89.74	2.18	86.31	5.16	1241	1.000
4	UM0107457	01	MAYBRIDGE	BTB 01468	61.37	74.20	6.39	64.29	37.02	71.62	3.89	51.50	-23.24	41.54	3.78	1030	0.768
5	UM0107465	01	MAYBRIDGE	BTB 01397	65.21	59.62	23.57	38.70	45.45	61.07	17.66	-13.30	-0.79	23.13	0.41	638	0.793
6	UM0107466	01	MAYBRIDGE	BTB 01470	62.68	87.76	7.51	84.96	32.04	89.40	5.59	80.38	-33.37	75.46	2.84	1005	0.315
7	UM0107535	01	MAYBRIDGE	BTB 02075	58.30	64.35	-7.01	66.17	26.54	47.47	-9.57	16.31	-35.15	6.23	4.37	811	0.703
8	UM0108212	01	MAYBRIDGE	BTB 04098	54.13	66.53	13.33	68.83	25.74	60.04	7.94	28.16	-18.98	25.97	2.89	1083	0.204
9	UM0108871	01	MAYBRIDGE	BTB 04426	56.34	77.93	-52.46	69.98	49.28	70.98	60.77	58.61	9.16	81.03	0.17	592	0.071
10	UM0024677	03	MAYBRIDGE	BTB 04957	67.60	93.78	36.31	85.60	42.67	93.83	-14.46	73.44	-11.38	63.73	4.31	1024	1.000
11	UM0111426	01	MAYBRIDGE	BTB 06019	46.60	76.68	1.34	48.60	29.73	77.70	-13.92	40.97	-10.42	53.70	7.54	910	0.309
12	UM0112103	01	MAYBRIDGE	BTB 06776	68.25	89.25	38.71	86.66	20.39	81.39	7.11	70.27	-12.51	63.67	4.08	1178	0.621
13	UM0112687	01	MAYBRIDGE	BTB 07155	76.83	85.35	51.15	84.05	61.45	79.50	68.19	66.43	18.66	83.95	3.70	612	0.040
14	UM0113547	01	MAYBRIDGE	BTB 08799	67.58	86.85	39.91	80.51	41.26	89.74	-38.11	59.00	-14.66	39.49	5.24	1125	0.985
15	UM0113559	01	MAYBRIDGE	BTB 08960	67.91	84.93	31.50	81.13	41.55	78.03	53.01	72.97	-4.59	86.14	0.81	823	0.017
16	UM0113574	01	MAYBRIDGE	BTB 09085	67.34	86.24	52.83	88.39	30.79	81.37	30.06	74.57	0.37	80.17	15.73	789	0.022
17	UM0114221	01	MAYBRIDGE	CD 03862	64.09	91.79	17.43	85.48	62.50	91.30	3.77	83.77	-44.82	79.36	9.70	843	0.772
18	UM0114440	01	MAYBRIDGE	CD 08589	45.55	78.56	8.26	87.41	42.59	81.79	60.17	69.60	7.13	87.60	0.54	458	0.047
19	UM0044124	02	MAYBRIDGE	DSHS 00690	77.53	88.69	42.12	87.63	36.81	92.52	10.92	83.45	-26.61	81.39	0.57	585	0.019
20	UM0114956	01	MAYBRIDGE	DSHS 00902	46.34	87.87	54.93	89.27	21.53	91.98	-17.19	86.96	-22.73	82.72	5.77	1281	0.977
21	UM0114962	01	MAYBRIDGE	DSHS 00691	45.76	78.45	23.02	53.92	23.26	69.35	-13.00	46.29	-15.44	29.17	5.77	1201	0.738
22	UM0115060	01	MAYBRIDGE	GK 00932	57.37	84.54	33.24	85.72	41.81	95.30	-1.29	89.97	-2.53	87.37	4.14	1226	1.000
23	UM0044370	02	MAYBRIDGE	GK 01179	78.41	82.19	-37.15	81.61	48.10	81.45	64.99	70.40	-65.02	83.59	2.63	797	0.131
24	UM0115252	01	MAYBRIDGE	GK 02719	68.10	88.13	54.62	87.02	36.96	93.36	5.68	86.41	5.36	88.72	5.59	1240	0.315
25	UM0115281	01	MAYBRIDGE	GK 03147	45.23	89.37	53.14	89.82	26.07	94.57	-8.10	92.30	-8.57	88.81	4.78	1116	0.543
26	UM0115294	01	MAYBRIDGE	GK 03068	52.76	87.32	36.28	78.36	40.41	87.86	-6.40	79.33	-8.86	73.75	3.92	1083	0.590
27	UM0115297	01	MAYBRIDGE	GK 03183	80.81	80.09	31.44	87.95	39.21	79.07	68.50	70.64	18.98	87.05	2.84	705	0.016
28	UM0044649	02	MAYBRIDGE	HAN 00116	45.62	78.12	9.04	88.52	50.48	80.87	64.88	70.41	9.73	87.46	2.87	764	0.168
29	UM0115430	01	MAYBRIDGE	HAN 00410	35.64	79.81	4.37	39.68	27.83	77.47	-13.59	29.30	-17.95	16.64	3.94	1057	1.000
30	UM0115588	01	MAYBRIDGE	HTS 01167	50.70	82.37	26.25	82.76	38.26	90.34	0.67	84.37	-16.55	76.63	4.46	1130	0.777
31	UM0045020	02	MAYBRIDGE	HTS 01731	59.16	85.22	35.04	72.15	53.24	87.56	-1.36	67.10	8.69	55.48	5.27	1336	0.437
32	UM0115695	01	MAYBRIDGE	HTS 01716	82.98	78.54	52.78	85.20	32.39	61.04	10.87	55.86	-15.05	34.17	6.85	1221	0.011
33	UM0045568	02	MAYBRIDGE	HTS 04586	46.30	21.41	9.33	66.80	25.44	52.76	3.17	40.96	-7.50	25.32	5.01	895	0.504
34	UM0116166	01	MAYBRIDGE	HTS 05440	47.36	37.79	35.79	59.50	32.78	64.09	9.76	34.37	4.03	13.92	5.87	1113	0.456
35	UM0116479	01	MAYBRIDGE	HTS 07461	47.28	41.69	13.69	17.06	33.04	51.36	-2.19	-5.46	4.61	-12.42	6.35	1023	1.000
36	UM0116537	01	MAYBRIDGE	HTS 07564	51.44	43.67	14.13	58.96	38.28	47.30	52.07	8.81	6.72	33.27	0.04	812	0.075
37	UM0046711	02	MAYBRIDGE	HTS 11170	45.24	80.82	30.49	82.80	26.10	87.33	-14.06	77.84	-17.55	62.37	5.51	1278	0.783
38	UM0117516	01	MAYBRIDGE	JFD 02414	51.01	63.14	12.78	19.29	43.35	69.42	0.03	6.66	3.22	2.00	4.51	1105	0.769
39	UM0117931	01	MAYBRIDGE	KM 01532	46.92	58.25	38.21	68.10	28.92	72.06	39.04	13.68	-28.02	51.27	1.85	915	0.081
40	UM0107759	01	MAYBRIDGE	KM 03640	57.91	59.70	3.04	65.18	33.00	64.89	28.30	40.55	15.17	70.46	0.76	711	0.234
41	UM0107915	01	MAYBRIDGE	KM 05478	48.25	76.91	9.15	82.74	22.85	91.45	-1.55	83.71	-0.82	89.16	0.65	752	0.023
42	UM0107958	01	MAYBRIDGE	KM 05637	46.57	80.72	33.05	84.16	25.51	75.79	-14.59	37.29	-5.11	13.72	4.64	1146	0.414
43	UM0109122	01	MAYBRIDGE	NRB 04070	54.41	82.81	6.29	68.33	29.95	84.93	35.79	79.11	10.00	84.92	0.38	961	0.400
44	UM0050240	02	MAYBRIDGE	NRB 04635	46.80	65.81	27.75	81.91	31.30	73.89	46.34	65.89	4.73	75.24	6.66	844	0.329

					Primary screen Plk1-Map (stable)		Primary screen CRAF-BRAF (transient)		Confirmation Plk1-Map (stable)		Confirmation Plk1-Map (transient)		Confirmation CRAF-BRAF (transient)		Mitotic index		Co-IP
Cpd #	ID	Batch	Supplier	Supplier ID	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	Mitotic index (%)	Nb of analyzed cells	Values relative to top in plate column = 1
45	UM0110305	01	MAYBRIDGE	RH 01618	45.20	71.08	1.10	51.49	33.67	85.08	-30.22	54.24	-9.24	31.93	5.38	1008	0.483
46	UM0110531	01	MAYBRIDGE	RJC 01628	53.18	82.37	16.40	63.54	30.35	91.04	-29.25	72.57	19.45	66.12	29.44	831	0.219
47	UM0111224	01	MAYBRIDGE	S 04076	45.29	83.95	5.83	80.16	25.06	89.54	-34.97	71.30	7.01	59.13	5.81	1150	0.233
48	UM0111382	01	MAYBRIDGE	S 09528	64.65	53.86	-17.19	85.32	33.05	76.00	66.76	64.63	-36.81	71.69	2.86	844	1.000
49	UM0111558	01	MAYBRIDGE	S 12721	47.04	83.98	26.90	82.37	22.76	94.87	-25.39	85.65	-4.13	80.84	5.42	1246	1.000
50	UM0053923	02	MAYBRIDGE	SEW 02094	50.27	81.17	30.34	74.95	21.62	89.00	-26.33	68.27	4.23	62.33	4.49	1043	0.801
51	UM0112278	01	MAYBRIDGE	SEW 04487	45.29	68.33	38.92	79.48	23.19	82.72	-21.49	67.09	-13.15	65.32	5.08	1140	0.748
52	UM0054719	02	MAYBRIDGE	SEW 06560	51.80	71.42	33.34	75.79	34.00	87.94	-7.60	68.31	1.45	68.36	5.54	1251	0.801
53	UM0054792	02	MAYBRIDGE	SP 00278	73.92	65.68	-13.36	70.12	60.53	70.21	60.20	19.84	-23.26	57.92	0.25	711	0.464
54	UM0112835	01	MAYBRIDGE	SP 01000	53.70	67.85	37.49	80.97	41.04	90.19	-32.83	72.33	-10.41	72.90	4.26	1083	0.824
55	UM0112874	01	MAYBRIDGE	SP 00932	65.38	72.24	36.23	82.49	28.65	80.09	-12.63	42.73	11.99	57.62	5.27	1126	0.684
56	UM0112902	01	MAYBRIDGE	SP 01131	67.10	77.26	28.17	81.60	50.83	90.14	-16.36	57.34	6.88	51.70	4.98	1253	0.619
57	UM0113055	01	MAYBRIDGE	SPB 01974	53.19	68.79	4.09	89.46	50.51	72.14	71.14	50.57	-10.70	69.51	0.54	716	0.017
58	UM0092330	01	SPECS	AE-842/33004023	65.44	83.13	31.26	89.59	53.82	91.81	30.67	78.98	-23.16	86.72	3.75	918	1.000
59	UM0098743	01	SPECS	AH-487/14754106	45.99	44.75	-7.85	49.56	29.09	69.72	-9.80	25.20	4.44	18.13	5.75	1110	0.671
60	UM0101058	01	SPECS	AK-968/13146122	35.23	63.07	11.13	67.69	32.12	85.23	-20.95	61.21	-6.53	54.44	5.92	1170	0.644
61	UM0096778	01	SPECS	AG-690/11634009	34.79	55.41	-3.90	28.94	28.84	66.72	-11.44	10.58	5.94	8.20	5.04	1218	0.607
62	UM0099861	01	SPECS	AK-778/40958947	57.06	92.90	34.60	89.55	25.31	91.53	-4.94	88.74	-31.50	86.36	8.40	1202	0.528
63	UM0093537	01	SPECS	AF-399/40772847	55.30	85.95	14.36	72.59	22.43	90.81	-31.39	74.12	-6.96	70.12	6.28	1310	0.534
64	UM0098333	01	SPECS	AG-690/40751449	54.01	89.49	23.13	74.11	26.54	78.16	-17.93	44.43	16.60	30.91	5.57	1198	0.316
65	UM0103856	01	SPECS	AM-814/41094541	61.32	62.29	18.29	63.96	36.24	55.25	66.45	5.21	9.99	52.39	1.27	757	0.017
66	UM0097001	01	SPECS	AG-690/11972457	50.52	77.46	22.17	64.38	37.47	87.78	-38.28	67.78	4.54	58.51	5.92	1233	0.617
67	UM0074324	02	SPECS	AK-968/37055100	31.29	37.41	6.83	16.36	22.95	61.79	1.09	8.90	10.69	2.61	4.67	1193	1.000
68	UM0007091	02	SPECS	AK-968/11165178	57.54	82.61	17.83	79.56	28.10	89.68	14.45	80.27	5.89	81.72	4.61	1032	0.381
69	UM0091821	01	SPECS	AA-768/34978021	77.38	86.93	18.32	72.42	49.85	91.91	-0.61	62.73	7.18	67.88	30.13	831	0.866
70	UM0100032	01	SPECS	AK-918/36937003	46.12	46.18	0.06	56.69	22.36	62.75	5.50	38.39	-28.95	58.72	5.83	1133	0.895
71	UM0007012	02	SPECS	AK-968/11164008	41.57	76.66	0.10	47.69	26.24	76.59	-2.74	22.97	-6.09	29.41	4.26	1195	0.620
72	UM0092622	01	SPECS	AE-848/32013001	59.39	89.58	12.98	64.73	40.47	93.63	-13.07	20.20	1.93	48.04	4.61	1133	0.374
73	UM0094671	01	SPECS	AG-205/33681021	63.62	64.67	17.70	53.74	61.07	86.87	14.84	43.80	12.53	60.89	4.74	1166	0.918
74	UM0012934	01	CHEMBRIDGE	5627239	57.14	79.85	-17.08	62.42	46.92	89.77	5.59	53.85	7.55	57.95	6.08	1066	0.726
75	UM0018211	01	CHEMBRIDGE	6104940	33.13	63.27	-20.12	46.85	24.23	69.66	0.50	24.45	-5.25	20.92	4.13	1028	0.913
76	UM0021111	01	CHEMBRIDGE	6444619	36.23	86.67	-25.66	71.84	28.13	79.68	2.98	24.94	1.10	21.27	4.63	1102	0.992
77	UM0028721	01	CHEMBRIDGE	7638598	34.94	70.82	11.65	66.48	23.10	68.69	0.66	31.04	-0.51	27.16	4.73	1061	1.000
78	UM0030219	01	CHEMBRIDGE	7661641	38.19	81.82	-6.39	47.50	20.02	52.04	5.19	-9.88	-1.19	-10.17	3.26	1158	0.714
79	UM0031037	01	CHEMBRIDGE	7677876	30.97	85.21	-33.90	85.58	33.17	75.37	14.03	16.72	0.90	17.29	4.67	1033	0.610
80	UM0033912	01	CHEMBRIDGE	7730387	58.62	83.13	-26.68	76.10	48.50	88.73	6.61	51.14	10.29	60.54	4.08	1079	0.293
81	UM0036576	01	CHEMBRIDGE	7783341	39.52	85.34	-49.88	66.67	35.26	82.85	1.40	43.11	2.04	36.38	4.88	943	0.540
82	UM0037150	01	CHEMBRIDGE	7797334	32.75	91.18	-7.30	87.22	20.83	87.73	-21.33	68.55	-4.05	66.89	5.46	1099	0.717
83	UM0035177	01	CHEMBRIDGE	7748948	65.55	95.99	31.78	73.09	67.35	91.06	2.84	47.17	8.72	47.08	5.53	1249	1.000
84	UM0006319	01	CHEMBRIDGE	5332028	49.16	88.80	42.57	87.02	25.03	87.25	-3.78	80.90	-15.37	79.45	6.12	906	0.901
85	UM0006445	01	CHEMBRIDGE	5335954	35.60	76.35	-55.12	85.87	23.46	82.55	0.97	72.99	-1.99	73.05	5.48	1057	0.997
86	UM0007481	01	CHEMBRIDGE	5356835	48.16	47.44	-19.74	57.89	22.94	62.97	-0.99	25.56	-5.93	22.95	5.12	1099	0.886
87	UM0021300	01	CHEMBRIDGE	6464887	61.67	64.02	-29.49	82.11	25.43	71.46	-2.68	11.01	-5.72	28.44	4.16	1175	0.798
88	UM0032114	01	CHEMBRIDGE	7693232	34.75	90.60	-36.49	84.35	24.24	91.70	-6.13	77.82	-9.37	82.92	4.57	1120	0.539
89	UM0024677	02	MAYBRIDGE	BTB 04957	65.89	83.32	-2.21	60.82	37.03	90.16	2.91	44.38	2.11	51.19	5.10	1080	0.576

Cpd #	ID	Batch	Supplier	Supplier ID	Primary screen Plk1-Map (stable)		Primary screen CRAF-BRAF (transient)		Confirmation Plk1-Map (stable)		Confirmation Plk1-Map (transient)		Confirmation CRAF-BRAF (transient)		Mitotic index		Co-IP
					% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	% Inhibition	% Quench RLuc	Mitotic index (%)
90	UM0026675	01	CHEMBRIDGE	7573944	31.54	93.69	-34.02	86.30	24.87	84.92	1.54	72.23	16.47	76.14	4.73	994	0.865
91	UM0040880	01	MAYBRIDGE	BTB 02165	31.42	80.31	-62.27	88.13	21.59	87.71	-6.44	81.17	-1.92	82.34	5.46	938	0.874
92	UM0024677	01	CHEMBRIDGE	7182303	31.73	89.49	-2.91	71.76	37.27	88.36	-7.35	35.36	-4.79	40.97	5.01	1114	1.000
93	UM0043471	01	MAYBRIDGE	CD 09636	45.16	88.06	-19.74	80.17	26.62	85.18	-8.24	58.20	-2.23	66.84	5.09	1026	0.841
94	UM0044029	01	MAYBRIDGE	DP 01660	51.69	94.69	3.78	88.15	44.21	87.91	35.12	87.25	17.09	86.44	4.83	1089	0.860
95	UM0045661	01	MAYBRIDGE	HTS 05195	45.78	92.59	22.52	68.30	24.98	86.69	-10.29	55.66	18.00	64.03	5.31	1028	0.688
96	UM0046655	01	MAYBRIDGE	HTS 10845	59.02	55.37	10.66	4.08	20.78	62.69	3.53	-18.74	11.95	-26.33	4.80	1011	0.422
97	UM0046929	01	MAYBRIDGE	HTS 12603	46.44	-15.66	-4.09	7.76	27.48	64.10	6.60	-1.90	10.94	-8.01	5.52	1064	0.924
98	UM0048554	01	MAYBRIDGE	KM 04938	61.42	73.36	9.17	35.83	30.73	72.21	11.50	-7.56	7.06	6.67	5.60	912	1.000
99	UM0053488	01	MAYBRIDGE	SCR 00880	64.82	96.93	-6.51	86.09	43.59	95.67	-1.55	69.68	-0.16	73.58	4.14	1035	0.535
100	UM0055888	01	MAYBRIDGE	SPB 06986	56.28	70.67	-5.98	49.29	21.68	81.71	36.75	83.83	10.77	85.00	3.46	1091	0.906
101	UM0057305	01	CHEMBRIDGE	7973027	58.20	71.17	11.87	24.48	25.59	81.25	-2.26	5.22	-1.25	8.14	4.90	1082	0.693
102	UM0057777	01	CHEMBRIDGE	7994213	53.01	86.51	-16.54	76.76	22.75	89.27	1.44	29.97	0.02	26.31	5.04	1156	0.955
103	UM0057973	01	CHEMBRIDGE	7998959	37.22	87.50	-4.16	82.45	23.14	87.15	-4.15	54.58	0.96	63.97	5.09	1269	0.873
104	UM0057989	01	CHEMBRIDGE	7999397	39.98	76.07	-14.54	44.82	21.84	74.15	3.12	3.89	2.23	7.27	4.60	1136	1.000
105	UM0061111	01	CHEMBRIDGE	9032660	34.88	91.43	-17.93	73.39	23.41	91.83	-7.42	66.08	-3.58	70.37	4.26	1151	0.913
106	UM0061877	01	CHEMBRIDGE	9037275	33.90	88.43	2.59	15.25	36.57	82.59	-5.15	17.65	4.63	4.64	4.67	1038	0.760
107	UM0063060	01	CHEMBRIDGE	9043703	35.84	92.37	-35.90	89.70	23.41	94.03	-6.99	88.59	3.94	89.06	5.67	1132	0.605
108	UM0064980	01	CHEMBRIDGE	9053935	49.87	77.46	1.34	75.29	27.61	91.79	-7.08	69.97	2.78	65.83	4.53	1115	0.353
109	UM0065245	01	CHEMBRIDGE	9055626	30.40	48.25	0.23	45.69	21.21	77.52	-4.20	43.98	-1.50	40.13	4.24	1303	0.457
110	UM0065246	01	CHEMBRIDGE	9055637	60.41	86.89	20.65	86.28	31.50	92.21	-16.97	73.64	1.99	74.92	4.62	1078	0.759
111	UM0067000	01	CHEMBRIDGE	5134355	57.76	73.12	25.91	66.49	36.70	62.66	11.76	35.42	17.99	39.64	4.74	1312	0.133
112	UM0067184	01	CHEMBRIDGE	5144173	57.92	71.84	0.00	55.88	32.29	74.81	7.79	10.94	5.72	32.74	5.62	1358	0.960
113	UM0074300	01	CHEMBRIDGE	5676580	67.59	86.55	27.85	82.94	50.01	93.10	15.89	81.59	14.86	87.48	5.03	1266	1.000
114	UM0126848	01	UdeM	UDM-000144	31.05	72.96	22.02	52.38	20.34	72.77	-4.60	17.86	-1.60	29.77	5.74	1089	0.824
115	UM0127131	01	UdeM	UDM-000770	62.96	77.03	35.58	89.67	22.78	80.34	-12.42	18.67	3.87	23.45	6.01	1152	0.644
116	UM0127134	01	UdeM	UDM-000775	73.20	82.73	20.93	83.07	45.22	85.88	-14.02	31.94	1.20	37.47	5.34	1046	0.427
117	UM0127135	01	UdeM	UDM-000776	65.86	88.45	29.57	86.80	42.50	88.81	-15.83	29.77	5.21	45.01	4.99	1161	0.526
118	UM0127153	01	UdeM	UDM-000933	45.02	65.77	-0.56	75.63	23.49	75.56	-12.10	25.09	-5.41	36.92	5.11	1229	0.926
119	UM0127154	01	UdeM	UDM-000934	69.73	78.13	30.28	81.67	28.42	83.23	-9.75	51.09	5.62	60.39	5.09	1122	0.917
120	UM0127191	01	UdeM	UDM-001223	47.30	72.72	25.58	69.47	25.29	76.88	-13.60	36.57	6.13	47.60	5.09	1163	1.000
121	UM0011080	01	CHEMBRIDGE	5541594	63.35	73.89	-6.16	36.21	38.10	75.39	-10.29	6.91	2.84	5.90	4.87	841	0.955
122	UM0052184	01	SPECS	AA-768/33245010	44.05	86.60	-8.30	44.16	30.57	79.14	-1.93	25.69	3.03	35.15	NT	NT	0.776

NT= Non tested

Table S3. Structure-Activity Relationship (SAR) analysis of Cpd 16. Chemical structure and IC<sub>50</sub> values obtained in FP for the different analogs of Cpd 16 are presented.

Cpd #	UM number	Molecular mass	Molecular formula	Structure	FP - IC <sub>50</sub> (μM)
16	UM0113574	407.79	C <sub>16</sub> H <sub>13</sub> ClF <sub>3</sub> NO <sub>4</sub> S		18.4 ± 10.2
123	UM0131803	373.35	C <sub>16</sub> H <sub>14</sub> F <sub>3</sub> NO <sub>4</sub> S		7.2
124	UM0131783	291.32	C <sub>14</sub> H <sub>13</sub> NO <sub>4</sub> S		68.6 ± 18.5
125	UM0131784	393.76	C <sub>15</sub> H <sub>11</sub> ClF <sub>3</sub> NO <sub>4</sub> S		11.9 ± 7.2
126	UM0131785	360.21	C <sub>14</sub> H <sub>11</sub> Cl <sub>2</sub> NO <sub>4</sub> S		4.3
127	UM0131786	325.77	C <sub>14</sub> H <sub>12</sub> ClNO <sub>4</sub> S		187.4
128	UM0131787	393.76	C <sub>15</sub> H <sub>11</sub> ClF <sub>3</sub> NO <sub>4</sub> S		8.4
129	UM0041779	359.32	C <sub>15</sub> H <sub>12</sub> F <sub>3</sub> NO <sub>4</sub> S		8.5
130	UM0113524	428.21	C <sub>15</sub> H <sub>10</sub> Cl <sub>2</sub> F <sub>3</sub> NO <sub>4</sub> S		7.8
131	UM0113530	360.21	C <sub>14</sub> H <sub>11</sub> Cl <sub>2</sub> NO <sub>4</sub> S		70.5
132	UM0113515	415.43	C <sub>19</sub> H <sub>20</sub> F <sub>3</sub> NO <sub>4</sub> S		26.1
133	UM0113549	370.76	C <sub>14</sub> H <sub>11</sub> ClN <sub>2</sub> O <sub>6</sub> S		5.4
15	UM0113559	350.78	C <sub>15</sub> H <sub>11</sub> ClN <sub>2</sub> O <sub>4</sub> S		5.7 ± 2.8
136	UM0131792	293.34	C <sub>14</sub> H <sub>15</sub> NO <sub>4</sub> S		No inhibition
137	UM0131779	338	C <sub>16</sub> H <sub>16</sub> ClNO <sub>3</sub> S		No inhibition

Cpd #	UM number	Molecular mass	Molecular formula	Structure	FP - IC <sub>50</sub> (μM)
138	UM0032816	387.28	C <sub>17</sub> H <sub>16</sub> Cl <sub>2</sub> O <sub>4</sub> S		No inhibition
139	UM0131782	391.23	C <sub>14</sub> H <sub>12</sub> Cl <sub>2</sub> N <sub>2</sub> O <sub>3</sub> S		No inhibition
140	UM0131780	337.82	C <sub>16</sub> H <sub>16</sub> ClNO <sub>3</sub> S		No inhibition
141	UM0131777	372.27	C <sub>16</sub> H <sub>15</sub> Cl <sub>2</sub> NO <sub>3</sub> S		No inhibition
142	UM0041694	392.78	C <sub>15</sub> H <sub>12</sub> ClF <sub>3</sub> N <sub>2</sub> O <sub>3</sub> S		No inhibition
143	UM0131778	429.31	C <sub>16</sub> H <sub>10</sub> F <sub>7</sub> NO <sub>3</sub> S		No inhibition
144	UM0113371	387.33	C <sub>15</sub> H <sub>12</sub> F <sub>3</sub> N <sub>3</sub> O <sub>4</sub> S		No inhibition
145	UM0109778	382.27	C <sub>16</sub> H <sub>16</sub> BrNO <sub>3</sub> S		No inhibition
146	UM0131761	237.24	C <sub>9</sub> H <sub>10</sub> F <sub>3</sub> NOS		No inhibition
147	UM0131763	375.79	C <sub>16</sub> H <sub>13</sub> ClF <sub>3</sub> NO <sub>2</sub> S		No inhibition
148	UM0131764	391.79	C <sub>16</sub> H <sub>13</sub> ClF <sub>3</sub> NO <sub>3</sub> S		No inhibition
149	UM0131772	425.35	C <sub>17</sub> H <sub>13</sub> F <sub>6</sub> NO <sub>3</sub> S		No inhibition
150	UM0131773	355.37	C <sub>17</sub> H <sub>16</sub> F <sub>3</sub> NO <sub>2</sub> S		No inhibition
151	UM0131774	457.34	C <sub>17</sub> H <sub>13</sub> F <sub>6</sub> NO <sub>5</sub> S		41.7



Cpd #	UM number	Molecular mass	Molecular formula	Structure	FP - IC <sub>50</sub> (μM)
152	UM0131775	387.37	C <sub>17</sub> H <sub>16</sub> F <sub>3</sub> NO <sub>4</sub> S		54.4
153	UM0131776	420.25	C <sub>16</sub> H <sub>13</sub> BrF <sub>3</sub> NO <sub>2</sub> S		No inhibition
154	UM0131790	452.24	C <sub>16</sub> H <sub>13</sub> BrF <sub>3</sub> NO <sub>4</sub> S		36.8
155	UM0131794	410.24	C <sub>16</sub> H <sub>12</sub> Cl <sub>2</sub> F <sub>3</sub> NO <sub>2</sub> S		No inhibition
156	UM0131795	442.24	C <sub>16</sub> H <sub>12</sub> Cl <sub>2</sub> F <sub>3</sub> NO <sub>4</sub> S		48.1
157	UM0131805	361.81	C <sub>16</sub> H <sub>15</sub> ClF <sub>3</sub> NOS		No inhibition
158	UM0131806	269.24	C <sub>9</sub> H <sub>10</sub> F <sub>3</sub> NO <sub>3</sub> S		No inhibition
159	UM0131811	393.81	C <sub>16</sub> H <sub>15</sub> ClF <sub>3</sub> NO <sub>3</sub> S		No inhibition
160	UM0131831	379.4	C <sub>16</sub> H <sub>20</sub> F <sub>3</sub> NO <sub>4</sub> S		75.0
161	UM0131842	251.23	C <sub>9</sub> H <sub>8</sub> F <sub>3</sub> NO <sub>2</sub> S		8.3 ± 2.6
164	UM0131845	347.33	C <sub>10</sub> H <sub>12</sub> F <sub>3</sub> NO <sub>5</sub> S <sub>2</sub>		6.0
168	UM0131878	347.4	C <sub>16</sub> H <sub>20</sub> F <sub>3</sub> NO <sub>2</sub> S		No inhibition
169	UM0132247	253.24	C <sub>9</sub> H <sub>10</sub> F <sub>3</sub> NO <sub>2</sub> S		No inhibition