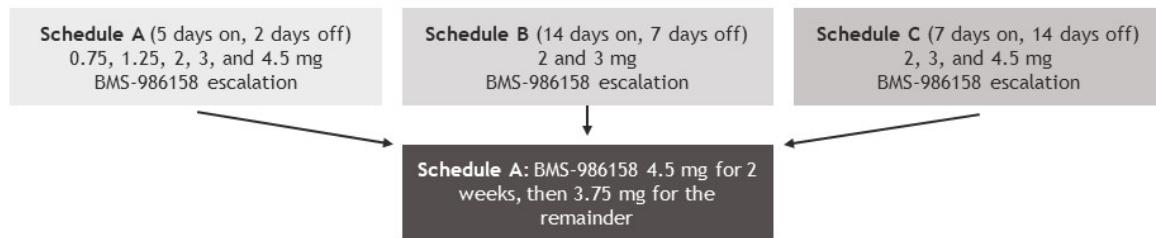

Supplemental Materials

Title: BMS-986158, a Small Molecule Inhibitor of the Bromodomain and Extraterminal Domain Proteins, in Patients With Selected Advanced Solid Tumors: Results From a Phase 1/2a Trial

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Part 1: dose escalation in patients with TNBC, ovarian cancer, SCLC, or other selected solid tumors



Part 2: dose expansion at RP2D

Initial Expansion

Fusion proteins: double hit lymphoma, NUT-midline carcinoma
BRD amplifications: TNBC
MYC amplifications: Non-GC-DLBL
AR amplifications: CRPC



Subsequent Expansion

Fusion proteins: double hit lymphoma, NUT-midline carcinoma, Ewing sarcoma, Burkitt's lymphoma
BRD amplifications: TNBC, ovarian cancer, neuroendocrine prostate cancer
MYC amplifications: Non-GC-DLBL, ovarian cancer, uterine carcinosarcoma
AR amplifications: CRPC
Other mutations: RCC,^a NSCLC,^b uveal melanoma^c

Figure S1. Study design. ^aRenal cell carcinoma with *SWI/SNF* mutations; ^bNon-small cell lung cancer with *SWI/SNF* or *KRAS/wtTK11*; ^cUveal melanoma with *Gnaq11*. AR, androgen receptor; BRD, bromodomain; CRPC, castration-resistant prostate cancer; Non-GC-DLBL, non-germinal center diffuse large B-cell lymphoma; NSCLC, non-small cell lung cancer; NUT, nuclear protein in testis; RCC, renal cell carcinoma; RP2D, recommended phase 2 dose; SCLC, small cell lung cancer; TNBC, triple-negative breast cancer.

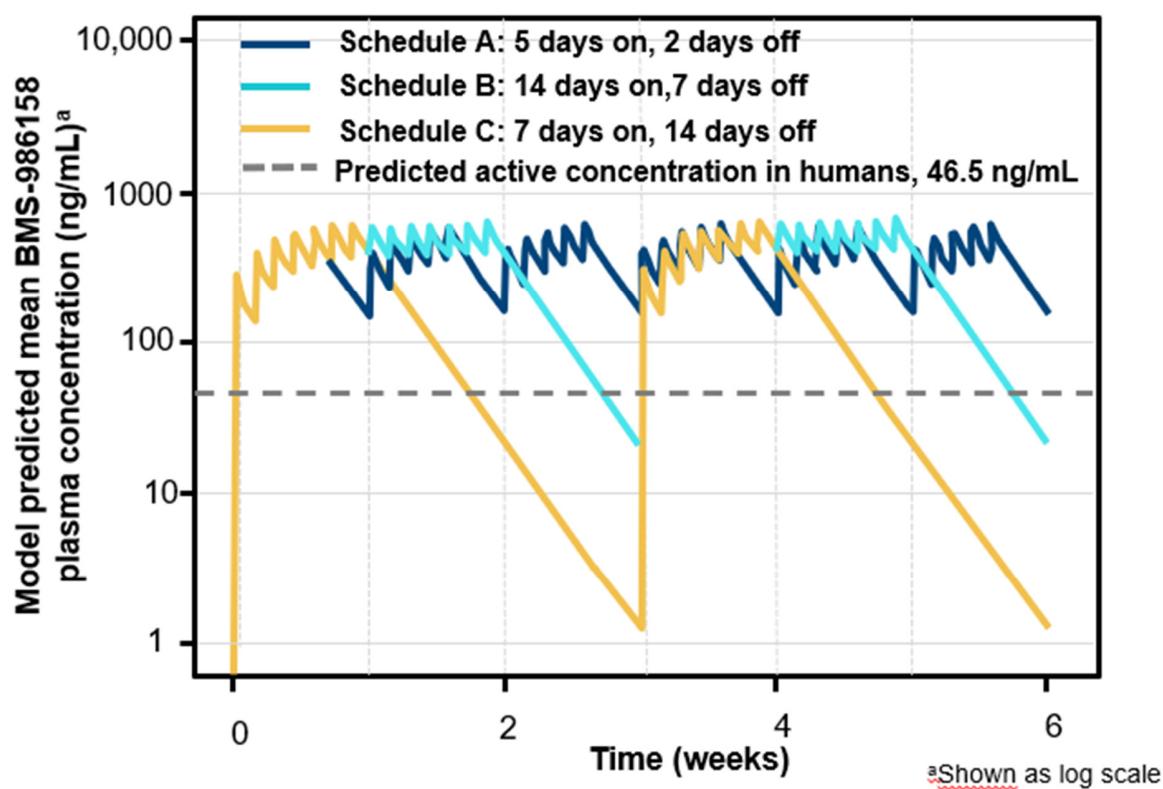
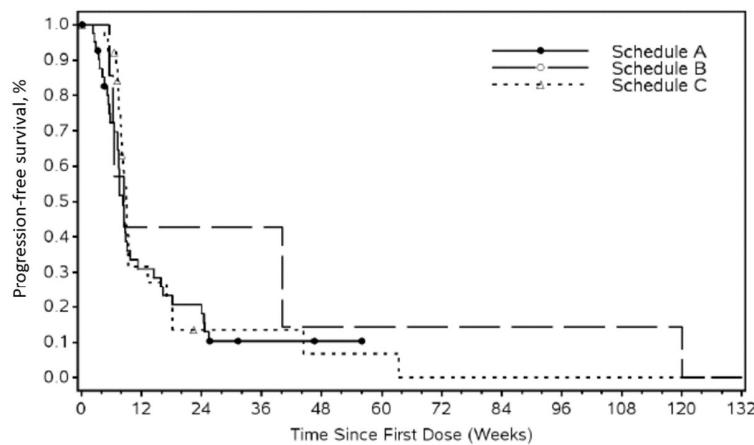


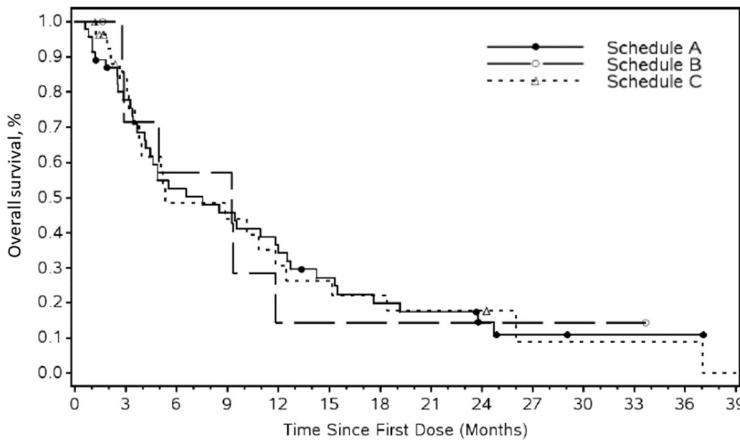
Figure S2. Predicted pharmacokinetic profile at BMS-986158 3-mg dose for each treatment schedule.

A)



Schedule	Number at risk												
A	46	12	8	2	1	0	0	0	0	0	0	0	0
B	8	3	3	3	1	1	1	1	1	1	1	1	0
C	29	7	2	2	1	1	0	0	0	0	0	0	0

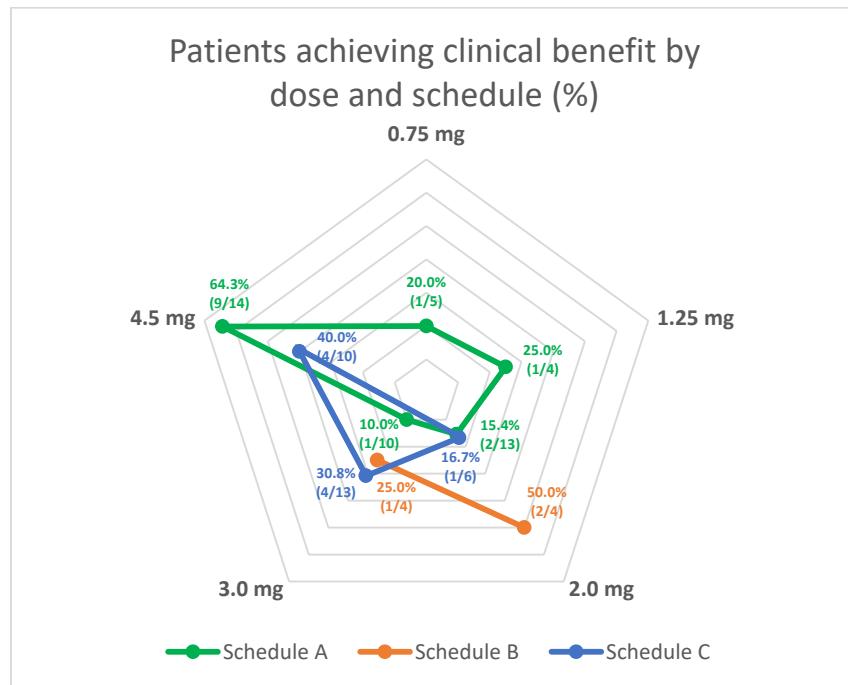
B)



Schedule	Number at risk												
A	46	34	23	20	15	11	8	7	4	2	1	1	1
B	8	5	4	4	1	1	1	1	1	1	1	1	0
C	29	19	11	10	7	6	5	4	4	1	1	1	0

Figure S3. Kaplan-Meier survival curves by treatment schedule for A) PFS and B) OS.

A)



B)

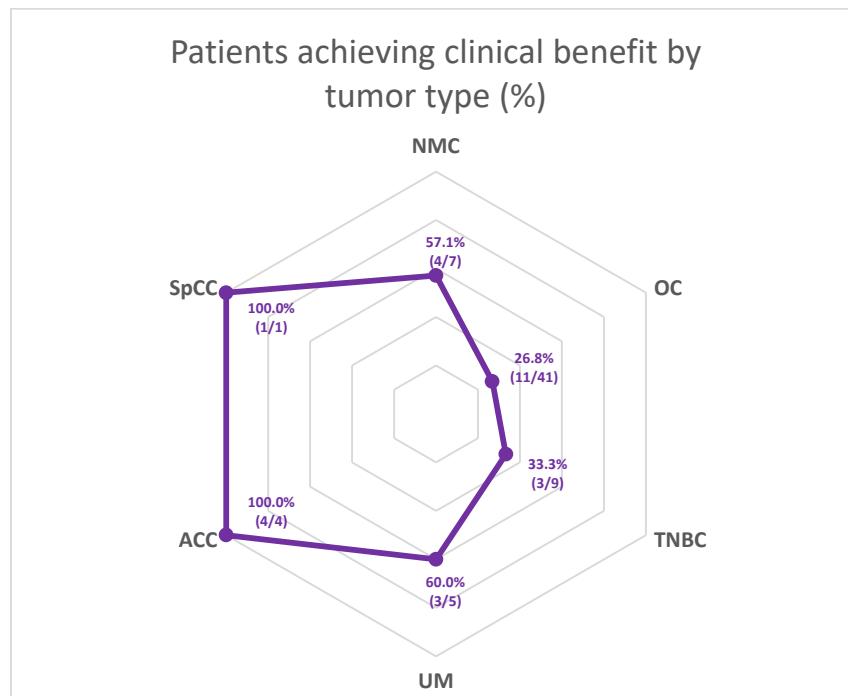


Figure S4. Patients who achieved clinical benefit (SD or better) by A) dose and schedule and B) tumor type.

Table S1. Pharmacokinetics of BMS-986158 and its metabolite at single dose by treatment schedule.

Pharmacokinetic parameter	Assigned to the cohort receiving:									
	Schedule A					Schedule B		Schedule C		
	0.75 mg n = 5	1.25 mg n = 3	2.0 mg n = 9	3.0 mg n = 10	4.5 mg n = 10	2.0 mg n = 1	3.0 mg n = 4	2.0 mg n = 6	3.0 mg n = 11	4.5 mg n = 7
BMS-986158										
C _{max} (ng/mL), GM (%CV)	68.8 (23)	175 (36)	260 (20)	328 (36)	478 (24)	207	481 (25)	295 (29)	370 (22)	567 (24)
T _{max} (h), median (range)	4.00 (2.00–4.02)	1.00 (1.00–2.00)	2.00 (1.00–4.03)	2.00 (0.98–6.15)	2.04 (1.00–4.03)	1.00 (1.00–4.03)	1.03 (1.00–4.03)	1.00 (0.50–2.03)	1.00 (0.50–2.02)	2.02 (1.00–4.00)
AUC _(0–24) (h×ng/mL), GM (%CV)	1027 (16)	2309 (13)	3610 (22)	4942 (43)	6786 (24)	2358	6921 (28)	3660 (27)	4468 (29)	7418 (43)
AUC _(0–T) (h×ng/mL), GM (%CV)	2150 (15)	6372 (14)	8564 (48)	10,452 (50)	19,124 (46)	5202	28,852 (41)	10,931 (43)	11,493 (54)	17,220 (84)
Effective T _{1/2} (h), mean (SD)	33.7 (1.4) ^a	48.7 (6.7)	41.4 (18.2) ^a	33.3 (7.6) ^b	48.3 (13.4) ^c	44.2	54.0 ^d	64.6 (17.5)	48.2 (23.8) ^e	35.7 (17.8) ^f
Vz/F (L), GM (%CV)	14.7 (21) ^a	12.5 (18)	14.6 (15) ^a	12.8 (25) ^b	15.7 (29) ^c	23.2	15.3 ^d	13.9 (29)	15.6 (28) ^e	11.6 (17) ^f
Metabolite of BMS-986158										
C _{max} (ng/mL), GM (%CV)	5.0 (30)	10.0 (29)	16.9 (32) ^g	22.5 (38) ^c	30.0 (41)	13.6	26.3 (63)	21.1 (35)	19.4 (44)	45.5 (32)
MR C _{max} , GM (%CV)	0.07 (14)	0.06 (9)	0.07 (30) ^g	0.07 (34) ^c	0.06 (45)	0.07	0.06 (42)	0.07 (48)	0.05 (34)	0.08 (25)
T _{max} (h), median (range)	24.0 (2.6–24.2)	2.0 (2.0–24.0)	5.1 (2.0–72.0) ^g	23.9 (1.0–48.0) ^c	15.1 (1.0–48.0)	1.00 (1.0–47.4)	1.52 (1.0–71.5)	14.9 (1.0–48.0)	2.02 (1.0–45.6)	6.27 (1.0–45.6)
MR AUC _(0–24) (h×ng/mL), GM (%CV)	0.10 (29)	0.08 (7)	0.08 (24) ^g	0.09 (44) ^c	0.08 (40)	0.08	0.06 (33)	0.10 (18)	0.08 (35)	0.10 (41)
MR AUC _(0–T) (h×ng/mL), GM (%CV)	0.15 (33)	0.12 (29)	0.12 (21) ^g	0.14 (38) ^c	0.14 (37)	0.08	0.10 (28)	0.17 (35)	0.11 (41)	0.15 (48)

^an = 4; ^bn = 7; ^cn = 9; ^dn = 1; ^en = 10; ^fn = 5; ^gn = 8.

AUC_(0–24), area under the plasma concentration-time curve from time 0 to time 24 hours post dose; AUC_(0–T), area under the plasma concentration-time curve from time 0 to time of the last quantifiable concentration; CL/F, apparent total body clearance; C_{max},

maximum observed plasma concentration; %CV, coefficient of variation; GM, geometric mean; MR, metabolite-to-parent; N/A, not available; $T_{1/2}$, half-life; T_{\max} , time to C_{\max} ; V_z/F , apparent volume of distribution.

Table S2. Baseline treatment assignments for BMS-986158 monotherapy and tumor type ^a.

Characteristic	Patients who achieved clinical benefit, n/N (%) ^a
All safety population	26/83 (31.3)
Schedule A, all doses	14/46 (30.4)
0.75 mg	1/5 (20.0)
1.25 mg	1/4 (25.0)
2.0 mg	2/13 (15.4)
3.0 mg	1/10 (10.0)
4.5 mg	9/14 (64.3)
Schedule B, all doses	3/8 (37.5)
2.0 mg	2/4 (50.0)
3.0 mg	1/4 (25.0)
Schedule C, all doses	9/29 (31.0)
2.0 mg	1/6 (16.7)
3.0 mg	4/13 (30.8)
4.5 mg	4/10 (40.0)
Tumor type, n (%)	
NUT carcinoma	4/7 (57.1)
Ovarian cancer	11/41 (26.8)
TNBC	3/9 (33.3)
UM	3/5 (60.0)
ACC	4/4 (100.0)
SpCC	1/1 (100.0)

^aPercentage of patients with SD or better calculated as number of patients with SD or better divided by number of patients in the safety population with the same characteristic multiplied by 100.

ACC, adenoid cystic carcinoma; NUT, nuclear protein in testis; QD, once daily; SpCC, spindle cell carcinoma; TNBC, triple-negative breast cancer, UM, uveal melanoma.