Supplementary Material

Analysis of a real-world progression variable and related endpoints for patients with five different cancer types

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Supplemental Table 1: rwEndpoint definitions

	Index date	Event date	Censor date
rwOS		Patient's date of death due to any cause	Last visit ^b or last abstracted oral episode of drug(s)
rwPFS		Date of first rwP event (abstracted) that occurs more than 14 days after index date or date of death, whichever occurs first.	Last unstructured clinic note assessed for rwP
rwTTP	Start of 1L therapy ^a	Date of first rwP event (abstracted) that occurs more than 14 days after index date.	Last unstructured clinic note assessed for rwP
rwTTD		Last administration or non-cancelled order (IV drugs) or last abstracted oral episode of drug(s) contained in the index therapy. Permanent discontinuation was defined as meeting one of the following conditions: start of index therapy, a gap in treatment after the end of index therapy or death.	Last administration or non-cancelled order (IV drugs) or last abstracted oral episode of drug(s)
rwTTNT		Date of the start of line of therapy subsequent to index therapy or date of death, whichever occurs first.	Last visit ^b

^aIndex date was 2L therapy for the aMel cohort

Abbreviations: 1L, first line; 2L, second line; aMel, advanced melanoma; EHR, electronic health record; IV, intravenous; rwOS, real-world overall survival; rwP, real-world progression; rwPFS, real-world progression-free survival; rwTTD, real-world time-to-discontinuation; rwTTP, real-world time-to-progression; rwTTNT, real-world time-to-next treatment.

bVisit categorized as vitals (any date on which a patient had a structured vitals measurement in the EHR); treatment (any date on which a patient had a structured medication administration in the EHR); lab (any date on which a patient had a structured lab in the EHR)

Supplemental Table 2: Baseline table for each analytic cohort by disease

Table 2a: mBC cohort

Characteristic	mBC cohort N=884
Age at metastatic diagnosis (years), median [IQR]	64.0 [54.0;74.0]
Age at metastatic diagnosis category (years), n (%)	
<35	8 (0.9)
35–44	127 (14.4)
45–54	315 (35.6)
55–64	228 (25.8)
≥65	206 (23.3)
Sex, n (%)	
Female	870 (98.4)
Male	14 (1.6)
Region, n (%)	
Northeast	201 (22.7)
Midwest	149 (16.9)
South	315 (35.6)
West	161 (18.2)
Other/Missing	58 (6.6)
Practice type, n (%)	
Academic	44 (5.0)
Community	840 (95.0)

Race, n (%)	
White	607 (68.7)
Black	86 (9.7)
Asian	12 (1.4)
Other Race	126 (14.3)
Unknown/missing	53 (6.0)
Stage at diagnosis, n (%)	
Stage I	84 (9.5)
Stage II	216 (24.4)
Stage III	180 (20.4)
Stage IV	282 (31.9)
Stage not reported	122 (13.8)
ER tested, n (%)	868 (98.2)
ER status among those tested, n (%)	
Negative	225 (25.9)
Positive	642 (74.0)
Results unknown/not documented	1 (0.1)
PR tested, n (%)	868 (98.2)
PR status among those tested, n (%)	1
Negative	344 (39.6)
Positive	485 (55.9)
Results unknown/not documented	39 (4.5)

HER2 tested, n (%)	841 (95.1)	
HER2 status among those tested, n (%)		
Negative	650 (77.3)	
Positive	148 (17.6)	
Equivocal	33 (4.0)	
Results unknown/not documented	10 (1.3)	
HR/HER2 status among those tested, n (%)	•	
HR- and HER2+	45 (5.1)	
HR+ and HER2-	508 (57.5)	
HR+ and HER2+	100 (11.3)	
Triple Negative	132 (14.9)	
HR/HER2 status unknown/not documented	99 (11.2)	
First-line therapy class, n (%)	,	
CDK4/6 inhibitor-based therapy	0 (0.0)	
Chemo + HER2 targeted-based therapy	95 (10.7)	
Chemotherapy	351 (39.7)	
Clinical study drug-based therapies	3 (0.3)	
Endocrine therapy	376 (42.5)	
HER2 targeted-based therapy	32 (3.6)	
mTOR inhibitor-based therapy	11 (1.2)	
Other therapies	16 (1.8)	

Abbreviations: CDK, cyclin-dependent kinase; ER, estrogen receptor, HER2, human epidermal growth factor receptor 2; HR, hormone receptor; IQR, interquartile range; mBC, metastatic breast cancer; mTOR, mammalian target of rapamycin; PR, progesterone receptor.

Table 2b: aMel cohort

	aMel cohort N = 152
Age at advanced diagnosis (years), median [IQR]	64.0 [56.0; 75.0]
Age at advanced diagnosis category (years), n (%)	
<35	7 (4.6)
35–44	16 (10.5)
45–54	56 (36.8)
55–64	32 (21.1)
≥65	41 (27.0)
Sex, n (%)	
Female	50 (32.9)
Male	102 (67.1)
Region, n (%)	
Northeast	24 (19.7)
Midwest	23 (15.8)
South	63 (41.4)
West	12 (7.9)
Other/Missing	30 (19.7)
Practice type, n (%)	•
Academic	29 (19.1)
Community	123 (80.9)
Race, n (%)	•
White	127 (83.6)

Black	1 (0.7)
Asian	0 (0.0)
Other Race	12 (7.9)
Unknown/Not Documented	12 (7.9)
Stage at diagnosis, n (%)	
Stage I	10 (6.6)
Stage II	33 (21.7)
Stage III	29 (19.1)
Stage IV	42 (27.6)
Group stage is not reported	38 (25.0)
BRAF tested, n (%)	147 (96.7)
BRAF status among those tested, n (%)	
Mutation negative	71 (48.3)
Mutation positive	75 (51.0)
Results unknown/not documented	1 (0.7)
KIT tested, n (%)	45 (29.6)
KIT status among those tested, n (%)	
Mutation negative	39 (86.7)
Mutation positive	5 (11.1)
Results unknown/not documented	1 (2.2)
PD-L1 tested, n (%)	10 (6.6)
PD-L1 status among those tested, n (%)	
PD-L1 negative/not detected	3 (30.0)
PD-L1 positive	2 (20.0)
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Results unknown/not documented	2 (20.0)
No interpretation given in report	3 (30.0)
NRAS tested, n (%)	43 (28.3)
NRAS status among those tested, n (%)	
Mutation negative	33 (76.7)
Mutation positive	9 (20.9)
Results unknown/not documented	1 (2.3)
First-line therapy class, n (%)	•
(Peg)interferon-based therapies	0 (0.0)
BRAF/MEK-based therapies	36 (23.7)
Chemotherapy only (single-agent or combos)	20 (13.2)
Clinical study drug-based therapies	2 (1.3)
Modified viral therapies	1 (0.7)
PD-1/PD-L1-based therapies	56 (36.8)
Yervoy + PD-1/PD-L1-based therapies	10 (6.6)
Yervoy-based therapies	22 (14.5)
Other therapies	5 (3.3)

Abbreviations: aMel, advanced melanoma; IQR, interquartile range; PD-1, programmed cell death protein 1; PD-L1, programmed death-ligand 1.

Table 2c: SCLC cohort

	SCLC cohort N=359
Age at initial diagnosis (years), median [IQR]	68.0 [60.0;74.5]
Age categories at initial diagnosis (years), n (%)	
<35	1 (0.28)
35–49	16 (4.5)
50–64	125 (34.8)
≥65	217 (60.4)
Sex, n (%)	
Female	178 (49.6)
Male	181 (50.4)
Region, n (%)	
Northeast	81 (22.6)
Midwest	78 (21.7)
South	132 (36.8)
West	45 (12.5)
Other/Missing	23 (6.4)
Practice type, n (%)	
Academic	22 (6.1)
Community	337 (93.9)
Race, n (%)	
White	250 (69.6)
Black	25 (7.0)
Asian	1 (0.3)
Other Race	35 (9.7)

Unknown/missing	48 (13.4)
Smoking status, n (%)	
History of smoking	352 (98.1)
No history of smoking	7 (1.9)
Unknown/not documented	0 (0.0)
Stage at diagnosis, n (%)	
Stage I	10 (2.8)
Stage II	11 (3.1)
Stage III	58 (16.2)
Stage IV	199 (55.4)
Stage not reported	81 (22.6)
SCLC-specific stage at diagnosis, n (%)	
Extensive disease	231 (64.3)
Limited disease	105 (29.2)
Unknown/not documented	23 (6.4)
Resection, n (%)	•
Yes	10 (2.8)
No/unknown	349 (97.2)
Upfront radiation therapy, n (%)	
Yes	123 (34.3)
No/unknown	236 (65.7)
First-line therapy class, n (%)	
Platinum + Etoposide	339 (94.4)
Other	19 (5.3)

Unknown/not documented	1 (0.3)
Abbreviations: IQR, interquartile range; SCLC, small cell lung cancer.	

Table 2d: mRCC cohort

	mRCC cohort N=323
Age at metastatic diagnosis (years), median [IQR]	66.0 [58.0;73.0]
Age categories at metastatic diagnosis (years), n (%)	•
<35	2 (0.6)
35–49	14 (4.3)
50–64	125 (38.7)
≥65	182 (56.3)
Sex, n (%)	•
Female	86 (26.6)
Male	237 (73.4)
Region, n (%)	
Northeast	56 (17.3)
Midwest	58 (18.0)
South	119 (36.8)
West	58 (18.0)
Other/Missing	32 (9.9)
Practice type, n (%)	·
Academic	25 (7.7)
Community	298 (92.3)

Race, n (%)	
White	215 (66.6)
Black or African American	23 (7.1)
Asian	4 (1.2)
Other Race	48 (14.9)
Unknown/missing	33 (10.2)
Smoking status, n (%)	•
History of smoking	202 (62.5)
No history of smoking	119 (36.8)
Unknown/Not documented	2 (0.6)
Stage at diagnosis, n (%)	•
Stage I–III	124 (38.4)
Stage IV	189 (58.5)
Stage not reported	10 (3.1)
Histology, n (%)	·
Clear Cell	225 (69.7)
Non-clear Cell	31 (9.6)
RCC, NOS	67 (20.7)
Nephrectomy, n (%)	·
Yes, pre-metastatic	125 (38.7)
Yes, post-metastatic	75 (23.2)
Yes, date unknown	2 (0.6)
No	121 (37.5)
First-line therapy class, n (%)	•

Anti-VEGF TKI	233 (72.1)
mTOR inhibitor	39 (12.1)
PD-1/PD-L1 therapy	14 (4.3)
Other therapies	37 (11.5)

Abbreviations: IQR, interquartile range; mRCC, metastatic renal cell carcinoma; mTOR, mammalian target of rapamycin; TKI, tyrosine kinase inhibitor; NOS, not otherwise specified; VEGF, vascular endothelial growth factor.

Table 2e: aGEC cohort

	aGEC cohort N=370	
Age at advanced diagnosis date (years), median [IQR]	67.0 [58.0;74.0]	
Age categories at advanced diagnosis date (years), n (%)		
<35	2 (0.5)	
35–49	28 (7.6)	
50–64	133 (35.9)	
≥65	207 (56.0)	
Sex, n (%)		
Female	89 (24.1)	
Male	281 (75.9)	
Region, n (%)		
Northeast	28 (7.6)	
Midwest	30 (8.1)	
South	152 (41.1)	
West	160 (43.2)	

Other/missing	0 (0.0)
Practice type, n (%)	
Academic	0 (0.0)
Community	370 (100.0)
Race, n (%)	
White	264 (71.4)
Black or African American	24 (6.5)
Asian	9 (2.4)
Other Race	28 (7.6)
Unknown/missing	45 (12.2)
Smoking status, n (%)	•
History of smoking	264 (71.4)
No history of smoking	106 (28.6)
Unknown/Not documented	0 (0.0)
Stage at diagnosis, n (%)	•
Stage I	18 (4.9)
Stage II	66 (17.8)
Stage III	75 (20.3)
Stage IV	176 (47.6)
Stage not reported	35 (9.5)
Disease site, n (%)	,
Esophageal	175 (47.3)
Gastric	112 (30.3)
Gastroesophageal Junction	83 (22.4)

Histology, n (%)	
Adenocarcinoma	306 (82.7)
Adenosquamous	3 (0.8)
Squamous Cell Carcinoma	58 (15.7)
Other	2 (0.5)
Unknown/Not Documented	1 (0.3)
Surgery status, n (%)	
Surgery on or before index	55 (14.9)
Unknown surgery date	0 (0.0)
No surgery on or before index	315 (85.1)
HER2 tested, n (%)	233 (63.0)
HER2 status among those tested, n (%)	<u>, </u>
Positive	42 (18.0)
Negative	178 (76.4)
Equivocal	4 (1.7)
Results unknown/not documented	9 (3.9)
First-line therapy class, n (%)	
Chemo mono or combination	317 (85.7)
Anti-VEGF +/- chemo-based	8 (2.2)
HER2 +/- chemo-based	28 (7.6)
Clinical study drug-based	12 (3.2)
Other therapies	5 (1.4)
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Abbreviations: aGEC, advanced gastric/esophageal cancer; HER2, human epidermal growth factor receptor 2; IQR, interquartile range; VEGF, vascular endothelial growth factor.