		Overall
Variable	Statistic	n (%)
Age (yrs)	n	189
	median (IQR)	67 (60-74)
Sex	n (%)	
Female		84 (44.4%)
Race Male	n (%)	105 (55.6%)
Asian	11 (70)	3 (1.6%)
Black or African American		10 (5.3%)
Other		3 (1.6%)
Unknown, Missing, or Not Reported		50 (26.5)
White or Caucasian		123 (65.1%)
Ethnicity	n (%)	
Hispanic or Latino		3 (1.6%)
Not Hispanic or Latino		124 (65.6%)
Unknown, Missing, or Not Reported Type of Therapy	n (%)	62 (32.8%)
anti-PD-(L)1	11 (70)	102 (54.0%)
Chemo + anti-PD-(L)1		87 (46.0%)
Line of Therapy	n (%)	2. (10.0.0)
1	` /	145 (76.7%)
2		35 (18.5%)
3+		9 (4.8%)
Primary Tumor Type	n (%)	
Bladder Cancer		13 (6.9%)
Breast Cancer		10 (5.3%)
Cancer of Unknown Primary Cervical Cancer		11 (5.8%) 2 (1.1%)
Esophagogastric Cancer		17 (9.0%)
Head and Neck Cancer		9 (4.8%)
Melanoma		2 (1.1%)
Non-Small Cell Lung Cancer		121 (64.0%)
Other Cancer		2 (1.1%)
Small Cell Lung Cancer		2 (1.1%)
Cohort	n (%)	00 (47 (0))
Discovery		90 (47.6%)
PD-L1 IHC Type	n (%)	99 (52.4%)
PD-L1 22C3 CPS	11 (70)	38 (20.1%)
PD-L1 SP142 IC		12 (6.3%)
PD-L1 22C3 TPS		139 (73.5%)
PD-L1 IHC Positive (>=1%)	n (%)	
Overall		157 (83.1%)
PD-L1 22C3 CPS		37 (97.4%)
PD-L1 SP142 IC		12 (100.0%)
PD-L1 22C3 TPS	median (IQR)	108 (77.7%)
PD-L1 IHC Score Overall	median (IQK)	3.5 (1.6-6.2)
PD-L1 22C3 CPS		6 (4-25)
PD-L1 SP142 IC		2.5 (1-8.75)
PD-L1 22C3 TPS		20 (2-75)
TMB Status	n (%)	
Low		140 (74.1%)
High	(0/)	49 (25.9%)
IRS Status	n (%)	106 (56.1%)
Low High		83 (43.9%)
Iligii		(.5.5 / 5)

Demographics and tumor types for all 189 patients treated with anti-PD-(L)1 monotherapy or chemotherapy (chemo) + anti-PD-(L)1 with available quantitative clinical PD-L1 IHC reported in accompanying pathology reports, including age (in years), sex, self reported race, self reported ethnicity, type of therapy, the anti-PD-(L)1 +/. chemo line of systemic therapy, tumor type, inclusion in discovery or separate validation cohorts, PD-L1 IHC type (and scoring method), % of overall and IHC type specific PD-L1 positivity (>=1%), overall and IHC type specific IHC score, TMB status (>=10 mutations/megabase as High) and IRS status (High vs. Low). n= sample size, %= percentage, IQR = interquartile range.