

Supplemental Material

***PD-L1* gene amplification and focality: relationship with protein expression**

Supplemental Table 1 – Cohort characteristics

Characteristic		Study Cohort (N=60,793)
Median Age		66.0 years
Gender (%)	Male	28177 (46)
	Female	32587 (54)
	Unknown	29
Genomic ancestry (%)	African American	6241 (10)
	American	4483 (7)
	East Asian	2095 (3)
	European	47411 (78)
	South-East Asian	532 (1)
	Unknown	31
Site of tissue specimen for CGP (%)	Local	26483 (44)
	Metastatic	23788 (39)
	Unknown	10522 (17)
Median tissue specimen tumor purity [95% CI]		49.7% [36.0% - 66.2%]
Most frequent histologies		N (%)
Lung Adenocarcinoma		11961 (20)
Colon adenocarcinoma		7344 (12)
Lung squamous cell carcinoma		4932 (8)
Ovary serous carcinoma		3004 (5)
Lung non-small cell lung carcinoma (nos)		2957 (5)
Prostate acinar adenocarcinoma		2929 (5)
Pancreas ductal adenocarcinoma		2560 (4)
Unknown primary adenocarcinoma		1808 (3)
Rectum adenocarcinoma		1626 (3)
Skin melanoma		1308 (2)

Abbreviations: CGP: comprehensive genomic profiling; nos: not other specified

Supplemental Table 2- Distribution of PDL1 IHC score category over CD274**focality thresholds at a CD274 gain threshold of 4**

Ploidy Category	Focality Category	PDL1 Status	N
<ploidy+4	<0.1Mb	TPS ≥ 50%	14
<ploidy+4	<0.1Mb	TPS 1% -49%	11
<ploidy+4	<0.1Mb	TPS < 1%	17
<ploidy+4	≥0.1Mb	TPS ≥ 50%	7716
<ploidy+4	≥0.1Mb	TPS 1% -49%	16210
<ploidy+4	≥0.1Mb	TPS < 1%	36087
≥ploidy+4	<0.1Mb	TPS ≥ 50%	18
≥ploidy+4	<0.1Mb	TPS 1% -49%	0
≥ploidy+4	<0.1Mb	TPS < 1%	0
≥ploidy+4	≥0.1Mb	TPS ≥ 50%	368
≥ploidy+4	≥0.1Mb	TPS 1% -49%	190
≥ploidy+4	≥0.1Mb	TPS < 1%	162
<ploidy+4	<4Mb	TPS ≥ 50%	317
<ploidy+4	<4Mb	TPS 1% -49%	357
<ploidy+4	<4Mb	TPS < 1%	620
<ploidy+4	≥4Mb	TPS ≥ 50%	7413
<ploidy+4	≥4Mb	TPS 1% -49%	15864
<ploidy+4	≥4Mb	TPS < 1%	35484
≥ploidy+4	<4Mb	TPS ≥ 50%	186
≥ploidy+4	<4Mb	TPS 1% -49%	45
≥ploidy+4	<4Mb	TPS < 1%	17
≥ploidy+4	≥4Mb	TPS ≥ 50%	200
≥ploidy+4	≥4Mb	TPS 1% -49%	145
≥ploidy+4	≥4Mb	TPS < 1%	145
<ploidy+4	<20Mb	TPS ≥ 50%	2416
<ploidy+4	<20Mb	TPS 1% -49%	3776
<ploidy+4	<20Mb	TPS < 1%	7064
<ploidy+4	≥20Mb	TPS ≥ 50%	5314
<ploidy+4	≥20Mb	TPS 1% -49%	12445
<ploidy+4	≥20Mb	TPS < 1%	29040
≥ploidy+4	<20Mb	TPS ≥ 50%	334
≥ploidy+4	<20Mb	TPS 1% -49%	134
≥ploidy+4	<20Mb	TPS < 1%	90
≥ploidy+4	≥20Mb	TPS ≥ 50%	52
≥ploidy+4	≥20Mb	TPS 1% -49%	56
≥ploidy+4	≥20Mb	TPS < 1%	72

Abbreviations: TPS, Tumor Proportion Score

Supplemental Table 3 –Distribution of PDL1 IHC score category over CD274**focality thresholds at a CD274 gain threshold of 2**

Ploidy Category	Focality Category	PDL1 Status	N
<ploidy+2	<0.1Mb	TPS ≥ 50%	8
<ploidy+2	<0.1Mb	TPS 1% -49%	6
<ploidy+2	<0.1Mb	TPS < 1%	16
<ploidy+2	≥0.1Mb	TPS ≥ 50%	6951
<ploidy+2	≥0.1Mb	TPS 1% -49%	15240
<ploidy+2	≥0.1Mb	TPS < 1%	34405
≥ploidy+2	<0.1Mb	TPS ≥ 50%	24
≥ploidy+2	<0.1Mb	TPS 1% -49%	5
≥ploidy+2	<0.1Mb	TPS < 1%	1
≥ploidy+2	≥0.1Mb	TPS ≥ 50%	1133
≥ploidy+2	≥0.1Mb	TPS 1% -49%	1160
≥ploidy+2	≥0.1Mb	TPS < 1%	1844
<ploidy+2	<4Mb	TPS ≥ 50%	220
<ploidy+2	<4Mb	TPS 1% -49%	290
<ploidy+2	<4Mb	TPS < 1%	569
<ploidy+2	≥4Mb	TPS ≥ 50%	6739
<ploidy+2	≥4Mb	TPS 1% -49%	14956
<ploidy+2	≥4Mb	TPS < 1%	33852
≥ploidy+2	<4Mb	TPS ≥ 50%	283
≥ploidy+2	<4Mb	TPS 1% -49%	112
≥ploidy+2	<4Mb	TPS < 1%	68
≥ploidy+2	≥4Mb	TPS ≥ 50%	874
≥ploidy+2	≥4Mb	TPS 1% -49%	1053
≥ploidy+2	≥4Mb	TPS < 1%	1777
<ploidy+2	<20Mb	TPS ≥ 50%	2005
<ploidy+2	<20Mb	TPS 1% -49%	3390
<ploidy+2	<20Mb	TPS < 1%	6610
<ploidy+2	≥20Mb	TPS ≥ 50%	4954
<ploidy+2	≥20Mb	TPS 1% -49%	11856
<ploidy+2	≥20Mb	TPS < 1%	27811
≥ploidy+2	<20Mb	TPS ≥ 50%	745
≥ploidy+2	<20Mb	TPS 1% -49%	520
≥ploidy+2	<20Mb	TPS < 1%	544
≥ploidy+2	≥20Mb	TPS ≥ 50%	412
≥ploidy+2	≥20Mb	TPS 1% -49%	645
≥ploidy+2	≥20Mb	TPS < 1%	1301

Abbreviations: TPS, Tumor Proportion Score

Supplemental Table 4 - Distribution of samples across focality (20 Mb cutoff) and PDL1 score (50 TPS cutoff) bins at a ploidy cutoff of 4, for Non Small Cell Lung Cancer (NSCLC), Colorectal cancer (CRC), and the rest of the cohort

NSCLC, \geq ploidy + 4 (n=321)	PDL1 TPS: <50	PDL1 TPS: \geq50
Focality category: <20 Mb	74	199
Focality category: \geq20 Mb	21	27
Two-sided Fisher's exact test odds ratio = 0.48[0.24-0.95], P Value = 0.02		
NSCLC, < ploidy + 4 (n=20241)	PDL1 TPS: <50	PDL1 TPS: \geq50
Focality category: <20 Mb	3815	1843
Focality category: \geq20 Mb	10802	3781
Two-sided Fisher's exact test odds ratio = 0.72[0.68-0.78], P Value = 6.56e-21		
CRC, \geq ploidy + 4 (n=72)	PDL1 TPS: <50	PDL1 TPS: \geq50
Focality category: <20 Mb	17	5
Focality category: \geq20 Mb	49	1
Two-sided Fisher's exact test odds ratio = 0.07[0.001-0.71], P Value = 0.009		
CRC, < ploidy + 4 (n=8922)	PDL1 TPS: <50	PDL1 TPS: \geq50
Focality category: <20 Mb	912	17
Focality category: \geq20 Mb	7901	92
Two-sided Fisher's exact test odds ratio = 0.62[0.37-1.12], P Value = 0.08		
Others, \geq ploidy + 4 (n=345)	PDL1 TPS: <50	PDL1 TPS: \geq50
Focality category: <20 Mb	133	130
Focality category: \geq20 Mb	58	24
Two-sided Fisher's exact test odds ratio = 0.42[0.24-0.74], P Value = 0.001		
Others, < ploidy + 4 (n=30892)	PDL1 TPS: <50	PDL1 TPS: \geq50
Focality category: <20 Mb	6113	556
Focality category: \geq20 Mb	22782	1441
Two-sided Fisher's exact test odds ratio = 0.70[0.63-0.77], P Value = 9.03e-12		