

PROJECTGENIE

Genomics Evidence Neoplasia Information Exchange

GENIE BPC NSCLC 2.0-PUBLIC

OVERVIEW

2022

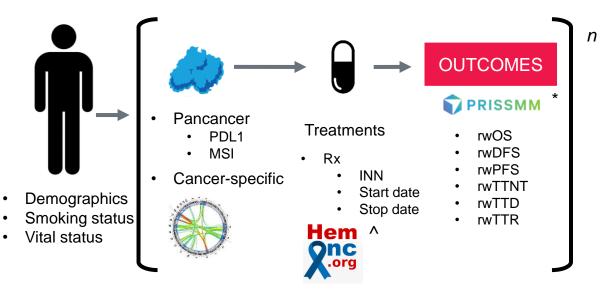
Release Notes



- PRISSMM™: the BPC NSCLC dataset uses the PRISSMM™ framework developed at the Dana-Farber Cancer Institute to determine outcomes from retrospective real-world data to ascertain cancer treatment responses in the real world. Additional information can be found in the <u>analytic data guide</u> and information about licensing PRISSMM™ can be obtained by emailing <u>PRISSMM@mskcc.org</u>
- **Pathologic information:** Each pathology specimen from diagnosis through death or last follow-up is curated with specimen type, site, and histology.
- **Imaging information:** Each CT, MRI, PET-CT scan from diagnosis through death or last follow-up is curated for the presence or absence of cancer and an evaluation of whether the cancer was stable, responding, or progressing.
- Medical oncologist's evaluations: Medical oncology notes (1/month) have been curated to ascertain the presence or absence of cancer and whether the cancer was stable, responding, or progressing.
- Additional relevant biomarkers: Information about select biomarkers not included on the NGS panels, including PDL1, PD1 and MSI, are also curated.
- NSCLC cancer diagnosis is considered the index tumor for this patient cohort. There are data about other cancer diagnoses antecedent to the NSCLC and subsequent to the NSCLC diagnosis.

GENIE BPC Data Model

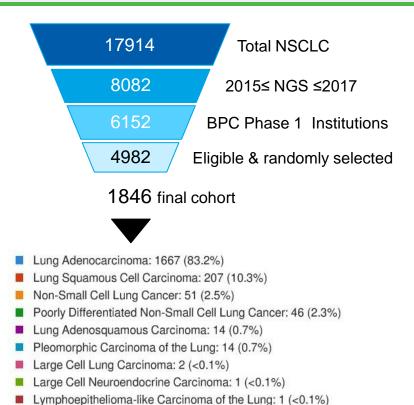


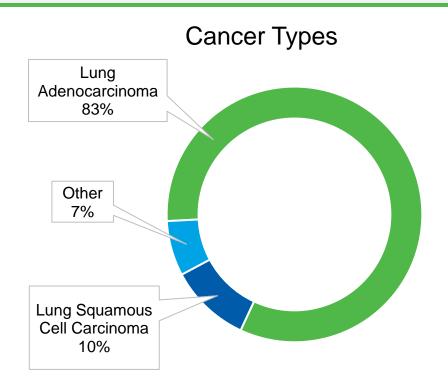


*PRISSMM is licensed from the DFCI ^HemOnc is available through a creative commons license





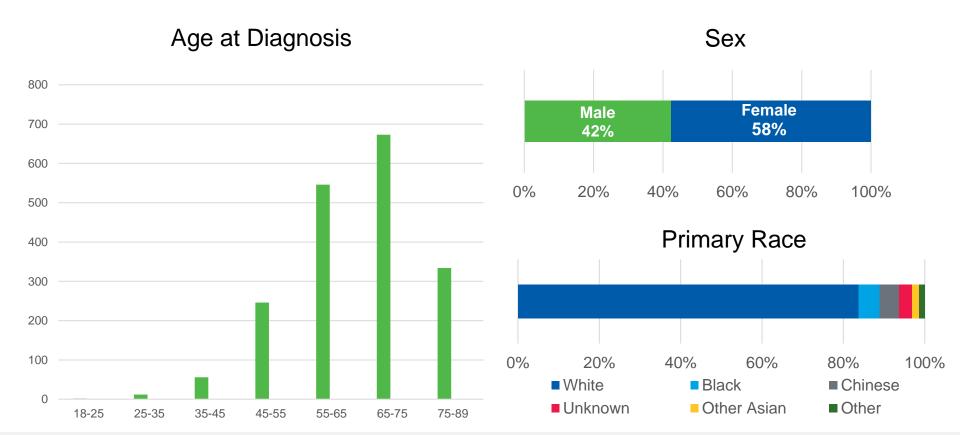




Sarcomatoid Carcinoma of the Lung: 1 (<0.1%)

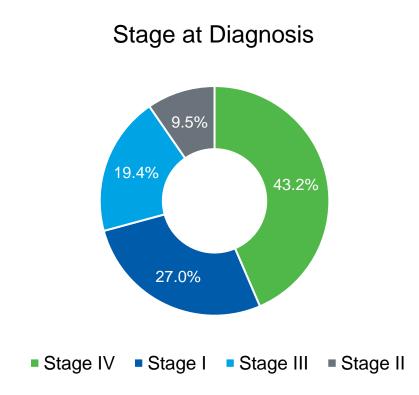
BPC NSCLC 2.0-public Demographics

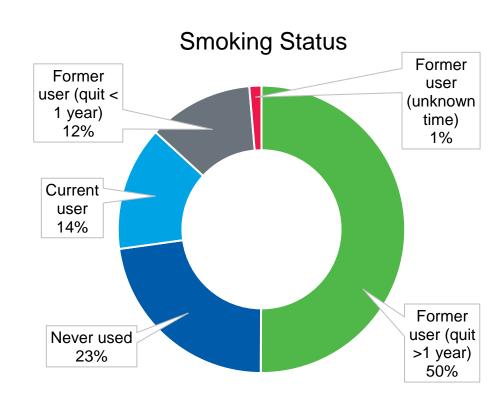




BPC NSCLC 2.0-public: Stage at Diagnosis and Smoking Status



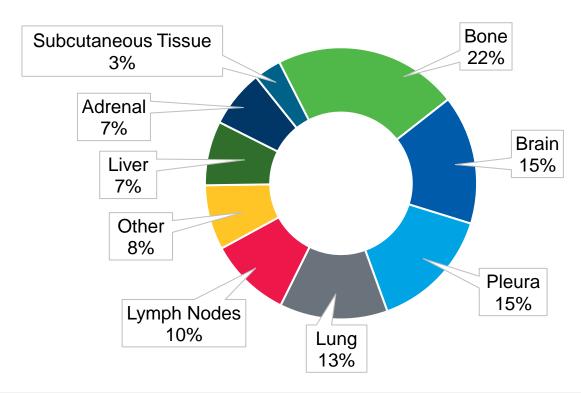




BPC NSCLC 2.0-public: Sites of Metastases at Diagnosis



Location of Metastases (Stage IV patients)



BPC NSCLC 2.0-public: Detailed Clinical Genomics



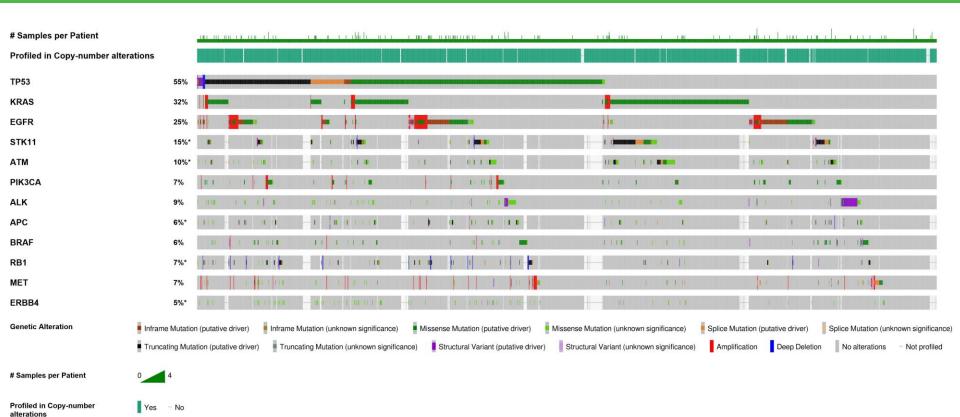
Mutated Genes (2004 profiled samples)			
▼ Gene	# Mut	#	Freq ▼
TP53	1,153	□ 1,074	53.6%
KRAS	607	□ 604	30.1%
LRP1B	30	_ 20	25.6%
EGFR	591	<u>481</u>	24.0%
SPTA1	15	□ 12	15.6%
STK11	276	□ 263	14.4%
COL7A1	44	<u> </u>	13.3%
KEAP1	249	235	13.3%
PRKDC	111	□ 95	11.7%
KMT2D	224	□ 186	10.5%
GRM3	9	8	10.4%
RBM10	146	□ 139	10.4%

Structural Variant Genes (1908 profiled samples)			
▼ Gene	# SV	#	Freq ▼
NRG1	11	□ 11	3.5%
ALK	67	□ 61	3.2%
EML4	52	□ 52	2.7%
STAT4	1	_ 1	2.4%
EGFR	31	□ 31	1.6%
RET	29	□ 27	1.4%
ROS1	28	□ 26	1.4%
KIF5B	19	□ 19	1.0%
CD74	18	□ 18	0.9%
MET	16	□ 16	0.8%
TP53	15	□ 15	0.8%

CNA Genes (1949 profiled samples)				
▼ Gene	Cytoband	CNA	#	Freq ▼
CDKN2A	9p21.3	HOMDEL	□ 147	8.3%
CDKN2B	9p21.3	HOMDEL	□ 135	7.6%
NKX2-1	14q13.3	AMP	□ 95	5.3%
MDM2	12q15	AMP	□ 95	5.3%
MTAP	9p21.3	HOMDEL	□ 16	5.3%
EGFR	7p11.2	AMP	□ 102	5.2%
GRM3	7q21.11-q	AMP	□ 3	3.9%
SPTA1	1q23.1	AMP	□ 3	3.9%
ARFRP1	20q13.33	AMP	□ 3	3.8%
FGF10	5p12	AMP	□ 3	3.8%
TERT	5p15.33	AMP	□ 63	3.7%
MYC	8q24.21	AMP	□ 65	3.7%

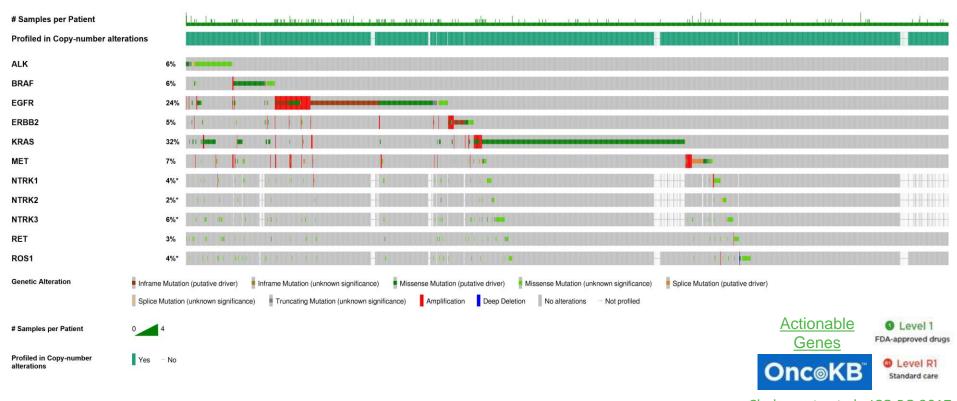


Top 12 Mutated Genes NSCLC 2.0-public



Clinically Actionable Genes NSCLC 2.0-public

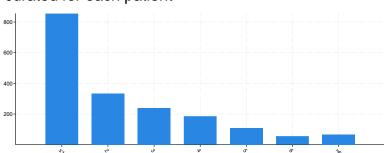




BPC NSCLC 2.0-public:Complete Treatment Histories



Number of cancer-directed drug regimens curated for each patient



Cancer-directed Drug Regimens (Includes regimens ever received for NSCLC irrespective of line or stage)

<u> </u>	
Regimen	# of patients
Carboplatin, Pemetrexed	615
Cisplatin, Pemetrexed	285
Carboplatin, Paclitaxel	207
Bevacizumab, Carboplatin, Pemetrexed	168

Treatments by Patient

Treatment	# ▼
Pemetrexed Disodium	□ 813
Carboplatin	□ 795
Cisplatin	□ 388
Investigational Drug	□ 341
Nivolumab	□ 279
Bevacizumab	□ 225
Paclitaxel	□ 218
Pembrolizumab	□ 204
Erlotinib Hydrochloride	□ 199
Gemcitabine Hydrochloride	□ 195
Docetaxel	□ 191

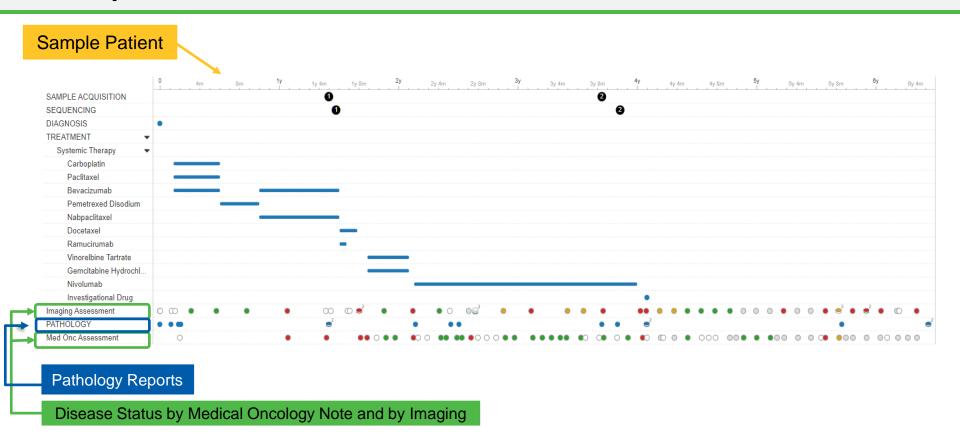
Treatments by Sample (pre- and post-)

Treatment	Pre / Post	# 🕶
Pemetrexed Disodium	Pre	□ 667
Pemetrexed Disodium	Post	235
Carboplatin	Pre	□ 663
Carboplatin	Post	□ 223
Investigational Drug	Pre	□ 303
Investigational Drug	Post	□ 83
Nivolumab	Pre	□ 263
Nivolumab	Post	_ 40
Cisplatin	Pre	□ 255
Cisplatin	Post	□ 166

- Individual regimens can be calculated from the raw data found <u>here</u> on Synapse
- Treatment history for non-index cancers can be found <u>here</u> on Synapse

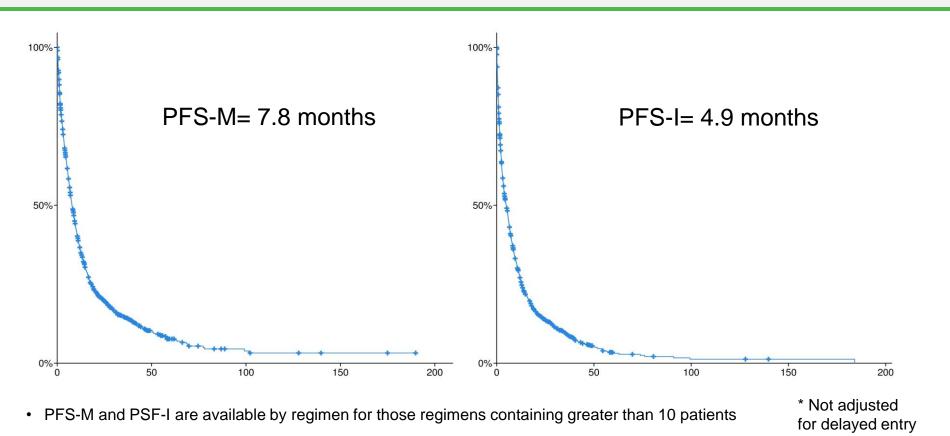
BPC NSCLC 2.0-public: Comprehensive Patient View





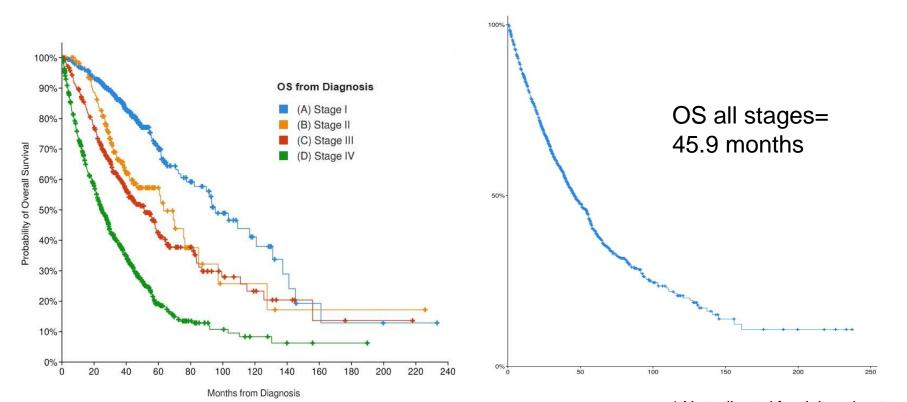
BPC NSCLC 2.0-public: High Quality Clinical Data





BPC NSCLC 2.0-public: OS by Stage





GENIE BPC Acknowledgements



FINDING CURES TOGETHER™



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Former BPC Team Members can be found <u>here</u>