

PROJECTGENIE

Genomics Evidence Neoplasia Information Exchange

GENIE BPC CRC 2.0-PUBLIC

OVERVIEW

2022

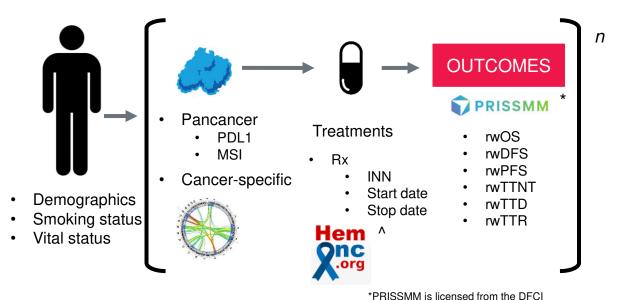
Release Notes



- PRISSMM™: the BPC CRC 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, MMR and MSI, are also curated.
- Colorectal cancer diagnosis is considered the index tumor for this patient cohort. There are data about other cancer diagnoses antecedent to the colorectal and subsequent to the colorectal diagnosis.

GENIE Data Model



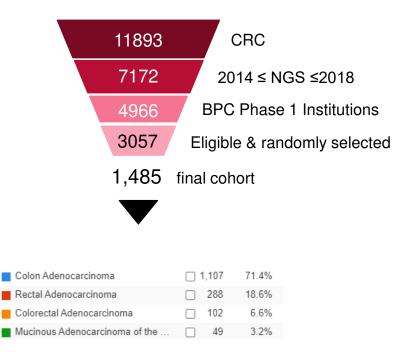


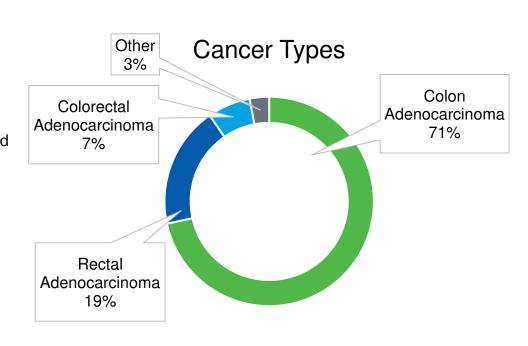
^HemOnc is available through a creative commons license

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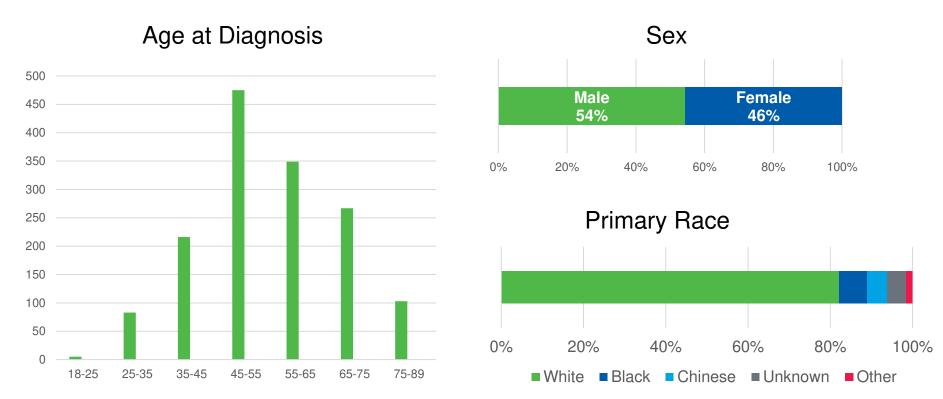




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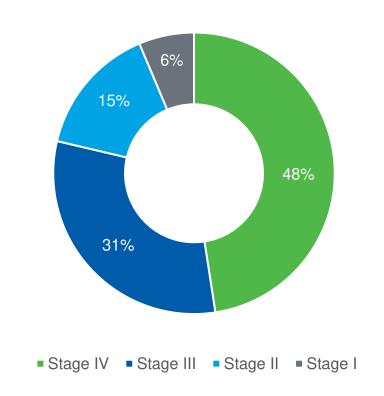




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BPC CRC 2.0-public: Stage at Diagnosis

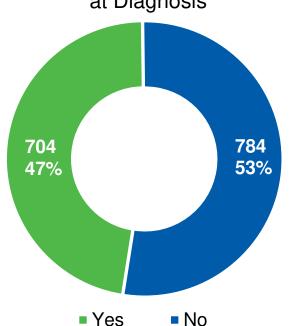


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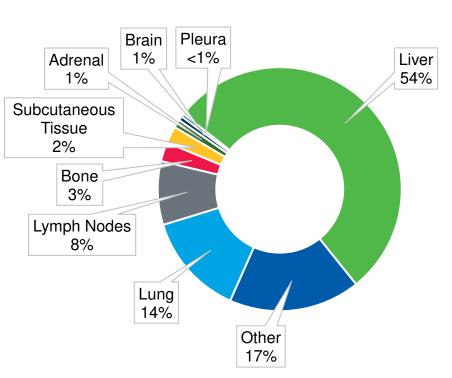
BPC CRC 2.0-public:Sites of Metastases at Diagnosis







Location of Metastasis



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BPC CRC 2.0-public: Detailed Clinical Genomics



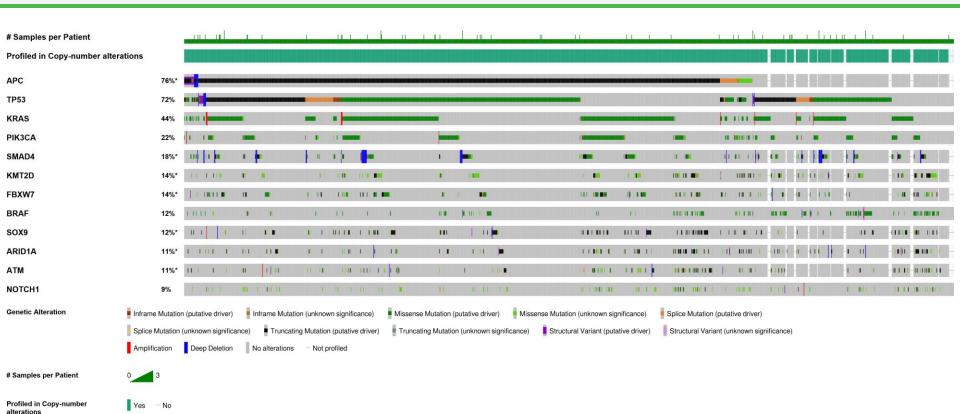
Mutated Genes (1551 profiled samples)			Structural Variant Genes (253 profiled samples)				CNA Genes (1487 profiled samples)					
▼ Gene	# Mut	#	Freq •	▼ Gene	# SV	#	Freq ▼	▼ Gene	Cytoband	CNA	#	Freq *
APC	1,810	1,114	74.9%	APC	19	□ 19	7.5%	ARFRP1	20q13.33	AMP	□ 19	10.4%
TP53	1,187	1,107	71.4%	TP53	14	□ 14	5.5%	FGF14	13q33.1	AMP	□ 16	8.8%
KRAS	678	□ 669	43.1%	CTNNB1	13	□ 13	5.1%	RTEL1	20q13.33	AMP	□ 20	5.3%
PIK3CA	367	□ 327	21.1%	BRAF	8	7	2.8%	FLT3	13q12.2	AMP	□ 73	4.9%
SMAD4	247	□ 224	15.1%	PRKN	6	□ 6	2.4%	BCL2L1	20q11.21	AMP	□ 67	4.5%
LRP1B	52	□ 27	14.8%	SMAD4	4	_ 4	1.6%	MYC	8q24.21	AMP	□ 67	4.5%
KMT2D	336	□ 208	14.0%	TCF7L2	4	_ 4	1.6%	SRC	20q11.23	AMP	□ 52	4.3%
FBXW7	226	□ 203	13.7%	PBRM1	4	_ 4	1.6%	FLT1	13q12.3	AMP	□ 61	4.1%
COL7A1	40	□ 35	12.5%	NTRK1	4	_ 4	1.6%	CDK8	13q12.13	AMP	□ 49	4.1%
BRAF	200	□ 190	12.3%	ATM	3	□ 3	1.2%	TOP1	20q12	AMP	□ 37	4.0%
SOX9	200	□ 174	11.7%	PTEN	3	□ 3	1.2%	AGO2	8q24.3	AMP	□ 15	4.0%
PTPRS	108	85	11.6%	NF1	3	_ 3	1.2%	DNMT3B	20q11.21	AMP	□ 28	3.8%

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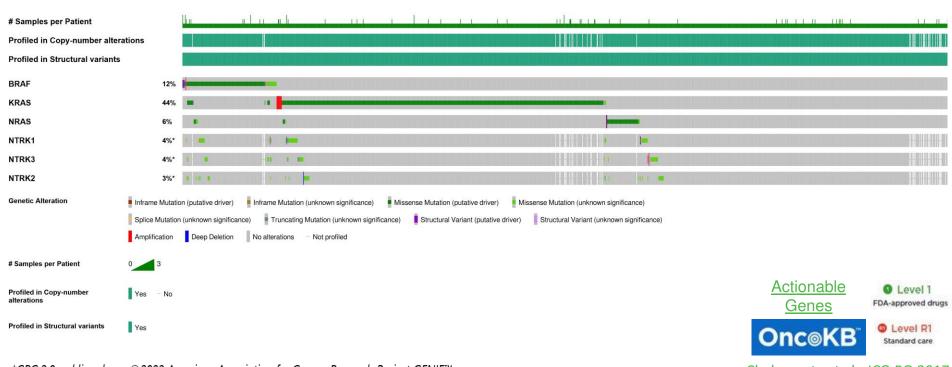
FINDING CURES TOGETHER™



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Clinically Actionable Genes CRC 2.0-public





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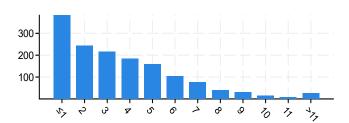
Chakravarty et al., JCO PO 2017

BPC CRC 2.0-public: Complete Treatment Histories



FINDING CURES TOGETHER™

Number of cancer-directed drug regimens curated for each patient



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

Regimen	# of patients
FOLFOX	1172
FOLFOX + Bevacizumab	702
FOLFIRI	853
FOLFIRI + Bevacizumab	665

Treatments by Patient

Treatments by Sample (pre- and post-)

Treatment per Patient					
Treatment	# 🕶				
Fluorouracil	_ 1,247				
Leucovorin Calcium	1,237				
Oxaliplatin	_ 1,209				
Irinotecan Hydrochloride	□ 866				
Bevacizumab	□ 746				
Capecitabine	□ 397				
Investigational Drug	□ 189				
Panitumumab	□ 171				
Cetuximab	□ 154				
Trifluridine and Tipiracil Hydrochloride	□ 153				

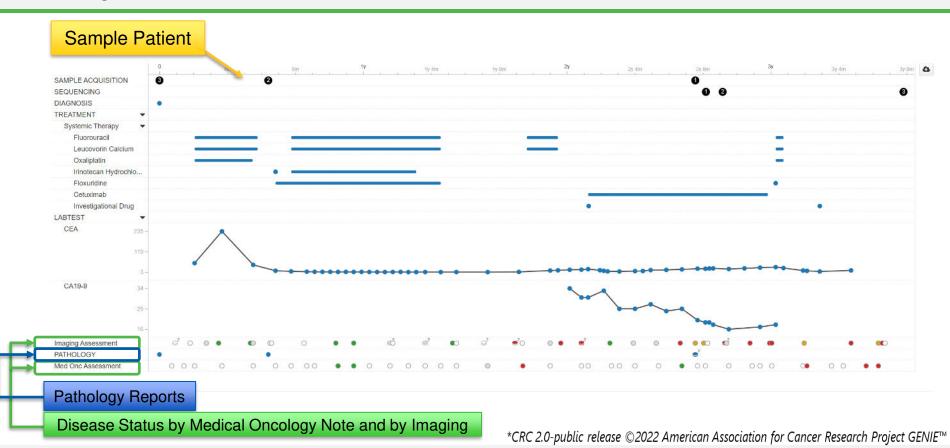
		•				
Treatment per Sample (pre/post)						
Treatment	Pre / Post	# ▼				
Leucovorin Calcium	Pre	□ 865				
Leucovorin Calcium	Post	□ 437				
Fluorouracil	Pre	□ 853				
Fluorouracil	Post	□ 459				
Oxaliplatin	Pre	□ 843				
Oxaliplatin	Post	_ 428				
Irinotecan Hydrochloride	Pre	□ 757				
Irinotecan Hydrochloride	Post	□ 167				
Bevacizumab	Pre	□ 630				
Bevacizumab	Post	□ 162				

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

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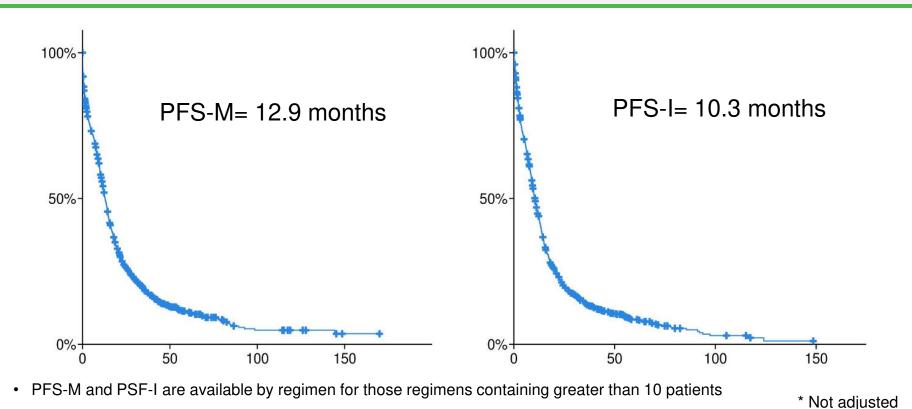
BPC CRC 2.0-public: Comprehensive Patient View





BPC CRC 2.0-public: High Quality Clinical Data



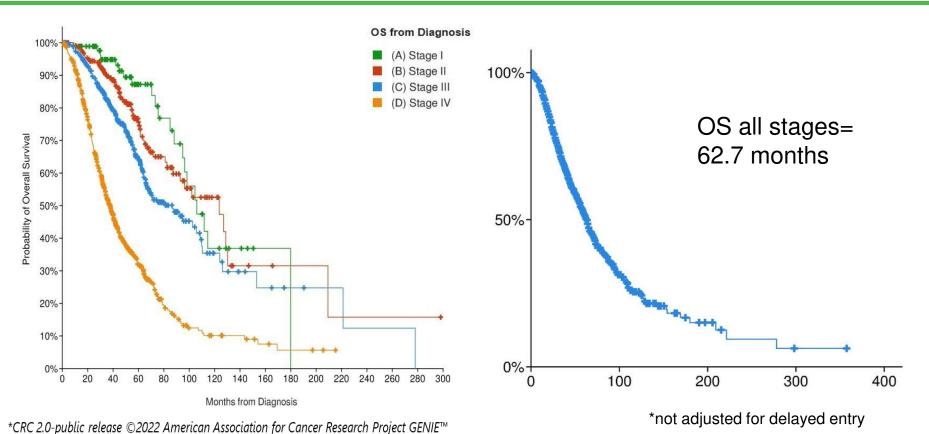


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for delayed entry

BPC CRC 2.0-public: OS by Stage





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GENIE BPC Acknowledgements



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Xindi Guo Thomas Yu Chelsea Nayan



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Deb Schrag
Julia Rudolph
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Cynthia Chu
Shirin Pillai

Statistical Core

Kathy Panageas Jessica Lavery Samantha Brown Axel Martin Michael Curry



Ken Kehl Asha Postle Ashley Newcomb Kevin Haigis John Orechia Daniel Quinn Simon Arango Baquero



Philippe Bedard
Celeste Yu
Samanta Del Rossi
Nitthusha Singaravelan
Demi Plagianakos
Alisa Nguyen
Nazish Qazi
Gunjan Srivastava
Sophie Cooke
Alisha Rizvi



Jeremy Warner
Michele Lenoue-Newton
Christine Micheel
Ben Ho Park
Sanjay Mishra
Daniel Fabbri
Marilyn Holt
Neha Jain
Protiva Rahman
Li Wen
Yuanchu James Yang
Kate Mittendorf

Former BPC Team Members can be found here