

AACR PROJECTGENIE

Genomics Evidence Neoplasia Information Exchange

GENIE BPC CRC 2.0-PUBLIC

OVERVIEW

2022

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Release Notes



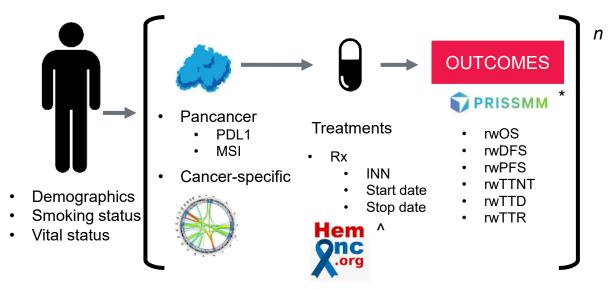
- PRISSMM™: the BPC CRC dataset uses the PRISSMM™ framework developed at the Dana-Farber Cancer Institute to determine outcomes from retrospective real-world data to ascertain cancer treatment responses in the real world. Additional information can be found in the <u>analytic data guide</u> and information 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



FINDING CURES TOGETHER





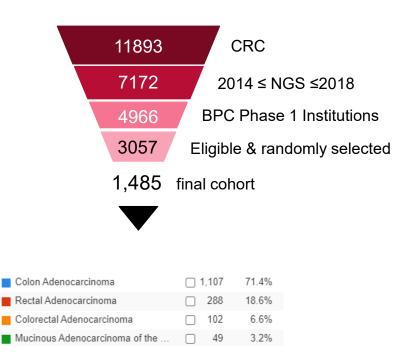
*PRISSMM is licensed from the DFCI
^HemOnc is available through a creative commons license

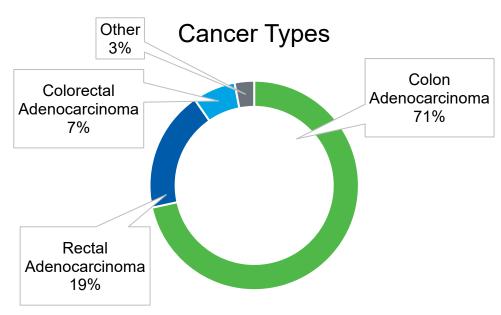
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BPC CRC 2.0-public Cohort







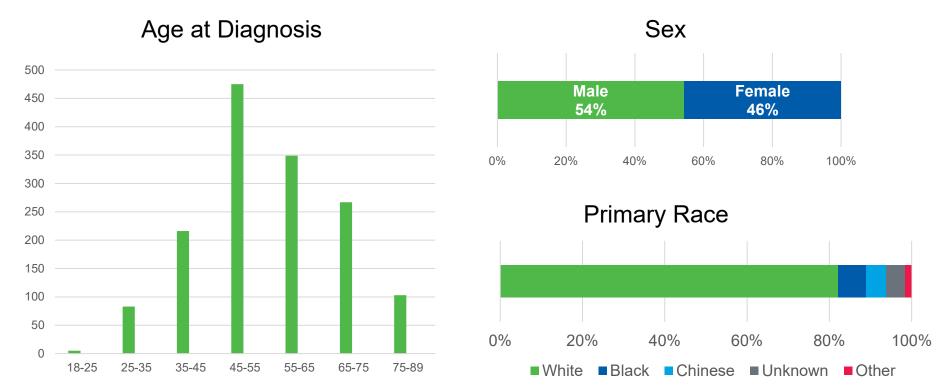


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BPC CRC 2.0-public Demographics





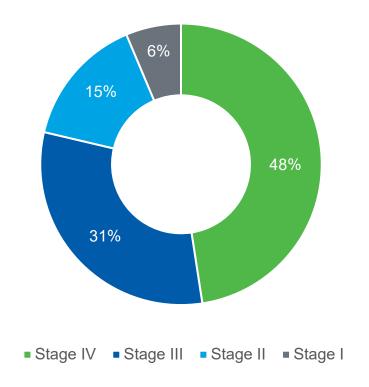


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BPC CRC 2.0-public: Stage at Diagnosis







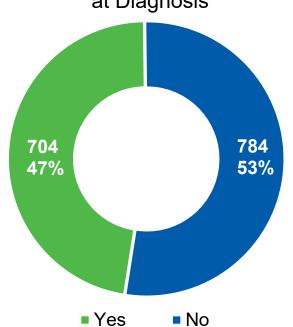
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BPC CRC 2.0-public: Sites of Metastases at Diagnosis

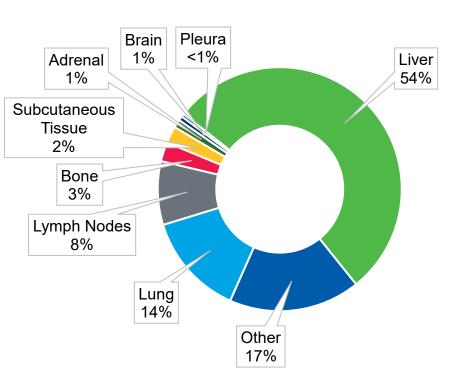








Location of Metastasis



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BPC CRC 2.0-public: Detailed Clinical Genomics



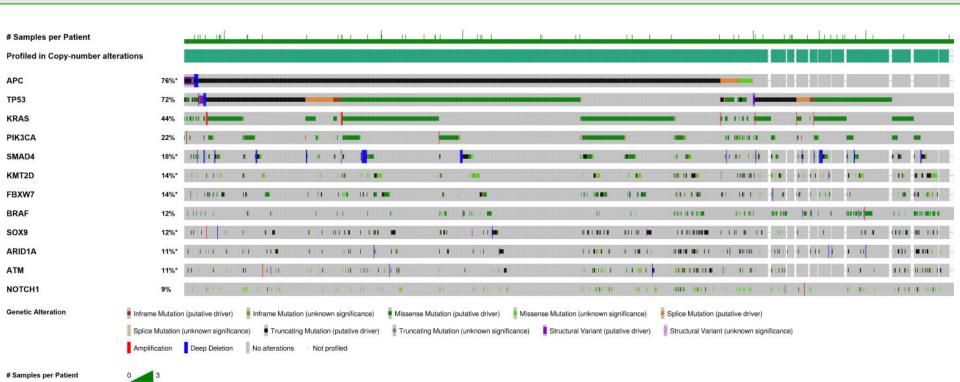


	CNA Genes (1487 profiled samples)			Structural Variant Genes (253 profiled samples)			Mutated Genes (1551 profiled samples)					
Freq *	#	CNA	Cytoband	▼ Gene	Freq ▼	#	# SV	▼ Gene	Freq ▼	#	# Mut	▼ Gene
10.4%	□ 19	AMP	20q13.33	ARFRP1	7.5%	□ 19	19	APC	74.9%	1,114	1,810	APC
8.8%	□ 16	AMP	13q33.1	FGF14	5.5%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

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Top 12 Mutated Genes CRC 2.0-public





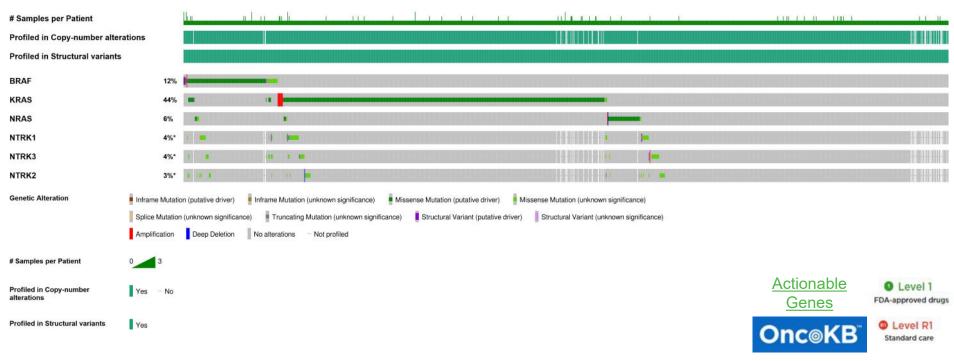
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Profiled in Copy-number

Clinically Actionable Genes CRC 2.0-public







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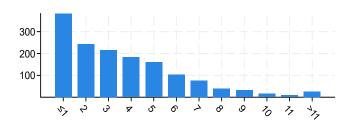
Chakravarty et al., JCO PO 2017

BPC CRC 2.0-public: Complete Treatment Histories





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					

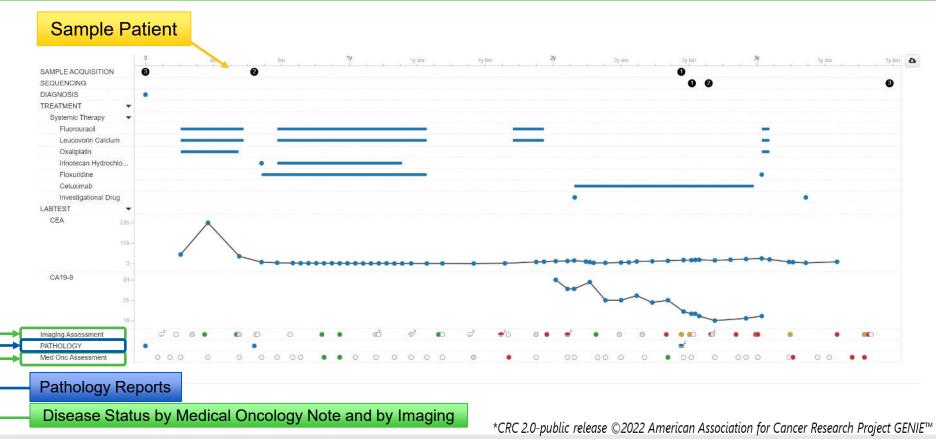
- Individual regimens can be calculated from the raw data found here on Synapse
- Treatment history for non-index cancers can be found here on Synapse

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BPC CRC 2.0-public:Comprehensive Patient View



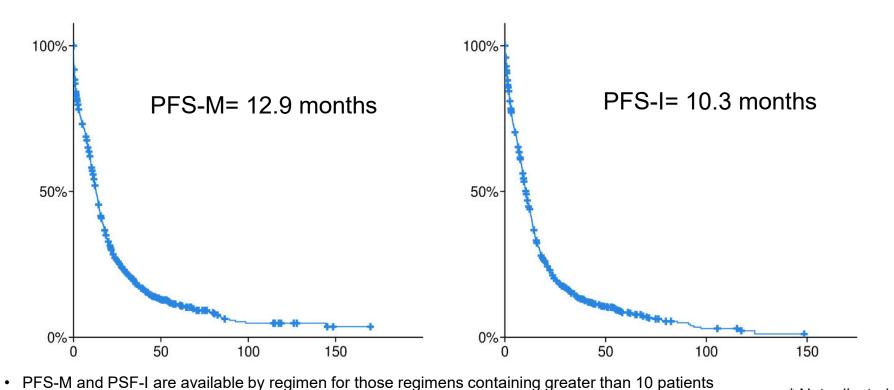




BPC CRC 2.0-public: High Quality Clinical Data





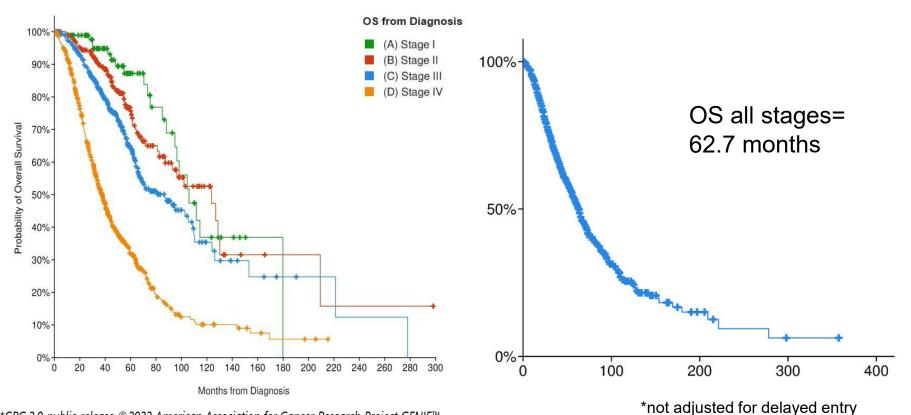


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* Not adjusted for delayed entry

BPC CRC 2.0-public: OS by Stage





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GENIE BPC Acknowledgements







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Former BPC Team Members can be found here