

Medical Laboratory  
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Leading a new era of precision oncology

# Oncofocus®

Precision Oncology

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## Patient demographics

ONC20	-	Requester	-
Surname	-	Contact details	-
Forename	-	Date requested	-
DOB	-		
Gender	Female		
Histology #	-	Tumour %	70%
Primary site	Right Eye	Tumour %	-
Tumour subtype	Spindle Cell Melanoma	(macrodissected)	
Tissue Type	Endorescection Right Eye		

## Comment

The DNA and RNA extracted from this sample were of optimal quality. The Oncofocus assay on which the sample was run met all assay specific quality metrics.

Oncofocus currently targets 505 genes covering oncogenes, fusion genes, genes susceptible to copy number variation and tumour suppressors. Actionable genetic variants detected by Oncofocus are currently linked to 738 anti-cancer targeted therapies/therapy combinations.

### The clinically significant bio-markers identified in this case are summarised on page 2

In addition to the variants listed below, a pathogenic variant in the SF3B1 gene was also detected, c.1874G>A, p.(Arg625His), however at this time it does not link to any therapies/clinical trials.

*Within the 'Current Clinical Trials Information' section of this report, starting on page 7, the NCT numbers are hyperlinks to the [clinicaltrials.gov](https://clinicaltrials.gov) webpages which should be accessed to gain further trial specific information*

## Clinically Significant Biomarkers

Indicated Contraindicated

Genomic Alteration	Alt Allele Freq	Relevant Therapies (In this cancer type)	Relevant Therapies (In other cancer type)	Clinical Trials
<b>GNA11 mutation</b> c.626A>T, p.(Gln209Leu)	42.26%	Clinical trials and/or off-label	Clinical trials and/or off-label	9
<b>BAP1 mutation</b> c.1366C>T, p.(Gln456Ter)	72.81%	Clinical trials and/or off-label	Clinical trials and/or off-label	7
<b>FANCD2 deletion</b> (copy number = 0.63)		Clinical trials and/or off-label	Clinical trials and/or off-label	5
<b>ATR deletion</b> (copy number = 0.76)		Clinical trials and/or off-label	Clinical trials and/or off-label	5
<b>SMO mutation</b> c.808G>A, p.(Val270Ile)	50.58%	Clinical trials and/or off-label	Clinical trials and/or off-label	3
<b>SETD2 deletion</b> (copy number = 0.84)		Clinical trials and/or off-label	Clinical trials and/or off-label	4

Sources included in relevant therapies: EMA1, ESMO, NCCN

Hotspot variants with >10% alternate allele reads are classified as 'detected' with an assay sensitivity and positive predictive value (PPV) of 99%. Copy number variants; amplifications of CN> 6 with the 5% confidence value of  $\geq 4$  after normalization and deletions with 95% CI  $\leq 1$  are classified as present when the tumour% >50% with a sensitivity of 80% and PPV 100%. Gene Fusions are reported when occurring in >40 counts and meeting the thresholds of assay specific internal RNA quality control with a sensitivity of 92% and PPV of 99%. Other mutations, copy number variations, or fusions that were detected but not classified by the Oncofocus Test as actionable by a known therapeutic targeted agent are not listed in the results section of this report. Supplementary technical information is available upon request.

## Biomarker Descriptions

### GNA11 (G protein subunit alpha 11)

**Background:** The GNA11 gene encodes an alpha subunit of heterotrimeric guanine nucleotide-binding proteins (G-proteins). G-protein alpha subunits bind guanine nucleotide, hydrolyze GTP, and interact with specific receptor and effector molecules. GNA11 is closely related to GNAQ, another G-protein alpha subunit.

**Alterations and prevalence:** Somatic activating mutations in GNA11 and GNAQ at amino acids R183 and Q209 are common in uveal melanoma and are mutually exclusive. These mutations render the G protein constitutively active leading to the stimulation of MAP kinases, PI3K/AKT, and protein kinase C, which promote tumor growth and proliferation<sup>1,2,3</sup>. Approximately 45% of uveal melanoma cases contain activating mutations in GNA11 and up to 50% of cases contain activating mutations in GNAQ<sup>4,5,6</sup>. By contrast, GNA11 and GNAQ mutations are infrequent in cutaneous melanoma, with a combined prevalence of approximately 1%, and are infrequently observed in other cancers<sup>5,6</sup>.

**Potential clinical relevance:** Currently, no therapies are approved for GNA11 aberrations. In a randomized phase II clinical trial of MEK inhibitor selumetinib versus chemotherapy, GNA11 and GNAQ positive uveal melanoma patients demonstrated a median progression-free survival (PFS) of 15.9 weeks versus 7 weeks, respectively<sup>7</sup>. However, no statistically significant improvement in overall survival (OS) was observed and the improvement in outcomes was associated with a high rate of adverse events<sup>7</sup>.

### SMO (smoothened, frizzled class receptor)

**Background:** The SMO gene encodes the smoothened, frizzled class receptor, a transmembrane G protein-coupled receptor that is part of the Hedgehog (Hh) signaling pathway<sup>8</sup>. SMO is negatively regulated by the tumor suppressor gene patched transmembrane receptor (PTCH). However, binding of the ligand sonic hedgehog (Shh) stops this inhibition thereby activating downstream genes such as glioma-associated (GLI) transcription factors<sup>9</sup>. Consequently, aberrations in SMO leading to constitutive activation have been identified to promote oncogenesis in certain cancer types including basal cell carcinoma (BCC)<sup>10,11,12</sup>.

**Alterations and prevalence:** Somatic mutations in SMO are observed in 10% of BCC and medulloblastoma, and in 5% of uterine cancer, 4% of stomach cancer, and 3% of lung adenocarcinoma<sup>5,6</sup>. SMO is amplified in up to 7% of ovarian cancer, 5% of glioma, and 4% of melanoma<sup>5,6</sup>.

**Potential clinical relevance:** Currently, no therapies are approved for SMO aberrations. However, FDA approved Hh pathway inhibitors that include SMO as a target include vismodegib (2012) and sonidegib (2015) for BCC and glasdegib (2018) for acute myeloid leukemia. Several missense mutations in SMO, including G497W and D473Y/H, have been associated with resistance to vismodegib in clinical cohorts of BCC patients<sup>13,14,15</sup>. Similarly, in a clinical trial of BCC patients treated with sonidegib, SMO W535L, Q477E, D473H, S533N, and D473G mutations demonstrated progressive disease<sup>16</sup>.

Lead Clinical Scientist: Keeda Hardisty

Clinical Scientist: Kaiya Chowdhary

Date:

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## Tier Criteria Met

Genomic Alteration	Tier Classification for Melanoma
<i>GNA11</i> mutation Tier: IIC	IIC: Biomarker is an inclusion criteria for clinical trials
<i>BAP1</i> mutation Tier: IIC	IIC: Biomarker is an inclusion criteria for clinical trials
<i>FANCD2</i> deletion Tier: IIC	IIC: Biomarker is an inclusion criteria for clinical trials
<i>ATR</i> deletion Tier: IIC	IIC: Biomarker is an inclusion criteria for clinical trials
<i>SETD2</i> mutation Tier: IIC	IIC: Biomarker is an inclusion criteria for clinical trials
<i>SMO</i> mutation Tier: IIC	IIC: Biomarker is an inclusion criteria for clinical trials

**Reference:** Li et al. *Standards and Guidelines for the Interpretation and Reporting of Sequence Variants in Cancer: A Joint Consensus Recommendation of the Association for Molecular Pathology, American Society of Clinical Oncology, and College of American Pathologists.* J Mol Diagn. 2017 Jan;19(1):4-23.

## Relevant Therapy Summary

● In this cancer type  
 ○ In other cancer type  
 ◐ In this cancer type and other cancer types  
 ⊘ Contraindicated  
 ⚠ Both for use and contraindicated  
 × No evidence

GNA11 mutation				
Relevant Therapy	EMA	ESMO	NCCN	Clinical Trials*
selumetinib, ulixertinib	×	×	×	● (II)
ASTX029	×	×	×	● (I/II)
cobimetinib	×	×	×	● (I/II)
lifirafenib, mirdametinib	×	×	×	● (I/II)
LXS-196	×	×	×	● (I/II)
abemaciclib, cetuximab, chemotherapy, encorafenib, LY3214996, midazolam	×	×	×	● (I)
everolimus + RO-5126766, RO-5126766	×	×	×	● (I)
LXH254	×	×	×	● (I)
RMC-4630	×	×	×	● (I)

\* Most advanced phase (IV, III, II/III, II, I/II, I) is shown and multiple clinical trials may be available.

ONC20:-  
Referring pathology dept: -

[www.oncologica.com](http://www.oncologica.com)

**Disclaimer:** The data presented here is a result of the curation of published data sources, but may not be exhaustive. The data version is 2019.12(005).

Lead Clinical Scientist: Keeda Hardisty

Clinical Scientist: Kaiya Chowdhary

Date:

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## Relevant Therapy Summary (continued)

● In this cancer type  
 ○ In other cancer type  
 ⓘ In this cancer type and other cancer types  
 ⊘ Contraindicated  
 ⚠ Both for use and contraindicated  
 × No evidence

### BAP1 mutation

Relevant Therapy	EMA	ESMO	NCCN	Clinical Trials*
atezolizumab	×	×	×	● (II)
durvalumab + olaparib	×	×	×	● (II)
niraparib	×	×	×	● (II)
olaparib	×	×	×	● (II)
BAY-1895344	×	×	×	● (I/II)
pamiparib, tislelizumab	×	×	×	● (I)

### FANCD2 deletion

Relevant Therapy	EMA	ESMO	NCCN	Clinical Trials*
durvalumab + olaparib	×	×	×	● (II)
olaparib	×	×	×	● (II)
prexasertib	×	×	×	● (II)
talazoparib	×	×	×	● (II)
BAY-1895344	×	×	×	● (I/II)

### ATR deletion

Relevant Therapy	EMA	ESMO	NCCN	Clinical Trials*
durvalumab + olaparib	×	×	×	● (II)
olaparib	×	×	×	● (II)
prexasertib	×	×	×	● (II)
talazoparib	×	×	×	● (II)
BAY-1895344	×	×	×	● (I/II)

\* Most advanced phase (IV, III, II/III, II, I/II, I) is shown and multiple clinical trials may be available.

ONC20:-  
Referring pathology dept: -

www.oncologica.com

Disclaimer: The data presented here is a result of the curation of published data sources, but may not be exhaustive. The data version is 2019.12(005).

Lead Clinical Scientist: Keeda Hardisty

Clinical Scientist: Kaiya Chowdhary

Date:

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## Relevant Therapy Summary (continued)

● In this cancer type  
 ○ In other cancer type  
 ● In this cancer type and other cancer types  
 ⊘ Contraindicated  
 ⚠ Both for use and contraindicated  
 × No evidence

### SETD2 mutation

Relevant Therapy	EMA	ESMO	NCCN	Clinical Trials*
atezolizumab	×	×	×	● (II)
olaparib	×	×	×	● (II)
BAY-1895344	×	×	×	● (I/II)
pamiparib, tislelizumab	×	×	×	● (I)

### SMO mutation

Relevant Therapy	EMA	ESMO	NCCN	Clinical Trials*
vismodegib	×	×	×	● (II)

\* Most advanced phase (IV, III, II/III, II, I/II, I) is shown and multiple clinical trials may be available.

## Relevant Therapy Details

### Current Clinical Trials Information

Clinical Trials information is current as of 2019-09-09. For the most up-to-date information regarding a particular trial, search [www.clinicaltrials.gov](http://www.clinicaltrials.gov) by NCT ID or search local clinical trials authority website by local identifier listed in 'Other identifiers'.

#### GNA11 mutation

**NCT03947385**

A Phase I/II Study of IDE196 in Patients With Solid Tumors Harboring GNAQ/11 Mutations or PRKC Fusions

**Cancer type:** Melanoma

**Variant class:** GNA11 mutation

**Other identifier:** IDE196-001

**Population segments:** Fourth line or greater, Second line, Stage III, Stage IV, Third line

**Exclusion criteria variant class:** Microsatellite instability-High

**Phase:** I/II

**Therapy:** LXS-196

**Locations:** Australia, United States

**US States:** PA, TN, TX

**Contact:** Dr. Julie Hambleton [650-262-3603; jhambleton@ideayabio.com]

**NCT02639546**

A Phase I/II, Multicenter, Open-Label, Dose-Escalation Study of The Safety And Pharmacokinetics of Cobimetinib In Pediatric And Young Adult Patients With Previously Treated Solid Tumors

**Cancer type:** Melanoma

**Variant class:** RAS/RAF/MEK/ERK pathway

**Other identifiers:** 15-524, 16-041, 2015-0929, CTX#15-0005, DRKS00010690, EudraCT Number: 2014-004685-25, GO29665, iMATRIX Cobi, iMATRIXcobi, IRAS ID: 174562, NCI-2016-00541, NL52503.078.16

**Population segments:** (N/A), Pediatric or Adolescent, Second line

**Phase:** I/II

**Therapy:** cobimetinib

**Locations:** Canada, France, Germany, Italy, United Kingdom, United States

**US States:** AZ, CA, PA

**Contact:** Reference Study ID Number: GO29665 [888-662-6728; global-roche-genentech-trials@gene.com]



## GNA11 mutation (continued)

### NCT03905148

A Phase Ib, Open-Label, Dose-escalation and Expansion Study to Investigate the Safety, Pharmacokinetics and Antitumor Activities of a RAF Dimer Inhibitor BGB-283 in Combination With MEK Inhibitor PD-0325901 in Patients With Advanced or Refractory Solid Tumors

**Cancer type:** Melanoma

**Variant class:** RAS/RAF/MEK/ERK pathway

**Other identifier:** BGB-283/PD-0325901-AU-001

**Population segments:** Locally advanced, Metastatic, Second line, Stage III, Stage IV, Unresectable

**Phase:** I/II

**Therapies:** lifirafenib, mirdametinib

**Location:** Australia

### NCT03155620

NCI-COG Pediatric MATCH (Molecular Analysis for Therapy Choice) Screening Protocol

**Cancer type:** Unspecified Solid Tumor

**Variant class:** RAS/RAF/MEK/ERK mutation

**Other identifiers:** 17-729, APEC1621SC, COGAPEC1621SC, N 55017 SC, NCI-2017-01251, NCI-COG Pediatric MATCH, Pediatric MATCH

**Population segments:** Aggressive, Indolent, Pediatric or Adolescent, Second line, Stage III, Stage IV

**Phase:** II

**Therapies:** selumetinib, ulixertinib

**Locations:** Puerto Rico, United States

**US States:** AL, AR, AZ, CA, CO, CT, DE, FL, GA, HI, IA, ID, IL, IN, KY, LA, MA, MD, ME, MI, MN, MO, MS, NC, NE, NH, NJ, NV, NY, OH, OK, OR, PA, RI, SC, TN, TX, UT, VA, VT, WA, WI

**Contact:** Multiple contacts: See [www.clinicaltrials.gov](http://www.clinicaltrials.gov) for complete list of contacts.

### NCT03520075

A Phase I/II Study of the Safety, Pharmacokinetics, and Activity of ASTX029 in Subjects With Advanced Solid Tumors

**Cancer type:** Unspecified Solid Tumor

**Variant class:** RAS/RAF/MEK/ERK pathway

**Other identifier:** ASTX029-01

**Population segments:** Second line, Stage III, Stage IV

**Phase:** I/II

**Therapy:** ASTX029

**Location:** United States

**US States:** CT, TX, VA

**Contact:** Richard J. Morishige [925-560-2882; [Richard.Morishige@astx.com](mailto:Richard.Morishige@astx.com)]

## GNA11 mutation (continued)

### NCT02407509

A Phase I Trial of RO5126766 (a Dual RAF/MEK Inhibitor) Exploring Intermittent, Oral Dosing Regimens in Patients With Solid Tumours or Multiple Myeloma, With an Expansion to Explore Intermittent Dosing in Combination With Everolimus

**Cancer type:** Unspecified Solid Tumor

**Variant class:** RAS/RAF/MEK/ERK mutation

**Other identifiers:** CCR3808, DDU RAF/MEK, EudraCT Number: 2012-001040-22, IRAS ID:102403

**Population segments:** Adenocarcinoma, Fourth line or greater, KRAS, Second line, Stage III, Stage IV, Third line

**Phase:** I

**Therapies:** everolimus + RO-5126766, RO-5126766

**Location:** United Kingdom

### NCT03634982

A Phase I, Open-Label, Multicenter, Dose-Escalation Study of RMC-4630 Monotherapy in Adult Participants with Relapsed/Refractory Solid Tumors

**Cancer type:** Unspecified Solid Tumor

**Variant class:** RAS/RAF/MEK/ERK mutation

**Other identifiers:** 18-559, 19683, NCI-2018-02064, RMC-4630-01, UCI-18-14

**Population segments:** KRAS, Second line, Stage III, Stage IV

**Phase:** I

**Therapy:** RMC-4630

**Location:** United States

**US States:** AZ, CA, CO, FL, OK, TN

**Contact:** Revolution Medicines [650-779-2300; CT-Inquiries@RevolutionMedicines.com]

### NCT02857270

A Phase I Study of an ERK1/2 Inhibitor (LY3214996) Administered Alone or in Combination With Other Agents in Advanced Cancer

**Cancer type:** Unspecified Solid Tumor

**Variant class:** RAS/RAF/MEK/ERK pathway

**Other identifiers:** 16419, 2016-0544, EudraCT Number: 2016-001907-21, F18132, I8S-MC-JUAB, I8S-MC-JUAB-c, JapicCTI-194728, JUAB, NCI-2017-00039

**Population segments:** Adenocarcinoma, BRAF, Fourth line or greater, Large Cell, Stage III, Stage IV

**Phase:** I

**Therapies:** abemaciclib, cetuximab, chemotherapy, encorafenib, LY3214996, midazolam

**Locations:** Australia, France, Japan, United States

**US States:** DC, FL, MA, NH, PA, TN, TX

**Contact:** Eli Lilly and Company [877-285-4559]

**GNA11 mutation (continued)****NCT02607813**

A Phase I Dose Finding Study of Oral LXH254 in Adult Patients With Advanced Solid Tumors Harboring MAPK Pathway Alterations

**Cancer type:** Unspecified Solid Tumor

**Variant class:** RAS/RAF/MEK/ERK pathway

**Other identifiers:** 16-225, 18-108, 2015-0913, CLXH254X2101, EudraCT Number: 2015-003421-33, JapicCTI-163224, NCI-2015-02280, NL55506.078.15, Nov RAFI (CLXH254X2101), REec-2016-2132, SNCTP000002708

**Population segments:** BRAF, Fourth line or greater, Second line, Stage I, Stage II, Stage III, Stage IV, Third line

**Phase:** I

**Therapy:** LXH254

**Locations:** Canada, France, Italy, Japan, Netherlands, Republic of Korea, Spain, Switzerland, United States

**US States:** MA, NY, TX

**Contact:** Novartis Pharmaceuticals [888-669-6682; Novartis.email@novartis.com]

**BAP1 mutation****NCT03207347**

A Phase II Trial of the PARP Inhibitor, Niraparib, in BAP1 and Other DNA Damage Response (DDR) Pathway Deficient Neoplasms (UF-STO-ETI-001)

**Cancer type:** Melanoma

**Variant class:** BAP1 mutation

**Other identifiers:** OCR15732, UF-ETG-001, UF-STO-ETI-001

**Population segments:** (N/A), Second line

**Exclusion criteria variant classes:** BRCA1 germline mutation, BRCA2 germline mutation

**Phase:** II

**Therapy:** niraparib

**Location:** United States

**US State:** FL

**Contact:** Project Management Office [352-273-6772; PMO@cancer.ufl.edu]

## BAP1 mutation (continued)

### NCT03925350

A Phase II Study of Niraparib in Patients With Advanced Melanoma With Genetic Homologous Recombination (HR) Mutation / Alteration

**Cancer type:** Melanoma

**Variant class:** BAP1 mutation

**Other identifier:** CPMC17-MEL01

**Population segments:** Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** niraparib

**Location:** United States

**US State:** CA

**Contact:** Peter Gasper [415-600-3472; gasperjp@sutterhealth.org]

### No NCT ID - see other identifier(s)

Single Arm, Open label, Signal Seeking, Phase IIa Trial Of The Activity Of Olaparib In Combination With Durvalumab In Patients With Tumours With Homologous Recombination Repair Defects

**Cancer type:** Unspecified Solid Tumor

**Variant class:** BAP1 mutation

**Other identifiers:** ACTRN12617001000392, MoST Addendum 3, U1111-1182-6652

**Population segments:** Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** durvalumab + olaparib

**Location:** Australia

### NCT03767075

Basket of Baskets: A Modular, Open-label, Phase II, Multicentre Study To Evaluate Targeted Agents in Molecularly Selected Populations With Advanced Solid Tumours

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair mutation

**Other identifiers:** (Basket of Baskets) (BoB), EudraCT number: 2017-005108-89, M039164, VHIO17002

**Population segments:** Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** atezolizumab

**Locations:** France, Spain, United Kingdom

## BAP1 mutation (continued)

### NCT03233204

NCI-COG Pediatric MATCH (Molecular Analysis for Therapy Choice)- A Phase II Subprotocol of Olaparib in Patients With Tumors Harboring Defects in DNA Damage Repair Genes

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair pathway

**Other identifiers:** 20170841, APEC1621H, NCI-2017-00766

**Population segments:** Aggressive, Indolent, Pediatric or Adolescent, Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** olaparib

**Locations:** Puerto Rico, United States

**US States:** AL, AR, AZ, CA, CO, DC, FL, GA, IA, ID, IL, IN, KY, LA, MD, ME, MI, MN, MO, MS, NC, NE, NJ, NY, OH, OK, OR, PA, SC, TN, TX, UT, VA, VT, WA, WI

**Contact:** Multiple contacts: See [www.clinicaltrials.gov](http://www.clinicaltrials.gov) for complete list of contacts.

### NCT03188965

An Open-label, First-in-human, Dose-escalation Study to Evaluate the Safety, Tolerability, Pharmacokinetics, Pharmacodynamics, and Maximum Tolerated Dose and / or Recommended Phase II Dose of the ATR Inhibitor BAY1895344 in Patients With Advanced Solid Tumors and Lymphomas

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair pathway

**Other identifiers:** 18-441, 18594, 2017-0186, BAY1895344/18594, EudraCT Number: 2016-004484-39, IRAS ID-218516, JapicCTI-183998, NCI-2018-00206

**Population segments:** Adenocarcinoma, Aggressive, Diffuse large B-cell lymphoma (DLBCL), Fourth line or greater, Hormone refractory, Indolent, Mantle cell lymphoma (MCL), Pulmonary, Second line, Squamous Cell, Stage III, Stage IV

**Phase:** I/II

**Therapy:** BAY-1895344

**Locations:** Canada, Japan, Singapore, Switzerland, United Kingdom, United States

**US States:** FL, GA, MA, NY, OH, PA, TX, UT

**Contact:** Bayer Clinical Trials Contact [888-842-2937; [clinical-trials-contact@bayer.com](mailto:clinical-trials-contact@bayer.com)]

### NCT02660034

A Phase I/Ib, Open Label, Multiple Dose, Dose Escalation and Expansion Study to Investigate the Safety, Pharmacokinetics and Antitumor Activity of the Anti-PD-1 Monoclonal Antibody BGB-A317 in Combination With the PARP Inhibitor BGB-290 in Subjects With Advanced Solid Tumors

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair mutation

**Other identifiers:** 16-183, 18-009, A317/290, BGB-A317/BGB-290, BGB-A317/BGB-290\_Study\_001, CT783, NCI-2018-00791, P 55217, VICCPHI1814

**Population segments:** Fourth line or greater, HER2 negative, Pulmonary, Second line, Stage III, Stage IV, Third line, Triple receptor negative

**Phase:** I

**Therapies:** pamiparib, tislelizumab

**Locations:** Australia, France, New Zealand, Spain, United Kingdom, United States

**US States:** AZ, CA, CO, FL, MA, TN, TX, VA

**Contact:** Rob Stewart [[clinicaltrials@beigene.com](mailto:clinicaltrials@beigene.com)]

## FANCD2 deletion

**No NCT ID - see other identifier(s)**  
Single Arm, Open label, Signal Seeking,  
Phase IIa Trial Of The Activity Of Olaparib  
In Combination With Durvalumab In  
Patients With Tumours With Homologous  
Recombination Repair Defects

**Cancer type:** Unspecified Solid Tumor

**Variant class:** FANCD2 deletion

**Other identifiers:** ACTRN12617001000392, MoST Addendum 3, U1111-1182-6652

**Population segments:** Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** durvalumab + olaparib

**Location:** Australia

### NCT03233204

NCI-COG Pediatric MATCH (Molecular  
Analysis for Therapy Choice)- A Phase  
II Subprotocol of Olaparib in Patients  
With Tumors Harboring Defects in DNA  
Damage Repair Genes

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair pathway

**Other identifiers:** 20170841, APEC1621H, NCI-2017-00766

**Population segments:** Aggressive, Indolent, Pediatric or Adolescent, Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** olaparib

**Locations:** Puerto Rico, United States

**US States:** AL, AR, AZ, CA, CO, DC, FL, GA, IA, ID, IL, IN, KY, LA, MD, ME, MI, MN, MO, MS, NC, NE, NJ, NY, OH, OK, OR, PA, SC, TN, TX, UT, VA, VT, WA, WI

**Contact:** Multiple contacts: See [www.clinicaltrials.gov](http://www.clinicaltrials.gov) for complete list of contacts.

### NCT02873975

A Phase II Study of the CHK1 Inhibitor  
LY2606368 in Patients With Advanced  
Solid Tumors Exhibiting Replicative  
Stress or Homologous Recombination  
Repair Deficiency

**Cancer type:** Unspecified Solid Tumor

**Variant class:** Fanconi anemia pathway

**Other identifiers:** 16-281, I4D-MC-E006, NCI-2016-01564

**Population segments:** Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** prexasertib

**Location:** United States

**US State:** MA

**Contact:** Dr. Geoffrey Shapiro [617-632-4942; [Geoffrey\\_S Shapiro@dfci.harvard.edu](mailto:Geoffrey_S Shapiro@dfci.harvard.edu)]

## FANCD2 deletion (continued)

### NCT02286687

Phase II Study of the PARP Inhibitor BMN 673 (Talazoparib Tosylate) in Advanced Cancer Patients With Somatic Alterations in BRCA1/2, Mutations/Deletions in PTEN or PTEN Loss, a Homologous Recombination Defect, Mutations/Deletions in Other BRCA Pathway Genes and Germline Mutation in BRCA1/2 (Not Breast or Ovarian Cancer)

**Cancer type:** Unspecified Cancer

**Variant class:** Fanconi anemia pathway

**Other identifiers:** 2013-0961, NCI-2014-02494

**Population segments:** Fourth line or greater, Stage III, Stage IV

**Phase:** II

**Therapy:** talazoparib

**Location:** United States

**US State:** TX

**Contact:** Dr. Sarina Piha-Paul [713-563-1930]

### NCT03188965

An Open-label, First-in-human, Dose-escalation Study to Evaluate the Safety, Tolerability, Pharmacokinetics, Pharmacodynamics, and Maximum Tolerated Dose and / or Recommended Phase II Dose of the ATR Inhibitor BAY1895344 in Patients With Advanced Solid Tumors and Lymphomas

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair pathway

**Other identifiers:** 18-441, 18594, 2017-0186, BAY1895344/18594, EudraCT Number: 2016-004484-39, IRAS ID-218516, JapicCTI-183998, NCI-2018-00206

**Population segments:** Adenocarcinoma, Aggressive, Diffuse large B-cell lymphoma (DLBCL), Fourth line or greater, Hormone refractory, Indolent, Mantle cell lymphoma (MCL), Pulmonary, Second line, Squamous Cell, Stage III, Stage IV

**Phase:** I/II

**Therapy:** BAY-1895344

**Locations:** Canada, Japan, Singapore, Switzerland, United Kingdom, United States

**US States:** FL, GA, MA, NY, OH, PA, TX, UT

**Contact:** Bayer Clinical Trials Contact [888-842-2937; clinical-trials-contact@bayer.com]

## ATR deletion

### No NCT ID - see other identifier(s)

Single Arm, Open label, Signal Seeking, Phase IIa Trial Of The Activity Of Olaparib In Combination With Durvalumab In Patients With Tumours With Homologous Recombination Repair Defects

**Cancer type:** Unspecified Solid Tumor

**Variant class:** ATR deletion

**Other identifiers:** ACTRN12617001000392, MoST Addendum 3, U1111-1182-6652

**Population segments:** Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** durvalumab + olaparib

**Location:** Australia



## ATR deletion (continued)

### NCT02286687

Phase II Study of the PARP Inhibitor BMN 673 (Talazoparib Tosylate) in Advanced Cancer Patients With Somatic Alterations in BRCA1/2, Mutations/Deletions in PTEN or PTEN Loss, a Homologous Recombination Defect, Mutations/Deletions in Other BRCA Pathway Genes and Germline Mutation in BRCA1/2 (Not Breast or Ovarian Cancer)

**Cancer type:** Unspecified Cancer

**Variant class:** ATR deletion

**Other identifiers:** 2013-0961, NCI-2014-02494

**Population segments:** Fourth line or greater, Stage III, Stage IV

**Phase:** II

**Therapy:** talazoparib

**Location:** United States

**US State:** TX

**Contact:** Dr. Sarina Piha-Paul [713-563-1930]

### NCT03233204

NCI-COG Pediatric MATCH (Molecular Analysis for Therapy Choice)- A Phase II Subprotocol of Olaparib in Patients With Tumors Harboring Defects in DNA Damage Repair Genes

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair pathway

**Other identifiers:** 20170841, APEC1621H, NCI-2017-00766

**Population segments:** Aggressive, Indolent, Pediatric or Adolescent, Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** olaparib

**Locations:** Puerto Rico, United States

**US States:** AL, AR, AZ, CA, CO, DC, FL, GA, IA, ID, IL, IN, KY, LA, MD, ME, MI, MN, MO, MS, NC, NE, NJ, NY, OH, OK, OR, PA, SC, TN, TX, UT, VA, VT, WA, WI

**Contact:** Multiple contacts: See [www.clinicaltrials.gov](http://www.clinicaltrials.gov) for complete list of contacts.

### NCT02873975

A Phase II Study of the CHK1 Inhibitor LY2606368 in Patients With Advanced Solid Tumors Exhibiting Replicative Stress or Homologous Recombination Repair Deficiency

**Cancer type:** Unspecified Solid Tumor

**Variant class:** Fanconi anemia pathway

**Other identifiers:** 16-281, I4D-MC-E006, NCI-2016-01564

**Population segments:** Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** prexasertib

**Location:** United States

**US State:** MA

**Contact:** Dr. Geoffrey Shapiro [617-632-4942; [Geoffrey\\_Shapiro@dfci.harvard.edu](mailto:Geoffrey_Shapiro@dfci.harvard.edu)]



## ATR deletion (continued)

### NCT03188965

An Open-label, First-in-human, Dose-escalation Study to Evaluate the Safety, Tolerability, Pharmacokinetics, Pharmacodynamics, and Maximum Tolerated Dose and / or Recommended Phase II Dose of the ATR Inhibitor BAY1895344 in Patients With Advanced Solid Tumors and Lymphomas

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair pathway

**Other identifiers:** 18-441, 18594, 2017-0186, BAY1895344/18594, EudraCT Number: 2016-004484-39, IRAS ID-218516, JapicCTI-183998, NCI-2018-00206

**Population segments:** Adenocarcinoma, Aggressive, Diffuse large B-cell lymphoma (DLBCL), Fourth line or greater, Hormone refractory, Indolent, Mantle cell lymphoma (MCL), Pulmonary, Second line, Squamous Cell, Stage III, Stage IV

**Phase:** I/II

**Therapy:** BAY-1895344

**Locations:** Canada, Japan, Singapore, Switzerland, United Kingdom, United States

**US States:** FL, GA, MA, NY, OH, PA, TX, UT

**Contact:** Bayer Clinical Trials Contact [888-842-2937; clinical-trials-contact@bayer.com]

## SETD2 mutation

### NCT03767075

Basket of Baskets: A Modular, Open-label, Phase II, Multicentre Study To Evaluate Targeted Agents in Molecularly Selected Populations With Advanced Solid Tumours

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair mutation

**Other identifiers:** (Basket of Baskets) (BoB), EudraCT number: 2017-005108-89, MO39164, VHIO17002

**Population segments:** Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** atezolizumab

**Locations:** France, Spain, United Kingdom

### NCT03233204

NCI-COG Pediatric MATCH (Molecular Analysis for Therapy Choice)- A Phase II Subprotocol of Olaparib in Patients With Tumors Harboring Defects in DNA Damage Repair Genes

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair pathway

**Other identifiers:** 20170841, APEC1621H, NCI-2017-00766

**Population segments:** Aggressive, Indolent, Pediatric or Adolescent, Second line, Stage III, Stage IV

**Phase:** II

**Therapy:** olaparib

**Locations:** Puerto Rico, United States

**US States:** AL, AR, AZ, CA, CO, DC, FL, GA, IA, ID, IL, IN, KY, LA, MD, ME, MI, MN, MO, MS, NC, NE, NJ, NY, OH, OK, OR, PA, SC, TN, TX, UT, VA, VT, WA, WI

**Contact:** Multiple contacts: See www.clinicaltrials.gov for complete list of contacts.

## SETD2 mutation (continued)

### NCT03188965

An Open-label, First-in-human, Dose-escalation Study to Evaluate the Safety, Tolerability, Pharmacokinetics, Pharmacodynamics, and Maximum Tolerated Dose and / or Recommended Phase II Dose of the ATR Inhibitor BAY1895344 in Patients With Advanced Solid Tumors and Lymphomas

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair pathway

**Other identifiers:** 18-441, 18594, 2017-0186, BAY1895344/18594, EudraCT Number: 2016-004484-39, IRAS ID-218516, JapicCTI-183998, NCI-2018-00206

**Population segments:** Adenocarcinoma, Aggressive, Diffuse large B-cell lymphoma (DLBCL), Fourth line or greater, Hormone refractory, Indolent, Mantle cell lymphoma (MCL), Pulmonary, Second line, Squamous Cell, Stage III, Stage IV

**Phase:** I/II

**Therapy:** BAY-1895344

**Locations:** Canada, Japan, Singapore, Switzerland, United Kingdom, United States

**US States:** FL, GA, MA, NY, OH, PA, TX, UT

**Contact:** Bayer Clinical Trials Contact [888-842-2937; clinical-trials-contact@bayer.com]

### NCT02660034

A Phase I/Ib, Open Label, Multiple Dose, Dose Escalation and Expansion Study to Investigate the Safety, Pharmacokinetics and Antitumor Activity of the Anti-PD-1 Monoclonal Antibody BGB-A317 in Combination With the PARP Inhibitor BGB-290 in Subjects With Advanced Solid Tumors

**Cancer type:** Unspecified Solid Tumor

**Variant class:** DNA repair mutation

**Other identifiers:** 16-183, 18-009, A317/290, BGB-A317/BGB-290, BGB-A317/BGB-290\_Study\_001, CT783, NCI-2018-00791, P 55217, VICCPHI1814

**Population segments:** Fourth line or greater, HER2 negative, Pulmonary, Second line, Stage III, Stage IV, Third line, Triple receptor negative

**Phase:** I

**Therapies:** pamiparib, tislelizumab

**Locations:** Australia, France, New Zealand, Spain, United Kingdom, United States

**US States:** AZ, CA, CO, FL, MA, TN, TX, VA

**Contact:** Rob Stewart [clinicaltrials@beigene.com]

## SMO mutation

**No NCT ID - see other identifier(s)**  
Single Arm, Open Label, Signal Seeking, Phase IIa Trial of The Activity of Vismodegib in Patients With Tumours Harboursing PTCH1 or SMO Mutations

**Cancer type:** Unspecified Solid Tumor

**Variant class:** SMO mutation

**Other identifiers:** ACTRN12618000281291, CTC0141-addendum 4, MoST Addendum 4, U1111-1182-6652

**Population segments:** Second line, Stage III, Stage IV

**Exclusion criteria variant classes:** SMO D473Y mutation, SMO G497W mutation, SUFU mutation

**Phase:** II

**Therapy:** vismodegib

**Locations:** Australia, New Zealand

## SMO mutation (continued)

### NCT02465060

Molecular Analysis for Therapy Choice (MATCH)

**Cancer type:** Unspecified Solid Tumor

**Variant class:** SMO mutation

**Other identifiers:** 15-7002, 16-750, AAP9159, CTSU/EAY131, EAY131, EAY131- K1, EAY131- K2, EAY131- Z1K, EAY131-A, EAY131-B, EAY131-C1, EAY131-C2, EAY131-E, EAY131-F, EAY131-G, EAY131-H, EAY131-I, EAY131-J, EAY131-L, EAY131-M, EAY131-MATCH, EAY131-N, EAY131-P, EAY131-Q, EAY131-R, EAY131-S1, EAY131-S2, EAY131-T, EAY131-U, EAY131-V, EAY131-W, EAY131-X, EAY131-Y, EAY131-Z1A, EAY131-Z1B, EAY131-Z1C, EAY131-Z1D, EAY131-Z1E, EAY131-Z1F, EAY131-Z1G, EAY131-Z1H, EAY131-Z1I, EAY131-Z1J, EAY131-Z1L, EAY131-Z1M, ECOG-ACRIN EAY131, ECOGEAY131-M, MATCH, NCI-2015-00054, NCI-MATCH

**Population segments:** Aggressive, BRAF, Classical, Fourth line or greater, HER2 positive, Indolent, Nodular lymphocyte-predominant, Pulmonary, Second line, Stage III, Stage IV, Third line, Unspecified

**Phase:** II

**Therapy:** vismodegib

**Locations:** Puerto Rico, United States

**US States:** AK, AL, AR, AZ, CA, CO, CT, DC, DE, FL, GA, HI, IA, ID, IL, IN, KS, KY, LA, MA, MD, ME, MI, MN, MO, MS, MT, NC, ND, NE, NH, NJ, NM, NV, NY, OH, OK, OR, PA, RI, SC, SD, TN, TX, UT, VA, VT, WA, WI, WV, WY

**Contact:** Multiple contacts: See [www.clinicaltrials.gov](http://www.clinicaltrials.gov) for complete list of contacts.

### NCT03297606

Canadian Profiling and Targeted Agent Utilization Trial (CAPTUR): A Phase II Basket Trial

**Cancer type:** Unspecified Solid Tumor

**Variant class:** SMO aberration

**Other identifiers:** CA209-9DL, CAPTUR, ESR-17-12831, ML39800, PM1, WI233446

**Population segments:** Aggressive, Diffuse large B-cell lymphoma (DLBCL), Extranodal marginal zone B-cell lymphoma (MALT), First line, Follicular lymphoma (FL), Indolent, Lymphoblastic lymphoma (LBL), Mantle cell lymphoma (MCL), Other subtype, Second line, Stage III, Stage IV, Waldenstrom's macroglobulinemia (WM)

**Phase:** II

**Therapy:** vismodegib

**Location:** Canada

## Evidence Summary by Variant Class

A variant class hierarchy was created to summarize gene variants with associated clinical evidence. Evidence items refers to citations across the different global data sources.

### GNA11 mutation

Variant Class	Evidence Items
RAS/RAF/MEK/ERK pathway	5
↳ RAS/RAF/MEK/ERK mutation	3
↳ GNA11 mutation	1

### BAP1 mutation

Variant Class	Evidence Items
DNA repair pathway	8
↳ DNA repair mutation	4
↳ BAP1 mutation	3

### FANCD2 deletion

Variant Class	Evidence Items
DNA repair pathway	8
↳ DNA repair deletion	0
↳ FANCD2 deletion	0
Fanconi anemia pathway	3
↳ FANC deletion	1
↳ FANCD2 deletion	0

## Evidence Summary by Variant Class (continued)

A variant class hierarchy was created to summarize gene variants with associated clinical evidence. Evidence items refers to citations across the different global data sources.

### ATR deletion

Variant Class	Evidence Items
DNA repair pathway	8
↳ DNA repair deletion	0
↳ ATR deletion	2
Fanconi anemia pathway	3
↳ ATR deletion	2

### SETD2 mutation

Variant Class	Evidence Items
DNA repair pathway	8
↳ DNA repair mutation	4
↳ SETD2 mutation	0

### SMO mutation

Variant Class	Evidence Items
SMO aberration	1
↳ SMO mutation	2

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Report Authorised by

Signed



printed

Keeda Hardisty

Clinical Scientist ☒

Pathologist ☐

Report reviewed by

Signed



printed

Kaiya Chowdhary

Clinical Scientist ☒

BMS ☐





**oncologica®**



Oncologica UK Ltd, Suite 2, The Newnham Building,  
Chesterford Research Park, Little Chesterford,  
Cambridge, CB10 1XL

**+44 (0) 1223 785 327 - [info@oncologica.com](mailto:info@oncologica.com)**

## Ireland

Bymac Centre, Northwest  
Business Park, Blanchardstown, Dublin 15

**+353 1 8604204**

## Italy

Parco Tecnologico della Sardegna  
Pula, Località Piscinamanna

**+39 02 808 88210**