- 1. Project title: Genomic Report and Data Interoperability with Epic
- 2. Submitter's name, title, and contact information

Michael Marchant, Director – Health Information Exchange – . 916 7340942 – mbmarchant@ucdavis.edu

3. Names of project leader(s) and team members

Willy Bansi, David Vaillancourt, Scott Nathanson, Marvin Asercion, Paul Robbins

4. A simple short statement (even just one sentence) summarizing what the project does

UC Davis Health built a context aware link in Epic giving providers access to discrete data generated from genomic reports provided by Foundation Health with live link access to clinical trial information embedded in the report.

5. A project narrative that:

• Describes the problem being solved and the project goals

Genomic reports are delivered to UC Davis Health by mail, fax or by hand and then are scanned into the EHR and placed in the media tab with other scanned documents. The project goal is to take those scanned documents and create a dynamic frame of discrete data and links in context in the patient's chart with access to information provided by the genomic reporting agency and allow real time access and retrieval of those reports.

• Emphasize the solution and innovation, rather than technical detail

This is an embedded link in Epic, which will render the Genomics data placed in an external data storage. The connection leverages a FHIR API along with Single Sign On (SSO) technology. SSO is used to authentication users between the two systems, and FHIR Resources are used to pass the Patient and Provider information between Epic and Healthshare. The Embedded viewer GUI (Graphical Unit Interface) mimics Epic's and gives the user an experience that they never left Epic.

• Tell how the solution has impacted customers/users

Providers and Genetic Counselors were spending time looking for scanned reports in Epic's media tab and finding reports were difficult due to the number of scanned documents found there. The integration saves them time as they can easily locate patient reports. The embedded integration adds interactivity and search capability. Providers can search for key words and it gives them the ability to quickly narrow down available clinical trials based on study locations, Genes Types or Bio Markers. The report includes links that take providers to the ClinicalTrials.gov site to get more detail on the specifics of the research study.

- Explains how project success was measured
 - o Budget & Project Schedule: We were on Budget and Project went live on schedule
 - Customer Satisfaction: Gathered requirements from our customers to make sure all the deliverables were met
 - Reporting: All transactions and activities are being logged and reports are in place showing the user activities.
 - Team Satisfaction: As we were developing a new capability and provided team members an opportunity to innovate, all team members were enthusiastic and engaged and their productivity showed the results
 - Quality of work: We had stakeholders who were able to perform quality audits to make sure their requirements were met.
- Highlighted collaboration with other locations, departments, or teams
 - Multiple Teams were involved in the collaboration Systems Integration, Epic Applications, Cancer Center Providers, Infrastructure Team, Foundation Medicine and our vendors (J2 and InterSystems).

- Timeframe of deployment
 - This project took roughly 4 months from start to finish.
- Describe the technology utilized
 - This integration uses Web Services, FHIR with oAuth, Single Sign On as well as the Epic Interconnect and InterSystems Healthshare platform.
- Optional: Relevant screenshots

The **Genomics Results** activity opens the **Genomics** tab. Available result reports are listed. Double-click the report link to open the desired report.

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Genomics			? ×
Date	Description	Status	Provider
Historic			
11/13/2019	FoundationOne CDX	Report Received	Ruskin, Rachel
11/12/2019	FoundationOne Heme	Report Received	Canter, Robert

At the top of the report, the header provides information about the test.

- Click the PDF hyperlink to view the Complete Report. The PDF report was previously viewed when scanned to the Chart Review Media tab.
- Click through the tabs to access the sections of the report for **Genomic Results**, **Therapies**, **Clinical Trials**, and **Variants of Unknown Significance**.

Report Date	Nov 13, 2019	Tumor Type	Uterus endometrial adenocarcinoma (NOS)	Specimen Site	Lymph Node
Case #	ORD-0685340-01-SIT	Ordering Facility	UC Davis Comprehensive Cancer Center	Specimen Type	Slide Deck
Test Type	FoundationOne CDX	Ordering Physician	Ruskin, Rachel	Specimen Received	Oct 29, 2019
Complete Report	PDF	Pathologist	Afify, Alaa	Date of Collection	Sep 17, 2019

View Genomic Results:

Genomic Results (9) Therapies (15) Clinical Trials (26) Variants of Unknown Significance	e (8)		Q	filter	Reset	
nic Results (9) Therapies (15) Clinical Trials (26) Variants of Unknown Significance TIA	í	ARID1A encodes the AT-rich interactive domain-cont Bat250a, a member of the SW//SNF chromatin remod inactivation of ARID1A has been reported in many ca tumor suppressor 18 19 180 121 12 10 14 15 18, ARID1A	nain-containing protein 1A, also known as tin remodeling complex. Mutation, loss, or 1 many cancers, and the gene is considered a 24, ARID1A mutations, which are mostly			
ERBB2 Alteration: amplification Therapies for other tumor types: Trastuzumab-pkrb, Lapatinib, Trastuzumab, Trastuzumab-qyyp, Ado-trastuzumab emtansine, Trastuzumab-anns, Pertuzumab, Afatinib, Trastuzumab-ditb, Trastuzumab-dist, Neratinib, Dacomitinib Clinical Trials		truncating, have been identified along the entire gen protein loss ¹¹⁹ ¹³⁰ ¹³¹ ¹³⁷ ¹³⁸ , whereas ARID1A missens uncharacterized, ARID1A alterations are particularly p (46-50%), ovarian and uterine endometrioid carcinom (27%): they are also reported in up to 27% of gastric adenocarcinoma. Waldenstrom macroglobulinemia, p hepatocellular carcinoma, colorectal carcinoma (CRC)	e and o e mutat revalen as (24 carcinor ediatric . and ur	ften correlate with AR tions are mostly t in ovarian clear cell e 44%), and cholangioci ma, esophageal c Burkitt lymphoma, rothelial carcinoma sai	ID1A arcinoma arcinoma mples	
PIK3CA Alteration: E545K Therapies for other tumor types: Alpelisib, Temsirolimus, Everolimus Clinical Trials	ï	analyzed (COSMIC, cBioPortal, 2019) 119 129 130 131 132 13 microsatellite instability in ovarian and endometrial e 134 135 139 137, CRC 138 139 149, and gastric cancer 135 138 no correlation between ARID1A loss and clinicopatho or endometrioid carcinomas or other endometrial car	ARID ndomet at taz ta logical ncers	1A loss is associated w trioid adenocarcinoma ¹ . Several studies have parameters in ovarian ⁴ ¹⁴⁵ ¹⁴⁶ ¹⁴⁷ , whereas ot	reported clear cell hers	
BCOR Alteration: N1425S		suggest that ARID1A loss is a negative prognostic fac approved to address the mutation or loss of ARID1A limited clinical and preclinical evidence, ARID1A inact	: factor ¹⁴⁸ ¹⁴⁹ . There are no therapies)1A in cancer. However, on the basis o nactivating mutations may lead to			
Microsatellite status Alteration: MS-Stable	A RID 1A encodes the AT-rich interactive domain-containing Bat250a, a member of the SWI/SNF chromatin remodeling i inactivation of ARID1A has been reported in many cancers, timer suppressor 19 19 10 tot tot 20 10 at 15 100, ARID1A mutation of ARID1A has been reported in many cancers, timer suppressor 19 19 10 tot 10 20 10 10 20 10 at 15 100, ARID1A missense mut uncharacterized. ARID1A alterations are particularly prevale (46-50%), ovarian and uterine endometrioid carcinoma (24 (25%); they are also reported in up to 27% of gastric carcina and uterine endometrioid carcinoma (24 (25%); they are also reported in up to 27% of gastric carcina and uterine endometrioid carcinoma (24 (25%); they are also reported in up to 27% of gastric carcina (24 (25%); they are also reported in up to 27% of gastric carcina (24 (25%); they are also reported in up to 27% of gastric carcina (24 (25%); they are also reported in up to 27% of gastric carcina (24 (25%); they are also reported in up to 27% of gastric carcina (24 (25%); they are also reported in up to 27% of gastric carcina (24 (25%); they are also reported in up to 27% of gastric carcina (24 (25%); they are also reported in up to 27% of gastric carcina (24 (25%); they are also reported in up to 27% of gastric carcina action ad endometrioid carcinoma (24 (25%); they are also reported in up to 27% of gastric carcina (25%); they are also reported in up to 27% of gastric carcina advective (25%); they are also reported in up to 27% of gastric carcina advective (25%); they are also reported in up to 27% of gastric carcina advective (25%); ovarian and uterine endometrioid carcinoma (26, and up to 19 10 19 19 19 19 19 19 19 19 19 19 19 19 19	with sm with M6 m studi	hall cell lung cancer ha 620 combined with to ies in ovarian cancer, A	potecan ARID1A		
PPP2R1A Alteration: \$256F		inactivation may predict sensitivity to inhibitors of EZ in clinical trials. Other studies have reported that loss pathway and be linked with sensitivity to inhibitors of expression has been associated with chemoresistance	of ARIE this pa	-: which are under inv 21A may activate the F ithway ¹⁵² ¹⁵⁴ ¹⁵⁵ , Loss inum-based therapy i	estigation PI3K-AKT of ARID1A n patients	
SMARCA4 Alteration: rearrangement exon 20		with ovarian clear cell carcinoma ¹⁴⁹ ¹⁵⁰ and to 5-fluo	ouracil	(5-FU) in CRC cell line	s 157	

View Therapies:



Therapies Tab

View Clinical Trials:

- Filter clinical trial results by location, Genes, or keyword(s)
- Click an NCTID hyperlink to go directly to the NLM listing on ClinicalTrials.gov

Genomic Results	(9) Therapies (15) Clinical Trials (26)		Q location		All Genes	;	Q	filter		Reset
IMPORTANT: Clinica every effort is made to complete list of avail below into the search	I be supported by gene and prioritized by: age range inclusion criteria for to ensure the accuracy of the information contained below, the information av able trials. In order to conduct a more thorough search, please go to www.clin h bar.	pediatric patio ailable in the icaltrials.gov o	ents, proximity to ordering medical facili public domain is continually updated a and use the search terms provided below	ity, later trial ; nd should be i w. For more in,	hase, and verification o nvestigated by the phys formation about a speci	f trial i ician o fic clini	informa r resear ical tria	tion within the la ch staff. This is n l, type the NCT la	ast two months. Iot meant to be D of the trial ini	While a dicated
Gene	Title	Phase	Targets	Loca	tions				NCTID	^
ARID1A (S617fs*6)	ATR Kinase Inhibitor VX-970 and Irinotecan Hydrochloride in Treating Patients With Solid Tumors That Are Metastatic or Cannot Be Removed by Surgery	PHASE 1	ATR	Califo Mich Penn	rnia, Connecticut, F gan, Missouri, Nort sylvania, Tennessee	lorida h Car	a, Mas olina,	sachusetts,	NCT02595	931
ARID1A (S617fs*6)	Ascending Doses of AZD6738 in Combination With Chemotherapy and/or Novel Anti Cancer Agents	PHASE 1/2	ATR, PARP, PD-L1	Califo Villeji of), S (Uniti Mano Kingo	rnia, New York, Sai uif (France), Seongn eoul (Korea, Republ ed Kingdom), Londo hester (United King Iom)	nt He am-si ic of), on (Ur idom)	rblain i (Kore . Caml nited H I, Sutte	(France), ea, Republic oridge Kingdom), on (United	NCT02264	678
ARID1A (S617fs*6)	Trial of Topotecan With VX-970, an ATR Kinase Inhibitor, in Small Cell Cancers	PHASE 1/2	ATR	Mary	and				NCT02487	095
ARID1A (S617fs*6)	Veliparib (ABT-888), an Oral PARP Inhibitor, and VX-970, an ATR Inhibitor, in Combination With Cisplatin in People With Refractory Solid Tumors	PHASE 1	PARP, ATR	Mary	and, Massachusett	s, Tex	as		NCT02723	864

Clinical Trials Tab



View Variants of Unknown Significance:

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Report Date Case # Test Type	Nov 13, 2019 ORD-0685340-01-SIT FoundationOne CDX	Tumor Type Ordering Facility Ordering Physician	Uterus endometrial adenocarcinoma (NOS) UC Davis Comprehensive Cancer Center Ruskin, Rachel	Specimen Site Specimen Type Specimen Received	Lymph Node Slide Deck Oct 29, 2019	×
Complete Report	PDF	Pathologist	Afify, Alaa	Date of Collection	Sep 17, 2019	_
Senomic Results (9	9) Therapies (15) Clir	nical Trials (26) Va	ariants of Unknown Significance (8)	Q filter	Re	set
erature at the time th at they become clinic Gene	is report was issued, and/or t ally meaningful in the future. Variant	ne genomic context of	Microsatellite status	e choose to include them	nere in the event	
			MSS			
RBM10	V238M		Tumor Mutation Burden			
NTRK2	M713K		intermediate; 6.3 mutations-per-m	egabase		
ARFRP1	A123V					
ASXL1	K85R					
PTEN	M35K					
CCND1	E279*					
			~			

Variants of Unknown Significance Tab