

# Large scale analysis in von Hippel-Lindau disease

Andreea Chioorean

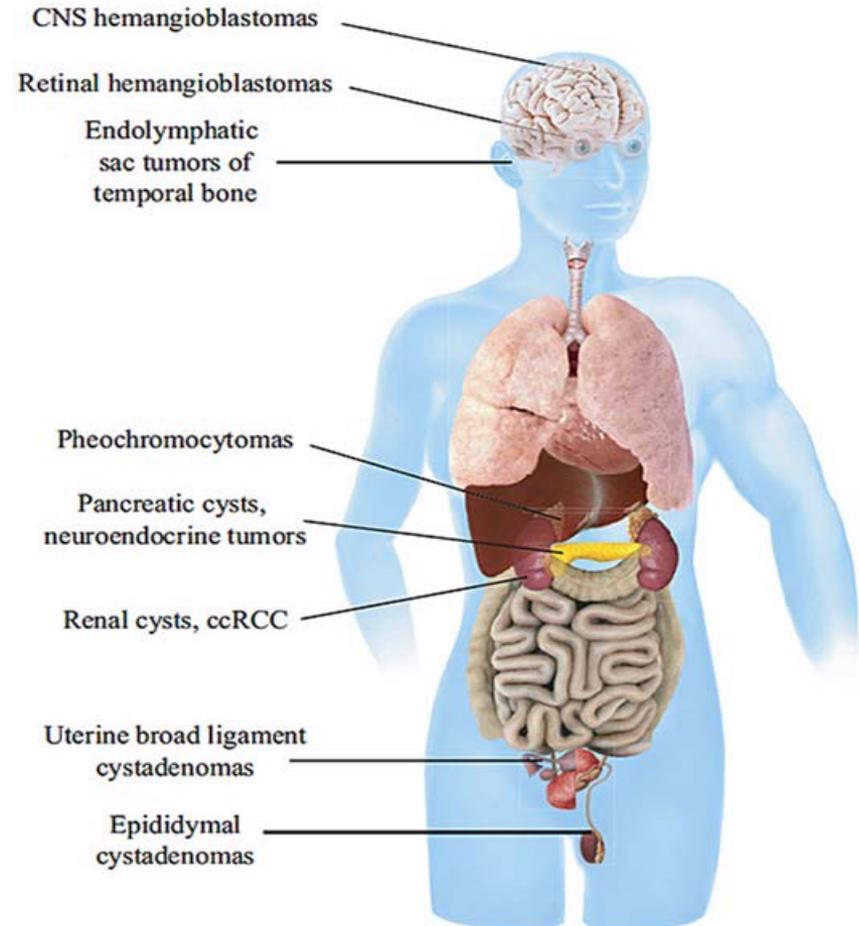
MD Candidate, University of Toronto

# Conflict of Interests

- None to declare

# Von Hippel Lindau

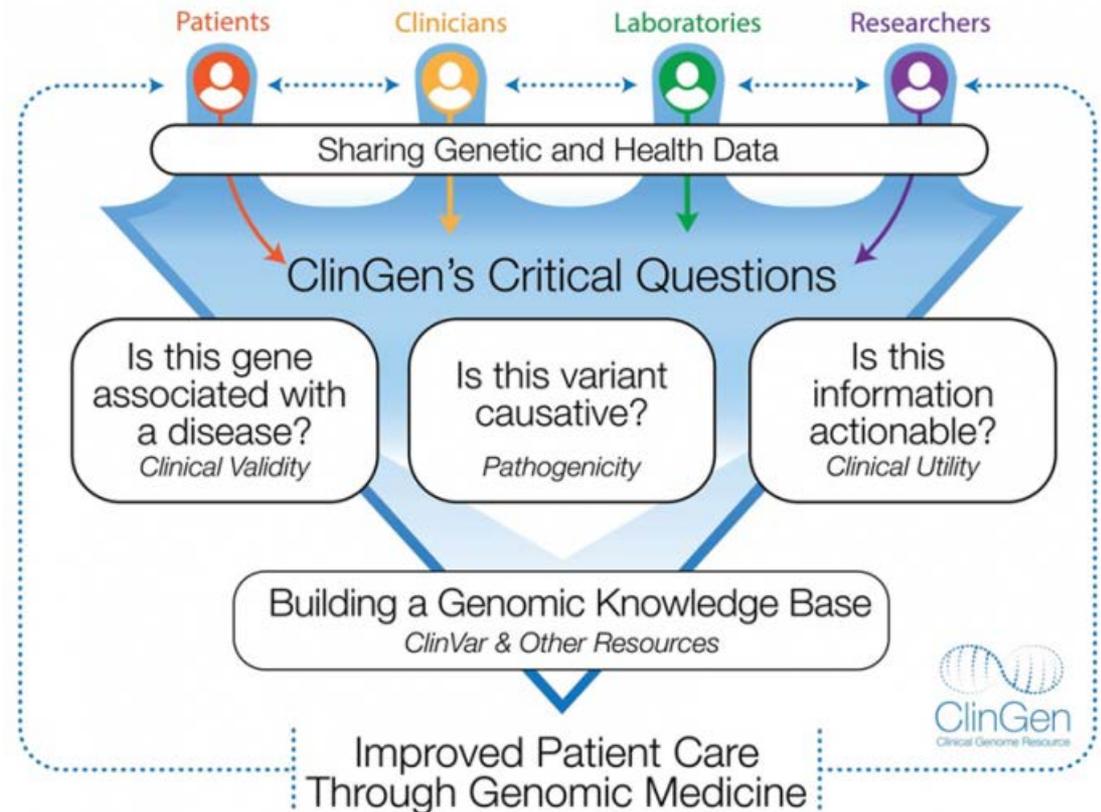
- Multi-system disorder
- Mutations in *VHL* gene responsible for degradation of hypoxia inducible factor 1-alpha (HIF1 $\alpha$ )
- Results in hemangioblastomas
  - Eye
  - Brain
  - Spine
- Deafness (endolymphatic sac tumours)
- Pancreatic cysts
- variable expressivity = not all mutation carriers develop all manifestations
- Frameshift mutations result in risk of renal cell carcinoma “type 1”
- Missense mutation result in pheochromocytoma “type 2”
- Genetics is not straightforward

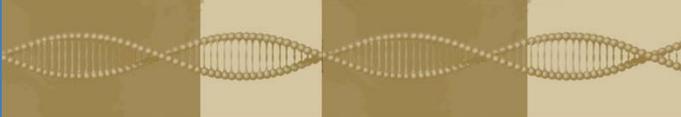


Adapted with permission from Ho & Jonasch (2014)

# VHL vs other genes....

- Large gene-based consortium
- BRCA1/2 (ENIGMA)
- Cystic fibrosis
- Hereditary Colon Cancer (INSIGHT)





# VHL Case Studies

## Scenario 1: Clinical Case Study

- ✦ 23 year old patient with renal cell carcinoma
- ✦ Gene panel testing reveals a germline *VHL* variant:
  - c.345C>G [p.H115Q missense mutation]
  - Lab classification: Variant of unknown significance
- ✦ Family history is negative for VHL disease
  - No VHL mutations in family members
- ✦ Does this patient have VHL disease?
- ✦ Should this patient receive life-long surveillance?

# Case study Patient Scenario: ClinVar gene database

- Assertion for (*VHL* c.345C>G) mutation is **inconclusive** on ClinVar, no phenotype described:

**NM\_000551.3(VHL):c.345C>G (p.His115Gln)**

## Interpretation ?

Go to:

Clinical significance: [Uncertain significance](#)

Last evaluated: Feb 5, 2016

Number of submission(s): 1

## Assertion and evidence details

Go to:

Clinical assertions

Summary evidence

Supporting observations

Submitter	Allele origin	Individuals	Phenotypes (Affected status)	Ethnicity	Geographic origin	Citations	Description
<a href="#">Integrated Genetics/Laboratory Corporation of America</a>	germline	not provided	not provided (unknown)	not provided	not provided	• <a href="#">PubMed</a>	not provided

# VISION

## VHL Information-Sharing International cONsortium

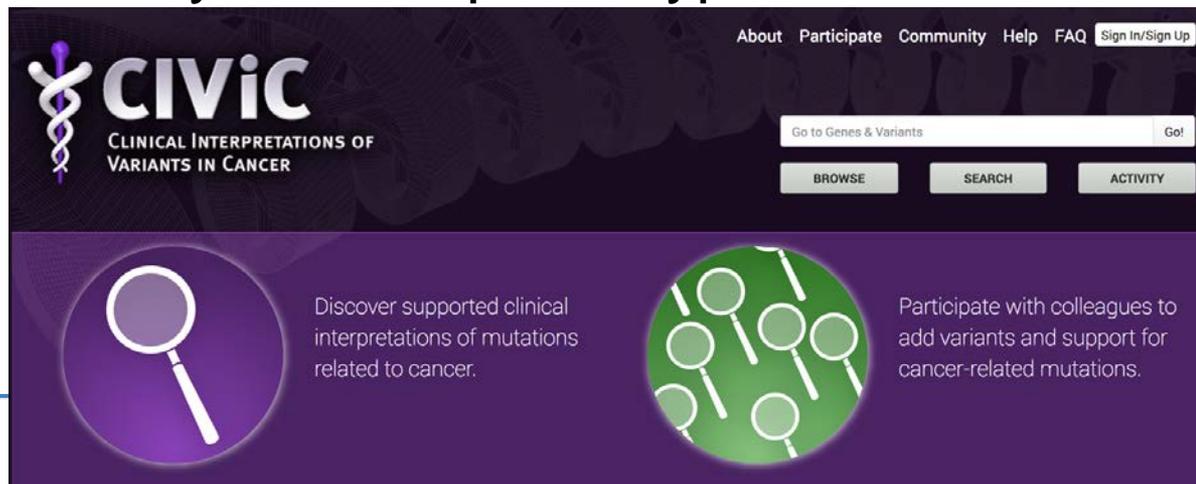
**Mandate:** Improve VHL disease understanding and treatment on a global scale by sharing genomic and clinical information.

### Projects:

- Curate a database of VHL variants
- Establish an Expert Panel to:
  1. Create rules for VHL variant interpretation
  2. Use VHL rules to identify pathogenic variants
- Freely share data with the VHL community around the world

# Housing *VHL* Variants in CIViC

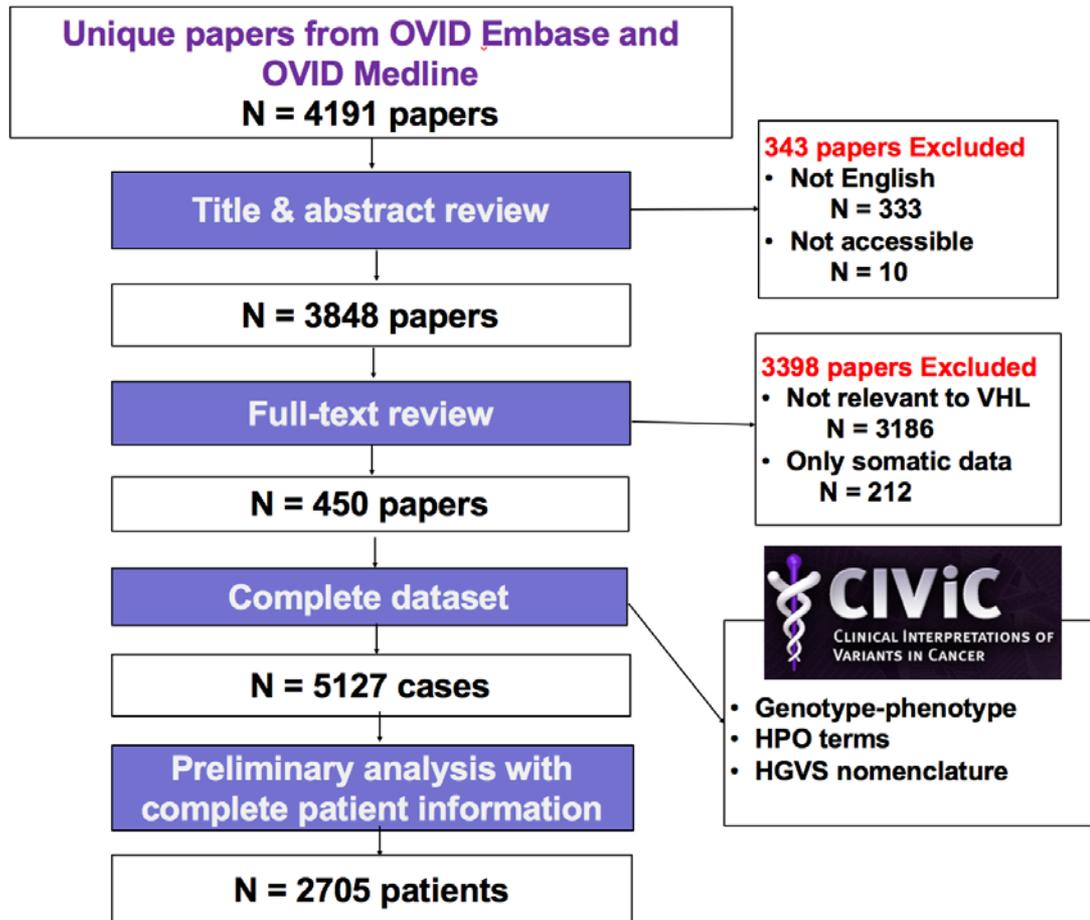
- *VHL* information is scattered and unorganized
- Open access, community-driven web resource for Clinical Interpretation of Variants in Cancer
- Standardized (Human Phenotype Ontology)
- Centralized, debated, and interpreted data of associations between specific mutations, phenotypes, and responses to a targeted therapy
- Search by variant, phenotype, disease, therapy



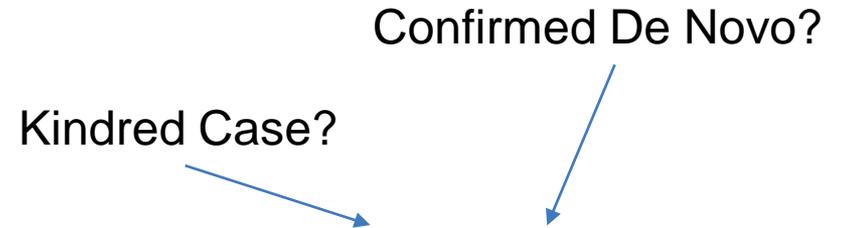
Discover supported clinical interpretations of mutations related to cancer.

Participate with colleagues to add variants and support for cancer-related mutations.

# Assembling all VHL cases ever published



# What does the data look like?



# What does the data look like?

Age Type?

- Evaluated
- Last known
- Death

Resolution?

- Patient
- Family
- Variant
- Tumor



fc	Phenotype	Reference	Age	Sex	Notes	Evidence Stat	Resolution	Input	Warnings	HGVS_transcrip	HGVS_intronic
ry	HP:0009711	(Alegret et al., 2011)	E14Y0M; lk15Y	M	resulting in in-f	A case study o	Tumour	NM_000551.3:c.227_229del		NM_000551.3:c.227_229del	
sc	HP:0010797; H	(AlFadhli et al., 2011)	unknown	F	index III-9	The germline V	Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	F	III-8		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	F	III		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	F	III		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	M	III		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	M	III		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	M	III		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	F	II		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	F	II		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	M	II		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	M	II		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	M	II		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
st	HP:0010797; H	(AlFadhli et al., 2011)	unknown	F	I		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
ry	HP:0002666; H	(Allen et al., 2011)	E11Y0M; lk22Y	unknown	patient #1 ; ph	Family history	Patient	NM_000551.3:c.292T>C		NM_000551.3:c.292T>C	
	none	(Allen et al., 2011)	lk24Y0M	unknown	patient #2; diagnosed at 24 w		Patient	NM_000551.3:c.292T>C		NM_000551.3:c.292T>C	
	none	(Allen et al., 2011)	lk39Y0M	unknown	patient #3; diagnosed at 39 w		Patient	NM_000551.3:c.292T>C		NM_000551.3:c.292T>C	

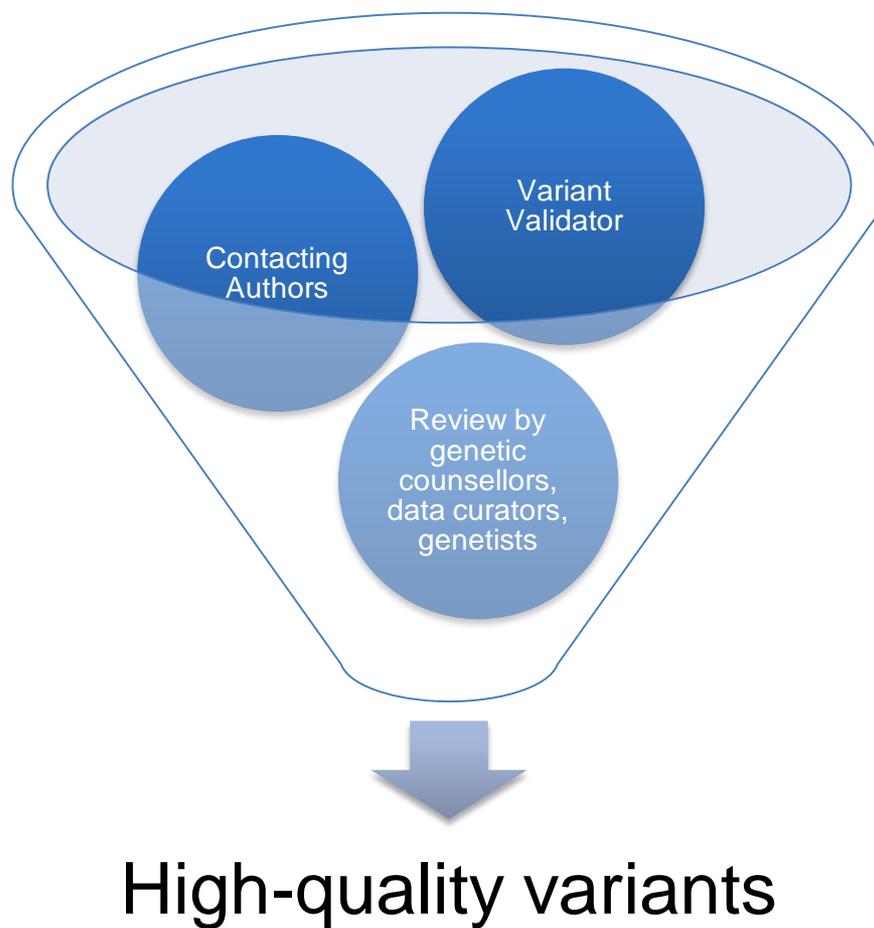
# What does the data look like?



Reference	Age	Sex	Notes	Evidence Stat	Resolution	Input	Warnings	HGVSc_transcript	HGVSc_intronic
(Alegret et al., 2015)	E14Y0M; 1k15Y0M	M	resulting in in-frame deletion	A case study of	Tumour	NM_000551.3:c.227_229del		NM_000551.3:c.227_229del	
; H (AlFadhli et al., 2015)	unknown	F	index III-9	The germline variant	Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	F	III-8		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	F	III		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	F	III		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	M	III		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	M	III		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	M	III		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	F	II		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	F	II		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	M	II		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	M	II		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	M	II		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (AlFadhli et al., 2015)	unknown	F	I		Patient	NM_000551.3:c.351G>T		NM_000551.3:c.351G>T	
; H (Allen et al., 2015)	E11Y0M; 1k22Y0M	unknown	patient #1 ; phenotype	Family history	Patient	NM_000551.3:c.292T>C		NM_000551.3:c.292T>C	
(Allen et al., 2015)	1k24Y0M	unknown	patient #2; diagnosed at 24 weeks		Patient	NM_000551.3:c.292T>C		NM_000551.3:c.292T>C	
(Allen et al., 2015)	1k39Y0M	unknown	patient #3; diagnosed at 39 weeks		Patient	NM_000551.3:c.292T>C		NM_000551.3:c.292T>C	

Wait, what's this section...?

# Validating the data



# Back to the case, c.345C>G

## CIViC: Search By Variant

Match  of the following conditions:

Gene Name
▼
contains
▼
VHL
✕

Variant Name
▼
contains
▼
c.345C>G
✕

Variant Origin
▼
is
▼
Germline Mutation
✕ +

Search

### Search Results 4 total items

Get Data

Help

EID	GENE...	VARIANT...	DESC	DIS	DRUGS	EL ▲	ET	ED	CS	VO	TR ▼	☰
5515	VHL	H115Q (c....	Study of 103 patients...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	ⓘ	⋮	4 ★	
5274	VHL	H115Q (c....	Screening of 92 unrel...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	ⓘ	⋮	3 ★	
5494	VHL	H115Q (c....	Two cases of VHL pat...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	ⓘ	⋮	3 ★	
5755	VHL	H115Q (c....	This study reports 1,5...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	ⓘ	⋮	2 ★	

# Case Study: Conclusion

- CIViC Evidence Supports c.345C>G Pathogenicity
- Ask lab to reclassify, pathogenic a severe variant

**Management decision:** Patient should undergo surveillance and likely has VHL disease

# CIViC vs Clinvar Search By Phenotype

The screenshot shows the CIViC search interface. At the top left is the CIViC logo. To the right are navigation links: About, Participate, Community, Help, FAQ, and Sign In/Sign Up. Below the navigation is a purple bar with buttons for 'Go to Genes & Variants', 'Go!', 'BROWSE', 'SEARCH', 'ACTIVITY', and 'ADD'. The main content area is titled 'Search Evidence' and has tabs for 'Evidence', 'Assertions', 'Variants', 'Genes', and 'Sources'. Under 'Example Searches', there are buttons for 'High Quality ALK Evidence', 'High Quality Predictive Evidence', 'High Quality Drug Predictions', and 'Alectinib Evidence'. The search criteria section is highlighted with a red box and includes a 'Match' dropdown set to 'all' and the text 'of the following conditions:'. Two conditions are listed: 'Phenotype HPO class contains renal cell carcinoma' and 'Gene Name contains VHL'. A 'Search' button is at the bottom right of the criteria section.

# CIViC: Search By Phenotype

Search Results 421 total items

Get Data

Help

EID	GENE	VARIANT	DESC	DIS	DRUGS	EL	ET	ED	CS	VO	TR
5129	VHL	N131T (c....	Genotype-phenotype ...	Von Hippel-Lindau Di...	N/A	B	⚠	👍	?	⋮	4★
6061	VHL	L188Q (c....	In a study of 114 unre...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	?	⋮	4★
5169	VHL	W88* (c.2...	Genotype-phenotype ...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	?	⋮	4★
5159	VHL	S65L (c.1...	Genotype-phenotype ...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	?	⋮	4★
5152	VHL	F136S (c....	Genotype-phenotype ...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	?	⋮	4★
5145	VHL	L153C (c....	Genotype-phenotype ...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	?	⋮	4★
5131	VHL	Q132P (c....	Genotype-phenotype ...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	?	⋮	4★
5130	VHL	N78H (c.2...	Genotype-phenotype ...	Renal Cell Carcinoma	N/A	C	⚠	👍	?	⋮	4★
1942	VHL	EXON 1-3 ...	Mutation detected in ...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	📄	⋮	4★
5866	VHL	Y112N (c....	This paper analyzes a...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	?	⋮	4★
6420	VHL	EXON 1 D...	Molecular analysis of...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	?	⋮	4★
4916	VHL	V62C (c.1...	In a study of 114 unre...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	?	⋮	4★
4918	VHL	Q73* (c.21...	In a study of 114 unre...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	?	⋮	4★
4919	VHL	F76del (c....	This deletion mutatio...	Von Hippel-Lindau Di...	N/A	C	⚠	👍	?	⋮	4★



Variants Associated with RCC

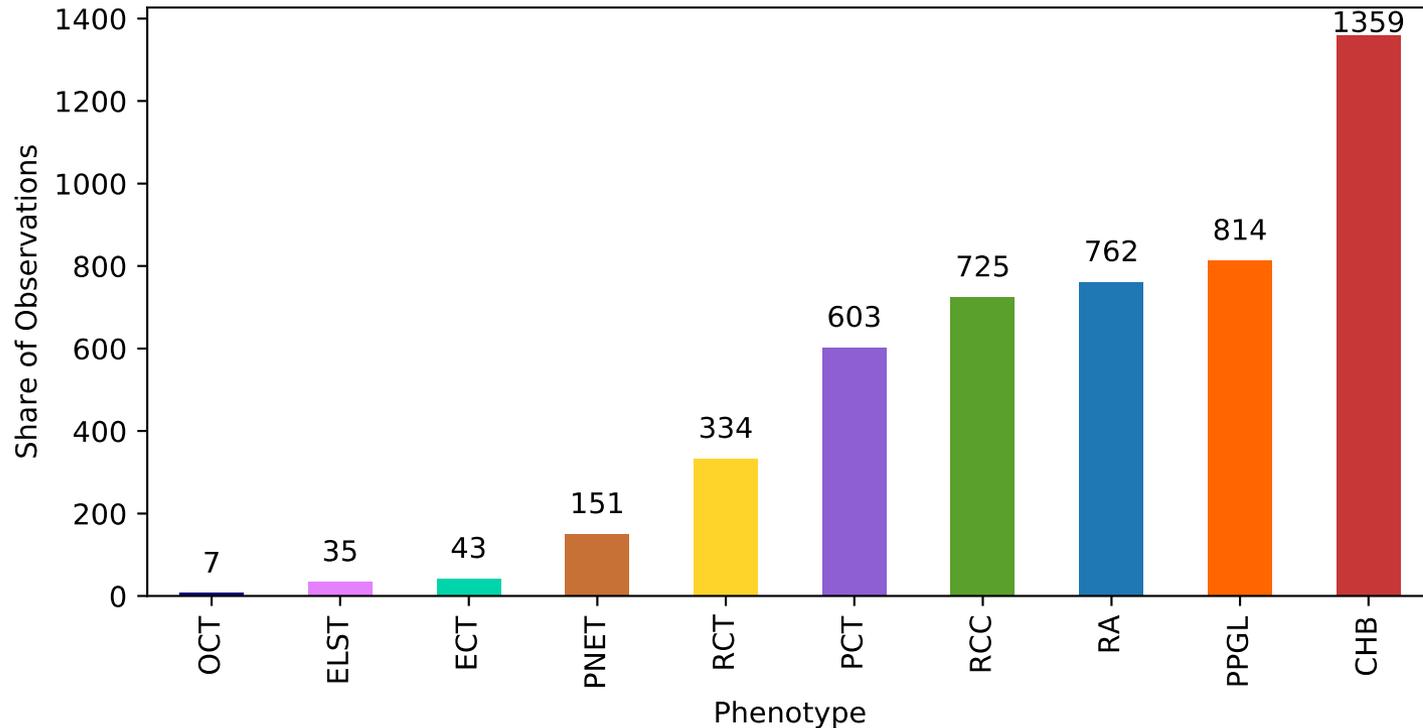


**SickKids<sup>®</sup>**



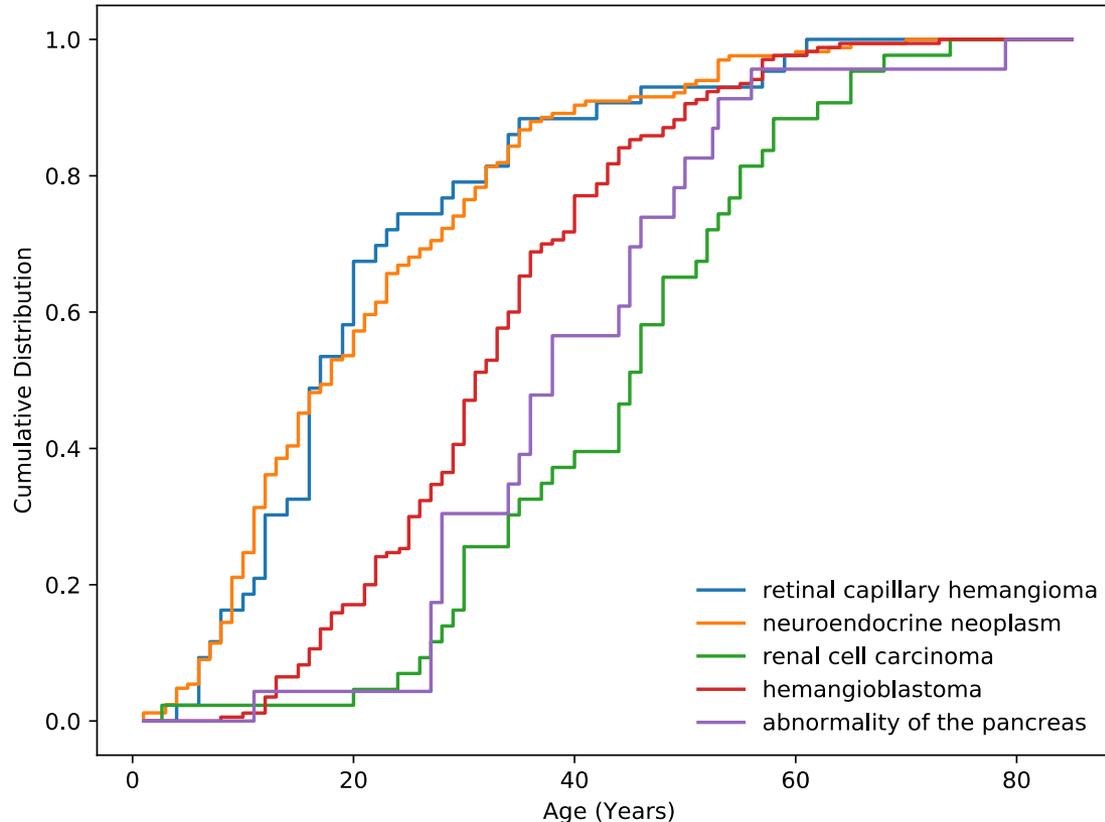
# Analyses on aggregate VHL database...

# Patient-level resolution: phenotype prevalence



Phenotype prevalence at patient-level data resolution showing highest reports of CNS hemangioblastoma.

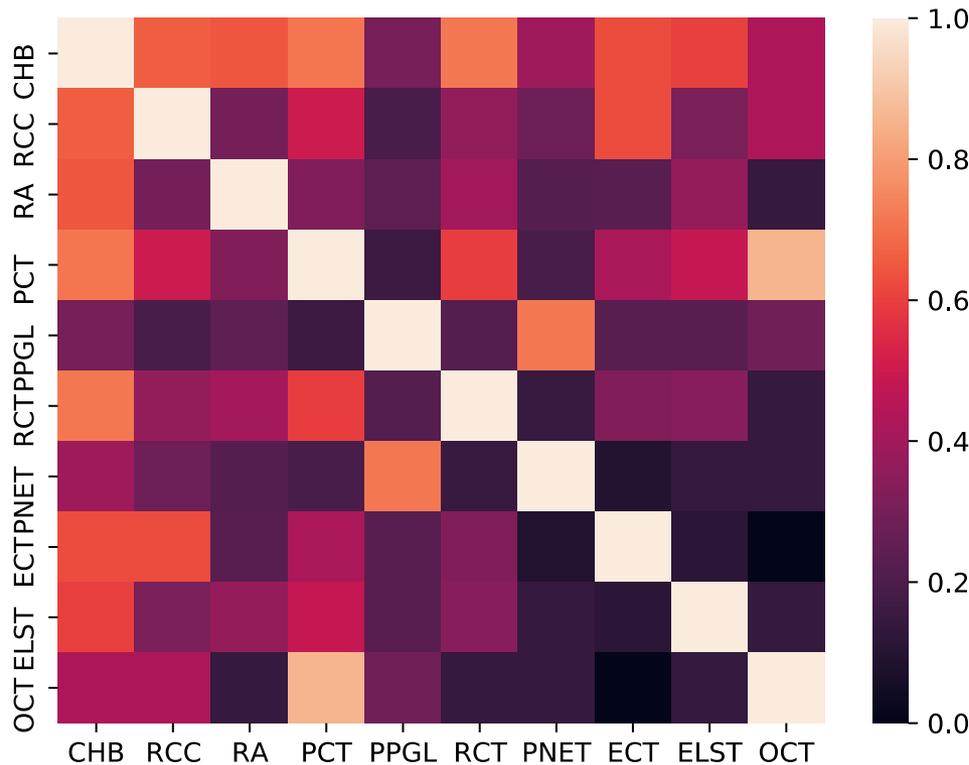
# Patient-level resolution: age-related penetrance



Why is N = 458?

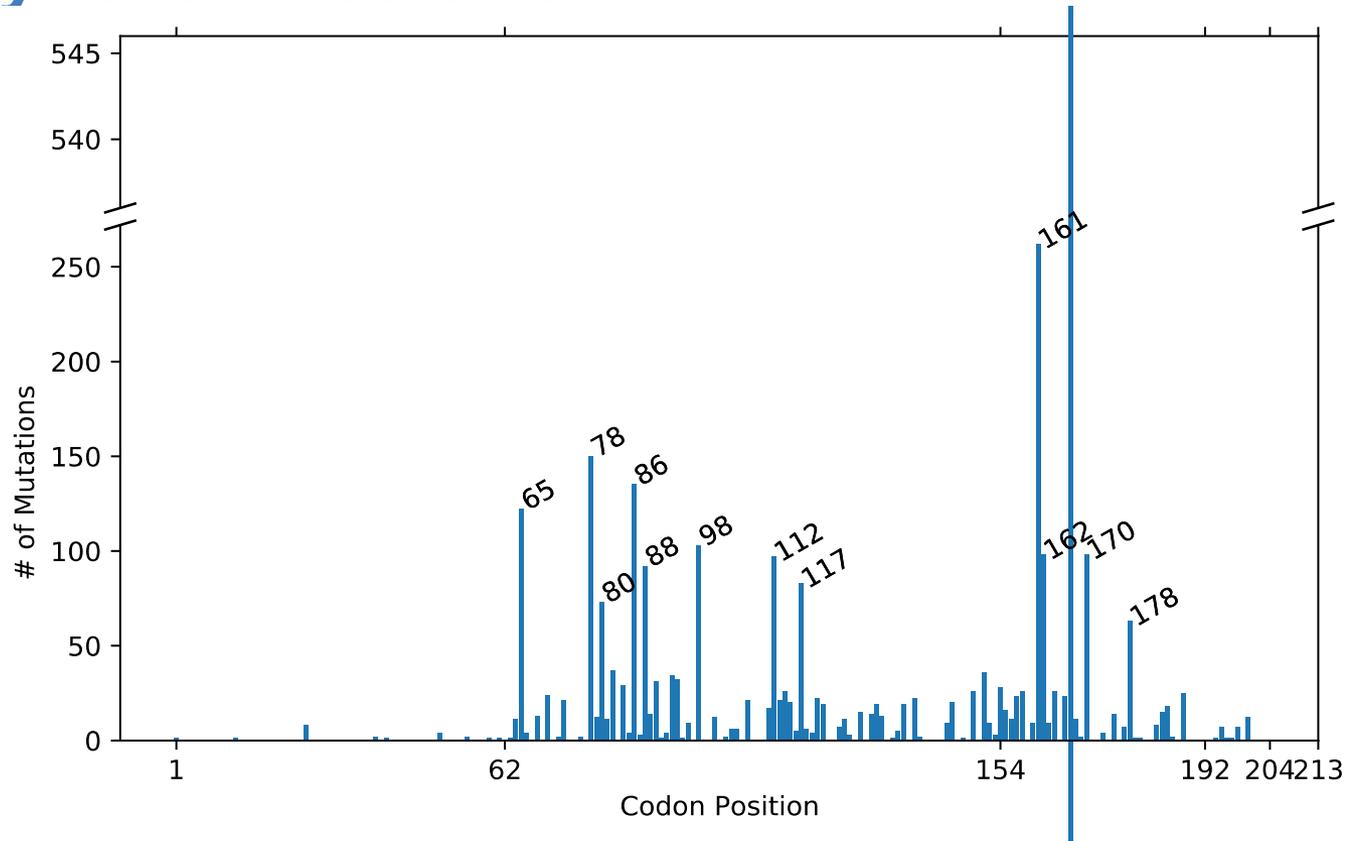
- Age of onset for patients with only one phenotype as a surrogate for age-related penetrance
- Suggests earlier onset with retinal angioma and pheochromocytoma/ paraganglioma.

# Patient-level resolution: phenotype correlation ratios



High correlation between pheochromocytoma / paraganglioma and pancreatic neuroendocrine tumors.

# Family-level resolution



Frequency of missense mutations in families supports potential hotspots at codon 167 and 161.

# Next steps and what we learned

- Duplicated cohorts
- Challenges with aggregating different data types
- Release complete dataset for collaborative researchers
- Other genes?

# Why was it not in ClinVar?

- ClinVar database quality

★ Single submitter

★★★ Expert Panel

**NM\_000551.3(VHL):c.345C>G (p.His115Gln)**

Interpretation ?

Go to: [v] [^]

Clinical significance: [Uncertain significance](#)

Last evaluated: Feb 5, 2016

Number of submission(s): 1

Assertion and evidence details

Go to: [v] [^]

- Clinical assertions
- Summary evidence
- Supporting observations

Submitter	Allele origin	Individuals	Phenotypes (Affected status)	Ethnicity	Geographic origin	Citations	Description
<a href="#">Integrated Genetics/Laboratory Corporation of America</a>	germline	not provided	not provided (unknown)	not provided	not provided	• <a href="#">PubMed</a>	not provided



# VHL ClinGen Expert Panel

## Creating Customized VHL-specific Rules



Eamonn  
Maher



Tina  
Pesaran



Carrie  
Horton



Jerry  
Machado



Deborah  
Ritter



Malachi  
Griffith



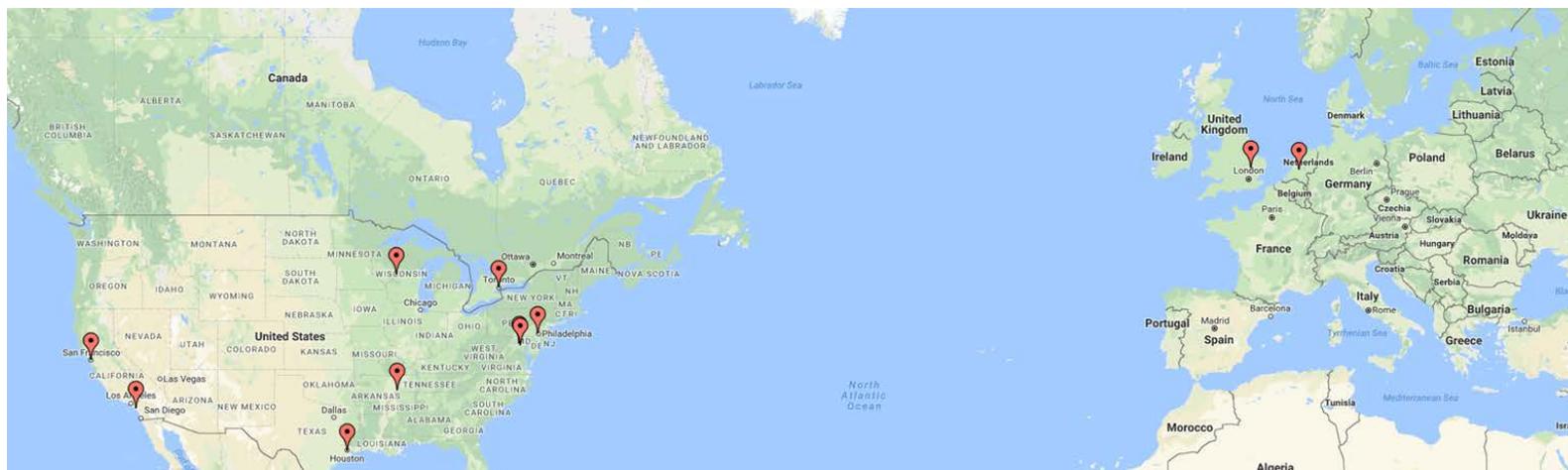
Raymond  
Kim



Minjie Luo



Chimene  
Kesserwan



Amit Tirosh



Sharon  
Plon



Kathleen  
Hruska



Hio Chung  
Kang



Ying Wang



Rachel Giles



Obi  
Griffith



Kilannin  
Krysiak

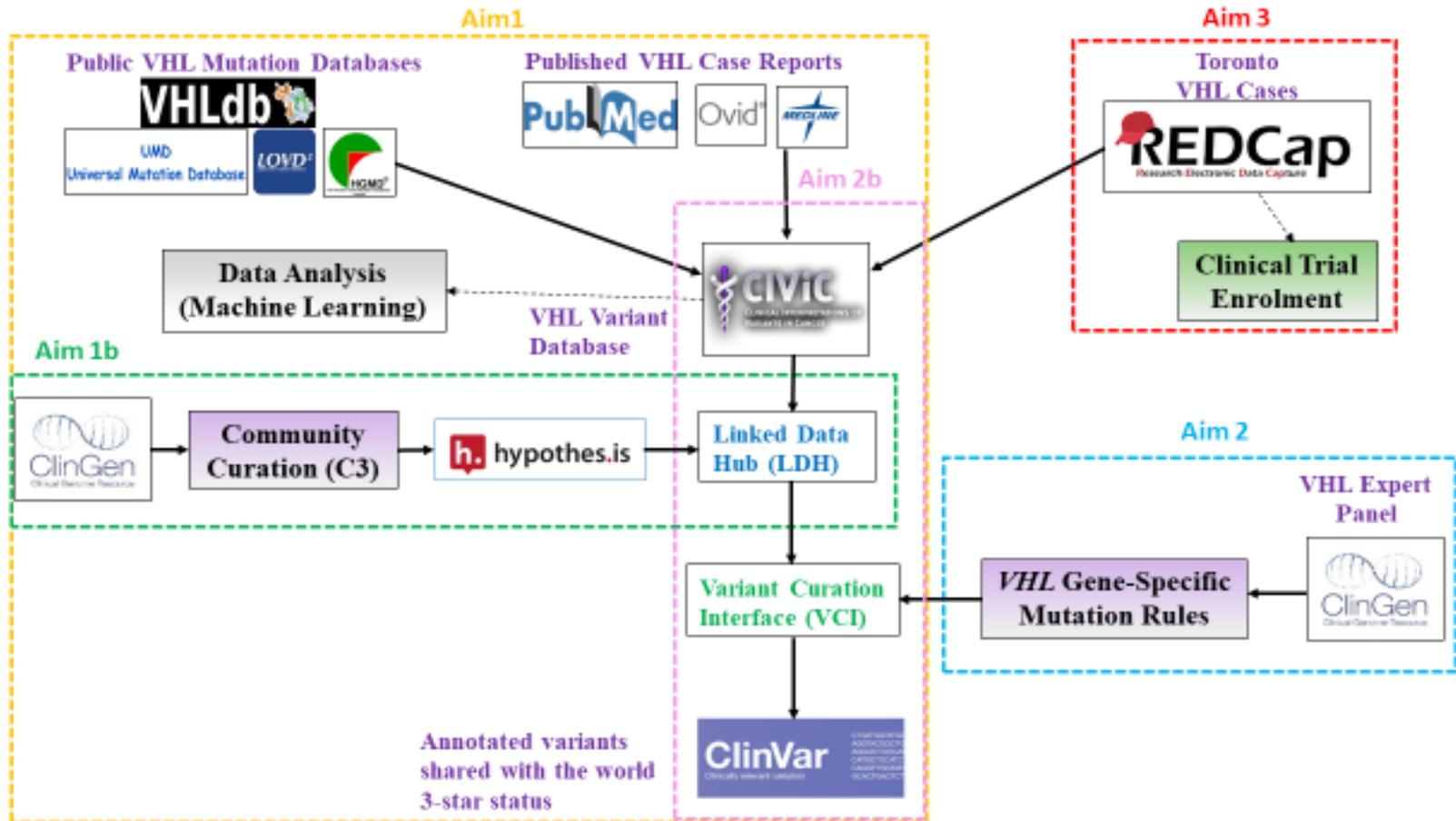


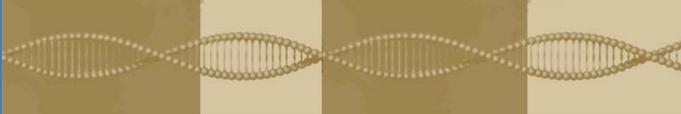
Michael  
Anderson



Kelly  
McGoldrick

# VISION Overview





# Acknowledgements

This project has been supported by a VHL Alliance Research Grant. Funding contributions were also made through the Starbucks Clinical Genetics/Genomics Research Studentship Award.

