


Clinical Reporting Made Easy



Golden Labs
 203 Enterprise Blvd
 Bozeman, 59718
 Phone: 406-587-8137
 Fax: 406-555-5555

Provider Information
 Physician: Dr. James McCoy
 Institution: Acme General Labs
 Case Id: 1234

Patient Information		Sample Information	
Name	Arthur Dent	Sample Site	Blood
Gender	Male	Sample Type	Blood
Date of Birth	10/11/1985	Collection Met...	Peripheral Draw
Id	42	Panel Coverage	85.96%
		Avg. Read Depth	3850x
		Collection Date	11/1/2016
		Receipt Date	11/3/2016
		Report Date	11/9/2016

Results

Positive: Mutations with an establish somatic link detected.

Affected Genes

ABL1 (0)	ASXL1 (0)	BRAF (1)	BRCA1 (0)	BRCA2 (0)	BUB1B (0)	CALR (0)	CBL (0)	CDH1 (0)	CDK4 (0)	CDKN1C (0)
CDKN2A (0)	CEBPA (0)	CEP57 (0)	CHEK2 (0)	CUX1 (0)	CYLD (0)	DDB2 (0)	DICER1 (0)	DIS3L2 (0)	KRAS (1)	NRAS (1)

Primary Findings

Gene	Zygoty	Variant	Exon	Pathogenicity
NRAS	Heterozygous	NM_002524.4:c.181C>A(NP_002515.1:p.Gln61Lys)	3	Pathogenic
BRAF	Heterozygous	NM_004333.4:c.1799T>A(NP_004324.2:p.Val600Glu)	15	Likely Pathogenic
KRAS	Heterozygous	NM_033360.2:c.38G>A(NP_203524.1:p.Gly13Asp)	2	Pathogenic

Interpretation Summary

Mutations in three known oncogene were detected.

Recommendations

Enrollment in a clinical trial testing drugs targeting the mutation should be considered.

February 15th, 2017

Steven Hystad
Field Application Scientist



1 Overview Golden Helix

2 Value of Clinical Reporting in VarSeq

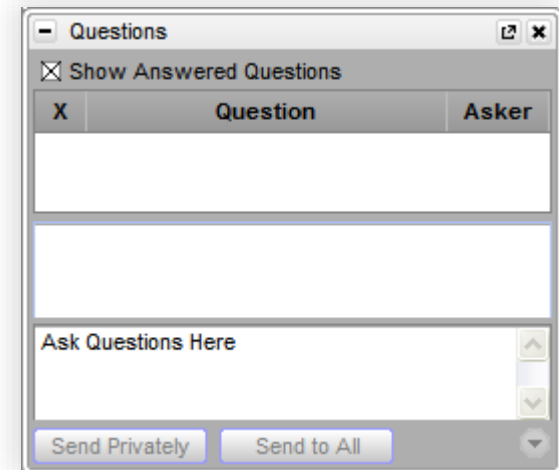
3 Live Demo

4 Q&A



Questions during the presentation

Use the Questions pane in your GoToWebinar window



Golden Helix – Who We Are



Golden Helix is a global bioinformatics company founded in 1998.



Filtering and Annotation
Single Sample CNV-Analysis
Clinical Reports
Pipeline: Run Workflows

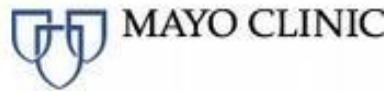


Variant Warehouse
Centralized Annotations
Hosted Reports
Sharing and Integration

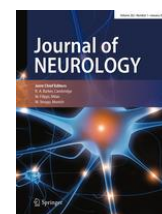
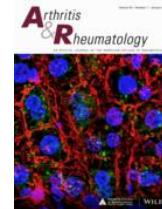
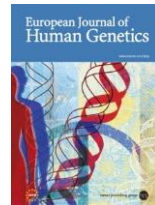
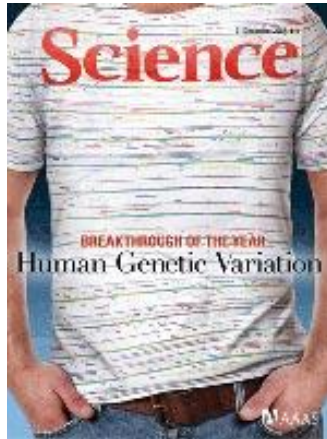


GWAS
Genomic Prediction
Large-N-Population Studies
RNA-Seq
Large-N CNV-Analysis

Over 350 customers globally



Cited in over 1000 peer-reviewed publications

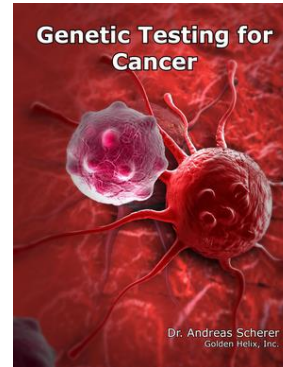


Golden Helix – Who We Are



When you choose a Golden Helix solution, you get more than just software

- REPUTATION
- TRUST
- EXPERIENCE



- INDUSTRY FOCUS
- THOUGHT LEADERSHIP
- COMMUNITY

- TRAINING
- SUPPORT
- RESPONSIVENESS



- TRANSPARENCY
- INNOVATION and SPEED
- CUSTOMIZATIONS



GOLDEN HELIX				
Patient Information				
Name	Jodie Doe	Gender	Female	Case #
Date of Birth	November 10, 2010	MI	1234	
Mother Information				
Name	Jane Doe	Date of Birth	September 28, 1988	MI
MI	1235			
Father Information				
Name	John Doe	Date of Birth	November 10, 1988	MI
MI	1236			
Reference Information				
Physician	Dr. Hyslop	Institution	Golden Labs	Case #
Case #	GL-1234			
Sample Information				
Sample Site	Blood	Sample Type	Blood	Collection Method
Collection Date	January 30, 2017	Receipt Date	January 30, 2017	Report Date
Report Date	January 31, 2017			
Results - Positive				
Mutations with an established link detected.				
Primary Findings				
Gene	Exon	Variant	Zygosity	Pathogenicity
LPIN2	20	NM_014648.2:c.73C>T	Heterozygous	Uncertain Significance
Interpretation Summary				
Patient possess a single Pathogenic de novo variants in the LPIN2 gene.				
Recommendations				
The variant could be associated with Chronic Recurrent Multifocal Osteomyelitis (CRMO). Treatment of CRMO typically involves administration of nonsteroidal anti-inflammatory drugs (NSAIDs) and physical therapy to avoid disease atrophy of muscles and contractures.				
Incidental Findings				
Gene	Exon	Variant	Zygosity	Pathogenicity
BRCA2	2	NM_000059.3:c.6513G>C	Heterozygous	Pathogenic
Interpretation Summary				
A pathogenic variant has been detected in the BRCA2 gene. High risk of breast and ovarian cancer is associated with germline BRCA2 mutations.				
Recommendations				
Referral to an oncologist. Preventive or "prophylactic" mastectomy or removal of both breasts, has been found to reduce the risk of breast cancer in high-risk women by about 90% and should be considered.				

Golden Labs		Provider Information		
203 Enterprise Blvd Bismarck, ND 58103 Phone: 406-637-8137 Fax: 406-555-5555		Physician: Dr. James McCoy Institution: Acme General Labs Case Id: 1234		
Patient Information		Sample Information		
Name	Arthur Dent	Sample Site	Blood	
Gender	Male	Sample Type	Blood	
Date of Birth	10/11/1985	Collection Met...	Peripheral Draw	
MI	42	Panel Coverage	85.96%	
		Avg. Read Depth	3550x	
		Collection Date	11/1/2016	
		Receipt Date	11/3/2016	
		Report Date	11/8/2016	
Results				
Positive: Mutations with an established genetic link detected.				
Affected Genes				
ABL1 (0)	ASXL1 (0)	BRAF (1)	BRCA1 (0)	
BRCA2 (0)	BUB1B (0)	CALR (0)	CBL (0)	
CDH1 (0)	CDK4 (0)	CDKN1C (0)	COKN2A (0)	
CEBPA (0)	CEP57 (0)	CHEK2 (0)	CLUX1 (0)	
CYLD (0)	DDI2 (0)	DICER1 (0)	DIS3L2 (0)	
KRAS (1)	NRAS (1)			
Primary Findings				
Gene	Zygosity	Variant	Exon	Pathogenicity
NRAS	Heterozygous	NM_002524.4:c.181C>A(NP_002515.1:p.G461Lys)	3	Pathogenic
BRAF	Heterozygous	NM_004333.4:c.1799T>A(NP_004324.2:p.Val600Gln)	15	Likely Pathogenic
KRAS	Heterozygous	NM_033360.2:c.38G>A(NP_203524.1:p.Gly13Asp)	2	Pathogenic
Interpretation Summary				
Mutations in three known oncogenes were detected.				
Recommendations				
Enrollment in a clinical trial testing drugs targeting the mutation should be considered.				
OncoMD Drug Summary				
Drug	Generic	Response Rate		
Trifluoromethine	Generic	86%		

- **Generate Clinical Grade Reports**

- ACMG Standard Germline Report

- **Reports are Fully Customizable**

- Configurable Global Settings
- Logo & Lab Information
- Test Description / Disclaimers

- **Customizable Sample Inputs**

- Patient Information & Test Results

- **Integrated with OMIM and OncoMD**

- Functional Description Genes and Phenotypes
- Drug Targeting Information
- Ongoing Clinical Trials



- **Illumina TruSight Myeloid Sequencing Panel**
 - Oncogenes & Tumor Suppressor Genes
 - Drug Targeting Information & Ongoing Clinical Trials

- **Exome Trio Workflow**
 - Casual Variants Associated with Phenotype
 - Incidental Findings

- **Illumina TruSight Cardio Sequencing Panel**
 - Cardiomyopathies, Arrhythmias, etc.
 - Integration with VSWarehouse

Sample Data for Cancer Gene Panels



■ Illumina TruSight Myeloid Sequencing Panel

- Comprehensive coverage of 54 genes designed to target exons of key tumor suppressor genes and frequently cited oncogenes.
- BAM and VCF files for each replicate are available

Chr:Pos	Ref/Alt	Identifier	Quality	Filter	Read Depth (DP)
1:36931696	T	?	100	PASS	1087
1:36931697	T	?	100	PASS	1085
1:36931698	T	?	100	PASS	1087
1:36931699	A	?	100	PASS	1090
1:36931700	C	?	100	PASS	1092
1:36931701	A	?	100	PASS	1092
1:36931703	T	?	100	PASS	1092
1:36931704	A	?	100	PASS	1091
1:36931705	C	?	100	PASS	1088
1:36931706	T	?	100	PASS	1084
1:36931707	G	?	100	PASS	1091
1:36931708	A	?	100	PASS	1086
1:36931709	A	?	100	PASS	1089
1:36931710	G	?	100	PASS	1093
1:36931711	T	?	100	PASS	1085
1:36931712	T	?	100	PASS	1084
1:36931713	A	?	100	PASS	1087
1:36931714	T	?	100	PASS	1089
1:36931715	A	?	100	PASS	1093

VSReports



Golden Helix
203 Enterprise Blvd
Bozeman, 59718
Phone: 406-587-8137
Fax: 406-555-5555

Provider Information
Physician: Dr. James McCoy
Institution: Golden Labs
Case Id: 42

Patient Information
Name: 50 Percent
Gender: Male
Date of Birth: 2/13/2017
Id: 1234

Sample Information
Sample Site: Blood
Sample Type: Blood
Collection Method: Venipuncture
Panel Coverage: 85.96%
Avg. Read Depth: 3850x
Collection Date: 2/6/2017
Receipt Date: 2/9/2017
Report Date: 2/13/2017

Results
Positive: Mutations with an establish somatic link detected.

Affected Genes

ABL1 (0)	ASXL1 (0)	BRAF (1)	BRCA1 (0)	BRCA2 (0)	BUB1B (0)	CALR (0)	CBL (0)	CDH1 (0)	CDK4 (0)	CDKN1C (0)
CDKN2A (0)	CEBPA (0)	CEP57 (0)	CHEK2 (0)	CUX1 (0)	CYLD (0)	DOB2 (0)	DICER1 (0)	DIS3L2 (0)	KRAS (1)	NRAS (0)

Primary Findings

Gene	Zygosity	Variant	Exon	Pathogenicity
BRAF	Heterozygous	NM_004333.4 c.1799T>A(NP_004324.2 p.Val600Glu)	15	Pathogenic
KRAS	Heterozygous	NM_033360.2 c.38G>A(NP_203524.1 p.Gly13Asp)	2	Pathogenic

Interpretation Summary
Mutations in known Oncogenes were detected.

Recommendations
Enrollment in a clinical trial testing drugs targeting the mutations should be considered.



Sample Data for Exome Trio Workflows



- **NA1278 17-year old Caucasian Women**

- Admitted for Symptoms of Acute Gastroenteritis.
- Serum testing revealed elevated pancreatic enzymes.
- Genetic testing revealed Heterozygosity for CFTR gene mutation Δ F508

- **Incidental Findings**

Patient Information

Name: ⓘ

Gender: ⓘ

Date of Birth: ⓘ

Id: ⓘ

Mother Information

Name: ⓘ

Date of Birth: ⓘ


Id: ⓘ

Father Information

Name: ⓘ

Date of Birth: ⓘ

Id: ⓘ



Golden Labs | 203 Enterprise dr Bozeman , 59718 | Phone: 406-581-8137 | Fax: 406-555-5555

Patient Information					
Name	Gender	Date of Birth	Id		
Jane Doe	Male	May 26, 1999	1234		
Mother Information					
Name	Date of Birth	Id			
Jane Doe	October 28, 1968	1235			
Father Information					
Name	Date of Birth	Id			
Steven Doe	November 10, 1968	1236			
Reference Information					
Physician	Institution	Case Id			
Dr James McCoy	Golden Labs	42			
Sample Information					
Sample Site	Sample Type	Collection Method	Collection Date	Receipt Date	Report Date
Blood	Blood	Venipuncture	November 15, 2016	November 17, 2016	November 24, 2016

Results - Positive

Mutations with an established link detected.

Primary Findings

Gene	Exon	Variant	Zygosity	Pathogenicity
CFTR	11	NM_000492.3:c.1521_1523delCTT (NP_000483.3:p.Phe508del)	Heterozygous	Pathogenic



Sample Data for Hereditary Gene Panels



■ TruSight Cardio Sequencing Kit from Illumina

- Comprehensive coverage of 174 genes with known associations to 17 inherited cardiac conditions (cardiomyopathies, arrhythmias, etc).



Golden Labs
303 University Blvd
Ipswich, MA 01938
Phone: 435-587-8137
Fax: 435-555-5555

Provider Information
Physician: Dr. James McCoy
Institution: Golden Labs
Case ID: 42

Patient Information
Name: 50 Percent
Gender: Male
Date of Birth: 2/13/2017
ID: 1234

Sample Information
Sample Site: Blood
Sample Type: Venipuncture
Collection Date: 2/8/2017
Collection Method: Venipuncture
Panel Coverage: 85.96%
Avg. Read Depth: 380x
Report Date: 2/8/2017

Results

Positive: Mutations with an established somatic link detected

Affected Genes

ABL1 (0)	ASXL1 (0)	BRAF (1)	BRC1A1 (0)	BRC1A2 (0)	SVB1B (0)	GALR (0)	CBL (0)	CDH1 (0)	CDK4 (0)	CDKN1C (0)
CDKN2A (0)	CEBPA (0)	CEP350 (0)	CHEK2 (0)	CLU1 (0)	CHLD (0)	DDR2 (0)	DICER1 (0)	DISL2 (0)	KRAS (1)	NRAS (0)

Primary Findings

Gene	Zygosity	Variant	Exon	Pathogenicity
BRAF	Heterozygous	NM_004333.4:c.1797T>A (NP_041242.2:p.M600G)	15	Pathogenic
KRAS	Heterozygous	NM_033962.2:c.385G>A (NP_203524.1:p.G13Asp)	2	Pathogenic

Interpretation Summary
Mutations in known Oncogenes were detected.

Recommendations
Enrollment in a clinical trial testing drugs targeting the mutations should be considered.

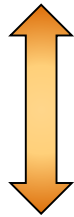
■ Have I ever seen this variant in my previous test samples?

■ Have I put this variant into a clinical report for any previous samples?

■ Did I provide a pathogenicity assessment for this variant? Has that changed?



WAREHOUSE



- **VarSeq can “Export to Warehouse”**
 - Create a new Project
 - Current project template is used
 - Add Variants to Existing (data only)
- **Annotate with Warehouse Counts**
 - Allele / Genotype counts computed on warehouse variants (or other sources in template).
- **Variant Catalogs/ Annotation**
 - Annotate against variant catalog
- **Remote Reports**
 - Central versioning of report
 - Queryable database of all saved reports, with all data in form ready for EMR/LIMS integration



Summary



GOLDEN HELIX

Patient Information					
Name	Gender	Date of Birth	Id		
Jane Doe	Female	November 10, 2010	1234		
Mother Information					
Name	Date of Birth	Id			
Jane Doe	September 28, 1988	1235			
Father Information					
Name	Date of Birth	Id			
John Doe	November 10, 1988	1236			
Reference Information					
Physician	Institution	Case Id			
Dr. Hystad	Golden Labs	GL-1234			
Sample Information					
Sample Site	Sample Type	Collection Method	Collection Date	Receipt Date	Report Date
Blood	Blood	Venipuncture	January 30, 2017	January 30, 2017	January 31, 2017

Results - Positive

Mutations with an established link detected.

Primary Findings

Gene	Exon	Variant	Zygoty	Pathogenicity
LPIN2	20	NM_014646.2:c.*3C>T	Heterozygous	Uncertain Significance

Interpretation Summary

Patient possess a a single Pathogenic de novo variants in the LPIN2 gene.

Recommendations

The variant could be associated with Chronic Recurrent Multifocal Osteomyelitis (CRMO). Treatment of CRMO typically involves administration of nonsteroidal anti-inflammatory drugs (NSAIDs) and physical therapy to avoid disuse atrophy of muscles and contractures.

Incidental Findings

Gene	Exon	Variant	Zygoty	Pathogenicity
BRCA2	2	NM_000059.3:c.6513G>C	Heterozygous	Pathogenic

Interpretation Summary

A pathogenic variant has been detected in the BRCA2 gene. High risk of breast and ovarian cancer is associated with germline BRCA2 mutations.

Recommendations

Referral to an oncologist. Preventive or "prophylactic" mastectomy or removal of both breasts, has been found to reduce the risk of breast cancer in high-risk women by about 90% and should be considered.

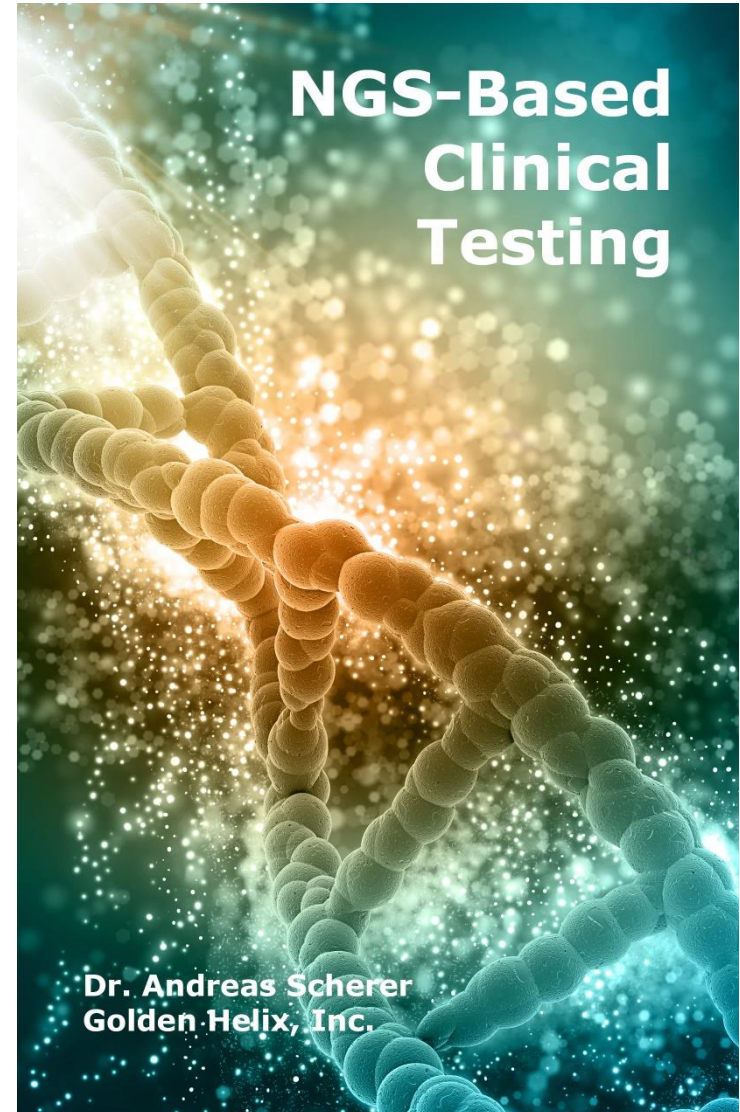
- **VSReports & VarSeq**
 - Powerful Filtering & Annotation Engine Integrated with Clinical Reporting.
- **Clinical Report Customization**
 - Parameters, Layout, and Logo.
 - Golden Helix Offers Client Customizations.
- **Integration with VSWarehouse**
 - Track Reports, Variant Classifications.

New eBook Tomorrow!



NGS-Based Clinical Testing by Dr. Andreas Scherer

- Will be released tomorrow
- Chapter dedicated to clinical reporting





VarSeq Power Pack



- *First 3 customers only!*
- 2 seats for 15 months with access to VarSeq, VSReports, VSPipeline & VSWarehouse
- Access to OMIM & CADD included
- **\$35,995!**
- Offer expires on March 31st, 2017 and is for new customers only
- Request a personal demo by emailing us info@goldenhelix.com

Special
OFFER



Questions or more info:

- Email info@goldenhelix.com
- Request an evaluation of the software at www.goldenhelix.com

