

ARCHER

MSN PTCH2

Archer™ Assay Designer

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Introduction

Product Description

The Archer FusionPlex[™] and VariantPlex[™] target-enrichment assays are used to create libraries for next generation sequencing. Libraries are created by using the FusionPlex assay in conjunction with the Archer Universal RNA Reagent Kits V2 (AK0040-8, AK0042-8), or VariantPlex assay in conjunction with the Archer Universal DNA Reagent Kit V2 for Illumina® (AK0037-8) and Archer[™] MBC Adapters for Illumina or Ion Torrent[™]. Please note that DNA custom design is not supported on the Ion Torrent platform at this time. **These products should never be used independently** (see Associated Products).

Once sequenced, Archer FusionPlex or VariantPlex libraries may be analyzed via <u>Archer Analysis</u> (<u>http://archer.archerdx.com/</u>) to detect and identify SNPs, InDels and fusion partners of designated genes. Copy number variation is not supported through Assay Designer at this time. **For Research Use Only.** Not for use in diagnostic procedures.

While Archer has a large and growing family of FusionPlex and VariantPlex assay panels, these panels cannot possibly provide detection for the vast combination of gene fusions and mutations possible in the human genome. For this reason, Archer has designed an online tool to facilitate panel customization. The Archer Assay **Designer** enables the user to pick and choose desired genes and gene exons. In the following pages, you will find detailed instructions for using and navigating this powerful tool.

Assay Type

Currently, the Archer Assay Designer offers RNA fusion detection, RNA full exon detection and DNA full exon detection. If you require Copy Number Variation detection, or DNA fusion detection, please contact tech@archerdx.com.

Overview

- 1.Register / login
- 2. Create a new project
- 3. Add gene targets
- 4. Specify target exons, fusion or tiling direction, and optional inclusion of UTRs
- 5.Submit design
- 6.Review completed design
- 7. Order assay





Getting Access

ARCHER	
WELCOME	TO THE ARCHER ASSAY DESIGNER
	_
	advanced custom assay design tool available. If you have an account, log in below. ise, register to start creating your custom designs.
Username*	
Password*	
	Forgot your password?
	Log in

http://assay.archerdx.com/

Registration

- First-time users must register before gaining access to the Archer Assay Designer. Follow the link circled to the left to go to the Registration Page.
- Enter your username, email address, institution, and password on the registration page, read the Terms and Conditions of Use, and click Register at the bottom.

	Postal code		
	Country	United States	\$
WELCOME TO THE ARCHER ASSAY DESIGNER	Password*		
—	Password (again)*		
The Archer Assay Designer is the most advanced custom assay design tool available. If you have an account, log in below.	By re	gistering you accept the terms of the E	ULA.
Other dese, register to stort creating your custom designs.	ArcherDX, Inc. Website Terms and Last Updated: Ma		
Password*	These Website Ter the ArcherDX, Inc.	ms and Conditions of Use (the "Terms (("ArcherDX") web site located at www.	ArcherDX.com,
Forgot your password?	subsidiaries and at	sites linked to www.ArcherDX.com by A filiates (collectively, the "Site"). The Site censors. BY USING THE SITE, YOU AGR	e is the property of
Log in		Register	





Confirmation

• Upon successful registration you will receive an activation email at the address provided. To activate your account simply follow the link in the e-mail. Once your account has been successfully activated, you will see the following message.

Your account has been activated.

Please Login to start your first assay design.







RC	CHER Dashboard \diamond New Project
	♦ New Project
	YOU CURRENTLY DO NOT HAVE ANY CUSTOM ASSAY DESIGNS STARTED
	Create a new project to start your first design.
	窗 View Removed Projects

New Project

- Click the "Create New Project" button to create a new project.
- After selecting Create New Project, Assay Designer will walk you through selecting and defining the target exons of interest.

ARCHER
Create New Project
<u>í</u>
YOU CURRENTLY DO NOT HAVE ANY CUSTOM ASSAY DESIGNS STARTED
Create a new project to start your first design.
Create New Project S View Removed Projects
Create New Project: Click here to start a new assay design

Design Name and Type

- 1. Enter a name for your project.
- 2. Select RNA or DNA
 - If you'd prefer to analyze DNA for fusion or CNV contact: tech@archerdx.com.
- 3. Click the green checkbox to start customizing your design.





RCHER		
	ame your project	
	● RNA ● DNA	
Enter a project name, select RNA or DNA then hit the green check box to proceed with your design	NO TARGETS IN DESIGN	

• After clicking on the green box above you will be prompted to add the target regions of interest.

Things to Consider

Before adding targets to the design, you may want to consider a few important features.

Target Gene Variants

- When choosing a gene target, please note that some genes have multiple transcript variants.
- It is important to consider transcript variants because exon numbers may vary between them.
- It is critical to make sure you are specifying exons numbers relative to the transcript you have selected

Fusion Exon

- Genes fusions may form at several different exon junctions.
- For this reason it is necessary to know the exact exon(s) by which this occurs.





Fusion Direction

• Fusion partners may bind at the 5' end, 3' end, or in some cases both ends of a gene exon. You will be given the option to designate which direction you expect the fusion partner to be located.



References

- If you are uncertain about any of the previously mentioned features, please visit the following sources for clarification:
 Quiver - http://archerdx.com/software/quiver
 - This curated database designed by Archer[™] contains entries from the sources listed below as well as several others, including manual entries from our inhouse fusion discoveries.
 - o **COSMIC**
 - http://cancer.sanger.ac.uk/cancergenome/projects/cosmic/
 - ChimerDB http://biome.ewha.ac.kr:8080/FusionGene/index.jsp
 - o TICdb http://www.unav.es/genetica/TICdb/





Add New Targets

• After entering in your project name, molecule type, and clicking on the green box you will be prompted to add the target regions of interest.

Add Target to Desi	gn		×
1 Upload Targets	Q Search Genes	🔥 Load Panel	
	S	ielect target file Drag & Drop Files	

- There are three methods for adding gene targets:
- 1. Upload a primer design file,
- 2. Upload all the targets from one of our commercially available panels, then edit
- 3. Search for the target regions of interest by gene name

1. Adding Targets by Uploading a Target File

• Click the indicated link to download a template file for adding targets.





Clicking on the "Help with custom target file" will display the following:

Adding Targets to Custom Target File

INSTRUCTIONS FOR GENERATING AND UPLOADING A CUSTOM TARGET FILE

Add Target to Desig	ţn		×
1 Upload Targets	Q Search Genes	📩 Load Panel	
	S	Select target file Drag & Drop Files Tile formats include .BED and Archer's custom format.	

1. Click on the "Download sample file" link to download a template file showing sample targets.

	🗋 prime	r_design_template.t	xt	
Target_name NCBI_re RET NM_020975 ALK NM_004304 ALK NM_004304 ROS1 NM_002944	eference_sequence Target_exon 8,9,10,11,12,13,14 5 19,20,21,22,23,24 5 25 3 fusion 31,32,33,34,35,36,37 5	s Direction fusion fusion fusion	Assay_type	Notes

2. By default, this template file is downloaded as a "txt" file, which may be difficult to add information to. Copy the template into an Excel spreadsheet, or similar application, to easily add gene targets and other necessary information. Below is an example of a correctly formatted custom target file.

	A	B	С	D	E	F
1	Target_name	NCBI_reference_sequence	Target_exons	Direction	Assay_type	Notes
2	RET	NM_020975	8,9,10,11,12,13,14	5	fusion	
3	ALK	NM_004304	19,20,21,22,23,24	5	fusion	
4	ALK	NM_004304	25	3	fusion	
5	ROS1	NM_002944	31,32,33,34,35,36,37	5	fusion	
6						





Each column name signifies the following:

Target Name: Canonical gene symbol for target gene.
NCBI_reference_sequence: The reference sequence associated with the target gene. Reference sequences must start with "NM".
Target Exon: The exons within the target gene, which will be amplified for gene fusions. Exons must be comma delimited.
Direction: The direction of primer amplification, designated by either the "5" or "3" prime end of the gene. If you would like to detect fusions in both directions you may do this by adding the target gene and gene exons twice but selecting opposing directions (see NTRK3 entries in example).

- Assay_type: Currently supported assay types include "Fusion" and "Tile".
- Notes: This section will not be considered by the assay designer. It is for your convenience to keep note of ideas while building your custom file.

÷

Format: Tab Delimited Text (.txt)

3. When you have finished adding targets to your custom file, be sure to save your spreadsheet file as a Tab Delimited Text (.txt) file.





4. Upload your file by dragging and dropping it in to the designated





area. Alternatively, you may select the file to be uploaded by navigating to its location on your computer.

- 2. Adding Targets by a Gene Name Search
 - Select the Search Genes tab circled below.

Add Target to Design		×
1 Upload Targets	rch Genes 🏟 Load Panel	
	Search by gene name or accession Search	

• Enter either the full or partial gene symbol and then click search.

Add Tar	Add Target to Design			
🌲 Uplo	ad Targets	Q Search Genes	🛉 Load Panel	
		ALK	Search	

 Select the correct transcript variant from the resulting list by clicking the corresponding ^(c) (see Things to Consider to find out more about transcript variants).





×

×

Add Target to Design

🌲 Upload	d Targets Q Sec	arch Genes 🏄 Load Panel		
		ALK		Search
	Gene Symbol	Accession	Quiver	Coordinates
0	ALK Homo sapiens anapla	NM_004304 Istic lymphoma receptor tyrosine ki	nase (ALK), mRNA.	chr2:29415639:30144477
•	ACVR1B Homo sapiens activin	NM_020328 A receptor, type IB (ACVR1B), trans	A script variant 3, mRNA.	chr12:52345450:52390863
•	ACVR1B Homo sapiens activin	NM_004302 A receptor, type IB (ACVR1B), trans	A script variant 1, mRNA.	chr12:52345450:52390863
•	ACVR1B Homo sapiens activin	NM_020327 A receptor, type IB (ACVR1B), trans	A script variant 2, mRNA.	chr12:52347163:52390863

• Verify you have the correct gene transcript by clicking the gene symbol link to view your selected target in the UCSC Genome Browser or click the accession number link to read about your selection at the NCBI GenBank.

Add Target to Design

Upload Targets	Q Search Genes	📩 Load Panel	
	ALK		Search
	ALK		Search
Gene Symbo	ol Accession	Quiver	Coordinates
Homo sapier	NM_004304 ns anaplastic lympnoma	receptor tyrosine kinase (ALK), mRNA	chr2:29415639:30144477 A.

Target Options

The software will walk you through the options for each target after you have selected it including the following:

Direction

- The direction of primer amplification is relative to the gene of interest and is based on the location or potential location of the fusion partner (see Things to Consider).
- DON'T FORGET: if you want fusion detection at both ends of a target exon, you must add that gene and select that target exon twice: once in the 3' direction and once in the 5' direction
- Click Next after making your selection





Add Ta	arget to Des	ign			×
~	ALK Homo sapien	NM_004304 s anaplastic lymphoma receptor t	₩ yrosine kinase (ALK), mRN	chr2:29415639:30144477 NA.	
		In which direction sho target gene? o	uld primers exter	nd relative to this	
		U		Next	

Untranslated Regions

• Specify whether or not to include untranslated regions in the design.

Add Ta	arget to Des	ign			×
~	ALK Homo sapien	NM_004304 s anaplastic lymphoma receptor t	n yrosine kinase (ALK), mR	chr2:29415639:30144477 NA.	
		Should primers target gene?	untranslated reg	ions of this target	
		No Yes			
		Previous		Next	

Exon Selection

- The assay designer provides the flexibility to choose exactly which exons in the target gene you want to sequence
- If you intend to detect fusion across all exons click Check All, otherwise, click the box next to the desired target exon(s).
- When the appropriate exons have been selected, exit by clicking $\underset{\ensuremath{\mathsf{Next}}}{\ensuremath{\mathsf{Next}}}$





~	ALK NM Homo sapiens anaplastic ly	_004304 mphoma recepto	or tyrosine kind	Aase (ALK), mRN	chr2:29415639:30144477 A.
	Please se	ect which o	exons you	would lik	e to target. o
	□ 1	9	17	25	ØAII
	2	□ 10	18	26	None
	3	□ 11	V 19	27	(None
	4	12	20	28	\searrow
	5	13	21	29	
	6	14	22		Check the boxes next to your
	7	15	23		exons of interest. You can also
	8	16	24		clear your selection by clicking none, or select all exons.
	Previous				Next

Exon Tiling or Fusion Detection

• Choose either exon tiling or fusion detection for the exons chosen for the target.



• After selecting finish, you will be directed back to the main Add Targets to Design screen. Choose more targets or proceed to submit your design by closing the Add Targets to Design window.

3. Adding Targets by Loading a Panel

Select from one of our commercially available panels

- This tab will give you the option of utilizing one of our commercially available panels. You may add targets, remove targets or combine panels.
- Select the desired panel(s) by clicking the corresponding 🔮





×

Add Target to Design

🌲 Upload	Targets	Q Search Genes	5 h- Load Panel	
	Panel Name		Version	Comments
\odot	FusionPlex Al	RR Panel	2.0	
0	FusionPlex N	TRK Panel	1.0	
0	FusionPlex H	eme Panel	1.0	
0	FusionPlex So	arcoma Panel	1.0	
0	FusionPlex FC	GFR Panel	1.0	

• After selecting a panel you may still add more targets or remove unwanted targets. Choose more targets, or proceed to submit for design by closing the Add Target to Design window.

	rget to Design	Genes 🔥 Load	When you are done adding panels and/or targets close the Add Target to Design	/`
L Opic	ad Targets Q Search	Genes h-Lodd	window by clicking on the grey x.	
	Panel Name	Version	Comments	
~	FusionPlex ARR Panel	2.0		
0	FusionPlex NTRK Panel	1.0		
0	FusionPlex Heme Panel	1.0		
0	FusionPlex Sarcoma Panel	1.0		
0	FusionPlex FGFR Panel	1.0	***	

Submit for Design

- Once you have uploaded all genes of interest and selected the correct directions and exons, click Submit for Design to initiate the design process.
- While the most recent project is being designed, another project may be created in the interim. However, only one project may be submitted for design at a time.



RCH	ER				nce finishe lecting tar bmit for D		Test U		
			Т	EST Project RN	A Target	S @		an be y clicking edited by n @1	
			O Add New T	arget 🕹 Download Target	F	Submit for Desig			
^ #	Query 2	Gene Name 😯	Accession 😯	Genomic Location ?	Direction	Assay Type 🕜	Target Exons	Include UTR 2 🖸	7
1	FusionPlex ARR Panel	ROS1	NM_002944	chr6:117609529-117747018	5'	Fusion	31, 32, 33, 34, 35, 36, 37	No	e i
2	FusionPlex ARR Panel	ALK	NM_004304	chr2:29415639-30144477	5'	Fusion	19, 20, 21, 22, 23, 24	No	ð

• Once the design has been submitted you will be able to see the status of the design. A status of Review Pending indicates that Archer Assay Designer has completed designing your primers and notified an Archer reviewer to review your design.

5'

Fusion

8, 9, 10, 11, 12, 13,

14

No

Ø 🖻

chr10:43572516-43625797

A	RCHER				Test User 👻
	Create New Project				
	Project	~ Created	Molecule	Status	
	Cest RNA Project	Jul 09 2015	RNA	Review Pending	42 ŵ

- After the design is reviewed, you will receive an e-mail with a quote, and a sales representative will contact you.
- Once the Design has been approved, the Dashboard status of the project will change to say "Assay Ready" and will provide you with new options. You will receive an additional email from the reviewer with any notes for your design.

ARCHER						Test User 👻
◆ Create New Project						
Project		Molecule	Status			
Test Project	Jul 09 2015	RNA	Assay Ready	🖃 Design Report	\$805.37	匬
	Create New F	Project 💼 V	ew Removed Projects			

3 FusionPlex ARR

Panel

RET

NM_020975





Review Order

• You may review your order by either following the link in the email or by clicking Design Report on the Dashboard.

Test Project RNA Design Report Assay Ready										
		لط Visu	alize Results 🕹 Downloo	ad Target File	` ≓ \$805	37				
		Overa	all Coverage: 100%	To	tal Targets:	21				
Coverage 🔺	Gene Symbol 🚱 🖨	Accession 🛛 🖨	Genomic Location 🛛 🔶	Direction \$	Assay Type 🖨	Target Exons 😧	Include UTR ¢			
100%	ROS1	NM_002944	chr6:117609529-117747018	5'	fusion	31, 32, 33, 34, 35, 36, 37	No			
100%	ALK	NM_004304	chr2:29415639-30144477	5'	fusion	19, 20, 21, 22, 23, 24	No			
100%	RET	NM_020975	chr10:43572516-43625797	5'	fusion	8, 9, 10, 11, 12, 13, 14	No			
100%	ALK	NM_004304	chr2:29415639-30144477	3'	fusion	25	No			

Coverage Key

 0% - 49%
 50% - 69%
 70% - 99%
 Full Coverage

DESIGN DOWNLOADS									
		Project created July							
primers_gsp1.bed primers_targets.bed		primers_primer_regions.bed		primers_gsp2_coverage.bed					

Check Coverage

- The coverage rating is an indication of how many of the designated targets will be amplified by the design.
- Unfortunately, the Assay Designer may not always produce an assay with 100% coverage due to complications with the target of interest.
 - Non-unique regions (contains gene homologues)
 - A-T rich regions
 - Low complexity regions (polynucleotide repeats)
- Overall coverage can quickly be assessed by comparing the coverage score to the color-coded coverage key below the target list.





• Each gene target, as well as the individual exons within that target is assigned a coverage rating. Clicking on the exon number will bring up a report. The Target Exon Report displays the percent coverage for each exon. If there was less than 100% coverage, the reason for design failure is specified. The cause of incomplete coverage will also be discussed in the email from the reviewer.

Coverage 🔺	Gene Symbol 🚱 🖨	Accession 😯 🖨	Genomic Location 😮	¢	Direction \$	Assay Type 🖨	Targe	et Exons 😧	Include UTR
100%	ROS1	NM_002944	chr6:117609529-1177470	18	5'	fusion	31, 32, 33	, 34, 35, 36, 37	No
100%	ALK	NM_004304	chr2:29415639-3014447	7	5'	fusion	19, 20, 3	21, 22, 23, 24	No
100%	RET	NM_020975	chr10:43572516-436257	97	5'	fusion	1 8, 9, 10,	11, 12, 13, 14	No
	ALK	NM_004304	chr2:29415639-3014447	7	3'	fusion		25	No
Target Exo	-			•		Click on th up the Tar	-	exon to bring Report	
Target		Genomic Loco	ation	2006	erage	L		1	_
31		chr6:117658334-11	7658503 1	.00%					
32		chr6:117650491-11	7650609	.00%					
33		chr6:117647386-11	7647577 1	.00%					
34		chr6:117645494-11	7645578 1	.00%					
35		chr6:117642421-11	7642557 1	.00%					
36		chr6:117641030-11	7641193 1	.00%					
37		chr6:117639350-11		.00%					

- View Coverage in Genome Browsers IGV
 - If you don't already have IGV on your computer, download and install the program.
 - IGV: http://www.broadinstitute.org/software/igv/download
 - IGV User guide: <u>http://www.broadinstitute.org/igv/UserGuide</u>
 - IGV compatible .bed files pertaining to your design may be downloaded by clicking any of the .bed files present at the bottom of the design report.

	DESIGN DOWNLOADS	
	Project created July 10, 2015	
primers_gsp1.bed primers_targets.bed	primers_primer_regions.bed	primers_gsp2_coverage.bed

• Upload .bed files in IGV to view coverage

File name	Description
primers_targets.bed	Position of the chosen target exons.
primers_gsp1.bed	Position of the first PCR gene specific





	primer.
primers_gsp2.bed	Position of the second PCR gene specific primer. Also, the Starting position of the amplicon
<pre>primers_gsp2_coverage.bed</pre>	The projected coverage resulting from the gene specific primer pair amplification.
Additional Files	
primers.gtf	A GTF formatted file, containing information necessary to analyze sample results with the Archer™ Analysis Pipeline.

• Alternatively you can open our Genome Browser containing the .bed files by clicking on Visualize Results.

CHER)						Test User
Modify Desig	n Design Rep	ort					
			Test Project	RNA Des	ign Report		
			Assay	Ready			
		Lill Visu	alize Results 🕹 ownload	d Target File	` ≓ \$805.37		
		Overa	all Coverage: 100%	Tot	tal Targets: 21		
	Gene Symbol 🛛 🖨	Accession 🛛 🖨	Genomic Location 🕄 🔶	Direction \$	Assay Type 🖨	Target Exons 😧	Include UTR ¢
Coverage 🖨				5'	fusion	31, 32, 33, 34, 35, 36, 37	
Coverage ¢	ROS1	NM_002944	chr6:117609529-117747018				No
	ROS1 ALK	NM_002944 NM_004304	chr2:29415639-30144477	5'	fusion	19, 20, 21, 22, 23, 24	No No
100%				5' 5'	fusion fusion	19, 20, 21, 22, 23, 24 8, 9, 10, 11, 12, 13, 14	

This link will bring up a searchable window containing ther .bed files





×

Genome Browser

Test Project		Search f symbol	or targets b nere.	y gene	
JBrowse File View Hel	p		v		Full-screen view
e	→ Q Q ⊕ ⊕ 117,650,000	chr6 chr6:117607189	.117743011 (135.8		1
Reference sequence sequence	Zoom i	n to see sequence		Zoom in	to see sequence
G dbSNP per 2,000 bp	 _ _				
TefGene ROS1					
BED_Primers_GSP1	ROS1_chr6_117639376_24_+_A1_GSP1 ROS1_chr6_117641152_26_+_A1_GSI ROS1_chr6_117642501_20_+_A1_G ROS1_chr6_1176425526_37_+_A ROS1_chr6_117647530_32_+ ROS1_chr6_117647530_32_+ ROS1_chr6_117650565_2	SP1 1_GSP1 _A1_GSP1			

View Coverage - What to Look For

• NOTE: This section is relevant to RNA fusion designs ONLY Nested Primers:

• The GSP2 should always be located (nested) inside of the GSP1



Exons Only:

• Primers should be positioned only on exons





the Coverage direction all match the direction specified in the design.

3' Directionality:



5' Directionality:



Junction Coverage

• Ensure that primer coverage extends through exon-exon junctions of all exons specified in design.



Ordering Your Assay

- "Order Assay" is accessible at two locations in the Assay Designer.
- 1. From the Assay Design Dashboard





ARCHER					Test User 👻
Create New Project				Click on the 🔚 box to order your design.	
Project	- Created	Molecule	Status		
Test Project	Jul 09 2015	RNA	Assay Ready	Design Repc 🛛 🗮 \$805.37	Ē
	💠 Create New F	Project	View Removed Projects		

2. From the Design Report

Test Project RNA Design Report order your design. Assay Ready 4							
		<u>اسا</u> Visu	ualize Results 🕹 Downloa	d Target Fi	≒ \$805.37		
		Overa	all Coverage: 100%	Tot	al Targets: 21		
Coverage 🔺	Gene Symbol Ø ¢	Overa Accession @+	Genomic Location 📀 💠	Tot Direction +	cal Targets: 21	Target Exons 😧	Include UTF
Coverage 100%	Gene Symbol �¢ ROS1		, in the second s				Include UTF
		Accession 🛛 🖨	Genomic Location 📀 💠	Direction \$	Assay Type 🗢	Target Exons 🛿	Include UTF No No
100%	ROS1	Accession ② ¢ NM_002944	Genomic Location ? + chr6:117609529-117747018	Direction \$	Assay Type 🗢 fusion	Target Exons 🛿	No

0% - 49% 50% - 69% 70% - 99% Full Coverage		Covera	ge Key	
	0% - 49%	50% - 69%	70% - 99%	Full Coverage

Complete Order Information

• Fill out the following fields and click Request Quote when done.





×

Request Quote

the details of your order. Ple need a Universal RNA Reag	request, a sales representative case note that in addition to yo gent Kit and platform-specific N oth items are available at the	our custom assay, you will MBC adapters in order to
Custom Assay 21 targets × 24 reactions	Quantity	Price Total \$805.37
Company* ArcherDX		
Phone *	tact information for future quot	tes?
Shipping Address	Stree	Billing Address
Street2	Stree	12
City *	City	
State Select	\$	
Select Other for non US address. Postal code *		t Other for non US address. Il code
Country *	Coun	try
United States	\$ Unit	ed States
placed through the Arch www.ArcherDX.com (the licensors. BY PLACING A YOU DO NOT AGREE, Dr By checl	2015	elocated at of ArcherDX and its SE PURCHASE TERMS; IF &
	Request Quote	\triangleright

• A confirmation message is displayed if the ordering information was entered correctly.





Thank you!

We have received your quote request for Project #640 – "Test Project".

A sales representative will be in touch shortly. Please note that in addition to the assay you've designed, you will also need a Universal RNA Reagent Kit and platform-specific MBC adapters in order to perform this experiment. Both items are available at the ArcherDX product page.

 NOTE: Unless you have already been given an order number, you must contact <u>orders@archerdx.com</u> and provide payment information before the order is processed.





FAQs

URL: http://archerdx.com/faqs

Associated Products

P/N	Product Description
AK0040-8	Archer™ Universal RNA Reagent Kit for Illumina®
AK0042-8	Archer™ Universal RNA Reagent Kit for Ion Torrent™ AK0037-8 Archer™ Universal DNA Reagent Kit for Illumina®
N/A	Archer™ Analysis Pipeline - http://analysis.archerdx.com

URL: http://archerdx.com/

Limitations of Use

For Research Use Only. Not for use in diagnostic procedures.

This product was developed, manufactured, and sold for in vitro use only. The product is not suitable for administration to humans or animals. SDS sheets relevant to this product are available upon request.

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For more information please visit http://www.archerdx.com

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ArcherDX, Inc. 2477 55th Street, Suite 202 Boulder, CO 80301 303-357-9001 www.archerdx.com