



Software for
Business Intelligence

BizInt Smart Charts

Patents & IP Sequences | Clinical Trials | Drug Pipelines



Creating IP Reports Integrating Sequence, Family, and Hit Structure data with BizInt Smart Charts

PIUG 2019 Biotechnology Conference, Boston MA

John Willmore

19 February 2019

www.bizint.com



THE JOURNEY BEGINS...

doxifun.com/puppies

Agenda

- IP Sequence reports
- Hit Structure reports
- Summary Record exports
- Reference Rows

BizInt.com/slides for other presentations

Agenda

- IP Sequence reports
- Hit Structure reports
- Summary Record exports
- Reference Rows



BizInt Smart Charts

VERSION

5

for Patents

IP Sequence Databases

Provide data on sequences filed in patents

- GQ Life Sciences GenomeQuest (Geneseq, GQ-PAT)
- STN (USGENE, DGENE, PCTGEN)

Sequence Databases on STN

- DGENE, USGENE, PCTGEN
- Jim Brown's workshop this morning provided step-by-step instructions for searching
- **D BIB SCORE ALIGN**
- Import your transcripts into BizInt Smart Charts
- Each row is a sequence from a patent pub.
- **Don't skip Validate Sequences step!**

Sequence Databases on STN

Combined: sequences2019					
	Title	Database	Sequence ID	Patent Sequence Location	Score
29	New vector comprising promoter and polynucleotide useful for producing fusion protein and target protein e.g. enzyme, blood protein, binding protein, hormone, synthetic protein and peptide.	Derwent GeneSeq	US20110151514-0005	Example; SEQ ID NO 5	44 2% of query self score 2022
30	Vectors, Methods, Systems and Kits for Protein Purification (PublishedApplication)	USGENE	US20110151514-0005	SEQ ID NO 5	44 2% of query self score 2022
31	New vector comprising promoter and polynucleotide useful for producing fusion protein and target protein e.g. enzyme, blood protein, binding protein, hormone, synthetic protein and peptide.	Derwent GeneSeq	US20110151514-0010	Example; SEQ ID NO 10	44 2% of query self score 2022
32	Vectors, Methods, Systems and Kits for Protein Purification (PublishedApplication)	USGENE	US20110151514-0010	SEQ ID NO 10	44 2% of query self score 2022
33	Genetic resistance prediction against antimicrobial drugs in microorganism using structural changes in the genome	PCTGEN	WO20170021529-0144		109 5% of query self score 2022
34	Determining structural variations in genome of microorganism particularly bacterial microorganism, comprising change in genome, by obtaining first data set gene sequences of clinical isolates, and correlating data with second data set.	Derwent GeneSeq	WO20170021529-0144	Disclosure; SEQ ID NO 144	109 5% of query self score 2022

Sequence Databases on STN

- CAplus
- Not supported ... yet
- Save both your transcript...
- **D BIB HITRN**
- ...and your Alignment report (.xss)

Sequence Databases on GenomeQuest

- GQPat, GENESEQ
- Results filters now apply to BizInt exports (e.g. top results per query)



Sequence Databases on GenomeQuest

- CAS Biosequences
- Watch this space!

Export your search results

The screenshot displays the BizInt Smart Charts website interface. At the top left is the logo for BizInt Smart Charts, with the tagline "Software for Business Intelligence". To the right of the logo is a search bar containing the word "search" and a magnifying glass icon. Further right, contact information is provided: "help: support@bizint.com" and "1.714.289.1000 [8am - 5pm Pacific]".

The main navigation bar includes links for "Home", "Products", "Support", "News & Events", "About Us", and "Contact Us". The "Support" dropdown menu is open, showing the following options:

- About Support
- Creating Reports from Databases and Hosts
- Tips for Using BizInt Smart Charts
- Installation and IT Issues
- Subscribe to Release Announcements
- Standard License Agreement

The "Creating Reports from Databases and Hosts" option is highlighted with a mouse cursor. To the right of the navigation bar, a sidebar contains several links with right-pointing arrows:

- Tips For Using BizInt Smart Charts
- Cookbook of Reports & Visualizations
- Software Download
- Download a Free Trial
- Upcoming Events
IC-SDV 2018
23-24 April 2018
Nice, France

The main content area features a large image of a brown dog (Remy) sitting on grass. Text overlaid on the image reads "Get [something] with tools and clin...". Below the image, a dark blue banner contains the text "Try it with your own data!" and a "DOWNLOAD" button with a right-pointing arrow. At the bottom left of the banner, there are five small square icons. At the bottom right, the text "BizInt Smart Charts Drug Development Suite" is visible.

Export from GenomeQuest

The screenshot shows the BizInt Smart Charts website. At the top, there is a navigation bar with the BizInt logo (Software for Business Intelligence) and the product name "BizInt Smart Charts". Contact information includes an email address (help: support@bizint.com) and a phone number (1.714.289.1000 [8am - 5pm Pacific]). A search bar is also present. The main navigation menu includes Home, Products, Support, News & Events, About Us, and Contact Us.

The "Support" section is highlighted, with a sub-section titled "Support: Creating Reports from Databases/Hosts". The main heading is "Creating from GenomeQuest". The text below states: "BizInt Smart Charts for Patents can build charts from search results from GenomeQuest. Additional details are provided in this pdf document."

A "Step by Step" section contains two numbered steps:

1. Create your results set on GenomeQuest.
2. Click on the "Applications" drop-down menu and select "BizInt Smart Charts", as shown below.

The screenshot also shows a screenshot of the software interface. It displays a menu with "Result", "Export", and "Applications" (with a help icon). The "Applications" menu is open, showing options: Assembly, EMBOSS, Geneious, Patent Statistics, To UCSC Genome Browser, BizInt Smart Charts (highlighted with a mouse cursor), Clustalw, and Sequence Search. Below the menu, there are input fields for "Match" (set to "all of the following"), "Application pub. date", "Abstract", and "Patent number".



- Added Unique Family Sequence ID
- Possible development: each unique sequence only appears once per family
- Added Sequence Listing Equivalents
- Useful in publication-level reports

Features of IP Sequences in BizInt reports

- Each row corresponds to a sequence in the context of a query.
- Columns contain bibliographic data, sequence details, and query results.
- You may see the same sequence more than once in a report.
- Family equivalents are not removed.

Sample sequence data report

GQPAT Proteins: Antibodies_GenomeQuest									
	Title	Patent Assignee	Seq. ID Number	Organism Species	Alignment				Percenta
1	Nucleic acid sequences relating to Bacteroides fragilis for diagnostics and therapeutics	OSCIENT PHARMACEUTICALS CORPORATION WALTHAM, MA	US7090973-6862	Bacteroides fragilis	Q:	1	KV--SNR-LY	7	70.00
					S:	340	KVDMSNRILY	349	
2	Expression of microbial proteins in plants for production of plants with improved properties	MONSANTO TECHNOLOGY, LLC ST. LOUIS, MO	US7314974-14121	Pseudomonas fluorescens	Q:	1	K-VS--NRLY	7	70.00
					S:	597	KLVSDDLNRLY	606	
3	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS INC. MORRIS PLAINS, NJ	US6962702-0008	Artificial Sequence	Q:	1	RSSQSIVHSNGNTYLQ	16	93.75
					S:	24	+ RSSQSIVHSNGNTYLE	39	
4	Chimeric, human and humanized anti-CSAP monoclonal antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	US7387772-0032	Murine sp.	Q:	1	RSSQSIVHSNGNTYLQ	16	93.75
					S:	24	+ RSSQSIVHSNGNTYLE	39	
5	Chimeric, human and humanized anti-CSAp monoclonal antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	US7414121-0032	Murine sp.	Q:	1	RSSQSIVHSNGNTYLQ	16	93.75
					S:	24	+ RSSQSIVHSNGNTYLE	39	
6	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	US7429381-0008	Artificial Sequence	Q:	1	RSSQSIVHSNGNTYLQ	16	93.75
					S:	24	+ RSSQSIVHSNGNTYLE	39	
7	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS INC. MORRIS PLAINS, NJ	US6962702-0012	Artificial Sequence	Q:	1	RSSQSIVHSNGNTYLQ	16	93.75
					S:	24	+ RSSQSIVHSNGNTYLE	39	
8	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	US7429381-0012	Artificial Sequence	Q:	1	RSSQSIVHSNGNTYLQ	16	93.75
					S:	24	+ RSSQSIVHSNGNTYLE	39	
9	Covalently reactive transition state analogs and methods of use thereof	BOARD OF REGENTS, THE UNIVERSITY OF TEXAS SYSTEM AUSTIN, TX	US6855804-0042	Mus musculus domesticus	Q:	1	RSSQSIVHSNGNTYLQ	16	93.75
					S:	24	+ RSSQSIVHSNGNTYLE	39	
	Covalently reactive transition	ADLER; BENJAMIN	US7524663-0042	Mus musculus	Q:	1	RSSQSIVHSNGNTYLQ	16	93.75

Sample sequence data report - features

GQPAT Proteins: Antibodies_GenomeQuest

	Title	Patent Assignee	Organism Species	Alignment	Percent
1	Nucleic acid sequences relating to Bacteroides fragilis for diagnostics and therapeutics	OSCIENT PHARMACEUTICALS CORPORATION WALTHAM, MA	Bacteroides fragilis	Q: 1 KV--SNR-LY 7 	70.00
			S: 340 KVDMSNRILY 349		
2	Expression of microbial proteins in plants for production of plants with improved properties	MONSANTO TECHNOLOGY, LLC ST. LOUIS, MO	Pseudomonas fluorescens	Q: 1 K-VS--NRLY 7 	70.00
			S: 597 KLVSDLNRLY 606		
3	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS INC. MORRIS PLAINS, NJ	Artificial Sequence	Q: 1 RSSQSIVHSNGNTYLQ 16 +	93.75
			S: 24 RSSQSIVHSNGNTYLE 39		
4	Chimeric, human and humanized anti-CSAp monoclonal antibodies	IMMUNIMEDICS, INC. MORRIS PLAINS, NJ	Murine sp.	Q: 1 RSSQSIVHSNGNTYLQ 16 +	93.75
			S: 24 RSSQSIVHSNGNTYLE 39		
5	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	Murine sp.	Q: 1 RSSQSIVHSNGNTYLQ 16 +	93.75
			S: 24 RSSQSIVHSNGNTYLE 39		
6	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	Artificial Sequence	Q: 1 RSSQSIVHSNGNTYLQ 16 +	93.75
			S: 24 RSSQSIVHSNGNTYLE 39		
7	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	Artificial Sequence	Q: 1 RSSQSIVHSNGNTYLQ 16 +	93.75
			S: 24 RSSQSIVHSNGNTYLE 39		
8	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	Artificial Sequence	Q: 1 RSSQSIVHSNGNTYLQ 16 +	93.75
			S: 24 RSSQSIVHSNGNTYLE 39		
9	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS INC. MORRIS PLAINS, NJ	Mus musculus domesticus	Q: 1 RSSQSIVHSNGNTYLQ 16 +	93.75
			S: 24 RSSQSIVHSNGNTYLE 39		
			Mus musculus domesticus	Q: 1 RSSQSIVHSNGNTYLQ 16 +	93.75

Bibliographic Data

Sample sequence data report - features

GQPAT Proteins: Antibodies_GenomeQuest						
	Title	Patent Assignee	Seq. ID Number	Organism Species	Alignment	Percentage
1	Nucleic acid sequences relating to <i>Bacteroides fragilis</i> for diagnostics and therapeutics	OSCIENT PHARMACEUTICAL CORPORATION WALTHAM, MA	US7090973-6862	<i>Bacteroides fragilis</i>	1 KV--SNR-LY 7 340 KVDMSNRILY 349	70.00
2	Expression of microbial proteins in plants for production of plants with improved properties	MONSANTO TECHNOLOGY, LLC ST. LOUIS, MO	US7314974-14121	<i>Pseudomonas fluorescens</i>	1 K-VS--NRLY 7 597 KLVSDLNRLY 606	70.00
3	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS INC MORRIS PLAINS, N.J.	US6962702-0008	Artificial Sequence	1 RSSQSIVHSNGNTYLQ 16 + 24 RSSQSIVHSNGNTYLE 39	93.75
4	Chimeric, human and humanized anti-CSAP monoclonal antibodies	IMMUNOMEDICS, INC MORRIS PLAINS, N.J.	US6962702-0008	Artificial Sequence	1 RSSQSIVHSNGNTYLQ 16 + 24 RSSQSIVHSNGNTYLE 39	93.75
5	Chimeric, human and humanized anti-CSAP monoclonal antibodies	IMMUNOMEDICS, INC MORRIS PLAINS, N.J.	US7397772-0032	Murine sp.	1 RSSQSIVHSNGNTYLQ 16 + 24 RSSQSIVHSNGNTYLE 39	93.75
6	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS, INC MORRIS PLAINS, N.J.	US7414121-0032	Murine sp.	1 RSSQSIVHSNGNTYLQ 16 + 24 RSSQSIVHSNGNTYLE 39	93.75
7	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS INC MORRIS PLAINS, N.J.	US7429381-0008	Artificial Sequence	1 RSSQSIVHSNGNTYLQ 16 + 24 RSSQSIVHSNGNTYLE 39	93.75
8	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS, INC MORRIS PLAINS, N.J.	US7429381-0008	Artificial Sequence	1 RSSQSIVHSNGNTYLQ 16 + 24 RSSQSIVHSNGNTYLE 39	93.75
9	Covalently reactive transition state analogs and methods of use thereof	BOARD OF REGENTS THE UNIVERSITY OF TEXAS SYSTEM AUSTIN, TX	US6962702-0012	Artificial Sequence	1 RSSQSIVHSNGNTYLQ 16 + 24 RSSQSIVHSNGNTYLE 39	93.75
	Covalently reactive transition state analogs and methods of use thereof	ADLER, BENJAMIN			1 RSSQSIVHSNGNTYLQ 16 +	93.75

Sequence Data

Sample sequence data report - features

GQPAT Proteins: Antibodies_GenomeQuest										
	Title	Patent Assignee	Seq. ID Num	Alignment						Percentage
1	Nucleic acid sequences relating to Bacteroides fragilis for diagnostics and therapeutics	OSCIENT PHARMACEUTICALS CORPORATION WALTHAM, MA	US7090973-6	Q:	1	KV--SNR-LY	7			70.00
				S:	340	KVDMSNRILY	349			
2	Expression of microbial proteins in plants for production of plants with improved properties	MONSANTO TECHNOLOGY, LLC ST. LOUIS, MO	US7314974-1	Q:	1	K-VS--NRLY	7			70.00
				S:	597	KLVS DLNRLY	606			
3	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS INC. MORRIS PLAINS, NJ	US6962702-0	Q:	1	RSSQSIVHSNGNTYLQ	16			93.75
				S:	24	RSSQSIVHSNGNTYLE	39	+		
4	Chimeric, human and humanized anti-CSAP monoclonal antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	US7387772-0	Q:	1	RSSQSIVHSNGNTYLQ	16			93.75
				S:	24	RSSQSIVHSNGNTYLE	39	+		
5	Chimeric, human and humanized anti-CSap monoclonal antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	US7414121-0	Q:	1	RSSQSIVHSNGNTYLQ	16			93.75
				S:	24	RSSQSIVHSNGNTYLE	39			
6	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	US7429381-0	Q:	1	RSSQSIVHSNGNTYLQ	16			93.75
				S:	24	RSSQSIVHSNGNTYLE	39	+		
7	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS INC. MORRIS PLAINS, NJ	US6962702-0	Q:	1	RSSQSIVHSNGNTYLQ	16			93.75
				S:	24	RSSQSIVHSNGNTYLE	39	+		
8	Production and use of novel peptide-based agents for use with bi-specific antibodies	IMMUNOMEDICS, INC. MORRIS PLAINS, NJ	US7429381-0	Q:	1	RSSQSIVHSNGNTYLQ	16			93.75
				S:	24	RSSQSIVHSNGNTYLE	39	+		
9	Covalently reactive transition state analogs and methods of use thereof	BOARD OF REGENTS, THE UNIVERSITY OF TEXAS SYSTEM AUSTIN, TX	US6855804-0	Q:	1	RSSQSIVHSNGNTYLQ	16			93.75
				S:	24	RSSQSIVHSNGNTYLE	39	+		
	Covalently reactive transition state analogs and methods of use thereof	ADLER; BENJAMIN	US7524663-0	Q:	1	RSSQSIVHSNGNTYLQ	16			93.75
				S:	24	RSSQSIVHSNGNTYLE	39	+		

Query Results

About the Alignment Column

- The Alignment column appears in the same default font as all columns
- You might be tempted to change the font to Courier New - DON'T!
- **Text menu | Fixed Width (e.g. Alignment)** is the correct technique
- Preserves runs of whitespace when exporting to HTML, Word, or Excel

Features of IP Sequences in BizInt reports

	Title	Alignment	Alignment Style
1	HEPATITIS C VIRUS GENE	Q: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSEERSQPRG 60 S: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSEERSQPRG 60 Q: 61 RRQPIPKDRRSTGKSWGKPGYPWPLYGNEGCGWAGWLLSPRGSRPTWGPTDPRHRSRNLG 120 + ++ + + [CONT.]	Default
2	HCV GENE	Q: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSEERSQPRG 60 S: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSEERSQPRE 60 Q: 61 RRQPIPKDRRSTGKSWGKPGYPWPLYGNEGCGWAGWLLSPRGSRPTWGPTDPRHRSRNLG 120 ++ + [CONT.]	Changed Font
3	HCV GENE	Q: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSEERSQPRG 60 S: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSEERSQPRE 60 Q: 61 RRQPIPKDRRSTGKSWGKPGYPWPLYGNEGCGWAGWLLSPRGSRPTWGPTDPRHRSRNLG 120 ++ + [CONT.]	Text Fixed Width

Features of IP Sequences in BizInt reports

GQPAT Gold+ Proteins: hepc_claimed_gqprt

	<i>Title</i>	<i>Alignment</i>	<i>Alignment Style</i>
1 Link	HEPATITIS C VIRUS GENE	Q: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSESRQPRG 60 S: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSESRQPRG 60 Q: 61 RRQIPKDRRSTGKSWGKPGYPWPLYGNEGCGWAGWLLSPRGSRPTWGPTDPRHRSRNLG 120 + ++ + [CONT.]	Default
2 Link	HCV GENE	Q: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSESRQPRG 60 S: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSESRQPRE 60 Q: 61 RRQIPKDRRSTGKSWGKPGYPWPLYGNEGCGWAGWLLSPRGSRPTWGPTDPRHRSRNLG 120 + ++ +	Changed Font
3 Link	HCV GENE	Q: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSESRQPRG 60 S: 1 MSTNPKPQRKTKRNTNRRPQDVKFPGGGQIVGGVYLLPRRGPRLGVRATRKTSESRQPRE 60 Q: 61 RRQIPKDRRSTGKSWGKPGYPWPLYGNEGCGWAGWLLSPRGSRPTWGPTDPRHRSRNLG 120 + ++ +	Text Fixed Width

Non-Patent Sequence Databases

- Can support as needed
- e.g. SWISSPROT, DrugBank

- Can be combined with IP sequences
- Cannot be grouped with patent families
- No tools for deduplication

Sequences in Literature

- Most collections are structured around sequences with multiple citations per seq
- CPlus is organized by document, with multiple sequences per doc

Improved Literature Support -- Project Goals

- Bring literature support up to the same level as drug pipelines, clinical trials, patents, and IP sequences
- Collecting user priorities to help set our development plans
- Focus of development for the next year.

Improved literature support - summary

- “Identify Common Citation” to allow Reference Rows operations and deduplication
- Literature support in all products
- Combine literature with patents OR trials
- Options for Citation formats, exports
- Support for additional sources
- Additional fields for analysis
- Improved hyperlink generation

Agenda

- IP Sequence reports
- **Hit Structure reports**
- Summary Record exports
- Reference Rows

Deliver reports in HTML, Word, Excel...

LifeQuest: Sample

	Title	Patent	Patent Family Kind	Date	Inventor(s)	CPC	Abstract	Claims	Pub. Status
1 Link	Drug delivery compositions	US7329638	B2	2008-02-12	Victor C. Yang Yoon Jeong Park Junfeng Liang	A61K2039/505 A61K47/48315 A61K47/48884 A61K47/4823 A61K47/48561	The present invention relates to multicomponent compositions and methods of administering these compositions, which specifically translocate	1. A composition, comprising: a. a first targeting component, wherein said first targeting component comprises i. a molecular recognition element, wherein said molecular recognition element has	Grant
2 Link	Drug delivery compos	<p>Link to record on source platform</p> <p>Link to web resource (like USPTO)</p> <p>ALL-TEXT AND IMAGE DATABASE</p> <p>Advanced Pat Num Help</p> <p>Bottom</p> <p>view Cart Add to Cart</p> <p>Images</p> <p>(1 of 1)</p> <p>United States Patent 7,329,638 Yang, et al. February 12, 2008</p> <p>Drug delivery compositions</p> <p>Abstract</p> <p>The present invention relates to multicomponent compositions and methods of administering these compositions, which specifically translocate therapeutic molecules (e.g., drugs or prodrugs) across biological membranes thus reducing potential toxic side effects on nontargeted cells and tissues.</p> <p>Inventors: Yang; Victor C. (Ann Arbor, MI), Park; Yoon Jeong (Seoul, KR), Liang; Junfeng (Westfield, NJ) Assignee: The Regents of the University of Michigan (Ann Arbor, MI) Family ID: 34198915 Appl. No.: 10/835,151 Filed: April 29, 2004</p> <p>Oncology drug innova</p> <p>Prior Publication Data</p> <p>Document Identifier: US 20050042753 A1 Publication Date: Feb 24, 2005</p> <p>Related U.S. Patent Documents</p>							
3 Link	POLYMORPHISMS OF REGION OF THE HUMAN HT1A GENE, ASSOCIATED WITH PROTEINS OF THE 5' HTR2A GENE AND A DIAGNOSTIC TEST FOR MAJOR DEPRESSION AND RELATED MENTAL ILLNESSES								
4 Link	Oncology drug innova								

Summary Record export in Word

3.	Basic Patent Number:	WO2012033858A2
	Title:	Boron-containing small molecules
	Inventor(s):	Hernandez, Vincent S.; Ding, Charles; Plattner, Jacob J.; Alley, Michael Richard Kevin; Rock, Fernando; Zhang, Suoming; Easom, Eric; Li, Xianfeng; Zhou, Ding
	Patent Assignee:	Anacor Pharmaceuticals, Inc., USA
	Hyperlinks:	Source WO2012033858A2



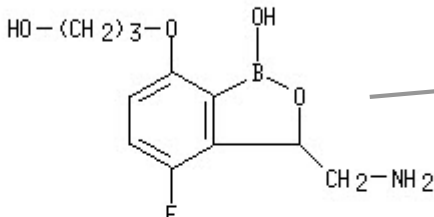
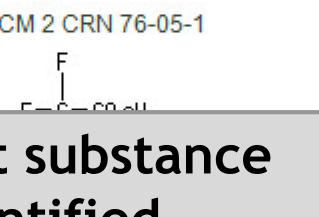
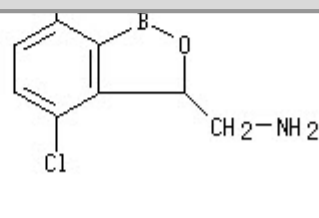
- Fields are the columns in chart
- Content, like hyperlinks, is included in the Summary Record

THE JOURNEY BEGINS...

Summary Record export with Hit Structures

3.	Basic Patent Number:	WO2012033858A2
	Title:	Boron-containing small molecules
	Inventor(s):	Hernandez, Vincent S.; Ding, Charles; Plattner, Jacob J.; Alley, Michael Richard Kevin; Rock, Fernando; Zhang, Suoming; Easom, Eric; Li, Xianfeng; Zhou, Ding
	Patent Assignee:	Anacor Pharmaceuticals, Inc., USA
	Hyperlinks:	Source WO2012033858A2

Text from CPlus record

Hit Structures:		
<p>1364682-96-1 (Cmpd. 2)</p> <p>1-Propanol, 3-[[3-(aminomethyl)-4-fluoro-1,3-dihydro-1-hydroxy-2,1-benzoxaborol-7-yl]oxy]-, 2,2,2-trifluoroacetate (1:2) (CA INDEX NAME)</p>	<p>CM 1 CRN 1364682-95-0</p>  <p>CM 2 CRN 76-05-1</p> 	<p>RL: PAC (Pharmacological activity); SPN (Synthetic preparation); TH (Therapeutic use); BIOL (Biological study); U (Preparation); U (Uses)</p> <p>prepn. of benzoxaborole useful for treating bacterial infections</p>
<p>1364683-03-3 (Cmpd. 3)</p> <p>1-Propanol, 3-[[3-(aminomethyl)-4-chloro-1,3-dihydro-1-hydroxy-2,1-benzoxaborol-7-yl]oxy]-, hydrochloride (1:1) (CA INDEX NAME)</p>	<p>CM 1 CRN 1364683-03-0</p>  <p>• HCl</p>	<p>RL: PAC (Pharmacological activity); SPN (Synthetic preparation); THU (Therapeutic use); BIOL (Biological study); U (Preparation); U (Uses)</p> <p>prepn. of benzoxaborole derivs. useful for treating bacterial infections</p>

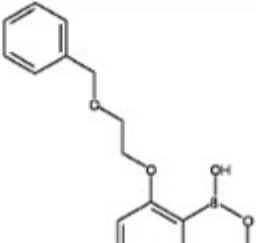
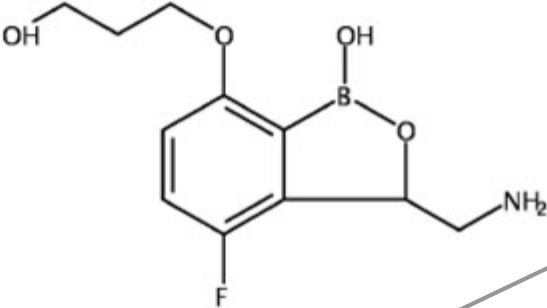
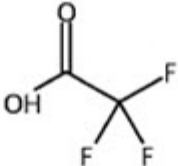
Structures with annotations including multi page images

Each hit substance identified

Index Terms

A structure oriented “Index of Hit Structures”

Index of Hit Structures

Substance	Structure
<p>1 1655492-02-6</p> <p>2,1-Benzoxaborole, 4-fluoro-1,3-dihydro-1-hydroxy-3-(nitromethyl)-7-[2-(phenylmethoxy)ethoxy]-</p>	 <p>Each hit substance identified</p>
<p>2 1364682-96-1</p> <p>1-Propanol, 3-[[3-(aminomethyl)-4-fluoro-1,3-dihydro-1-hydroxy-2,1-benzoxaborol-7-yl]oxy]-, 2,2,2-trifluoroacetate (1:2)</p>	 <p>CM2 CRN 76-05-1</p> 

Structures with annotations

prepn. and biol. applications of tricyclic benzoxaborole compds. [Reference 1](#)

prepn. of benzoxaborole derivs. useful for treating bacterial infections [Reference 2](#)

prepn. of benzoxaborole derivs. useful for treating bacterial infections [Reference 3](#)

Links to references for each structure

Multiple images (continuation) or mixtures

Option: Index of Hit Structures

Summary Record Export Options

The Summary Record export shows the columns (fields) visible in your chart.

- Number the records
- Start each record on new page
- Include Links section
- Include editable Notes section
- Include Index of Hit Structures

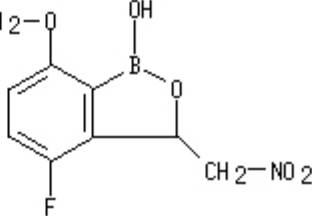
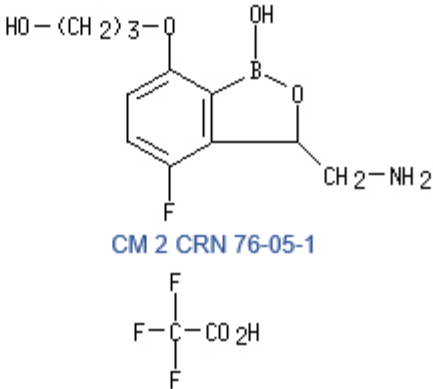
You may also include the following information for the record:

- Include Claims
- Include Alignment
- Include Hit Structures
- Include Index Terms

OK Cancel Help

- Include Editable Notes Section
- Include Index of Hit Structures

Index of Hit Structures

	Substance	Structure	Reference
1	<p>1655492-02-6</p> <p>2,1-Benzoxaborole, 4-fluoro-1,3-dihydro-1-hydroxy-3-(nitromethyl)-7-[2-(phenylmethoxy)ethoxy]- (CA INDEX NAME)</p>	<p>Ph-CH₂-O-CH₂-CH₂-O</p> 	<p>prepn. and antimycobacterial activity of benzoxaborole compds. Reference 1</p> <p>prepn. and biol. applications of tricyclic benzoxaborole compds. Reference 2</p>
2	<p>1364682-96-1</p> <p>1-Propanol, 3-[[[3-(aminomethyl)-4-fluoro-1,3-dihydro-1-hydroxy-2,1-benzoxaborol-7-yl]oxy]-, 2,2,2-trifluoroacetate (1:2) (CA INDEX NAME)</p>	<p>CM 1 CRN 1364682-95-0</p>  <p>CM 2 CRN 76-05-1</p>	<p>prepn. of benzoxaborole derivs. useful for treating bacterial infections Reference 3</p>

Option: Hit Structures

Summary Record Export Options

The Summary Record export shows the columns (fields) visible in your chart.

Number the records
 Start each record on new page
 Include Links section
 Include editable Notes section
 Include Index of Hit Structures

You may also include the following information for the record:

Include Claims
 Include Alignment
 Include Hit Structures
 Include Index Terms

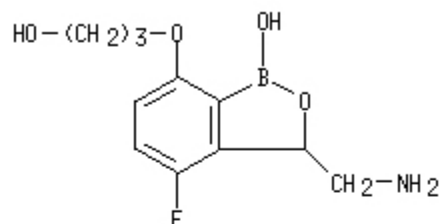
- Include Alignment
- Include Hit Structures
- Include Index Terms

Hit Structures:

1364682-96-1 [\(Cmpd. 2\)](#)

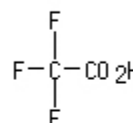
1-Propanol, 3-[[3-(aminomethyl)-4-fluoro-1,3-dihydro-1-hydroxy-2,1-benzoxaborol-7-yl]oxy]-, 2,2,2-trifluoroacetate (1:2) (CA INDEX NAME)

CM 1 CRN 1364682-95-0



RL: PAC (Pharmacological activity); SPN (Synthetic preparation); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); USES (Uses)

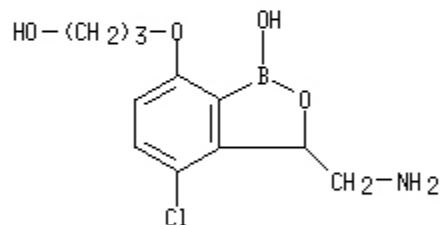
CM 2 CRN 76-05-1



prepn. of benzoxaborole derivs. useful for treating bacterial infections

1364683-03-3 [\(Cmpd. 3\)](#)

1-Propanol, 3-[[3-(aminomethyl)-4-chloro-1,3-dihydro-1-hydroxy-2,1-benzoxaborol-7-yl]oxy]-, hydrochloride (1:1) (CA INDEX NAME)



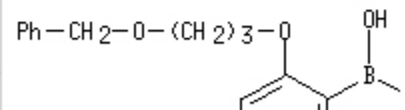
RL: PAC (Pharmacological activity); SPN (Synthetic preparation); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); USES (Uses)

• HCl

prepn. of benzoxaborole derivs. useful for treating bacterial infections

1364684-69-4 [\(Cmpd. 4\)](#)

2,1-Benzoxaborole, 4-fluoro-1,3-dihydro-1-hydroxy-3-



RL: RCT (Reactant); SPN (Synthetic preparation); PREP (Preparation); RACT

Option: Index Terms

Summary Record Export Options


The Summary Record export shows the columns (fields) visible in your chart.

- Number the records
- Start each record on new page
- Include Links section
- Include editable Notes section
- Include Index of Hit Structures

You may also include the following information for the record:

- Include Claims
- Include Alignment
- Include Hit Structures
- Include Index Terms

OK Cancel Help



Include Hit Structures

Include Index Terms

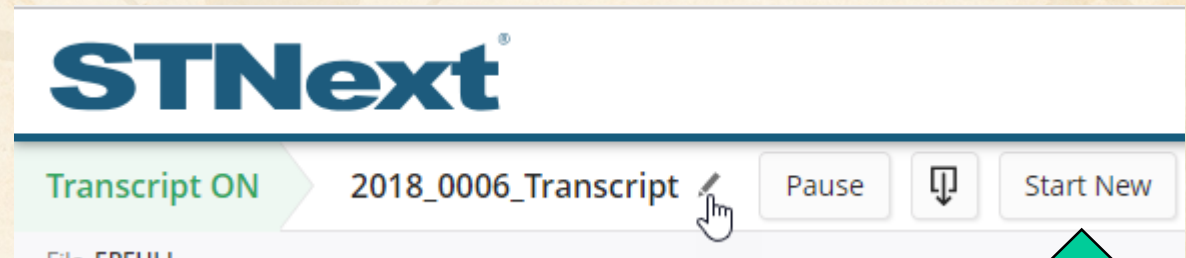
Index Terms:

1364682-96-1P ([Cmpd. 2](#)) 1364683-03-3P ([Cmpd. 3](#)) PAC (Pharmacological activity); SPN (Synthetic preparation); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); USES (Uses) (prepn. of benzoxaborole derivs. useful for treating bacterial infections)

1364684-69-4P ([Cmpd. 4](#)) 1364684-75-2P ([Cmpd. 5](#)) RCT (Reactant); SPN (Synthetic preparation); PREP (Preparation); RACT (Reactant or reagent) (prepn. of benzoxaborole derivs. useful for treating bacterial infections)

Importing transcripts with hit structures (STNext)

- Make sure that Classic Display is on
- Start new transcript before displaying.



BIB vs. IBIB

- We recommend using tagged (BIB), rather than indented (IBIB), display formats
- Some field contents (table headings) appear before the label in IBIB
- Indent levels in RTF are more reliably detected in BIB

Agenda

- IP Sequence reports
- Hit Structure reports
- **Summary Record exports**
- Reference Rows

Summary Record export in Word

3.	Basic Patent Number:	WO2012033858A2
	Title:	Boron-containing small molecules
	Inventor(s):	Hernandez, Vincent S.; Ding, Charles; Plattner, Jacob J.; Alley, Michael Richard Kevin; Rock, Fernando; Zhang, Suoming; Easom, Eric; Li, Xianfeng; Zhou, Ding
	Patent Assignee:	Anacor Pharmaceuticals, Inc., USA
	Hyperlinks:	Source WO2012033858A2

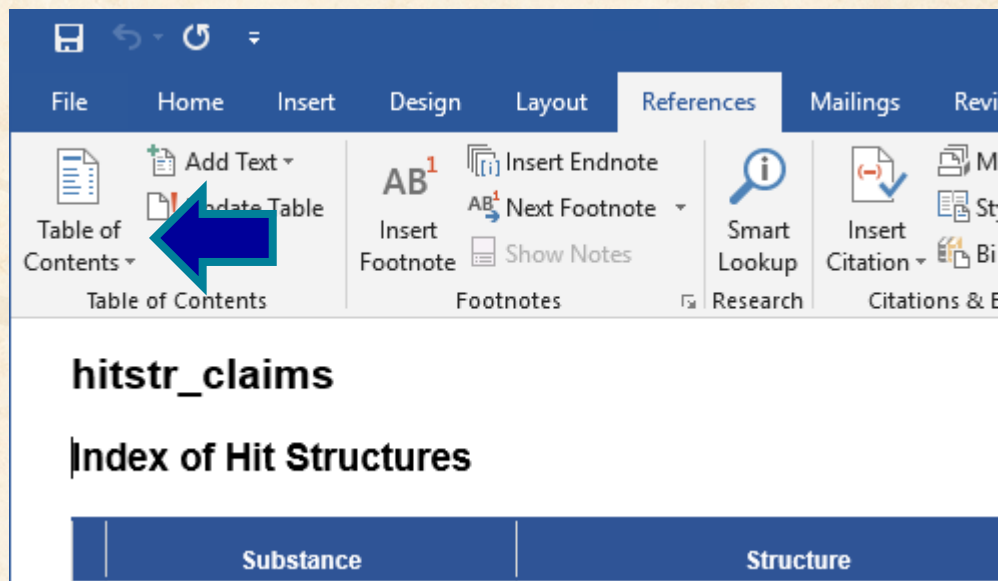


- Fields are the columns in chart
- Content, like hyperlinks, is included in the Summary Record

THE JOURNEY BEGINS...

Summary Records - Table of Contents

- Add a table of contents in Word



Summary Records - Table of Contents

- Sections are marked up for headings
- First column in chart is used as TOC entry

hitstr claims

Contents

[Index of Hit Structures](#)

[References](#)

[Tricyclic benzoxaborole compounds and uses thereof](#)

[Boron-containing small molecules](#)

[Benzoxaborole compounds and uses thereof](#)

Index of Hit Structures

Substance	Structure
-----------	-----------

Summary Records - Table of Contents

- Can collapse the Index of Hit Structures (in recent versions of Word)

▷ **Index of Hit Structures**

References

1.	Title:	Tricyclic benzoxaborole compounds and uses thereof		
	Common Family:	EP 3030519		
	Database:	Chemical Abstracts EP Patents Fulltext		
	Patent Family:	Patent	Kind	Date
		WO 2015021396	A2	20150212
		WO 2015021396	A3	20151029

F

Option: Claims + Hit Structures

Summary Record Export Options

The Summary Record export shows the columns (fields) visible in your chart.

Number the records
 Start each record on new page
 Include Links section
 Include editable Notes section
 Include Index of Hit Structures

You may also include the following information for the record:

Include Claims
 Include Alignment
 Include Hit Structures
 Include Index Terms

You may also include the following information for the record:

Include Claims

Include Alignment

10. An in vitro method of:

(A) inhibiting an enzyme, comprising: contacting the enzyme with the compound of any of claims 1 to 5, thereby inhibiting the enzyme;

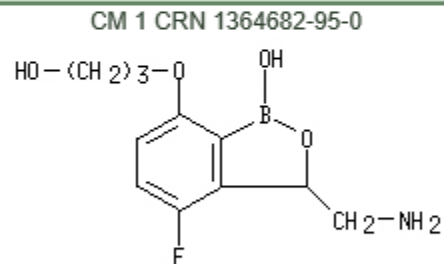
(B) killing and/or preventing the growth of a microorganism, comprising: contacting the microorganism with an effective amount of the compound of any of claims 1 to 5, thereby killing and/or preventing the growth of the microorganism; or

(C) inhibiting the editing domain of a t-RNA synthetase, comprising: contacting the synthetase with an effective amount of a compound of any of claims 1 to 5, or a pharmaceutically-acceptable salt thereof, thereby inhibiting the synthetase.

Hit Structures:

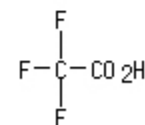
1364682-96-1 ([Cmpd. 2](#))

1-Propanol, 3-[[3-(aminomethyl)-4-fluoro-1,3-dihydro-1-hydroxy-2,1-benzoxaborol-7-yl]oxy]-, 2,2,2-trifluoroacetate (1:2) (CA INDEX NAME)



RL: PAC
(Pharmacological activity); SPN (Synthetic preparation); THU (Therapeutic use); BIOL (Biological study); PREP (Preparation); USES (Uses)

CM 2 CRN 76-05-1



prepn. of [benzoxaborole derivs.](#) useful for treating bacterial infections

1364683-03-3 ([Cmpd. 3](#))



RL: PAC

Link sequence results with hit structures

Follow Cookbook recipe to create a summary of sequence hits for each

1.	Title: Selective high-affinity polydentate ligands and methods of making such																																
	Database: GQPAT Gold+ Proteins GQPAT Gold+ Proteins GQPAT Gold+ Proteins GQPAT Gold+ Proteins GQPAT Gold+ Proteins GQPAT Gold+ Proteins GQPAT Gold+ Proteins GQPAT Gold+ Proteins Chemical Abstracts Chemical Abstracts																																
	Organism Species: Homo sapiens (human)																																
Sequence Summary:	<table border="1"><thead><tr><th>Seq. ID Number</th><th>Length</th><th>% Identity</th><th>Location</th></tr></thead><tbody><tr><td>US20180008621-0003</td><td>237</td><td>100.00</td><td>probable disclosure (not found by automated parsing)</td></tr><tr><td>US9884070-0003</td><td>237</td><td>100.00</td><td>probable disclosure (not found by automated parsing)</td></tr><tr><td>US20180008622-0003</td><td>237</td><td>100.00</td><td>probable disclosure (not found by automated parsing)</td></tr><tr><td>JP5623384-0003</td><td>237</td><td>100.00</td><td>probable disclosure (not found by automated parsing)</td></tr><tr><td>JP2014122234-0003</td><td>237</td><td>100.00</td><td>probable disclosure (not found by automated parsing)</td></tr><tr><td>US20110144065-0003</td><td>237</td><td>100.00</td><td>probable disclosure (not found by automated parsing)</td></tr><tr><td>CA2721980-0003</td><td>237</td><td>100.00</td><td>probable disclosure (not found by automated parsing)</td></tr></tbody></table>	Seq. ID Number	Length	% Identity	Location	US20180008621-0003	237	100.00	probable disclosure (not found by automated parsing)	US9884070-0003	237	100.00	probable disclosure (not found by automated parsing)	US20180008622-0003	237	100.00	probable disclosure (not found by automated parsing)	JP5623384-0003	237	100.00	probable disclosure (not found by automated parsing)	JP2014122234-0003	237	100.00	probable disclosure (not found by automated parsing)	US20110144065-0003	237	100.00	probable disclosure (not found by automated parsing)	CA2721980-0003	237	100.00	probable disclosure (not found by automated parsing)
Seq. ID Number	Length	% Identity	Location																														
US20180008621-0003	237	100.00	probable disclosure (not found by automated parsing)																														
US9884070-0003	237	100.00	probable disclosure (not found by automated parsing)																														
US20180008622-0003	237	100.00	probable disclosure (not found by automated parsing)																														
JP5623384-0003	237	100.00	probable disclosure (not found by automated parsing)																														
JP2014122234-0003	237	100.00	probable disclosure (not found by automated parsing)																														
US20110144065-0003	237	100.00	probable disclosure (not found by automated parsing)																														
CA2721980-0003	237	100.00	probable disclosure (not found by automated parsing)																														

Option: Alignments + Hit Structures

Summary Record Export Options

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You may also include the following information for the record:

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- Include Alignment
- Include Hit Structures
- Include Index Terms

- Include Claims
- Include Alignment
- Include Hit Structures

Alignment:

```
Q:      1  GDTRPRFLEEVKFECHFFNGTIERVRLLERRVHNOEEYARYSDVGEYRAVTELGRPDAEY  60
      |
S:      1  GDTRPRFLEEVKFECHFFNGTIERVRLLERRVHNOEEYARYSDVGEYRAVTELGRPDAEY  60
      |
Q:     61  WNSQKDLLERRRAAVDTYCRHNYGVGESFTVQRRVQPKVTIVYPSKTQPLQHHNLLVCSVN  120
      |
S:     61  WNSQKDLLERRRAAVDTYCRHNYGVGESFTVQRRVQPKVTIVYPSKTQPLQHHNLLVCSVN  120
      |
Q:    121  GFYPGSIEVRWFRNGQEEKTGCVSTGLIQNGDWTFOILVMLETVPOSGEVYTCQVEHPSV  180
      |
S:    121  GFYPGSIEVRWFRNGQEEKTGCVSTGLIQNGDWTFOILVMLETVPOSGEVYTCQVEHPSV  180
      |
Q:    181  MSPLTVEWRARSESAQSKMLSGVGGFVLGLLVLGAGLFYIFRNQKGHSGLPPTGFLS  237
      |
S:    181  MSPLTVEWRARSESAQSKMLSGVGGFVLGLLVLGAGLFYIFRNQKGHSGLPPTGFLS  237
```

Option: Alignments + Hit Structures

Summary Record Export Options

The Summary Record export shows the columns (fields) visible in your chart.

- Number the records
- Start each record on new page
- Include Links section
- Include editable Notes section
- Include Index of Hit Structures

You may also include the following information for the record:

- Include Claims
- Include Alignment
- Include Hit Structures
- Include Index Terms

OK Cancel Help

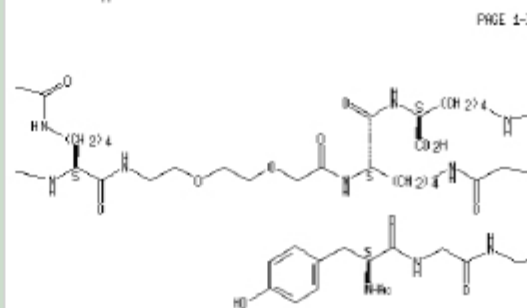
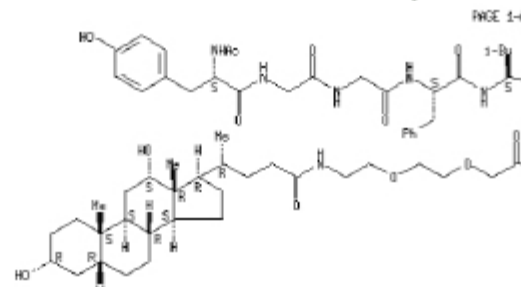
- Include Alignment
- Include Hit Structures

Hit Structures:

1045703-23-8 ([Cmpd. 1](#))

L-Lysine, N2,N6-bis[N6-(N-acetyl-L-tyrosylglycylglycyl-L-phenylalanyl-L-leucyl)-N2-[2-[2-[[[(3-oxocholan-24-yl)amino]ethoxy]ethoxy]acetyl]-L-lysyl-2-[2-(2-aminoethoxy)ethoxy]acetyl]-L-lysyl-N6-[5-[(3aS,4S,6aR)-hexahydro-2-oxo-1H-thieno[3,4-d]imidazol-4-yl]-1-oxopentyl]- (CA INDEX NAME)

Absolute stereochemistry.



RL: BSU (Biological study, unclassified); DGN (Diagnostic use); PAC (Pharmacological activity); THU (Therapeutic use); BIOL (Biological study); USES (Uses)

HLA-DR10 selective high-affinity polydentate ligand; selective high-affinity polydentate ligands of target mols. and methods of making such and uses for diagnosis and therapeutics in relation to delivery of effectors

Agenda

- IP Sequence reports
- Hit Structure reports
- Summary Record exports
- **Reference Rows**



BizInt Smart Charts

VERSION

5

for Patents

Patent Databases

Provide data on patents filed worldwide

- STN - Classic, STNext, & New STN
- Questel Orbit.com
- Minesoft PatBase
- Innovation, Cortellis IP, Integrity Patents
- LexisNexis TotalPatent
- GQ LifeSciences LifeQuest

Tools for integrating patent data

- **Combine charts** using **File | Combine** command
- **Identify related records** using the **“Identify Common Patent Family”** tool.

BizInt Smart Charts

for Patents

Tools for integrating patent data

- **Combine charts** using **File | Combine** command
- **Identify related records** using the “**Identify Common Patent Family**” tool.
- Use **BizInt Smart Charts Reference Rows** to summarize related records in a single row.

BizInt Smart Charts

for Patents

BizInt Smart Charts

Reference Rows™

Combining Charts

- You can combine two or more charts into a single report
- BizInt Smart Charts usually removes duplicate rows when combining
- Combining different queries? Use the “Combine without removing duplicates” option in the Combine wizard
- See 2013 PIUG Biotech presentation for more details (@ bizint.com/slides)

Combining multiple queries

Create Combined Chart Wizard

Step 3 - Select options for new combined chart:

Enter new chart title:

Sequence results - three queries

Select the operation you would like to perform:

- Combine charts from different databases.**
Build a report from different sources, aligning common fields.
- Add additional results from same search.**
Create one report from results saved in several files.

• Combine without removing duplicates.
Useful for gene sequence charts from multiple queries.

Advanced...

Finish

< Back

Cancel

Help

Preserving Multiple Queries

- First combine non-sequence charts with standard options (remove duplicates)
- Then combine the resulting non-sequence chart with the sequence results (without removing duplicates)
- Only use “without removing duplicates” option when you want to see variations on a row “side by side”

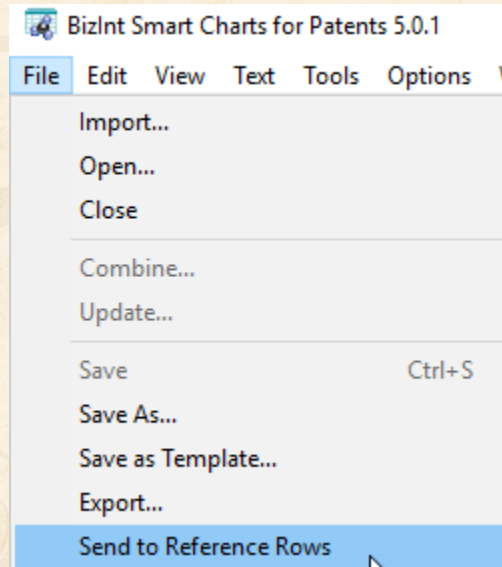
Common Patent Family

Database	Common Family	Patent Family		
		Patent	Kind	Date
Derwent World Patents Index	US 2014356956	US20140356959	A1	20141204
Derwent World Patents Index	US 2014356956	US20140356956 WO2014197568 WO2014197568 CA2914638	A1 A2 A3 A1	20141204 20141211 20150312 20141211
FAMPAT	US 2014356956	US 2014356956 US 2014356959 US 9267135	A1 A1 B2	2014-12-04 2014-12-04 2016-02-23
GQPAT Gold+ Proteins	US 2014356956	US20140356959		20141204
GQPAT Gold+ Proteins	US 2014356956	US20140356956		20141204
PatBase	US 2014356956	US 2014356959 US 2014356956 AU 2014274939 WO 14197568 WO 14197568 CA2914638 KR 20160014036	A A AA A2 A3 AA A	2014-12-04 2014-12-04 2014-12-11 2014-12-11 2015-03-12 2015-12-04 2016-02-05

“Identify Common Patent Families” tool

Minimum effort with Reference Rows

- Save chart in BizInt Smart Charts for Patents
- Send to Reference Rows



Minimum effort with Reference Rows

- Generate Common Patent Family (if needed)

Create Reference Rows (1 of 3)

Welcome to the Create Reference Rows Wizard

BizInt Smart Charts Reference Rows offers the ability to create a "Reference Row" which combines information from a set of related records into a single row. The Create Reference Rows Wizard will help you create and set up rules for Reference Rows.

NOTE: Reference Rows are based on the Common Patent Family column in your report. You can generate this column now.

Click Next to continue.

Generate

< Back Next > Finish Cancel



Minimum effort with Reference Rows

- Simply “Finish” on step two

Create Reference Rows (2 of 3)

Database Ranking


Data in cells will be chosen according to the Database Ranking if no other rule is present or if there is a tie in the rules.

Rank the databases in your preferred order

CAS REGISTRY Chemical Abstracts	Move Up
	Move Down
	Properties

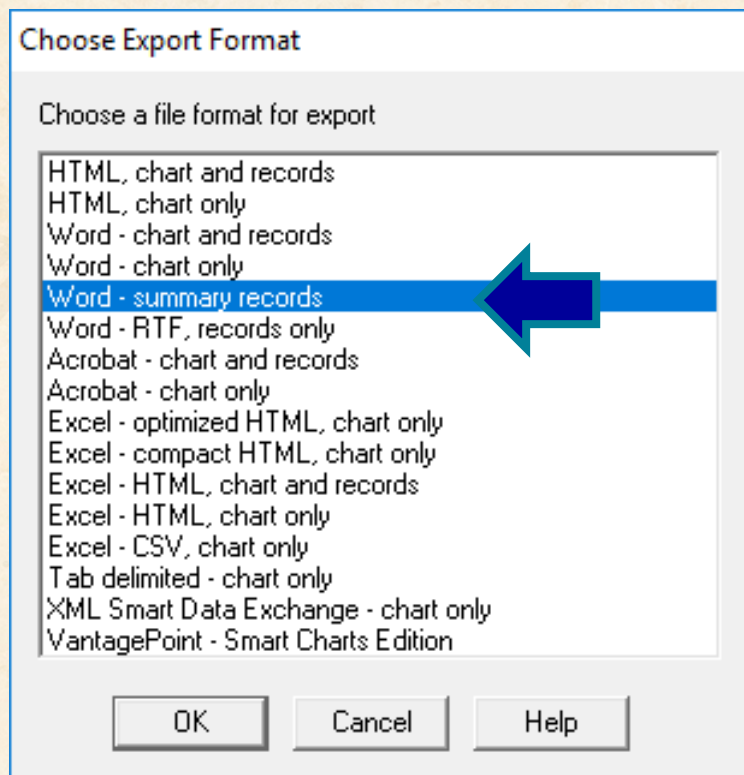
Rules Template

Database rankings and rules based on:



Minimum effort with Reference Rows

- File | Export ... Word - summary records



Reference Rows: Selection View

Unique fields are easily integrated in BizInt Smart Charts Reference Rows

Enhanced Title	Indications	Patent Type	Classifications	Family Status			
				Pub No.	State	Status	Expiry
Monoclonal antibodies and vaccines against epitopes on the Ebola virus glycoprotein ✓	Ebola virus infection ✓	Product ✓	Anti-Infectives Biologicals and Immunologicals ✓				
				WO200116183	DEAD	LAPSED	2006-03-26 ✓
				AU7089600	DEAD	LAPSED	2006-03-26
				US6630144	ALIVE	GRANTED	2020-08-29
Monoclonal antibodies against glycoprotein of Ebola Sudan Boniface (ESB) virus - useful in the diagnosis and treatment of ESB virus infection. ✓	Ebola virus infection ✓	Diagnostic, Analysis and Assay Product (Macromolecule) ✓	Anti-Infectives Biologicals and Immunologicals Diagnostics ✓				
				WO2011071574	ALIVE	PENDING	2030-09-01 ✓
				EP2473525	DEAD	LAPSED	2014-08-27
				US2012164153	ALIVE	PENDING	2030-09-01
Ebola virus liposome vaccines - useful in eliciting immune responses against Ebola virus infection. ✓	Ebola virus infection ✓	Formulation ✓	Anti-Infectives Biologicals and Immunologicals Pharmaceutics ✓				
				WO2012050193	DEAD	LAPSED	2013-12-03 ✓
				JP2014005205	ALIVE	PENDING	2030-10-14

Reference Rows: HTML exports

As seen in the fully integrated view

	Enhanced Title	Indications	Patent Type	Classifications	Family Status				Database
					Pub No.	State	Status	Expiry	
2.	Monoclonal antibodies and vaccines against epitopes on the Ebola virus glycoprotein	Ebola virus infection	Product	Anti-Infectives Biologicals and Immunologicals	WO200116183	DEAD	LAPSED	2006-03-26	2.1 CORTP link 2.2 FAMPAT link
					AU7089600	DEAD	LAPSED	2006-03-26	
					US6630144	ALIVE	GRANTED	2020-08-29	
	2.1 CORTP	2.1 CORTP	2.1 CORTP	2.1 CORTP					2.2 FAMPAT
3.	Monoclonal antibodies against glycoprotein of Ebola Sudan Boniface (ESB) virus - useful in the diagnosis and treatment of ESB virus infection.	Ebola virus infection	Diagnostic, Analysis and Assay Product (Macromolecule)	Anti-Infectives Biologicals and Immunologicals Diagnostics	WO2011071574	ALIVE	PENDING	2030-09-01	3.1 CORTP link 3.2 FAMPAT link
					EP2473525	DEAD	LAPSED	2014-08-27	
					US2012164153	ALIVE	PENDING	2030-09-01	
	3.1 CORTP	3.1 CORTP	3.1 CORTP	3.1 CORTP					3.2 FAMPAT
4.	Ebola virus liposome vaccines - useful in eliciting immune responses against Ebola virus infection.	Ebola virus infection	Formulation	Anti-Infectives Biologicals and Immunologicals Pharmaceutics	WO2012050193	DEAD	LAPSED	2013-12-03	4.1 CORTP link 4.2 FAMPAT link
					JP2014005205	ALIVE	PENDING	2030-10-14	
	4.1 CORTP	4.1 CORTP	4.1 CORTP	4.1 CORTP					4.2 FAMPAT
5.	Chimeric filovirus glycoproteins useful in vaccines against Ebola and Marburg virus infections	Marburg virus infection Ebola virus infection	Product	Anti-Infectives Biologicals and Immunologicals	WO02079239	DEAD	LAPSED	2006-03-29	5.1 CORTP link 5.2 FAMPAT link
					US7731975	DEAD	LAPSED	2014-06-08	
	5.1 CORTP	5.1 CORTP	5.1 CORTP	5.1 CORTP					5.2 FAMPAT

Integrate data from related records

Enhanced Title	Database	Patent Family			Family Status				Alignment	% Identity
		Patent	Kind	Date	Pub No.	State	Status	Expiry		
5. Methods for detecting the presence of isolated attenuated hEbola virus - useful as vaccines.	5.1 FAMPAT link	WO 201048615	A2	2010-04-29	WO2010048615	ALIVE	PENDING	2029-10-26	Q: 1 SFKAALSSL 9 100.00	
	5.2 CORTP link	CA 2741523	A1	2010-04-29	AU2009308422	ALIVE	PENDING	2029-10-26		
	5.3 GPATPRT link	AU 2009308422	A1	2010-04-29	CA2741523	ALIVE	PENDING	2029-10-26	S: 279 SFKAALSSL 287	
	5.4 GPATPRT link	WO 201048615	A3	2010-11-25	EP2350270	ALIVE	PENDING	2029-10-26		
	5.5 GPATNUC link	EP 2350270	A2	2011-08-03	IN3817/DELNP/2011	ALIVE	PENDING	2029-10-26		
	5.6 GPATNUC link	EP 2350270	A4	2012-04-11	US2012251502	ALIVE	PENDING	2029-10-26		
	5.7 GENESEQ link	US 20120251502	A1	2012-10-04						
		IN 2011DN03817	A	2013-09-27						
6. Recombinant biologically compatible filovirus	6.1 FAMPAT link	WO 2009128867	A2	2009-10-22	WO2009128867	DEAD	LAPSED	2010-08-08	Q: 1 SFKAALSSL 9 100.00	
	6.2 GENESEQ link	WO 2009128867	A3	2010-03-25						
									S: 1 SFKAALSSL 9	
7. Nucleic acid comprising a polynucleotide encoding a modified filovirus glycoprotein - useful as vaccines against filovirus infections, specifically Ebola virus.	7.1 FAMPAT link	WO 200637038	A1	2006-04-06	WO2006037038	ALIVE	PENDING	2025-09-27	Q: 1 HNTFPVYKLDISEATQVE 17 100.00	
	7.2 CORTP link	CA 2581840	A1	2006-04-06	AU2005289439	ALIVE	GRANTED	2025-09-27		
	7.3 GPATPRT link	AU 2005289439	A1	2006-04-06	CA2581840	ALIVE	GRANTED	2025-09-27	S: 389 HNTFPVYKLDISEATQVE 405	
	7.4 GPATPRT link	WO 200637038	A9	2006-05-26	EP1797113	ALIVE	GRANTED	2025-09-27		
	7.5 GPATPRT link	WO 200637038	B1	2006-08-03	IL182225	DEAD	LAPSED	2012-09-20		
	7.6 GPATPRT link	EP 1797113	A1	2007-06-20	IN2674/DELNP/2007	ALIVE	GRANTED	2025-09-27		
	7.7 GPATPRT link	IN 2007DN02674	A	2007-08-03	JP2008514203	ALIVE	GRANTED	2025-09-27		
	7.8 GENESEQ link	IL 182225	D0	2007-09-20	US2009232841	ALIVE	GRANTED	2027-06-07		
	7.9 GENESEQ link	JP 2008514203	A	2008-05-08	US8101739	ALIVE	GRANTED	2027-06-07		
	7.10 GENESEQ link	US 20090232841	A1	2009-09-17	US2012156239	ALIVE	PENDING	2025-09-27		
		AU 2005289439	B2	2011-12-01						
		US 8101739	B2	2012-01-24						
		US 20120156239	A1	2012-06-21						
		JP 5046941	B2	2012-10-10						
		IN 259912	B	2014-04-04						
		CA 2581840	C	2014-08-05						
		EP 1797113	B1	2014-11-26						

Reference Rows: user-defined rules

Cell Selection Rule - Title

Title
Choose how Reference Rows will select data for this column.

Selection Rule: Use database ranking

Match column:

i Use the database select.

Database Ranking for this column:

- Derwent World Patents Index
- Thomson Innovation + DWPI
- TotalPatent
- PatBase
- MicroPatent
- FAMPAT

Cell Selection Rule - Patent Assignee

Patent Assignee
Choose how Reference Rows will select data for this column.

Selection Rule: Most Recently Updated

Match column: Use database ranking, Earliest Date, Latest Date, Most Content (characters), Most Content (lines), Highest Development Phase, **Most Recently Updated**, Match Column

Database Ranking for this column:

- Derwent World Patents Index
- MicroPatent
- TotalPatent
- PatBase
- FAMPAT
- Thomson Innovation + DWPI

Move Up

Move Down

OK Cancel

Summarize data from related records

Title	Database	Patent Assignee	Query ID	Sequence Locations				
				Seq. ID Number	% Identity	Length	Location	
1. PRODUCTION OF PEPTIDES IN PLANTS AS VIRAL COAT PROTEIN FUSION	1.1 Patbase link	LARGE SCALE BIOLOGY CORP.	query2	WO20050108564-0101	100.00	17	Example 6; SEQ ID NO 101;	1.2
	1.2 GENESEQ link							
1.1 Patbase		1.2 GENESE						
2. Chimeric ebola virus envelopes and uses therefor	2.1 Patbase link	UNIV PENNSYLVANIA.	query2	US20050255123-0001	100.00	17	claim: 17	2.2
	2.2 GPATPRT link			WO03092582-0009	100.00	498	claim: 17	2.3
	2.3 GPATPRT link			WO03092582-0001	100.00	17	claim: 17	2.4
	2.4 GPATPRT link			US20050255123-0009	100.00	498	claim: 17	2.5
	2.5 GPATPRT link			WO20030092582-0001	100.00	17	Claim 17; SEQ ID NO 1; 107pp; English.	2.6
	2.6 GENESEQ link			WO20030092582-0009	100.00	498	Claim 17; SEQ ID NO 9; 107pp; English.	2.7
	2.7 GENESEQ link							
2.1 Patbase		2.6 GENESE						
3. ANTIGEN FRAGMENT AND TRUNCATION BASED ON EBOLA VIRUS ENVELOPE PROTEIN AS WELL AS APPLICATION	3.1 Patbase link	BIOENGINEERING RES INST ACAD MEDICAL SCI.	query2	CN103864904-0008	100.00		Example 1; SEQ ID NO 8; 28pp; Chinese.	3.2
	3.2 GENESEQ link			CN103864904-0002	100.00	17	Example 1; SEQ ID NO 2; 28pp; Chinese.	3.3
	3.3 GENESEQ link							
3.1 Patbase		3.2 GENESE						
4. HUMAN EBOLA VIRUS SPECIES AND COMPOSITIONS AND METHODS THEREOF	4.1 Patbase link	US DEPT HEALTH & HUMAN SERVICES.	query7 query5	US20120251502-0011	100.00	9	claim: 8; 11; 12	4.2
	4.2 GPATPRT link			EP2350270-0011	100.00	9	TBD (information not in GQ-Pat)	4.3
	4.3 GPATPRT link			US20120251502-0027	100.00	20	probable disclosure (not found by automated parsing)	4.4
	4.4 GPATNUC link			EP2350270-0027	100.00	20	TBD (information not in GQ-Pat)	4.5
	4.5 GPATNUC link			WO20100048615-0027	100.00	20	Claim 30; SEQ ID NO 27; 98pp; English.	4.6
	4.6 GENESEQ link							
4.1 Patbase		4.6 GENESE						

Sequence Summary Recipe

- Recipe for creating the sequence summary table at bizint.com/cookbook
- Create Subtable from any columns you want
NOTE: alignment loses fixed width formatting
- In Reference Rows, choose Summarize All Values column rule
- Export chart

Summary Record export

1.	Title:	Modulating expression of a target nucleic acid comprises providing to the cell a guide RNA including a transcriptional activator or repressor domain as a fusion protein, and providing to the cell a nuclease null Cas9 protein			
	Database:	Derwent World Patents Index Derwent World Patents Index GQPAT Gold+ Proteins GQPAT Gold+ Proteins PatBase FAMPAT			
	Patent Family:	Patent	Kind	Date	
		US 2014356959	A	2014-12-04	
		US 2014356956	A	2014-12-04	
		AU 2014274939	AA	2014-12-11	
		WO 14197568	A2	2014-12-11	
		WO 14197568	A3	2015-03-12	
		CA 2914638	AA	2015-12-04	
		KR 20160014036	A	2016-02-05	
	Family Status:	Pub No.	State	Status	Expiry
		US 20140356956 A1	ALIVE	PENDING	2034-06-04
		US 9267135 B2	ALIVE	GRANTED	2034-06-04
	Probable Assignee:	PRESIDENT AND FELLOWS OF HARVARD COLLEGE			
	Sequence Locations:	Seq. ID Number	% Identity	Length	Location
		US20140356959-0001	100.00	1368	probable disclosure (not found by automated parsing)
		US20140356956-0001	100.00	1368	probable disclosure (not found by automated parsing)

Notes

Alignment:

```

Q:          1  MDKKYSIGLDIGTNSVGWAVITDEYKVPSSKKFKVLGNTDRHSIKKNLIGALLFDSGETAE  60
  | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | | |
S:          1  MDKKYSIGLDIGTNSVGWAVITDEYKVPSSKKFKVLGNTDRHSIKKNLIGALLFDSGETAE  60

```

Summarize at Patent or Sequence Level

- Always start with Tools | Identify Common Patent Family to create the Common Family column (this is a “magic” column)
- Replace contents of the Common Family column with the data you want to group by
Select column, copy
Select Common Family, paste
- Patent level: Patent Number
- Sequence level: Sequence ID (pub+seqidno)



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