

Patents & IP Sequences | Clinical Trials | Drug Pipelines

Building Patent Reports from Orbit.com and GenomeQuest

BizInt Smart Charts eLearning Webinar

27 May 2020

John Willmore, VP Product Development

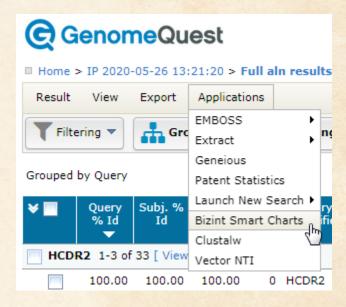
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Today's Agenda

- Creating reports from GenomeQuest
- Transferring publication numbers to Orbit
- Combining reports
- Identify Common Patent Family
- Reference Rows
- Creating a Summary Table of Sequence Hits
- Exports, including Summary Records

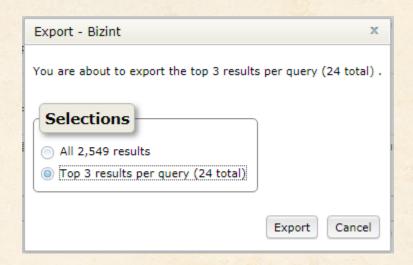
IP Sequences on GenomeQuest

- GQPAT
- GeneSeq
- CAS Biosequences (new in version 5.3.2)
- Use the BizInt application to export
- Discovery Browser export available (beta)



Improvements to GenomeQuest Exports

Filters and selections now carry over to exports



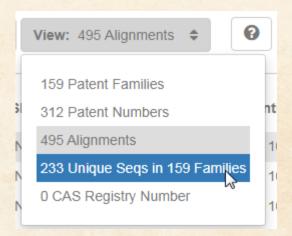


BizInt exports now available in Discovery Browser

- Great way to refine your results before export
- Filter to only claimed sequences

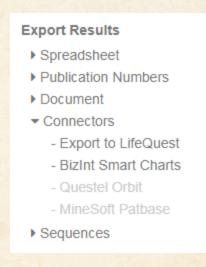
▼ Patent Sequence Location☑ Claim (495)

 Reduce results to unique family sequences



BizInt exports now available in Discovery Browser

 BizInt Smart Charts connector is available today in a beta version of Discovery Browser







Three Classes of Content in GQPAT

- Bibliographic data (title, family, inventors)
- Sequence data (sequence, organism, location)
- Query data (alignment, query ID, pct. Identity)

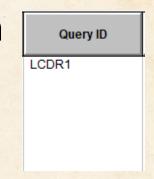
GQPAT Content in BizInt Smart Charts

Ebola .	Antibodies	(May	(2020)	

	Title	Title Query ID Patent Family Patent Ass		Patent Assignee	Inventor(s)	Alignment	Claimed Seg ID	Seq. ID Num		
	Title	Query ID	Patent	Kind	Date	Patent Assignee	ilivelitor(s)	Allylillielit	Claimed Seq ib	Seq. ID Wall
1	BROADLY NEUTRALIZING ANTIBODY TARGETING THE EBOLAVIRUS GLYCOPROTEIN INTERNAL FUSION LOOP	HCDR2	US20200040065 CN110087677 EP3525813 WO2018071345		20200206	INTEGRATED BIOTHERAPEUTICS, INC.; THE UNIVERSITY OF MARYLAND	AMAN MOHAMMAD JAVAD HOWELL KATIE A HOLTSBERG FREDERICK WAYNE ZHAO XUELIAN LI YUXING	Q: 1 GNIDNSASTNYNPSLKT 17	1-10	US2020004006
2	BROADLY NEUTRALIZING ANTIBODY TARGETING THE EBOLAVIRUS GLYCOPROTEIN INTERNAL FUSION LOOP	HCDR2	WO2018071345 CN110087677 EP3525813 US20200040065		20180419	INTEGRATED BIOTHERAPEUTICS; UNIVERSITY OF MARYLAND	AMAN, Mohammad Javad HOWELL, Katie A. HOLTSBERG, Frederick Wayne ZHAO, Xuelian	Q: 1 GNIDNSASTNYNPSLKT 17	1-15 17 20-27	WO2018071345
3	BROADLY NEUTRALIZING ANTIBODY TARGETING THE EBOLAVIRUS GLYCOPROTEIN INTERNAL FUSION LOOP	HCDR2	US20200040065 CN110087677 EP3525813 WO2018071345		20200206	INTEGRATED BIOTHERAPEUTICS, INC.; THE UNIVERSITY OF MARYLAND	AMAN MOHAN JAVAD HOWELL KATIE A HOLTSBERG FREDERICK WAYNE ZHAO XUELIAN LI YUXING	17 NPSLKT S: 50 GNIDNSASTNYNPSLKT	1-10	US2020004006
4	BROADLY NEUTRALIZING ANTIBODY TARGETING THE EBOLAVIRUS GLYCOPROTEIN INTERNAL FUSION LOOP	HCDR2	WO2018071345 CN110087677 EP3525813 US20200040065		20180419	INTEGRATED BIOTHERAPEUTICS; UNIVERSITY OF MARYLAND	AMAN, Mohammad Javad HOWELL, Katie A. HOLTSBERG, Frederick Wayne ZHAO, Xuelian LI, Yuxing	Q: 1 GNIDNSASTNYNPSLKT 17	1-15 17 20-27	WO2018071345
5	BROADLY NEUTRALIZING ANTIBODY TARGETING THE EBOLAVIRUS GLYCOPROTEIN INTERNAL FUSION LOOP	LCDR3	US20200040065 CN110087677 EP3525813 WO2018071345		20200206	INTEGRATED BIOTHERAPEUTICS, INC.; THE UNIVERSITY OF MARYLAND	AMAN MOHAMMAD JAVAD HOWELL KATIE A HOLTSBERG FREDERICK WAYNE	Q: 1 QQHNTLPLT 9 S: 1 QQHNTLPLT 9	1-10	US2020004006

Multi-query searches on GenomeQuest

• Query labels are available in the QueryID column e.g. LC, LCDR1/2/3...

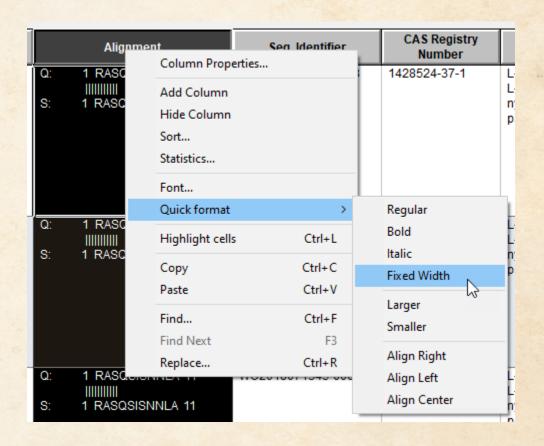


- If you run each query as a separate search, fill the Query ID in each chart by selecting the column and pasting in the new value.
- Then combine.

Formatting for Alignments

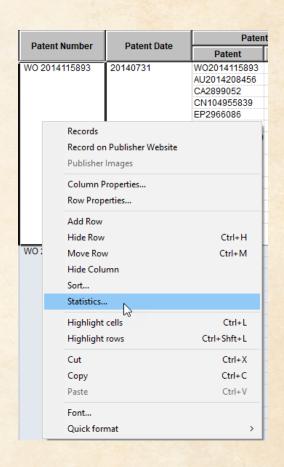
- Choose the Fixed Width option for Quick Format
- Don't just change the font for the column ... that looks ok in BizInt Smart Charts, but loses spaces when exported!

		Alignmen	ıt
Q:	1	RASQSISNNLA	11
s:	1	RASQSISNNLA	11



Transferring publication numbers to Orbit

- In a small set like this, collect all publication numbers
- Tools | Statistics creates an Excel sheet
- Copy the publication numbers
- An alternative that works with Orbit is to simply Copy the Patent Number column
- Paste the numbers into Orbit's "Number Search" panel



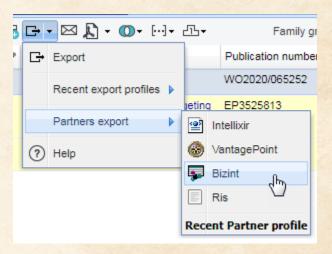
Searching Orbit Collections

- BizInt Smart Charts supports all three search collections on Orbit.com:
- Family (FAMPAT)
- Publication (FULLPAT)
- Full Text (definitions updated in 5.3.2)



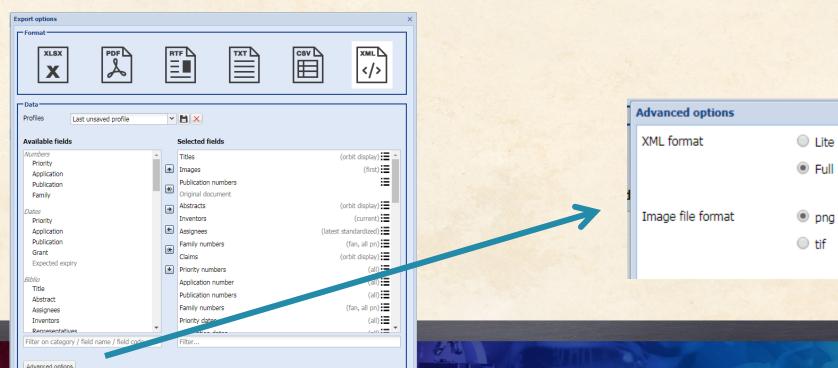
Exporting from Orbit

- There are two ways to export results from Orbit
- Easiest is the BizInt partner export
- Standard set of bibliographic and status data
- Independent Claims only



Exporting from Orbit - XML option

- You can select fields using Export and XML format
- Make sure you include family number (FAN)
- Allows you to choose full claims



Download Email

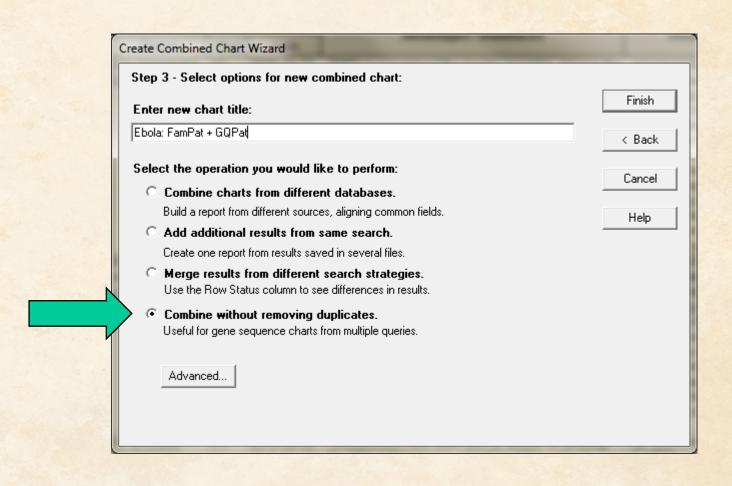
Orbit - export results, Family Status, Normalized Assignee

 Creating Reports from Databases and Hosts for instructions on each platform

Indonendent Claims	Family Status				Family Patent Assignee	
Independent Claims	Pub No.	State	Status	Expiry	Normalized	
(WO 202065252 A1 GB 201815629 D0	ALIVE DEAD	PENDING LAPSED	2022-03-25 2020-03-25	КҮМАВ	
An isolated antibody or antigen-binding fragment thereof comprising a binding domain that specifically binds to an orthologous epitope in the internal fusion loop of an ebolavirus glycoprotein, wherein the binding domain specifically binds to the epitope on two or more ebolavirus species or strains.	EP 3525813 A1 WO 201871345 A1 US 20200040065 A1 CN 110087677 A	ALIVE DEAD ALIVE ALIVE	PENDING LAPSED PENDING PENDING	2037-10-09 2020-04-11 2037-10-09 2037-10-09	INTEGRATED BIOTHERAEPUTICS UNIVERSITY OF MARYLAND	
(4. 5.5555)	EP 2760889 A1	DEAD	LAPSED	2018-10-09	TEVA PHARMACEUTICALS	

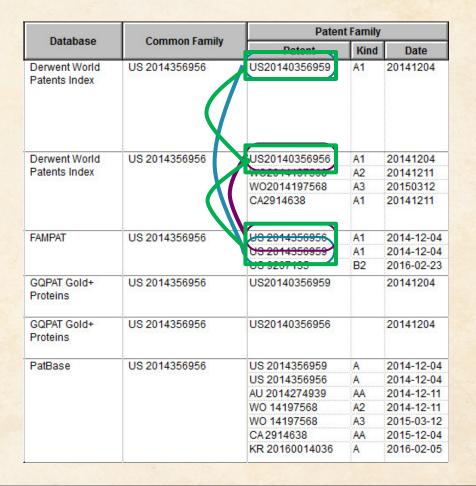
Combine Charts

- Both charts open
- File | Combine



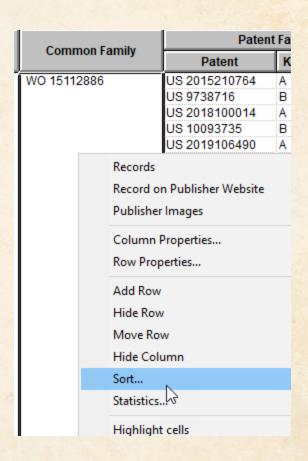
Match records by Tools | Identify Common Patent Families

- Links rows in the chart based on publication numbers in families
- Remember: Common Family is a sort key



Common Patent Family

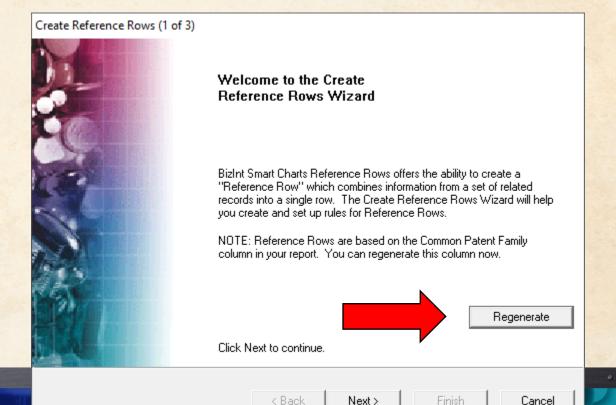
- Remember that Common Family is simply a sort key
- You can edit the assigned values
- You can paste another field into Common Family
- Patent Number group by publication
- Sequence ID group by sequence



Send to Reference Rows

- Save the combined chart first
- File | Send to Reference Rows
- On the first step of the wizard, do NOT select "Regenerate" if you have modified Common Family





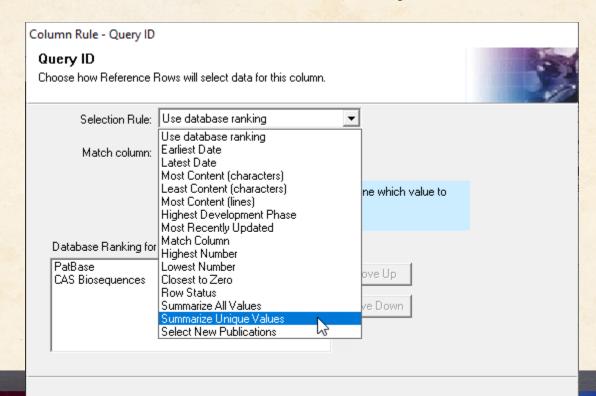
Standard behavior - fill in the blanks

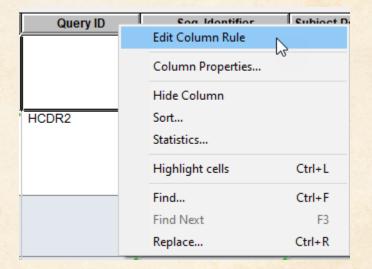
Check marks show representative values for each group

	4										
2	Broadly neutralizing antibody targeting the ebolavirus	EP 3525813	FAMPAT	INTEGRATED BIOTHERAEPUTICS	EP 3525813 A1 WO 201871345 A1	ALIVE DEAD	PENDING LAPSED	2037-10-09 			
.1	glycoprotein internal fusion loop			UNIVERSITY OF MARYLAND	US 20200040065 A1	ALIVE	PENDING	2037-10-09			
	1				CN 110087677 A	ALIVE	PENDING	2037-10-09			
2	BROADLY NEUTRALIZING ANTIBODY TARGETING THE	EP 3525813	GQPAT Gold+ Proteins						HCDR2 ✓	Q: 1	GNIDNSASTNYN
	EBOLAVIRUS GLYCOPROTEIN INTERNAL FUSION LOOP									S: 1	GNIDNSASTNYN
2	BROADLY NEUTRALIZING ANTIBODY TARGETING THE	EP 3525813	GQPAT Gold+ Proteins						HCDR2	Q: 1	GNIDNSASTNYN
.2	EBOLAVIRUS GLYCOPROTEIN INTERNAL FUSION LOOP									S: 1	GNIDNSASTNYN
2	Wide neutralizing antibodies targeted to internal fusion rings of	EP 3525813	GQPAT Platinum Proteins						HCDR2	Q: 1	GNIDNSASTNYN
.4	Ebola virus glycoproteins		Troteins							S: 1	GNIDNSASTNYN
2	BROADLY NEUTRALIZING ANTIBODY TARGETING THE	EP 3525813	GQPAT Gold+ Proteins						LCDR3	Q: 1	QQHNTLPLT 9
.?	EBOLAVIRUS GLYCOPROTEIN INTERNAL FUSION LOOP		, retemb							S: 1	QQHNTLPLT 9
2	BROADLY NEUTRALIZING ANTIBODY TARGETING THE	EP 3525813	GQPAT Gold+ Proteins						LCDR3	Q: 1	QQHNTLPLT 9
.0	EBOLAVIRUS GLYCOPROTEIN INTERNAL FUSION LOOP		1 1000113							S: 1	QQHNTLPLT 9
2	Wide neutralizing antibodies	EP 3525813	GQPAT Platinum						LCDR3	Q: 1	QQHNTLPLT 9
.7	targeted to internal fusion rings of Ebola virus glycoproteins		Proteins							S: 1	QQHNTLPLT 9

Show all Query IDs for a Family

- Edit Column Rule for Query ID
- Choose Summarize Unique Values





0K

Cancel

Summarize Unique Values

- Cell Glyph Changes
- Export or Statistics to see value

_					
C					
9					
)					
	Q:	1	RASQSISNNLA 11 💞	LCDR1 ##	WO2018071345-0006
			11111111111		* I
	S:	1	RASQSISNNLA 11		
	Q:	1	DPGFTIFGVVITSWSGLDS	HCDR3 ##	WO2018071345-0005
	19				
			111111111111111111111111111111111111111		
	S:	1	DPGFTIFGVVITSWSGLDS		
	19				
	Q:	1	DPGFTIFGVVITSWSGLDS	HCDR3 ##	WO2018071345-0001
	19				
			1111111111111111111111		
	S:	99	DPGFTIFGVVITSWSGLDS		
	117				
	Q:	1		HCDR2 ##	WO2018071345-0004
			1111111111111111111		
	S:	1	GNIDNSASTNYNPSLKT 17		
-					W00040074045 0004
	Q:	1		HCDR2 ##	WO2018071345-0001
	_		111111111111111111111111111111111111111		
	S:	50	GNIDNSASTNYNPSLKT 66		
-	٠.		COUNTED TO S	LODD3 ###	W02010071245 0000

LCDR1 HCDR3 HCDR2 LCDR3 HCDR1 LCDR2 LC-Ebola HC-Ebola

Create Subtable to Summarize Sequence Results

 But that solution only gives a list of Query IDs associated with the family - it doesn't tell you which sequence was returned with each query.

LCDR1 ##	WO2018071345-0006	100.00 🗸
HCDR3 ##	WO2018071345-0005	100.00
HCDR3 ##	WO2018071345-0001	14.84
HCDR2 ##	WO2018071345-0004	100.00
HCDR2 ##	WO2018071345-0001	13.28
	WO2018071345-0008	100.00
LCDR3 ##	WO2018071345-0002	8.49

Create Subtable to Summarize Sequence Results

 The answer is to create a table showing data from different sequence hits.

- Complete steps are available in a recipe at bizint.com/piugbio:
- Create subtable from columns
- Change rule to Summarize All Values
- Export

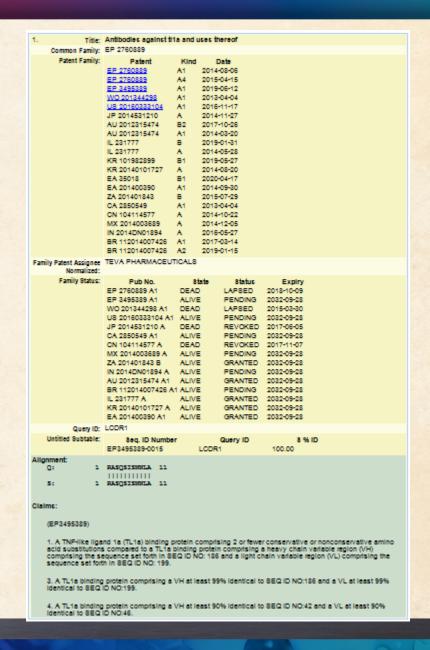
W 2018071345-0006	LCDR1	100.00	5.2
WO2018071345-0005	HCDR3	100.00	5.3
WO2018071345-0001	HCDR3	14.84	5.4
WO2018071345-0004	HCDR2	100.00	5.5
WO2018071345-0001	HCDR2	13.28	5.6
WO2018071345-0008	LCDR3	100.00	5.7
WO2018071345-0002	LCDR3	8.49	5.8
WO2018071345-0003	HCDR1	100.00	5.9
WO2018071345-0007	LCDR2	100.00	5.10
WO2018071345-0002	LC-Ebola	100.00	5.11
WO2018071345-0001	HC-Ebola	100.00	5.12

Key Tips of Summarized Subtable

- Rename columns before creating subtable you can't change them after the fact
- It is best to include a column which helps identify the data in each row (Sequence ID in the example)
- Fixed Width property of the Alignment column doesn't work in a subtable

Summary Records export

- A Word export containing content of the chart, full claims set, full alignment from selected records
- Yellow section contains columns of the chart
- Green section can contain one alignment, one full set of claims



Resources

- bizint.com/tips for links to key documentation
- BizInt Smart Charts for Patents Mini Guide
- bizint.com/piugbio for the recipe handout



THE JOURNEY CONTINUES...

BizInt Smart Charts

for Patents



Thank you!

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MARPAT Design Workshop

Wednesday, June 3 8:30 PT