



Paul:

That more or less say here's my person.
Here's the start and end of the exposure.

A non-related source value.

I have a dose table and then I have the NDC source code and this is important, so if we go back,

here you'll see that the the biggest box on the row standard table is the NDC, which is the national drug compound or something like that. I think it is that what the acronym stands for but,

Another just a set of terminology. I don't know if any of you on this call are more familiar with NDC than I am.

Especially the mimic table, although he have.

First, no worries, I'll push forward.

But in any case, we do have a non-standard representation.

Of those NDC codes in Cmpg and they do then have linkages to.

RX norm. So for instance, I can let the code up.

And one thing that is a bit weird with the way the data is represented here is the source code.

It's truncated or chopped off those the leading zeros of these terms. So normally NDC codes are supposed to be 11 digits.

With four padding of zeros, in this case I had to do that manually, just as an FYI.

I search this what I didn't find is have a my NDC code on Adhena.

And then I see that it has a single relationship, which is just. It's nonstandard to standard map into an RX norm code, and that then RX norm code has all of its awesome relationships that we talked about before.

And so at a high level, what our my code will do and I guess there it is when the NDC representation with all the other information and in this case it is inside, right?

That was the. Yeah. Yeah. Inside let go.

And when we're not gonna write about this too much for now. Where it was is depicted.

But we have then this kind of view on how much results it was at the source level.

Another thing to note, and maybe there, you and I can touch base on this later, but the quantity at least in the demo was the field represented quantity.

I feel like it's there seems to be an issue with that.

So we can maybe talk about offset because there is a lot of 0 quantities and I'm not.

I.

I'm not sure on the exact interpretation of that, but.

To mimic specific thing.

In any event, so you have this quantity column and I'll focus on the ones where the quantity is one.

But then in some cases these data are represented either directly and.

To some extent, as you're joining your source tables, whether it be from Epic or Cerner or anything else, you're going to end up with this type of a kind of a transient view where you have a subject ID, you have their admission, you have a start and.

As well as some drug known drug exposure.

With the way that that drug was administered, what the drug actually was and maybe it's an NDC or maybe you have it in RX norm or some other terminology.

With the description and so in our case, for mimic, all we need to do is just like what I did with Adhena. We can use that map to relationship.

To again, this is the relationship that's being shown. If I come back here, this nonstandard to standard map, that's a map to relationship. And so we're joining on this padded NDC code.

With our maps to relationship to either going on RX norm.

Equivalent to that NDC.

And I'll do that.

I've written the code to advance this time, so I'm not writing it for you on the fly, but.

What you'll see is exactly what I looked upon Adhena, right?

We have this now concept ID of 401. Whatever it is 66274.

That's all good.

We should hopefully see some other stuff here.

Yeah. So then we have some medication's preparations.

We have again, we have oxy codone and extemporaneous.

We will have homogeneous or very heterogeneous representation of I want to use this directly, which is currently how the logic is written the way we're mapping things is going to be mapped to different levels.

What I haven't done though necessarily and this is something that is worth checking when you do any sort of map maps to relationships into RX norm is to check the concept class because I can see already just from the results that these are all like the.

And when we have a branded drug, we have the brand of the drug included in this.

And in most cases I should say, maybe that's not always the case, but.

It's a fairly high level.

But I would recommend when you're anytime you're going into RX norm to take a look at your concept codes and what we can do carry concept class and I can add that on here which is.

Just directly beside my.

Just so you're.

You understand where your mapping ended up in the.

NDC branded drugs pretty much all the way down. And sometimes you have clinical drug form in case you don't have an appropriate match.

But it's a good to keep an eye on where you are in that, that brand drug class.

Kind of things.

OK, and what you see that also we're in mimic is that it is.

We will have homogeneous or very heterogeneous representation of I want to use this directly, which is currently how the logic is written the way we're mapping things is going to be mapped to different levels.

Depending on the drug in question, so it's not always going to be to a branded drug. It's gonna be sometime to a branded drug. Sometimes the clinical drug form, sometimes clinical drug thing like that. And depending on the case in the study and what you're trying to.

Do you want to try to standardize that, or at least bring everything to the same level.

That's a bit of a discussion to have internally.

We also should.

Provide convention for a minimum level for certain some point.

That will, like we will need effectively to support some of the pharmacokinetic, especially studies that we're looking at.

But in any in any event, definitely keep an eye on the concept class of your standard code when you're doing these mappings.

OK, now.

So that now we've taken our representation of drugs and mimic and we've taken that NDC code and we've combined them with.

With the CDM vocabulary relationships to get our RX norm code equivalent.

And now we can take that and use essentially this table that I'm showing. You can be used to populate the drug exposure table.

I mean, these things drug missing.

You have a unit that you run want to capture.

You would then.

Route that you would still wanna capture that were kind of skimming over here for the moment.

But I'll leave to you like you're you're on your way to having a drug exposure.

For drug use?

So let's say we say like you're on your way to having a drug exposure.

We've done this.

We made our mappings and we've populated drug exposure, drug use.

You don't.

You get for the essentially you can just by running.

The drug-exposure script.

So if you come down here.

And I'll send a link to this in the chat.

I didn't know I can.

So the script here you all haven't seen it.

You can run this pretty much out of the box.

You might have to tweak it slightly to update the schema.

You know what?

Actually, maybe now is a good time to show you this if I go.

So there's a package developed by Chinyee called SQL Bender and you actually have a SQL Bender developer where you can take code like this that has.

These like a parameters. So it's CDM schema that you can replace with your own. If you place this this script in this SQL Bender developer.

I mean actually just as a at with folks at Emory where they brought up so.

This is something that that could be used in this case. This will tell you.

Which parameters are included in this query?

In this case, it's only a single parameter which is your CDM schema.

Let's say it's a CDM.

And my temp evaluation schema. I'm going to put this temp.

And in my case I want to put it into datasets which is not on here.

Let's say we'll OK, but then that's a way next to do it in the workspace.

You can select that workspace and you end up with your query that has your.

Your CDM schema and you can copy and paste this into any sort of SQL execution.

So it's useful to know about if you didn't already know it.

It's also why I'm here if you're in the workspace, epic or otherwise, that we start to share between institutions. You can use this tool to connect it between.

Frames of SQL.

So doing this to know about.

I'll also post this in the chat.

I mean, who didn't know about it.

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OK.

Yeah. So thank you for that.
I guess the one the one topic that I could kind of point to here at the ends, although I must say I'm a little bit hesitant to do so because it's not been fully fleshed out yet.
Is this idea of capturing?
Drug infusions and what that could look like and maybe what I'll say to be for now, is that there is an ongoing discussion on the GitHub discussions about this.
With our recommendations.
And kind of more generally, what's being discussed in the Ordinary community?
I don't think yet that we have settled on an official convention for how we will capture drug infusions within Chorus.
Unlike there is an SOP or something published that I wasn't aware of. But I don't think that we have come to a conclusion on that at the moment, nor have I don't think Ordinary as a community has not decided on a single way to do this because I think there is a bit of complexity.
I don't know, Andrew, if you have anything
Thank you.

Williams, Andrew T 10:15
Say this is an opportunity for us to help to define what's a really exciting potential for capturing these kinds of data.
So you've mentioned pharmacokinetics studies in a few times.
This is not typically something you do in large observational data sets, and the ability to really examine questions about pharmacokinetics outside of a very controlled environment at all is a very novel and important thing, and to do so requires some stuff that hasn't been figured out, so for the easy to do kinds of things are.
Medication possession ratios over days for or months, and that kind of a thing.
And that's super valuable also.
But really understanding kind of what's going into somebody's body and all the other physiological responses to the drug being in someone at a very specific point in time is a new thing and really exciting thing.
So we I guess I'm coming after you ask, which is certainly true that there is an established conventions on his head and saying so. Really important thing we're all going to contribute to it is defining how this can be done at scale for the first time in a way that will be really impactful. And so it's an exciting thing I think some of the what we're just talking about with flow sheets is similar to you know drug pumps or other information where we have various information. There's.
There's a lot there.
One of the key elements that we haven't figured out yet is how to change the overall scale of time from a unit of a day to something that's within a day.
A day, and it's getting back to things like start and end times.
Either times of exposure or times plus amounts of exposure getting to very, you know, precise indications of of those things and so.
All the standardization work that has been that gets us pretty far, but doesn't get all the way to where we want to go.
So that's we should look at that as an exciting thing to to help you know, to figure out together. And we'll be getting back to you on that.

Hughes, Jared 10:20
Same, thank you.
And indeed, just one more plug to.
Use the data time fields like, it's something that we expect to be done within Chorus, it says here it is not required from cruds and it is officially from the not required. But within Hirus, especially within the context of ICU settings, we do need you to fill in these fields.
So again, I've mentioned it on other calls.
I will continue to mention it on future calls, but the date time is critical.
For for many of the analyses things that we want to do, like, downstream.
Unless but yeah, so I guess all that save out infusions is a place we have an opportunity to define the Convention or or the best practice for capturing this and the scale of time is I think.
Where we've given need to be to work, to transition from this typical day time into minutes and potentially seconds level of of detail of when those things are being administered and how long they're being administered for.
Assume.
OK.

So that's that's mine.
That's a very crude drug transformation also with some some dose information and again referring you to this nice vignette on different scenarios for calculating drug dose. One thing I didn't really go into is if I were to do this, is a production like it is in an ETL where I would to get a final drug or dose error table.
I would establish.
A kind of a scenario or a case based logic where I would first group my drugs into.
At least three cases.
Based on probably the combination of their drug form.
And the drug itself, or maybe some other relationships related to it.
So I would first kind of stratify my set of drugs into tablets or intratecs or.
Or.
Drugs.
And then based on that stratification use the drug strength table to create the dosage calculation.
So and like I said, start with the easy ones first.
I think tablets are pretty straightforward.
They typically have a given weight per tablet and you have a quantity of tablets, so you can calculate the amount given over time.
And then work your way up into more complex things where you have to deal with the denominator. You have drops.
I mean you have conversions.
This is another thing we didn't even get into, but if you have units.
In your drug exposure that are different than what appears in the drug strength table, you need to convert those units because the drug strength table of course is.
A representation of some set of multiples for a single unit and you need to make sure that your quantity refers to that unit.
So this is, as I would argue one of the most complex points of logic in an ETL is to do this and to do this correctly.
And I think it would be more than happy to take help on calls or or at least discuss this in more detail and the more we see cases you can bring.
I think the the the more valuable it will be because of course this mimic one is, is it OK to see and like we have some tablets but if you have challenging examples of source drugs that you don't know how to calculate a dose for.
Maybe infusions things and.
That would be very please. Please bring them forward and we can go through how how you might do that.
I think that would be a very nice.
Kind of question, answer topic or or future session that we could walk through some different examples of of those calculations.
Alright, that was a lot.
But hopefully this is a useful view on drugs.
And I don't think it's gonna be the last time we will. We will talk about them for sure.
But we will be able to then share this content with you all together with the SQS scripts.
I should note as well the the script for the mimic, is already available online and I can post a link to the chat for that as well.
In our repository within course, so I'll link that here, but I'll also upload this together with the values into the Google Drive.
After this call.
So thank you all for sticking with me.
And please reach out if you have any questions on drugs or really anything else related to what we talked about today.
So thanks.

Williams, Andrew T 10:17
Great job again, Jared.
Thanks for for doing this. You know for a second time without the epic stuff was really clear exposition of really complex and important topic. And thanks so much.
Yeah. Do you have a minute to stay on afterwards?

Hughes, Jared 10:30
Yes, I have you're OK with me joining like a minute or two late.
See, maybe Brian, Joe, is it alright? Alright then.

Brian Caw 10:41
Yeah, sure.

Hughes, Jared 10:43
Yeah, good to go.

Alvarez, Marta stopped transcription

