WEBVTT

NOTE duration: "00:12:25.6320000"

NOTE language:en-us

NOTE Confidence: 0.801072

00:00:03.590 --> 00:00:05.274 Hello I'm Florida carino,

NOTE Confidence: 0.801072

00:00:05.274 --> 00:00:08.320 I'm a professor at their child study

NOTE Confidence: 0.801072

 $00{:}00{:}08.320 \dashrightarrow 00{:}00{:}10.500$ center and Department of Neuroscience

NOTE Confidence: 0.801072

 $00{:}00{:}10.500 \dashrightarrow 00{:}00{:}14.316$ and today I am sharing with you latest

NOTE Confidence: 0.801072

 $00:00:14.316 \longrightarrow 00:00:17.298$ highlights of research from my laboratory.

NOTE Confidence: 0.801072

 $00:00:17.300 \longrightarrow 00:00:20.036$ We are focusing on two projects.

NOTE Confidence: 0.801072

 $00{:}00{:}20.040 \dashrightarrow 00{:}00{:}22.854$ One is an induced pluripotent stem

NOTE Confidence: 0.801072

 $00:00:22.854 \longrightarrow 00:00:25.421$ cells as models of developmental

NOTE Confidence: 0.801072

 $00{:}00{:}25.421 \dashrightarrow 00{:}00{:}29.579$ disorders and the 2nd is on somatic

NOTE Confidence: 0.801072

 $00:00:29.579 \longrightarrow 00:00:32.540$ genomic mosaicism in the human brain.

NOTE Confidence: 0.801072

 $00:00:32.540 \longrightarrow 00:00:36.488$ So the first part.

NOTE Confidence: 0.801072

 $00:00:36.490 \longrightarrow 00:00:38.830$ Is about induced proponents themselves.

NOTE Confidence: 0.801072

 $00:00:38.830 \longrightarrow 00:00:40.698$ You can see here.

NOTE Confidence: 0.801072

 $00{:}00{:}40.698 \dashrightarrow 00{:}00{:}43.510$ You probably know this, says R.

 $00:00:45.920 \longrightarrow 00:00:48.632$ Clearly Button says mean means immortal

NOTE Confidence: 0.80069906

 $00:00:48.632 \longrightarrow 00:00:51.859$ cell lines that are derived from a

NOTE Confidence: 0.80069906

00:00:51.859 --> 00:00:54.115 living person, typically from his

NOTE Confidence: 0.80069906

00:00:54.115 --> 00:00:56.390 small biopsy of fibroblast cells.

NOTE Confidence: 0.80069906

 $00{:}00{:}56.390 \dashrightarrow 00{:}00{:}59.694$ But it could be also other cells obtained

NOTE Confidence: 0.80069906

 $00:00:59.694 \longrightarrow 00:01:03.135$ from the adult human body and they are

NOTE Confidence: 0.80069906

 $00:01:03.135 \longrightarrow 00:01:05.965$ expanded in vitro and we differentiate

NOTE Confidence: 0.80069906

 $00{:}01{:}05.965 \dashrightarrow 00{:}01{:}09.127$ them in different type of neurons.

NOTE Confidence: 0.80069906

00:01:09.130 --> 00:01:11.662 So they undergo recapitulation

NOTE Confidence: 0.80069906

00:01:11.662 --> 00:01:14.194 of the neuronal development.

NOTE Confidence: 0.80069906

 $00:01:14.200 \longrightarrow 00:01:16.310$ Over several days and they

NOTE Confidence: 0.80069906

 $00:01:16.310 \longrightarrow 00:01:18.940$ can be used in various ways.

NOTE Confidence: 0.80069906

 $00{:}01{:}18.940 \dashrightarrow 00{:}01{:}21.100$ We develop them into organoids.

NOTE Confidence: 0.80069906

 $00:01:21.100 \longrightarrow 00:01:23.000$ I'll show you later,

NOTE Confidence: 0.80069906

 $00:01:23.000 \longrightarrow 00:01:25.850$ but typically they can be used

 $00:01:25.945 \longrightarrow 00:01:28.457$ in screening and discovery.

NOTE Confidence: 0.80069906

00:01:28.460 --> 00:01:29.744 Of, for example,

NOTE Confidence: 0.80069906

00:01:29.744 --> 00:01:32.312 genes that are important in development

NOTE Confidence: 0.80069906

 $00:01:32.312 \longrightarrow 00:01:35.000$ or disease or for drug screening,

NOTE Confidence: 0.80069906

 $00:01:35.000 \longrightarrow 00:01:38.608$ or they can be used as models of

NOTE Confidence: 0.80069906

00:01:38.608 --> 00:01:41.223 human development in both normal

NOTE Confidence: 0.80069906

 $00:01:41.223 \longrightarrow 00:01:43.928$ development and these orders.

NOTE Confidence: 0.80069906

00:01:43.930 --> 00:01:47.434 And various sizes have been applied to them,

NOTE Confidence: 0.80069906

 $00{:}01{:}47.440 \dashrightarrow 00{:}01{:}51.016$ so we started this project about 10 years

NOTE Confidence: 0.80069906

00:01:51.016 --> 00:01:54.264 ago when we started recruiting first

NOTE Confidence: 0.80069906

 $00:01:54.264 \longrightarrow 00:01:57.648$ patients of the Child Study Center.

NOTE Confidence: 0.80069906

00:01:57.650 --> 00:02:00.520 With various new psychiatric diseases,

NOTE Confidence: 0.80069906

 $00:02:00.520 \longrightarrow 00:02:03.390$ and since then, we've acquired,

NOTE Confidence: 0.80069906

 $00:02:03.390 \longrightarrow 00:02:05.626$ we've developed about 600.

NOTE Confidence: 0.80069906

00:02:05.626 --> 00:02:10.280 I PS lines from more than 100 people,

NOTE Confidence: 0.80069906

00:02:10.280 --> 00:02:13.150 including both patients with autism,

00:02:13.150 --> 00:02:14.452 Tourette syndrome,

NOTE Confidence: 0.80069906

 $00{:}02{:}14.452 \dashrightarrow 00{:}02{:}17.056$ and other developmental disorders

NOTE Confidence: 0.80069906

 $00:02:17.056 \longrightarrow 00:02:18.358$ and controls.

NOTE Confidence: 0.80069906

 $00:02:18.360 \longrightarrow 00:02:21.426$ And so we can grow these organoids

NOTE Confidence: 0.80069906

 $00:02:21.426 \longrightarrow 00:02:24.019$ in vitro over several days,

NOTE Confidence: 0.80069906

 $00:02:24.020 \longrightarrow 00:02:27.100$ and you can see the increase in

NOTE Confidence: 0.80069906

 $00:02:27.100 \longrightarrow 00:02:30.627$ size we grow them by the hundreds.

NOTE Confidence: 0.80069906

 $00:02:30.630 \longrightarrow 00:02:33.682$ We have a highly efficient protocol for

NOTE Confidence: 0.80069906

 $00:02:33.682 \longrightarrow 00:02:36.300$ developing them into these structures.

NOTE Confidence: 0.80069906

 $00{:}02{:}36.300 \dashrightarrow 00{:}02{:}39.492$ You can see here if you cross section

NOTE Confidence: 0.80069906

00:02:39.492 --> 00:02:42.898 them and stained with various markers,

NOTE Confidence: 0.80069906

 $00:02:42.900 \longrightarrow 00:02:45.260$ you identify substructure within them.

NOTE Confidence: 0.80069906

 $00:02:45.260 \longrightarrow 00:02:47.930$ These are new epithelial progenitor

NOTE Confidence: 0.80069906

 $00:02:47.930 \longrightarrow 00:02:51.389$ cells that are staying in red for.

NOTE Confidence: 0.80069906

00:02:51.390 --> 00:02:53.885 Assess undergoing cell division and

 $00:02:53.885 \longrightarrow 00:02:57.123$ in green for a neuronal progenitor

NOTE Confidence: 0.80069906

 $00:02:57.123 \longrightarrow 00:03:00.368$ marker which is expressing expressing

NOTE Confidence: 0.80069906

 $00:03:00.368 \longrightarrow 00:03:02.315$ the cerebral cortex.

NOTE Confidence: 0.80069906

 $00:03:02.320 \longrightarrow 00:03:05.148$ And if you going more higher magnification

NOTE Confidence: 0.80069906

 $00:03:05.148 \longrightarrow 00:03:08.269$ in in in one of these organized,

NOTE Confidence: 0.80069906

 $00{:}03{:}08.270 \dashrightarrow 00{:}03{:}10.574$ you can see that they express

NOTE Confidence: 0.80069906

 $00{:}03{:}10.574 \dashrightarrow 00{:}03{:}12.678$ various cell types that are

NOTE Confidence: 0.80069906

00:03:12.678 --> 00:03:14.650 proper for normal development.

NOTE Confidence: 0.80069906

 $00{:}03{:}14.650 \dashrightarrow 00{:}03{:}18.038$ Normal human development in red you see

NOTE Confidence: 0.80069906

 $00:03:18.038 \longrightarrow 00:03:19.974$ ventricular zone progenitors thankful

NOTE Confidence: 0.80069906

 $00:03:19.974 \longrightarrow 00:03:22.816$ pack six and cortical layer one neuron

NOTE Confidence: 0.80069906

 $00:03:22.816 \longrightarrow 00:03:25.267$ stain for a gene called TVR one.

NOTE Confidence: 0.80069906

 $00:03:25.270 \longrightarrow 00:03:27.400$ They are positive for Foxy,

NOTE Confidence: 0.80069906

 $00:03:27.400 \longrightarrow 00:03:31.927$ One which is expressed in the hole for brain.

NOTE Confidence: 0.80069906

 $00:03:31.930 \longrightarrow 00:03:34.996$ Here you see a marker City 2,

NOTE Confidence: 0.80069906

 $00:03:35.000 \longrightarrow 00:03:38.600$ Four layer 5 and here down here in red

 $00{:}03{:}38.600 \dashrightarrow 00{:}03{:}42.434$ and marker for layer 23 neurons in red.

NOTE Confidence: 0.80069906

 $00:03:42.440 \longrightarrow 00:03:45.068$ So they like capitulate fairly faithfully.

NOTE Confidence: 0.80069906

00:03:45.070 --> 00:03:48.136 Early stages of human in this case.

NOTE Confidence: 0.80069906

00:03:48.140 --> 00:03:49.448 Human cortical development.

NOTE Confidence: 0.7614749

 $00:03:52.310 \longrightarrow 00:03:55.565$ And they can be stained with viruses.

NOTE Confidence: 0.7614749

00:03:55.570 --> 00:03:59.298 And then in this way you can visualize

NOTE Confidence: 0.7614749

00:03:59.298 --> 00:04:02.179 their morphology in finer detail and

NOTE Confidence: 0.7614749

00:04:02.179 --> 00:04:04.975 even down to showing early synaptic

NOTE Confidence: 0.7614749

 $00:04:05.059 \longrightarrow 00:04:08.202$ spines and we we can do electrical

NOTE Confidence: 0.7614749

 $00{:}04{:}08.202 \dashrightarrow 00{:}04{:}11.137$ recording on these cells by Patch

NOTE Confidence: 0.7614749

00:04:11.137 --> 00:04:14.311 clamp and they actually have synaptic

NOTE Confidence: 0.7614749

 $00:04:14.311 \longrightarrow 00:04:17.189$ currents develop synaptic currents.

NOTE Confidence: 0.7614749

 $00{:}04{:}17.190 \dashrightarrow 00{:}04{:}19.525$ Overtime we've used them for

NOTE Confidence: 0.7614749

 $00{:}04{:}19.525 \dashrightarrow 00{:}04{:}20.926$ studying various disorders.

NOTE Confidence: 0.7614749

 $00:04:20.930 \longrightarrow 00:04:23.722$ This is a paper we published in 2015

00:04:23.722 --> 00:04:26.164 on Autism Spectrum Disorder where

NOTE Confidence: 0.7614749

 $00:04:26.164 \longrightarrow 00:04:28.934$ we identified an imbalance between

NOTE Confidence: 0.7614749

00:04:28.934 --> 00:04:31.298 excitatory and inhibitory early

NOTE Confidence: 0.7614749

 $00:04:31.298 \longrightarrow 00:04:34.228$ developing neurons in these patients.

NOTE Confidence: 0.7614749

 $00:04:34.230 \longrightarrow 00:04:37.242$ And now we're in the middle

NOTE Confidence: 0.7614749

 $00:04:37.242 \longrightarrow 00:04:39.250$ of an ongoing study.

NOTE Confidence: 0.7614749

 $00{:}04{:}39.250 \dashrightarrow 00{:}04{:}41.760$ Larger study of ASD families,

NOTE Confidence: 0.7614749

 $00:04:41.760 \longrightarrow 00:04:44.844$ which comprises eleven families in which

NOTE Confidence: 0.7614749

00:04:44.844 --> 00:04:48.790 we have one problem and one control pair,

NOTE Confidence: 0.7614749

 $00:04:48.790 \longrightarrow 00:04:50.850$ typically an effective father,

NOTE Confidence: 0.7614749

 $00:04:50.850 \longrightarrow 00:04:54.310$ and they are grouped into microcephalic ASD,

NOTE Confidence: 0.7614749

 $00:04:54.310 \longrightarrow 00:04:57.316$ meaning people that have large brains.

NOTE Confidence: 0.7614749

00:04:57.320 --> 00:04:59.830 An normal cephalic ASD individuals,

NOTE Confidence: 0.7614749

 $00:04:59.830 \longrightarrow 00:05:03.344$ an excitingly we find differences among them.

NOTE Confidence: 0.7614749

 $00:05:03.350 \longrightarrow 00:05:04.637$ Here, you see.

NOTE Confidence: 0.7614749

00:05:04.637 --> 00:05:07.211 Then see in mapping of single

00:05:07.211 --> 00:05:09.478 cell phenotypes by irony,

NOTE Confidence: 0.7614749

 $00{:}05{:}09.480 \dashrightarrow 00{:}05{:}11.930$ single cell RNA sequencing in

NOTE Confidence: 0.7614749

 $00:05:11.930 \longrightarrow 00:05:14.865$ these families in the whole data

NOTE Confidence: 0.7614749

 $00:05:14.865 \longrightarrow 00:05:17.644$ set we can see that for example,

NOTE Confidence: 0.7614749

 $00{:}05{:}17.650 \dashrightarrow 00{:}05{:}20.590$ in patients with microcephaly we have an

NOTE Confidence: 0.7614749

 $00:05:20.590 \longrightarrow 00:05:23.548$ imbalance in the distribution of sales.

NOTE Confidence: 0.7614749

 $00:05:23.550 \longrightarrow 00:05:26.358$ So this is up here in in Blue Excel

NOTE Confidence: 0.7614749

 $00:05:26.358 \longrightarrow 00:05:29.582$ Group that we identify as deep cortical

NOTE Confidence: 0.7614749

 $00{:}05{:}29.582 \rightarrow 00{:}05{:}32.236$ plate excitatory neurons and in

NOTE Confidence: 0.7614749

 $00:05:32.236 \longrightarrow 00:05:34.976$ macrocephalic individuals patients versus.

NOTE Confidence: 0.7614749

 $00:05:34.980 \longrightarrow 00:05:38.085$ Others you see the day there is an increase

NOTE Confidence: 0.7614749

00:05:38.085 --> 00:05:40.908 in a subgroup of excitatory neuron,

NOTE Confidence: 0.7614749

 $00:05:40.910 \longrightarrow 00:05:42.382$ shown here in red,

NOTE Confidence: 0.7614749

 $00:05:42.382 \longrightarrow 00:05:44.590$ and a decrease in another subtype

NOTE Confidence: 0.7614749

 $00:05:44.670 \longrightarrow 00:05:47.596$ of excitatory neuron here and also a

 $00:05:47.596 \longrightarrow 00:05:49.988$ decrease of inhibitory neuron as well.

NOTE Confidence: 0.7614749

 $00:05:49.990 \longrightarrow 00:05:52.608$ So so at higher and higher the

NOTE Confidence: 0.7614749

 $00:05:52.608 \longrightarrow 00:05:54.682$ different type of resolution when

NOTE Confidence: 0.7614749

 $00:05:54.682 \longrightarrow 00:05:57.298$ we look at gene expression with

NOTE Confidence: 0.7614749

 $00:05:57.298 \longrightarrow 00:06:00.044$ this sub within each of these sub

NOTE Confidence: 0.7614749

 $00:06:00.044 \longrightarrow 00:06:02.702$ group of cells we can also identify

NOTE Confidence: 0.7614749

 $00:06:02.702 \longrightarrow 00:06:05.288$ certain imbalances you see here they.

NOTE Confidence: 0.7614749

00:06:05.290 --> 00:06:06.646 Differential gene expression

NOTE Confidence: 0.7614749

 $00:06:06.646 \longrightarrow 00:06:08.454$ in two cellular subproof,

NOTE Confidence: 0.7614749

 $00:06:08.460 \longrightarrow 00:06:10.675$ this the deep cortical plate

NOTE Confidence: 0.7614749

00:06:10.675 --> 00:06:12.890 excitatory neuron have an increase

NOTE Confidence: 0.7614749

 $00:06:12.968 \longrightarrow 00:06:15.824$ in markers for jeans that are typical

NOTE Confidence: 0.7614749

00:06:15.824 --> 00:06:17.970 of excitatory neuron development,

NOTE Confidence: 0.7614749

 $00:06:17.970 \longrightarrow 00:06:19.329$ such as emx.

NOTE Confidence: 0.7614749

 $00:06:19.329 \longrightarrow 00:06:21.594$ One an in the other,

NOTE Confidence: 0.7614749

 $00{:}06{:}21.600 \dashrightarrow 00{:}06{:}24.141$ in another sub group of cells you

 $00:06:24.141 \longrightarrow 00:06:26.943$ have a decrease in jeans that

NOTE Confidence: 0.7614749

 $00:06:26.943 \longrightarrow 00:06:29.119$ are characteristic of inhibitory

NOTE Confidence: 0.7614749

 $00:06:29.119 \longrightarrow 00:06:30.207$ neuron development,

NOTE Confidence: 0.7614749

 $00:06:30.210 \longrightarrow 00:06:32.814$ suggesting again that there is an

NOTE Confidence: 0.7614749

 $00:06:32.814 \longrightarrow 00:06:35.320$ imbalance between excited or inhibited.

NOTE Confidence: 0.7614749

00:06:35.320 --> 00:06:37.894 Keep inhibitory neurons in in ASD

NOTE Confidence: 0.7614749

00:06:37.894 --> 00:06:41.214 and even more exciting we find that

NOTE Confidence: 0.7614749

 $00:06:41.214 \longrightarrow 00:06:43.238$ normal cephalic and microcephalic

NOTE Confidence: 0.7614749

 $00:06:43.238 \longrightarrow 00:06:45.529$ individuals are not the same,

NOTE Confidence: 0.7614749

 $00:06:45.530 \longrightarrow 00:06:48.308$ suggesting that using IP's season organoid.

NOTE Confidence: 0.7614749

 $00{:}06{:}48.310 \longrightarrow 00{:}06{:}51.292$ Perhaps we can identify finer differences

NOTE Confidence: 0.7614749

 $00:06:51.292 \longrightarrow 00:06:54.250$ between group of patients that can

NOTE Confidence: 0.7614749

 $00:06:54.250 \longrightarrow 00:06:56.920$ be useful for clinical phenotype and

NOTE Confidence: 0.7614749

 $00:06:56.920 \longrightarrow 00:06:59.450$ drug screening and things like that.

NOTE Confidence: 0.7614749

 $00:06:59.450 \longrightarrow 00:07:02.698$ We've also done studies in Tourette syndrome.

 $00:07:02.700 \longrightarrow 00:07:05.430$ This is my graduate student.

NOTE Confidence: 0.7614749

 $00:07:05.430 \longrightarrow 00:07:06.258$ Johnny Brady.

NOTE Confidence: 0.7614749

 $00{:}07{:}06.258 \dashrightarrow 00{:}07{:}08.328$ She's spearheaded a project where

NOTE Confidence: 0.7614749

 $00:07:08.328 \longrightarrow 00:07:10.621$ she developed basal ganglia organoid

NOTE Confidence: 0.7614749

 $00:07:10.621 \longrightarrow 00:07:12.597$ rather than cortical organoids.

NOTE Confidence: 0.7614749

 $00:07:12.600 \longrightarrow 00:07:15.192$ There they develop many neurons that

NOTE Confidence: 0.7614749

 $00:07:15.192 \longrightarrow 00:07:17.970$ are characteristic of the basal ganglia,

NOTE Confidence: 0.7614749

 $00:07:17.970 \longrightarrow 00:07:20.840$ and she asked the question of whether

NOTE Confidence: 0.7614749

 $00{:}07{:}20.840 \dashrightarrow 00{:}07{:}23.067$ this development was affected in

NOTE Confidence: 0.7614749

 $00:07:23.067 \longrightarrow 00:07:25.367$ Tourette syndrome because in another

NOTE Confidence: 0.7614749

00:07:25.367 --> 00:07:28.727 earlier study on adult brain with Tourettes,

NOTE Confidence: 0.7614749

 $00:07:28.730 \longrightarrow 00:07:32.174$ we found a decrease in certain types

NOTE Confidence: 0.7614749

 $00{:}07{:}32.174 \dashrightarrow 00{:}07{:}35.348$ of interneurons in the basal ganglia.

NOTE Confidence: 0.7614749

 $00:07:35.350 \longrightarrow 00:07:37.639$ In so she asked the question whether

NOTE Confidence: 0.7614749

 $00:07:37.639 \longrightarrow 00:07:39.544$ this degrees was a developmental

NOTE Confidence: 0.7614749

 $00:07:39.544 \longrightarrow 00:07:41.699$ type decrease by growing organized

 $00:07:41.699 \longrightarrow 00:07:42.992$ from these patients

NOTE Confidence: 0.8114302

00:07:43.062 --> 00:07:45.526 and basically making a Long story short,

NOTE Confidence: 0.8114302

 $00:07:45.530 \longrightarrow 00:07:47.275$ she developed basal ganglia organoid

NOTE Confidence: 0.8114302

 $00:07:47.275 \longrightarrow 00:07:49.725$ down here and found that indeed there

NOTE Confidence: 0.8114302

 $00{:}07{:}49.725 \dashrightarrow 00{:}07{:}51.651$ is an early imbalance in certain

NOTE Confidence: 0.8114302

00:07:51.651 --> 00:07:53.838 genes that are characteristic of

NOTE Confidence: 0.8114302

00:07:53.838 --> 00:07:55.326 inhibitory neuron development.

NOTE Confidence: 0.8114302

 $00:07:55.330 \longrightarrow 00:07:58.260$ You can see here NCX 2.1 is one of the

NOTE Confidence: 0.8114302

00:07:58.339 --> 00:08:01.171 earliest jeans that develop in the

NOTE Confidence: 0.8114302

00:08:01.171 --> 00:08:04.000 basal ganglia, and as you can see,

NOTE Confidence: 0.8114302

 $00:08:04.000 \longrightarrow 00:08:05.970$ while is very prevalent in.

NOTE Confidence: 0.8114302

 $00{:}08{:}05.970 \dashrightarrow 00{:}08{:}08.430$ Basal ganglia from control is much

NOTE Confidence: 0.8114302

 $00{:}08{:}08.430 \dashrightarrow 00{:}08{:}10.530$ decrease in basal ganglia organoid

NOTE Confidence: 0.8114302

 $00{:}08{:}10.530 \dashrightarrow 00{:}08{:}12.828$ from patients with threats and you

NOTE Confidence: 0.8114302

 $00:08:12.828 \longrightarrow 00:08:15.474$ can see this quantified here on the

00:08:15.474 --> 00:08:18.280 right where you see a summary of five

NOTE Confidence: 0.8114302

 $00{:}08{:}18.280 \dashrightarrow 00{:}08{:}20.290$ patient with red and tank controls

NOTE Confidence: 0.8114302

 $00:08:20.290 \longrightarrow 00:08:22.269$ with highly significant degrees.

NOTE Confidence: 0.8114302

 $00:08:22.270 \longrightarrow 00:08:24.825$ Foreign players 2.1 which is in the

NOTE Confidence: 0.8114302

 $00:08:24.825 \longrightarrow 00:08:27.309$ middle and Lonnie Kalmenson also DLX.

NOTE Confidence: 0.8114302

00:08:27.310 --> 00:08:29.445 Another change which is expressed

NOTE Confidence: 0.8114302

 $00{:}08{:}29.445 \dashrightarrow 00{:}08{:}31.580$ throughout the basal ganglia and

NOTE Confidence: 0.8114302

 $00{:}08{:}31.650 \dashrightarrow 00{:}08{:}33.744$ this is also imbalance is also

NOTE Confidence: 0.8114302

 $00{:}08{:}33.744 \dashrightarrow 00{:}08{:}35.967$ evident in the preoptic area where

NOTE Confidence: 0.8114302

00:08:35.967 --> 00:08:37.475 again you see decreases.

NOTE Confidence: 0.8114302

00:08:37.480 --> 00:08:39.400 In inhibitory interneurons,

NOTE Confidence: 0.8114302

 $00:08:39.400 \longrightarrow 00:08:41.320$ an cholinergic interneuron,

NOTE Confidence: 0.8114302

00:08:41.320 --> 00:08:43.330 impatience versus control.

NOTE Confidence: 0.8114302

 $00:08:43.330 \longrightarrow 00:08:48.020$ But this is not evident in the

NOTE Confidence: 0.8114302

 $00:08:48.142 \longrightarrow 00:08:50.728$ in cortical organoids.

NOTE Confidence: 0.8114302

 $00:08:50.730 \longrightarrow 00:08:53.634$ So moving on a second project I was

00:08:53.634 --> 00:08:57.229 going to talk to you about is about

NOTE Confidence: 0.8114302

00:08:57.229 --> 00:09:00.528 semantic mosaicism and this is a

NOTE Confidence: 0.8114302

 $00:09:00.528 \longrightarrow 00:09:03.488$ phenomenon that is attracted recently.

NOTE Confidence: 0.8114302

 $00{:}09{:}03.490 \dashrightarrow 00{:}09{:}06.450$ A lot of attention because.

NOTE Confidence: 0.7337406

 $00:09:08.700 \longrightarrow 00:09:11.855$ Deals with mutations that are

NOTE Confidence: 0.7337406

 $00:09:11.855 \longrightarrow 00:09:15.720$ developed in the body in each.

NOTE Confidence: 0.7337406

00:09:15.720 --> 00:09:17.905 Organism basically from the time

NOTE Confidence: 0.7337406

00:09:17.905 --> 00:09:19.653 of fertilization on throughout

NOTE Confidence: 0.7337406

 $00:09:19.653 \longrightarrow 00:09:21.467$ the life of that person.

NOTE Confidence: 0.7337406

 $00:09:21.470 \longrightarrow 00:09:24.144$ And here you see that mutations can

NOTE Confidence: 0.7337406

00:09:24.144 --> 00:09:26.728 occur at anytime and the earlier

NOTE Confidence: 0.7337406

 $00:09:26.728 \longrightarrow 00:09:28.939$ they developed, the more sales.

NOTE Confidence: 0.7337406

 $00:09:28.939 \longrightarrow 00:09:31.104$ Of course they involve typically

NOTE Confidence: 0.7337406

 $00:09:31.104 \longrightarrow 00:09:33.389$ however they occur at any stage.

NOTE Confidence: 0.7337406

 $00:09:33.390 \longrightarrow 00:09:35.856$ In the later they do develop.

 $00:09:35.860 \longrightarrow 00:09:37.915$ The smaller the part of

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 $00:09:37.915 \longrightarrow 00:09:39.970$ the body that harbors them,

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 $00:09:39.970 \longrightarrow 00:09:42.436$ and they're very difficult to detect.

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 $00:09:42.440 \longrightarrow 00:09:44.180$ As you can imagine.

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 $00:09:44.180 \longrightarrow 00:09:46.790$ So you have to develop particular

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 $00{:}09{:}46.880 \dashrightarrow 00{:}09{:}49.834$ protocols in order to Geno type the.

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 $00:09:49.840 \longrightarrow 00:09:52.458$ This is of high resolution in order

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00:09:52.458 --> 00:09:54.406 to identify and characterize them

NOTE Confidence: 0.7337406

 $00{:}09{:}54.406 \dashrightarrow 00{:}09{:}57.398$ an in the past three years ago we

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 $00:09:57.477 \longrightarrow 00:10:00.117$ developed we developed a method for.

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 $00{:}10{:}00.120 \dashrightarrow 00{:}10{:}02.704$ Assessing this mutation and

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 $00:10:02.704 \longrightarrow 00:10:05.934$ we use them to reconstruct.

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 $00{:}10{:}05.940 \dashrightarrow 00{:}10{:}07.748$ Reconstruct the cellular mutation

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 $00:10:07.748 \longrightarrow 00:10:10.460$ and history of three individuals in

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 $00:10:10.526 \longrightarrow 00:10:13.302$ the reason you can do that is because

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 $00:10:13.302 \longrightarrow 00:10:15.438$ these mutations are actually markers,

00:10:15.440 --> 00:10:17.500 indelible marker of every cell,

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 $00:10:17.500 \longrightarrow 00:10:19.570$ division in the human body,

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 $00:10:19.570 \longrightarrow 00:10:20.704$ and more recently,

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 $00:10:20.704 \longrightarrow 00:10:23.350$ in an unpublished study we found a

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 $00:10:23.424 \longrightarrow 00:10:26.088$ way to actually map this mutation

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 $00:10:26.088 \longrightarrow 00:10:27.420$ in living individuals.

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 $00:10:27.420 \longrightarrow 00:10:30.327$ The way we do that is we take six

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00:10:30.327 --> 00:10:32.942 small skin biopsy biopsies from a

NOTE Confidence: 0.7337406

 $00:10:32.942 \longrightarrow 00:10:36.140$ person an we develop this fibroblast.

NOTE Confidence: 0.7337406

00:10:36.140 --> 00:10:37.950 Into I PS lines an.

NOTE Confidence: 0.7337406

 $00:10:37.950 \longrightarrow 00:10:40.122$ We genotyped each line and compare

NOTE Confidence: 0.7337406

 $00:10:40.122 \longrightarrow 00:10:41.570$ the genomes of these.

NOTE Confidence: 0.7337406

00:10:41.570 --> 00:10:43.874 I PS lines each of them is a

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 $00:10:43.874 \longrightarrow 00:10:45.954$ descendant of a single cell and

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 $00:10:45.954 \longrightarrow 00:10:47.739$ so any difference between them

 $00:10:47.739 \longrightarrow 00:10:50.298$ are clearly due to mutations that

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 $00{:}10{:}50.298 \dashrightarrow 00{:}10{:}52.428$ developed during the lifetime of

NOTE Confidence: 0.7337406

00:10:52.430 --> 00:10:54.656 that person and then we genotyped

NOTE Confidence: 0.7337406

00:10:54.656 --> 00:10:57.138 this mutation found in Ipas in blood,

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 $00:10:57.140 \longrightarrow 00:10:57.471$ saliva,

NOTE Confidence: 0.7337406

 $00:10:57.471 \longrightarrow 00:10:59.788$ and urine and that is enough to

NOTE Confidence: 0.7337406

 $00{:}10{:}59.788 \dashrightarrow 00{:}11{:}01.322$ reconstruct the ancestry tree

NOTE Confidence: 0.7337406

 $00:11:01.322 \longrightarrow 00:11:02.930$ of that particular person.

NOTE Confidence: 0.7337406

00:11:02.930 --> 00:11:05.730 And you can see an example here in

NOTE Confidence: 0.7337406

00:11:05.730 --> 00:11:08.350 a patient with Tourette syndrome.

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 $00{:}11{:}08.350 \dashrightarrow 00{:}11{:}11.062$ Where we could map the early

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 $00:11:11.062 \longrightarrow 00:11:12.870$ lineages of that person,

NOTE Confidence: 0.7337406

 $00:11:12.870 \longrightarrow 00:11:16.272$ starting from the very first cell division

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 $00:11:16.272 \longrightarrow 00:11:19.800$ up to about the 5th cell division.

NOTE Confidence: 0.7337406

00:11:19.800 --> 00:11:21.585 And one remarkable finding of

NOTE Confidence: 0.7337406

00:11:21.585 --> 00:11:23.882 this mapping is that you often

 $00:11:23.882 \longrightarrow 00:11:26.504$ find that there is a dominant

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00:11:26.504 --> 00:11:27.815 l'imaginaire recessive leaner,

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 $00:11:27.820 \longrightarrow 00:11:30.476$ and by that I mean one lineages that

NOTE Confidence: 0.7337406

 $00:11:30.476 \longrightarrow 00:11:32.821$ is over represented in the tissue

NOTE Confidence: 0.7337406

 $00:11:32.821 \longrightarrow 00:11:35.793$ in the body of that persons versus

NOTE Confidence: 0.7337406

 $00:11:35.793 \longrightarrow 00:11:38.248$ one that is less representative.

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00:11:38.250 --> 00:11:40.777 So this is very short but just

NOTE Confidence: 0.7337406

 $00{:}11{:}40.777 \dashrightarrow 00{:}11{:}43.550$ wanted you to give a brief overview

NOTE Confidence: 0.7337406

 $00:11:43.550 \longrightarrow 00:11:46.481$ and in closing I would like to

NOTE Confidence: 0.7337406

00:11:46.481 --> 00:11:48.666 acknowledge people in my lab,

NOTE Confidence: 0.7337406

 $00:11:48.670 \longrightarrow 00:11:50.790$ particularly Jessica Mariani, who developed.

NOTE Confidence: 0.7337406

 $00{:}11{:}50.790 \dashrightarrow 00{:}11{:}53.844$ Organized protocol and Alex to down

NOTE Confidence: 0.7337406

 $00:11:53.844 \longrightarrow 00:11:56.961$ and finance who were involved very

NOTE Confidence: 0.7337406

00:11:56.961 --> 00:11:59.907 much so in their recent project

NOTE Confidence: 0.7337406

00:11:59.907 --> 00:12:03.078 with ASD and also our knowledge,

 $00:12:03.080 \longrightarrow 00:12:06.146$ our collaborator at the Trustor Dissenter,

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 $00{:}12{:}06.150 \dashrightarrow 00{:}12{:}08.074$ including various clinicians that

NOTE Confidence: 0.7337406

 $00:12:08.074 \longrightarrow 00:12:10.479$ have been instrumental in patients

NOTE Confidence: 0.7337406

00:12:10.479 --> 00:12:12.290 recruitment and characterization,

NOTE Confidence: 0.7337406

 $00{:}12{:}12.290 \dashrightarrow 00{:}12{:}15.362$ which is of course essential to

NOTE Confidence: 0.7337406

00:12:15.362 --> 00:12:17.410 finally put everything together.

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 $00:12:17.410 \longrightarrow 00:12:21.506$ And thank you very much for your attention.