WEBVTT

NOTE duration: "00:50:24.8000000"

NOTE language:en-us

NOTE Confidence: 0.86679626

00:00:16.910 --> 00:00:18.786 Alright guys, why don't we go ahead

NOTE Confidence: 0.86679626

 $00:00:18.790 \longrightarrow 00:00:20.674$ and get started so we're having a

NOTE Confidence: 0.86679626

 $00:00:20.674 \longrightarrow 00:00:22.560$ zoom webinar format for this talk so

NOTE Confidence: 0.86679626

00:00:22.560 --> 00:00:24.314 Doctor Sakamoto will give her talk and

NOTE Confidence: 0.86679626

 $00:00:24.314 \longrightarrow 00:00:26.330$ then at the end we'll have questions.

NOTE Confidence: 0.86679626

00:00:26.330 --> 00:00:28.207 I think you're going to have to

NOTE Confidence: 0.86679626

 $00{:}00{:}28.207 \dashrightarrow 00{:}00{:}29.821$ put your questions either in the

NOTE Confidence: 0.86679626

00:00:29.821 --> 00:00:32.238 Q&A or in the chat and then I can

NOTE Confidence: 0.86679626

 $00{:}00{:}32.240 \dashrightarrow 00{:}00{:}34.160$ read them to her just given

NOTE Confidence: 0.86679626

 $00:00:34.160 \longrightarrow 00:00:35.860$ the format of this talk.

NOTE Confidence: 0.86679626

 $00{:}00{:}35.860 \dashrightarrow 00{:}00{:}37.318$ So I'm Jeannie Hendrickson.

NOTE Confidence: 0.86679626

00:00:37.318 --> 00:00:38.410 I'm representing the

NOTE Confidence: 0.86679626

 $00:00:38.410 \longrightarrow 00:00:39.499$ Yellow Cooperative Center

NOTE Confidence: 0.83495444

 $00:00:39.500 \longrightarrow 00:00:40.952$ of Excellence in hematology

 $00:00:40.952 \longrightarrow 00:00:42.404$ for the enrichment program,

NOTE Confidence: 0.83495444

 $00{:}00{:}42.410 \dashrightarrow 00{:}00{:}44.595$ and we're extremely happy to have

NOTE Confidence: 0.83495444

00:00:44.595 --> 00:00:46.783 Doctor Sakamoto who's giving a talk,

NOTE Confidence: 0.83495444

 $00:00:46.783 \longrightarrow 00:00:50.053$ a talk to us all the way from Stanford.

NOTE Confidence: 0.83495444

00:00:50.053 --> 00:00:51.144 Virtually, of course,

NOTE Confidence: 0.83495444

 $00:00:51.144 \longrightarrow 00:00:52.964$ she's a professor in pediatric

NOTE Confidence: 0.83495444

00:00:52.964 --> 00:00:54.420 hematology oncology at Stanford,

NOTE Confidence: 0.83495444

 $00:00:54.420 \longrightarrow 00:00:56.246$ and her research involves signaling

NOTE Confidence: 0.83495444

 $00{:}00{:}56.246 \dashrightarrow 00{:}00{:}58.062$ pathways and gene regulation in

NOTE Confidence: 0.83495444

 $00:00:58.062 \longrightarrow 00:00:59.514$ normal and aberrant masterpieces,

NOTE Confidence: 0.83495444

 $00:00:59.520 \longrightarrow 00:01:00.975$ including bone marrow failure.

NOTE Confidence: 0.83495444

 $00:01:00.975 \longrightarrow 00:01:04.255$ And today she's going to focus her talk on

NOTE Confidence: 0.83495444

 $00:01:04.255 \longrightarrow 00:01:07.190$ this as it relates to Diamond Black fan.

NOTE Confidence: 0.83495444

00:01:07.190 --> 00:01:09.145 Nina, so I will turn it over

NOTE Confidence: 0.83495444

 $00:01:09.145 \longrightarrow 00:01:11.005$ to her and again thank you so

 $00:01:11.005 \longrightarrow 00:01:12.330$ much for joining us today.

NOTE Confidence: 0.8300267

 $00{:}01{:}13.800 --> 00{:}01{:}15.270$ Thank you so much Gene.

NOTE Confidence: 0.8300267

 $00{:}01{:}15.270 \dashrightarrow 00{:}01{:}16.730$ I really appreciate the invitation.

NOTE Confidence: 0.8300267

00:01:16.730 --> 00:01:19.226 I also want to thank Pat and Diane

NOTE Confidence: 0.8300267

 $00:01:19.226 \longrightarrow 00:01:21.087$ especially for this kind of rotation

NOTE Confidence: 0.8300267

 $00{:}01{:}21.087 \dashrightarrow 00{:}01{:}23.691$ and what I'm going to do today for the

NOTE Confidence: 0.8300267

 $00{:}01{:}23.691 \dashrightarrow 00{:}01{:}26.114$ next 45 minutes or so is to talk about

NOTE Confidence: 0.8300267

 $00{:}01{:}26.114 \dashrightarrow 00{:}01{:}28.193$ some of the recent work that we've

NOTE Confidence: 0.8300267

 $00{:}01{:}28.193 \dashrightarrow 00{:}01{:}30.454$ been doing on signaling pathways in

NOTE Confidence: 0.8300267

 $00:01:30.454 \longrightarrow 00:01:32.660$ the pathogenesis and treatment of TBI.

NOTE Confidence: 0.8300267

 $00{:}01{:}32.660 \to 00{:}01{:}34.748$ So just to say I have no good disclosure,

NOTE Confidence: 0.8300267

 $00:01:34.750 \longrightarrow 00:01:36.470$ I have no disclosures that

NOTE Confidence: 0.8300267

 $00:01:36.470 \longrightarrow 00:01:38.190$ are related to this work.

NOTE Confidence: 0.8300267

 $00:01:38.190 \longrightarrow 00:01:40.297$ So Diamond life and anemia is a

NOTE Confidence: 0.8300267

00:01:40.297 --> 00:01:42.339 very rare boomer failure syndrome,

NOTE Confidence: 0.8300267

 $00:01:42.340 \longrightarrow 00:01:45.252$ and I'll discuss some of the background

 $00:01:45.252 \longrightarrow 00:01:47.669$ and clinical features of this disease.

NOTE Confidence: 0.8300267

 $00:01:47.670 \longrightarrow 00:01:50.274$ Also, the role of an email like

NOTE Confidence: 0.8300267

 $00:01:50.274 \longrightarrow 00:01:53.268$ high knees in DBA models and then

NOTE Confidence: 0.8300267

 $00:01:53.268 \longrightarrow 00:01:55.950$ targeting and OK for the treatment

NOTE Confidence: 0.8300267

 $00:01:56.035 \dashrightarrow 00:01:59.095$ of DBA and the last part of the talk.

NOTE Confidence: 0.8300267

 $00:01:59.100 \longrightarrow 00:02:02:214$ I will talk about a new project on 71

NOTE Confidence: 0.8300267

 $00:02:02.214 \longrightarrow 00:02:05.758$ in erythropoiesis and EPA pathogenesis.

NOTE Confidence: 0.8300267

 $00{:}02{:}05.760 \longrightarrow 00{:}02{:}08.168$ So DPA is a very rare congenital

NOTE Confidence: 0.8300267

00:02:08.168 --> 00:02:10.378 bone marrow failure syndrome that's

NOTE Confidence: 0.8300267

 $00:02:10.378 \longrightarrow 00:02:12.586$ associated with macrocytic anemia,

NOTE Confidence: 0.8300267

00:02:12.590 --> 00:02:15.146 congenital defects, Anna risk of cancer.

NOTE Confidence: 0.8300267

 $00{:}02{:}15.150 \dashrightarrow 00{:}02{:}17.712$ This disease is generally diagnosed in

NOTE Confidence: 0.8300267

 $00{:}02{:}17.712 \dashrightarrow 00{:}02{:}21.126$ early childhood, less than a year of age.

NOTE Confidence: 0.8300267

00:02:21.130 --> 00:02:24.441 The incidence is seven in a million

NOTE Confidence: 0.8300267

 $00:02:24.441 \longrightarrow 00:02:27.473$ and approximately 20 to 40 new cases

00:02:27.473 --> 00:02:30.100 are diagnosed per year in the US,

NOTE Confidence: 0.8300267

 $00:02:30.100 \longrightarrow 00:02:30.950$ in Canada.

NOTE Confidence: 0.76300555

 $00:02:34.190 \longrightarrow 00:02:35.879$ Download iPad anemia.

NOTE Confidence: 0.76300555

 $00:02:35.879 \longrightarrow 00:02:39.257$ Clinical features are quite diverse and

NOTE Confidence: 0.76300555

00:02:39.257 --> 00:02:42.392 there are many patients who present

NOTE Confidence: 0.76300555

 $00:02:42.392 \longrightarrow 00:02:45.525$ with short stature shown here and

NOTE Confidence: 0.76300555

 $00:02:45.525 \longrightarrow 00:02:48.340$ these Twins upper limb abnormalities

NOTE Confidence: 0.76300555

00:02:48.340 --> 00:02:51.734 including thumb and facial and pallet

NOTE Confidence: 0.76300555

 $00{:}02{:}51.734 \dashrightarrow 00{:}02{:}55.136$ malformations which can occur with certain

NOTE Confidence: 0.76300555

 $00:02:55.136 \longrightarrow 00:02:57.790$ ribosomal protein subunit mutations.

NOTE Confidence: 0.76300555

00:02:57.790 --> 00:03:00.358 Patients can present with small eyes,

NOTE Confidence: 0.76300555

00:03:00.360 --> 00:03:03.356 kidney defects, but only for 30% of

NOTE Confidence: 0.76300555

00:03:03.356 --> 00:03:05.486 all patients have physical findings,

NOTE Confidence: 0.76300555

 $00:03:05.490 \longrightarrow 00:03:07.505$ which makes it very challenging

NOTE Confidence: 0.76300555

00:03:07.505 --> 00:03:10.943 when you have a patient who may not

NOTE Confidence: 0.76300555

00:03:10.943 --> 00:03:13.178 manifest significant anemia and may

 $00:03:13.178 \longrightarrow 00:03:15.767$ be identified by their adult intern,

NOTE Confidence: 0.76300555

 $00:03:15.770 \longrightarrow 00:03:16.582$ internist, physician,

NOTE Confidence: 0.76300555

 $00:03:16.582 \longrightarrow 00:03:19.018$ who notices that they have microcytosis

NOTE Confidence: 0.76300555

00:03:19.018 --> 00:03:21.329 but no other physical features,

NOTE Confidence: 0.76300555

 $00{:}03{:}21.330 \dashrightarrow 00{:}03{:}25.098$ and we've had a few of these patients

NOTE Confidence: 0.76300555

 $00:03:25.098 \longrightarrow 00:03:28.879$ come to clinic in our at Stanford.

NOTE Confidence: 0.76300555

 $00:03:28.880 \longrightarrow 00:03:31.165$ The treatment for Diamondback anemia

NOTE Confidence: 0.76300555

 $00:03:31.165 \longrightarrow 00:03:33.456$ is typically steroids, chronic ritzel,

NOTE Confidence: 0.76300555

 $00:03:33.456 \longrightarrow 00:03:35.746$ transfusions and stem cell transplantation.

NOTE Confidence: 0.76300555

 $00:03:35.750 \longrightarrow 00:03:38.498$ For patients who are steroid refractory

NOTE Confidence: 0.76300555

00:03:38.498 --> 00:03:40.330 or chronically transfusion dependent,

NOTE Confidence: 0.76300555

 $00{:}03{:}40.330 \dashrightarrow 00{:}03{:}42.615$ and these are all associated

NOTE Confidence: 0.76300555

 $00{:}03{:}42.615 \dashrightarrow 00{:}03{:}43.986$ with significant morbidities,

NOTE Confidence: 0.76300555

00:03:43.990 --> 00:03:44.906 including immunosuppression,

NOTE Confidence: 0.76300555

 $00:03:44.906 \longrightarrow 00:03:47.654$ iron overload, graph versus host disease,

 $00{:}03{:}47.660 \dashrightarrow 00{:}03{:}50.402$ there are newer the rapies that have

NOTE Confidence: 0.76300555

00:03:50.402 --> 00:03:51.773 come been proposed.

NOTE Confidence: 0.76300555

 $00:03:51.780 \longrightarrow 00:03:54.986$ For example, L, leucine, the amino acid.

NOTE Confidence: 0.76300555

 $00:03:54.990 \longrightarrow 00:03:57.445$ Many of you are familiar

NOTE Confidence: 0.76300555

 $00:03:57.445 \longrightarrow 00:03:59.409$ with the clinical trial.

NOTE Confidence: 0.76300555

00:03:59.410 --> 00:04:02.056 It's been directed by Jeff Lipton,

NOTE Confidence: 0.76300555

 $00:04:02.060 \longrightarrow 00:04:03.383$ an Adreno blocos.

NOTE Confidence: 0.76300555

 $00:04:03.383 \longrightarrow 00:04:05.147$ They showed a modest

NOTE Confidence: 0.76300555

 $00{:}04{:}05.147 \dashrightarrow 00{:}04{:}06.470$ improvement with leucine,

NOTE Confidence: 0.76300555

 $00:04:06.470 \longrightarrow 00:04:08.234$ although the doses were

NOTE Confidence: 0.76300555

 $00:04:08.234 \longrightarrow 00:04:10.439$ much lower and for safety,

NOTE Confidence: 0.76300555

 $00:04:10.440 \longrightarrow 00:04:12.756$ but there were some patients who

NOTE Confidence: 0.76300555

00:04:12.756 --> 00:04:14.300 did experience improvement in

NOTE Confidence: 0.76300555

 $00{:}04{:}14.364 \dashrightarrow 00{:}04{:}16.170$ the transfusion requirements.

NOTE Confidence: 0.76300555

 $00:04:16.170 \longrightarrow 00:04:18.696$ So Tatter septis Eligant wrapped that

NOTE Confidence: 0.76300555

 $00:04:18.696 \longrightarrow 00:04:21.359$ inhibits the TGF beta signaling pathways

 $00:04:21.359 \longrightarrow 00:04:24.059$ that also has been shown recently

NOTE Confidence: 0.76300555

 $00{:}04{:}24.059 \dashrightarrow 00{:}04{:}26.750$ to improve with police in patients,

NOTE Confidence: 0.76300555

 $00:04:26.750 \longrightarrow 00:04:30.038$ for example with Milo dysplastic syndrome.

NOTE Confidence: 0.76300555

 $00:04:30.040 \longrightarrow 00:04:33.190$ But most of them are being

NOTE Confidence: 0.76300555

 $00:04:33.190 \longrightarrow 00:04:34.240$ cronian investigated.

NOTE Confidence: 0.7915295

 $00:04:36.550 \longrightarrow 00:04:38.986$ Why does summer deficiency in bone

NOTE Confidence: 0.7915295

 $00:04:38.986 \longrightarrow 00:04:41.049$ marrow failure syndrome is become

NOTE Confidence: 0.7915295

 $00:04:41.049 \longrightarrow 00:04:43.281$ more and more common in diseases

NOTE Confidence: 0.7915295

00:04:43.281 --> 00:04:45.441 such as Shockman Diamond syndrome

NOTE Confidence: 0.7915295

 $00:04:45.441 \longrightarrow 00:04:47.866$ which is another congenital bone

NOTE Confidence: 0.7915295

 $00:04:47.866 \longrightarrow 00:04:50.180$ marrow failure syndrome with other

NOTE Confidence: 0.7915295

 $00{:}04{:}50.180 \dashrightarrow 00{:}04{:}52.305$ physical findings, deletion 5 Q.

NOTE Confidence: 0.7915295

 $00:04:52.310 \longrightarrow 00:04:54.234$ Milo dysplastic syndromes which

NOTE Confidence: 0.7915295

 $00:04:54.234 \longrightarrow 00:04:56.639$ is associated with our PS14

NOTE Confidence: 0.7915295

 $00:04:56.639 \longrightarrow 00:04:58.279$ haploin sufficiency that's been described

00:04:58.279 --> 00:05:00.827 by Ben Ebras group several years ago.

NOTE Confidence: 0.7915295

 $00:05:00.830 \longrightarrow 00:05:02.790$ These are all considered diseases

NOTE Confidence: 0.7915295

 $00:05:02.790 \longrightarrow 00:05:04.750$ that are ribosome opathy's with

NOTE Confidence: 0.7915295

 $00:05:04.811 \longrightarrow 00:05:06.620$ defective ribosome Biogenesis.

NOTE Confidence: 0.7915295

 $00:05:06.620 \longrightarrow 00:05:08.460$ And function.

NOTE Confidence: 0.7915295

00:05:08.460 --> 00:05:11.166 Over 80\% of DBA patients have

NOTE Confidence: 0.7915295

 $00{:}05{:}11.166 \dashrightarrow 00{:}05{:}12.970$ mutations in ribosomal protein.

NOTE Confidence: 0.7915295

 $00:05:12.970 \longrightarrow 00:05:15.676$ Some units resulting in ribosome dysfunction,

NOTE Confidence: 0.7915295

 $00{:}05{:}15.680 \dashrightarrow 00{:}05{:}18.968$ an impaired protein translation.

NOTE Confidence: 0.7915295

 $00:05:18.970 \longrightarrow 00:05:21.514$ This is just a pie chart that shows

NOTE Confidence: 0.7915295

 $00:05:21.514 \longrightarrow 00:05:23.764$ the various percentages of patients

NOTE Confidence: 0.7915295

 $00:05:23.764 \longrightarrow 00:05:25.816$ who have ribosomal mutations.

NOTE Confidence: 0.7915295

 $00:05:25.820 \longrightarrow 00:05:28.550$ You can see that among the most

NOTE Confidence: 0.7915295

 $00:05:28.550 \longrightarrow 00:05:30.820$ common is RPS 1925% RP 11.

NOTE Confidence: 0.7915295

 $00{:}05{:}30.820 \dashrightarrow 00{:}05{:}33.762$ About 5% in RP 11 and five have been

NOTE Confidence: 0.7915295

 $00:05:33.762 \longrightarrow 00:05:36.697$ associated more with craniofacial defects,

 $00:05:36.700 \longrightarrow 00:05:38.785$ and there are several other

NOTE Confidence: 0.7915295

 $00{:}05{:}38.785 \dashrightarrow 00{:}05{:}40.870$ mutations have been identified but

NOTE Confidence: 0.7915295

 $00:05:40.938 \longrightarrow 00:05:42.716$ 30% have non ribosomal mutations.

NOTE Confidence: 0.7915295

00:05:42.716 --> 00:05:45.560 In an example is what Vijay Sankar

NOTE Confidence: 0.7915295

 $00{:}05{:}45.560 \dashrightarrow 00{:}05{:}47.580$ had reported that God would.

NOTE Confidence: 0.7915295

00:05:47.580 --> 00:05:49.368 Mutations can also contribute

NOTE Confidence: 0.7915295

 $00:05:49.368 \longrightarrow 00:05:50.709$ to this disease.

NOTE Confidence: 0.7915295

 $00:05:50.710 \longrightarrow 00:05:52.024$ But there are several more mutations

NOTE Confidence: 0.7915295

 $00{:}05{:}52.024 \dashrightarrow 00{:}05{:}53.529$ that have yet to be identified.

NOTE Confidence: 0.77043605

 $00:05:55.650 \longrightarrow 00:05:57.942$ This is a schematic that basically

NOTE Confidence: 0.77043605

 $00:05:57.942 \longrightarrow 00:06:00.329$ summarizes some of the defects that

NOTE Confidence: 0.77043605

 $00{:}06{:}00.329 \dashrightarrow 00{:}06{:}02.699$ have been identified in the ribosomes

NOTE Confidence: 0.77043605

 $00{:}06{:}02.699 \dashrightarrow 00{:}06{:}05.226$ are made synthesis pathway primarily

NOTE Confidence: 0.77043605

00:06:05.226 --> 00:06:07.887 the Treacher Collins syndrome, which,

NOTE Confidence: 0.77043605

00:06:07.887 --> 00:06:09.635 Interestingly does not usually

 $00:06:09.635 \longrightarrow 00:06:11.820$ manifest with bone marrow findings.

NOTE Confidence: 0.77043605

 $00{:}06{:}11.820 \dashrightarrow 00{:}06{:}14.150$ We have just keratosis congenita

NOTE Confidence: 0.77043605

 $00:06:14.150 \longrightarrow 00:06:16.480$ which most people are familiar

NOTE Confidence: 0.77043605

 $00:06:16.558 \longrightarrow 00:06:18.820$ with that can cause fibrosis as

NOTE Confidence: 0.77043605

 $00:06:18.820 \longrightarrow 00:06:21.000$ well as bone marrow failure.

NOTE Confidence: 0.77043605

 $00:06:21.000 \longrightarrow 00:06:23.215$ Cartilage hair hypoplasia has been

NOTE Confidence: 0.77043605

 $00{:}06{:}23.215 \dashrightarrow 00{:}06{:}24.987$ associated with macrocytic anemia.

NOTE Confidence: 0.77043605

 $00:06:24.990 \longrightarrow 00:06:26.770$ And then of course, TBA,

NOTE Confidence: 0.77043605

 $00:06:26.770 \longrightarrow 00:06:29.080$ which can involve the small subunit

NOTE Confidence: 0.77043605

00:06:29.080 --> 00:06:31.400 or large subunit 60 S or 40th,

NOTE Confidence: 0.77043605

 $00:06:31.400 \longrightarrow 00:06:33.180$ and then shockman diamond syndrome.

NOTE Confidence: 0.7897504

 $00:06:35.860 \longrightarrow 00:06:37.965$ It's still very interesting that

NOTE Confidence: 0.7897504

 $00:06:37.965 \longrightarrow 00:06:40.076$ this disease, which is germline

NOTE Confidence: 0.7897504

00:06:40.076 --> 00:06:41.764 mutation of these Robertson,

NOTE Confidence: 0.7897504

00:06:41.770 --> 00:06:44.400 will proteins in other mutations

NOTE Confidence: 0.7897504

 $00:06:44.400 \longrightarrow 00:06:46.504$ involved primarily the erythroid

 $00:06:46.504 \longrightarrow 00:06:49.366$ lineages and what I thought is in

NOTE Confidence: 0.7897504

 $00:06:49.366 \longrightarrow 00:06:51.641$ this very simplified version of Rip

NOTE Confidence: 0.7897504

 $00:06:51.641 \longrightarrow 00:06:53.729$ Oasis is that the cells increase

NOTE Confidence: 0.7897504

 $00:06:53.729 \longrightarrow 00:06:55.918$ in number as they become Earth,

NOTE Confidence: 0.7897504

 $00{:}06{:}55.918 \dashrightarrow 00{:}06{:}58.724$ will glass and somehow this results in

NOTE Confidence: 0.7897504

00:06:58.724 --> 00:07:00.760 increased requirement for ribosomes,

NOTE Confidence: 0.7897504

 $00:07:00.760 \longrightarrow 00:07:02.870$ most likely due to increase.

NOTE Confidence: 0.7897504

 $00:07:02.870 \longrightarrow 00:07:04.980$ Protein translation is required for

NOTE Confidence: 0.7897504

 $00:07:04.980 \longrightarrow 00:07:06.668$ cell proliferation and differentiation.

NOTE Confidence: 0.6911545

 $00:07:09.430 \longrightarrow 00:07:10.980$ The normal hematopoietic treated Mary.

NOTE Confidence: 0.6911545

00:07:10.980 --> 00:07:12.840 Are you familiar with already from?

NOTE Confidence: 0.6911545

00:07:12.840 --> 00:07:14.604 He might have put stem cells to

NOTE Confidence: 0.6911545

00:07:14.604 --> 00:07:16.545 come in my local gender Mega

NOTE Confidence: 0.6911545

 $00{:}07{:}16.545 \dashrightarrow 00{:}07{:}18.420$ Carey Service Rd Ripper genders,

NOTE Confidence: 0.6911545

 $00:07:18.420 \longrightarrow 00:07:19.970$ and then the birth funding

 $00:07:19.970 \longrightarrow 00:07:21.210$ isn't calling from Eunice.

NOTE Confidence: 0.6911545

 $00:07:21.210 \dashrightarrow 00:07:24.010$ Very earlier it'll bus stage all lead to

NOTE Confidence: 0.6911545

 $00:07:24.010 \longrightarrow 00:07:26.557$ eventually the maturing of red blood cells.

NOTE Confidence: 0.6911545

 $00:07:26.560 \longrightarrow 00:07:29.620$ In DBA, these very subunit mutations

NOTE Confidence: 0.6911545

 $00:07:29.620 \longrightarrow 00:07:32.086$ that have been described seemed

NOTE Confidence: 0.6911545

 $00:07:32.086 \longrightarrow 00:07:34.822$ to be a result in a block in

NOTE Confidence: 0.6911545

 $00{:}07{:}34.822 \dashrightarrow 00{:}07{:}37.428$ early committed with blaster be.

NOTE Confidence: 0.6911545

 $00:07:37.430 \longrightarrow 00:07:40.148$ If you be a few istage,

NOTE Confidence: 0.6911545

 $00{:}07{:}40.150 \to 00{:}07{:}42.170$ mostly resulting from haploin sufficiency

NOTE Confidence: 0.6911545

00:07:42.170 --> 00:07:45.200 inducing mutation and we know that

NOTE Confidence: 0.6911545

 $00{:}07{:}45.276 \dashrightarrow 00{:}07{:}46.812$ homozygous mutations result in

NOTE Confidence: 0.6911545

 $00:07:46.812 \longrightarrow 00:07:49.660$ mouse models as well as in humans.

NOTE Confidence: 0.6911545

 $00:07:49.660 \longrightarrow 00:07:51.524$ Embryonic lethality so helpful

NOTE Confidence: 0.6911545

 $00{:}07{:}51.524 \dashrightarrow 00{:}07{:}52.922$ insufficiency or heterozygous

NOTE Confidence: 0.6911545

 $00:07:52.922 \longrightarrow 00:07:55.100$ mutations are typically what results.

NOTE Confidence: 0.6911545

 $00:07:55.100 \longrightarrow 00:07:57.710$ What results in this disease.

 $00:07:57.710 \longrightarrow 00:08:00.518$ Ultimately leading to anemia.

NOTE Confidence: 0.6911545

 $00:08:00.520 \longrightarrow 00:08:03.130$ So we started out looking at

NOTE Confidence: 0.6911545

00:08:03.130 --> 00:08:04.870 variety of signaling molecules,

NOTE Confidence: 0.6911545

 $00:08:04.870 \longrightarrow 00:08:07.606$ one that was of interest to us initially

NOTE Confidence: 0.6911545

 $00{:}08{:}07.606 \dashrightarrow 00{:}08{:}09.883$ was a transcription factor MYB and

NOTE Confidence: 0.6911545

 $00:08:09.883 \longrightarrow 00:08:13.550$ we know that this is a very important

NOTE Confidence: 0.6911545

 $00:08:13.550 \longrightarrow 00:08:16.178$ protein that regulates erythropoiesis.

NOTE Confidence: 0.6911545

 $00:08:16.180 \longrightarrow 00:08:19.225$ It's also associated with my little keemia.

NOTE Confidence: 0.6911545

 $00:08:19.230 \longrightarrow 00:08:21.600$ When Aberrantly expressed and it's been

NOTE Confidence: 0.6911545

 $00:08:21.600 \longrightarrow 00:08:24.695$ reported that one of the kinases that

NOTE Confidence: 0.6911545

 $00{:}08{:}24.695 \dashrightarrow 00{:}08{:}26.615$ activates or phosphorylates Mibiz,

NOTE Confidence: 0.6911545

00:08:26.620 --> 00:08:27.600 NIMAL, Iconis,

NOTE Confidence: 0.6911545

 $00{:}08{:}27.600 \dashrightarrow 00{:}08{:}30.050$ and this kinese is Assyrian

NOTE Confidence: 0.6911545

 $00{:}08{:}30.050 --> 00{:}08{:}31.030$ threatening kinese.

NOTE Confidence: 0.6911545

 $00:08:31.030 \longrightarrow 00:08:34.096$ That is a revolutionary and evolutionary

 $00:08:34.096 \longrightarrow 00:08:36.140$ conserved margin activated protein

NOTE Confidence: 0.6911545

 $00:08:36.211 \longrightarrow 00:08:38.864$ kinase in the map kinase family member.

NOTE Confidence: 0.6911545

 $00:08:38.870 \longrightarrow 00:08:41.026$ Originally described when mutated

NOTE Confidence: 0.6911545

 $00:08:41.026 \longrightarrow 00:08:43.721$ result results in I development

NOTE Confidence: 0.6911545

00:08:43.721 --> 00:08:46.448 defects in just saffola is highly

NOTE Confidence: 0.6911545

 $00:08:46.448 \longrightarrow 00:08:48.553$ expressed in neural tissues and

NOTE Confidence: 0.6911545

00:08:48.632 --> 00:08:51.880 plays a critical role in a number of

NOTE Confidence: 0.6911545

 $00:08:51.880 \longrightarrow 00:08:53.978$ important cellular functions through

NOTE Confidence: 0.6911545

 $00{:}08{:}53.978 \dashrightarrow 00{:}08{:}57.173$ the regulation of various transcription

NOTE Confidence: 0.6911545

 $00:08:57.173 \longrightarrow 00:08:59.699$ transcription factors such as nib.

NOTE Confidence: 0.6911545

 $00{:}08{:}59.700 \dashrightarrow 00{:}09{:}01.700$ An OK also regulate signaling

NOTE Confidence: 0.6911545

 $00:09:01.700 \longrightarrow 00:09:03.700$ pathways involving wind beta catenin

NOTE Confidence: 0.6911545

 $00:09:03.763 \longrightarrow 00:09:05.395$ active in aisle 6 an notch,

NOTE Confidence: 0.6911545

 $00:09:05.400 \longrightarrow 00:09:08.365$ which are all very critical

NOTE Confidence: 0.6911545

 $00:09:08.365 \longrightarrow 00:09:10.144$ for normal hematopoiesis.

NOTE Confidence: 0.6911545

00:09:10.150 --> 00:09:12.518 So this is work done by Mark Walsh,

 $00:09:12.520 \longrightarrow 00:09:13.995$ who is a former postdoctoral

NOTE Confidence: 0.6911545

 $00:09:13.995 \longrightarrow 00:09:15.175$ fellow in my lab,

NOTE Confidence: 0.6911545

 $00:09:15.180 \longrightarrow 00:09:18.659$ and now is an instructor in Pediatrics.

NOTE Confidence: 0.6911545

 $00:09:18.660 \longrightarrow 00:09:21.412$ We used a model that had been used

NOTE Confidence: 0.6911545

 $00:09:21.412 \longrightarrow 00:09:24.313$ prior in my lab where we would

NOTE Confidence: 0.6911545

00:09:24.313 --> 00:09:27.000 transduced human CD 34 positive cells,

NOTE Confidence: 0.6911545

00:09:27.000 --> 00:09:28.188 hematopoietic stem progenitor

NOTE Confidence: 0.6911545

00:09:28.188 --> 00:09:30.168 cells with small hairpin RNAs,

NOTE Confidence: 0.6911545

 $00:09:30.170 \longrightarrow 00:09:34.463$ then knocked down either RPS 19 or RPO 11.

NOTE Confidence: 0.6911545

 $00:09:34.470 \longrightarrow 00:09:36.410$ Making lentiviral constructs with

NOTE Confidence: 0.6911545

 $00:09:36.410 \longrightarrow 00:09:39.320$ transduced and then show that we

NOTE Confidence: 0.6911545

 $00:09:39.397 \longrightarrow 00:09:41.861$ could get help low or half the

NOTE Confidence: 0.6911545

 $00:09:41.861 \longrightarrow 00:09:44.125$ expression at the protein level by

NOTE Confidence: 0.6911545

 $00{:}09{:}44.125 \dashrightarrow 00{:}09{:}46.750$ this Western blot or M RNA levels.

NOTE Confidence: 0.6911545

 $00:09:46.750 \longrightarrow 00:09:49.470$ And this is the model that we used.

 $00:09:49.470 \longrightarrow 00:09:51.549$ And when Mark looked at the expression

NOTE Confidence: 0.6911545

 $00{:}09{:}51.549 \dashrightarrow 00{:}09{:}53.950$ of an OK throughout, he matter.

NOTE Confidence: 0.6911545

00:09:53.950 --> 00:09:56.170 Police is we didn't see any

NOTE Confidence: 0.6911545

00:09:56.170 --> 00:09:57.630 significant differences in our bar.

NOTE Confidence: 0.6911545

 $00:09:57.630 \longrightarrow 00:10:00.955$ RPS 19 RP 11 knock down now.

NOTE Confidence: 0.6911545

 $00:10:00.960 \longrightarrow 00:10:01.810$ In contrast,

NOTE Confidence: 0.6911545

 $00:10:01.810 \longrightarrow 00:10:03.935$ looking at phosphorylation or activation

NOTE Confidence: 0.6911545

00:10:03.935 --> 00:10:07.609 of an OK when Mark noticed was that

NOTE Confidence: 0.6911545

 $00{:}10{:}07.609 \dashrightarrow 00{:}10{:}09.469$ taking three different substrates,

NOTE Confidence: 0.6911545

00:10:09.470 --> 00:10:12.606 either an OK itself this is auto

NOTE Confidence: 0.6911545

 $00{:}10{:}12.606 \mathrel{--}{>} 00{:}10{:}14.497$ phosphorylates, nib or wrapped,

NOTE Confidence: 0.6911545

 $00{:}10{:}14.497 \dashrightarrow 00{:}10{:}17.371$ or that there is significant increase

NOTE Confidence: 0.6911545

00:10:17.371 --> 00:10:19.994 in the activation of this kinese

NOTE Confidence: 0.6911545

 $00:10:19.994 \longrightarrow 00:10:23.006$ in the 1st 10 days of hematopoiesis

NOTE Confidence: 0.6911545

 $00:10:23.006 \longrightarrow 00:10:26.058$ and so this we decided to focus

NOTE Confidence: 0.6911545

 $00{:}10{:}26.058 \mathrel{--}{>} 00{:}10{:}29.496$ on an OK and looking at mid target

00:10:29.496 --> 00:10:31.989 genes with RPS 19 lockdown.

NOTE Confidence: 0.6911545

 $00:10:31.990 \longrightarrow 00:10:32.863$ Again, HSBC is,

NOTE Confidence: 0.6911545

 $00:10:32.863 \longrightarrow 00:10:34.318$ we see that their expression

NOTE Confidence: 0.6911545

 $00{:}10{:}34.318 \dashrightarrow 00{:}10{:}36.677$ of Alamo two in California were

NOTE Confidence: 0.6911545

00:10:36.677 --> 00:10:37.537 significantly decreased.

NOTE Confidence: 0.6911545

 $00:10:37.540 \longrightarrow 00:10:37.887$ However,

NOTE Confidence: 0.6911545

00:10:37.887 --> 00:10:40.663 with the expression or knockdown of an OK,

NOTE Confidence: 0.7975294

00:10:40.670 --> 00:10:43.183 we see that that we could partially

NOTE Confidence: 0.7975294

 $00:10:43.183 \longrightarrow 00:10:45.519$ rescue the expression of these two genes.

NOTE Confidence: 0.7975294

 $00:10:45.520 \longrightarrow 00:10:46.908$ Obviously there other Trump,

NOTE Confidence: 0.7975294

00:10:46.908 --> 00:10:48.990 you know the regulators of nib,

NOTE Confidence: 0.7975294

 $00{:}10{:}48.990 \dashrightarrow 00{:}10{:}52.113$ so that makes a lot of sense to us.

NOTE Confidence: 0.7975294

 $00{:}10{:}52.120 --> 00{:}10{:}53.158$ I'm sorry, Kathy,

NOTE Confidence: 0.7975294

00:10:53.160 --> 00:10:56.880 can you go back one slide?

NOTE Confidence: 0.7975294

 $00:10:56.880 \longrightarrow 00:10:59.320$ What is one more? I'm sorry.

 $00:10:59.320 \longrightarrow 00:11:02.576$ What are you showing on the PC MIB?

NOTE Confidence: 0.7975294

 $00:11:02.580 \longrightarrow 00:11:04.610$ One forward PC MIB an?

NOTE Confidence: 0.7975294

 $00:11:04.610 \longrightarrow 00:11:07.058$ What's the PC MIB? There is

NOTE Confidence: 0.71488774

00:11:07.060 --> 00:11:07.872 middling. Phosphorylated

NOTE Confidence: 0.71488774

 $00:11:07.872 \longrightarrow 00:11:09.496$ yeah phosphorylated. MIB is

NOTE Confidence: 0.71488774

00:11:09.500 --> 00:11:11.124 that active? Is phosphorylated

NOTE Confidence: 0.71488774

00:11:11.124 --> 00:11:12.750 MIB active? Yes yes

NOTE Confidence: 0.71488774

 $00:11:12.750 \longrightarrow 00:11:16.285$ so. So that is a key phosphorylation

NOTE Confidence: 0.71488774

 $00:11:16.285 \longrightarrow 00:11:19.110$ site that's recognized by in OK. OK,

NOTE Confidence: 0.7562662

00:11:19.110 --> 00:11:21.360 so any OK is phosphorylating

NOTE Confidence: 0.7562662

 $00{:}11{:}21.360 \dashrightarrow 00{:}11{:}23.610$ in activating them? Yes OK

NOTE Confidence: 0.7562662

 $00:11:23.610 \longrightarrow 00:11:26.370$ just clarifying yes and I'll talk

NOTE Confidence: 0.7562662

 $00:11:26.370 \longrightarrow 00:11:29.460$ more about that in a few slides.

NOTE Confidence: 0.7562662

 $00:11:29.460 \longrightarrow 00:11:31.710$ Yeah and so so now.

NOTE Confidence: 0.7562662

 $00:11:31.710 \longrightarrow 00:11:34.830$ How does an OK regulate nib actually what

NOTE Confidence: 0.7562662

 $00:11:34.830 \longrightarrow 00:11:38.195$ happens is that activated and OK then

00:11:38.195 --> 00:11:40.705 phosphorylates mid to ubiquitinated nib,

NOTE Confidence: 0.7562662

 $00{:}11{:}40.710 \dashrightarrow 00{:}11{:}43.860$ an result in 26 S proteasome degradation.

NOTE Confidence: 0.7562662

 $00:11:43.860 \longrightarrow 00:11:49.180$ So if we have an OK and we knock it down.

NOTE Confidence: 0.7562662

00:11:49.180 --> 00:11:51.798 RPS 19 meter put extent progenitor cells.

NOTE Confidence: 0.7562662

 $00:11:51.800 \longrightarrow 00:11:54.784$ What we actually see is stabilization of MIB.

NOTE Confidence: 0.7562662

 $00:11:54.790 \longrightarrow 00:11:57.022$ So you see that the protein

NOTE Confidence: 0.7562662

 $00:11:57.022 \longrightarrow 00:11:59.279$ levels are higher and Konan OK.

NOTE Confidence: 0.7562662

 $00:11:59.280 \longrightarrow 00:12:01.150$ It's knocked down and this

NOTE Confidence: 0.7562662

 $00:12:01.150 \longrightarrow 00:12:03.020$ again this is in RPS.

NOTE Confidence: 0.7562662

 $00{:}12{:}03.020 \dashrightarrow 00{:}12{:}05.548$ 19 knockdown cells so you can see that

NOTE Confidence: 0.7562662

 $00{:}12{:}05.548 {\:\raisebox{--}{\text{--}}}{\:\raisebox{--}{\text{--}}}{\:\raisebox{--}{\text{--}}} 00{:}12{:}08.627$ if we again look at phosphorylated nip,

NOTE Confidence: 0.7562662

 $00:12:08.630 \longrightarrow 00:12:10.681$ that compared to when we knock down

NOTE Confidence: 0.7562662

 $00{:}12{:}10.681 \dashrightarrow 00{:}12{:}13.595$ in OK we see higher levels of mid

NOTE Confidence: 0.7562662

 $00:12:13.595 \longrightarrow 00:12:15.560$ phosphorylation as we would expect

NOTE Confidence: 0.7562662

 $00{:}12{:}15.630 \dashrightarrow 00{:}12{:}17.930$ because MIB results in degradation

 $00:12:17.930 \longrightarrow 00:12:20.230$ and OK phosphorylation results in.

NOTE Confidence: 0.7562662

00:12:20.230 --> 00:12:23.488 Amid degradation and also here only

NOTE Confidence: 0.7562662

 $00:12:23.488 \longrightarrow 00:12:26.250$ milk is ubiquitinated without an OK.

NOTE Confidence: 0.7562662

00:12:26.250 --> 00:12:28.760 So you see higher levels,

NOTE Confidence: 0.7562662

 $00:12:28.760 \longrightarrow 00:12:32.274$ so without an OK it's not degraded.

NOTE Confidence: 0.82151514

 $00:12:34.370 \longrightarrow 00:12:37.762$ So we also looked at C71 expression in

NOTE Confidence: 0.82151514

 $00:12:37.762 \longrightarrow 00:12:40.857$ this schematic schema that you see to

NOTE Confidence: 0.82151514

00:12:40.857 --> 00:12:44.208 indicate Orthop Oasis will Mark did was to

NOTE Confidence: 0.82151514

 $00{:}12{:}44.208 \mathrel{--}{>} 00{:}12{:}47.200$ isolate the C 71 positive cells and then

NOTE Confidence: 0.82151514

00:12:47.200 --> 00:12:50.070 to perform a Western blot with antibody

NOTE Confidence: 0.82151514

 $00{:}12{:}50.070 --> 00{:}12{:}52.849$ specific for three nine, 298 on in.

NOTE Confidence: 0.82151514

 $00:12:52.849 \longrightarrow 00:12:56.574$ OK so here we see that only in C 71

NOTE Confidence: 0.82151514

 $00:12:56.574 \longrightarrow 00:12:59.745$ positive cells do we see an OK activation

NOTE Confidence: 0.82151514

 $00{:}12{:}59.745 \dashrightarrow 00{:}13{:}04.551$ phosphorylation at this site but not in C 71.

NOTE Confidence: 0.82151514

 $00:13:04.560 \longrightarrow 00:13:07.424$ Get of cells and that's whether you have

NOTE Confidence: 0.82151514

00:13:07.424 --> 00:13:10.392 herpes 19 knocked down or control. Again,

00:13:10.392 --> 00:13:13.626 if we immunoprecipitated and OK in cells,

NOTE Confidence: 0.82151514

00:13:13.630 --> 00:13:16.878 either control or RPS 19 knockdown cells,

NOTE Confidence: 0.82151514

00:13:16.880 --> 00:13:19.666 we see that the phosphorylation in CD

NOTE Confidence: 0.82151514

 $00:13:19.666 \longrightarrow 00:13:22.611$ 71 cells of these three substrates

NOTE Confidence: 0.82151514

 $00:13:22.611 \longrightarrow 00:13:24.276$ is is upregulated,

NOTE Confidence: 0.82151514

 $00:13:24.280 \longrightarrow 00:13:27.514$ and that's what we would expect because

NOTE Confidence: 0.82151514

 $00:13:27.514 \longrightarrow 00:13:30.768$ we get hyper activation of an OK.

NOTE Confidence: 0.82151514

00:13:30.770 --> 00:13:34.322 So this is only again in C 171671

NOTE Confidence: 0.82151514

 $00{:}13{:}34.322 \dashrightarrow 00{:}13{:}37.709$ positive cells but not in other lineages.

NOTE Confidence: 0.7884901

 $00:13:39.920 \longrightarrow 00:13:42.195$ Furthermore, we looked at the

NOTE Confidence: 0.7884901

00:13:42.195 --> 00:13:45.120 activation of MLK in mouse models,

NOTE Confidence: 0.7884901

 $00{:}13{:}45.120 \dashrightarrow 00{:}13{:}47.964$ so we collaborated with new on

NOTE Confidence: 0.7884901

 $00{:}13{:}47.964 \dashrightarrow 00{:}13{:}50.800$ Flickr at Lund University in Sweden,

NOTE Confidence: 0.7884901

 $00:13:50.800 \longrightarrow 00:13:54.576$ who provided to us stem cells from RPS.

NOTE Confidence: 0.7884901

 $00:13:54.580 \longrightarrow 00:13:55.996$ 19 knockdown mice.

00:13:55.996 --> 00:13:57.884 It's catcher cycling inducible.

NOTE Confidence: 0.7884901

 $00:13:57.890 \longrightarrow 00:14:01.674$ Similarly, we examine the stem cells from RP.

NOTE Confidence: 0.7884901

00:14:01.680 --> 00:14:05.343 11 flox mice, which is provided to us by

NOTE Confidence: 0.7884901

 $00:14:05.343 \longrightarrow 00:14:08.322$ Manuel Serrano from Barcelona, Spain.

NOTE Confidence: 0.7884901

 $00:14:08.322 \longrightarrow 00:14:09.798$ This is also.

NOTE Confidence: 0.7884901

00:14:09.798 --> 00:14:12.750 Tamoxifen induced to lose one allele

NOTE Confidence: 0.7884901

 $00:14:12.842 \longrightarrow 00:14:15.862$ of RP 11 gene and what we see is in

NOTE Confidence: 0.7884901

00:14:15.953 --> 00:14:19.215 these mice with which have anemia that

NOTE Confidence: 0.7884901

00:14:19.215 --> 00:14:22.570 the activity of in OK is much higher

NOTE Confidence: 0.7884901

 $00:14:22.570 \longrightarrow 00:14:25.569$ than our control mice or minus docs.

NOTE Confidence: 0.7884901

 $00:14:25.570 \longrightarrow 00:14:28.618$ In both the RPS 19 and RPL 11

NOTE Confidence: 0.7884901

00:14:28.618 --> 00:14:30.510 knockdown mice mouse cells.

NOTE Confidence: 0.7884901

 $00:14:30.510 \longrightarrow 00:14:32.778$ We also examined patient samples and

NOTE Confidence: 0.7884901

 $00:14:32.778 \longrightarrow 00:14:35.257$ this is a collaboration with Hannah

NOTE Confidence: 0.7884901

 $00:14:35.257 \longrightarrow 00:14:37.507$ Gosda from Boston Children's Hospital.

NOTE Confidence: 0.7884901

 $00:14:37.510 \longrightarrow 00:14:39.645$ She provided to us three

 $00:14:39.645 \longrightarrow 00:14:40.926$ different patient samples.

NOTE Confidence: 0.7884901

 $00:14:40.930 \longrightarrow 00:14:43.982$ With these mutations in our PS 19

NOTE Confidence: 0.7884901

 $00:14:43.982 \longrightarrow 00:14:47.525$ Mark then examined the relative in OK

NOTE Confidence: 0.7884901

00:14:47.525 --> 00:14:50.210 kindness activity compared to healthy

NOTE Confidence: 0.7884901

 $00{:}14{:}50.210 \dashrightarrow 00{:}14{:}54.150$ control cells and show that that the

NOTE Confidence: 0.7884901

 $00:14:54.150 \longrightarrow 00:14:56.362$ activity was significantly increased.

NOTE Confidence: 0.7884901

00:14:56.370 --> 00:14:57.114 And finally,

NOTE Confidence: 0.7884901

00:14:57.114 --> 00:14:58.974 in collaboration with here Mitsunaga,

NOTE Confidence: 0.7884901

 $00:14:58.980 \longrightarrow 00:15:00.468$ which he had Stanford,

NOTE Confidence: 0.7884901

00:15:00.468 --> 00:15:01.584 his postdoctoral fellow,

NOTE Confidence: 0.7884901

 $00:15:01.590 \longrightarrow 00:15:02.670$ to she developed,

NOTE Confidence: 0.7884901

 $00{:}15{:}02.670 \dashrightarrow 00{:}15{:}05.190$ I PS cells from two different patients

NOTE Confidence: 0.7884901

 $00{:}15{:}05.259 \dashrightarrow 00{:}15{:}07.189$ who are diagnosed at Stanford,

NOTE Confidence: 0.7884901

 $00:15:07.190 \longrightarrow 00:15:09.055$ who had the typical clinical

NOTE Confidence: 0.7884901

 $00:15:09.055 \longrightarrow 00:15:10.920$ and physical features of DVA.

 $00:15:10.920 \longrightarrow 00:15:14.268$ And one of them had an RPS 26 mutation.

NOTE Confidence: 0.7884901

 $00{:}15{:}14.270 \dashrightarrow 00{:}15{:}16.888$ The other one had an unknown mutation,

NOTE Confidence: 0.7884901

 $00:15:16.890 \longrightarrow 00:15:19.098$ but in all three clones that

NOTE Confidence: 0.7884901

 $00:15:19.098 \longrightarrow 00:15:21.360$ we analyzed from I PS cells,

NOTE Confidence: 0.7884901

 $00:15:21.360 \longrightarrow 00:15:24.041$ we see hyperactivation of an OK using

NOTE Confidence: 0.7884901

00:15:24.041 --> 00:15:26.069 the substrates that I mentioned.

NOTE Confidence: 0.7884901

 $00:15:26.070 \longrightarrow 00:15:27.978$ And OK leban rector.

NOTE Confidence: 0.8358886

 $00:15:30.530 \longrightarrow 00:15:32.616$ So given this in all of the

NOTE Confidence: 0.8358886

 $00{:}15{:}32.616 \dashrightarrow 00{:}15{:}34.789$ models that we examined for DBA,

NOTE Confidence: 0.8358886

 $00:15:34.790 \longrightarrow 00:15:36.258$ regardless of the mutation,

NOTE Confidence: 0.8358886

 $00{:}15{:}36.258 \dashrightarrow 00{:}15{:}39.570$ we were able to see an OK activation.

NOTE Confidence: 0.8358886

 $00{:}15{:}39.570 \dashrightarrow 00{:}15{:}42.354$ And so we wanted to test the hypothesis

NOTE Confidence: 0.8358886

 $00:15:42.354 \longrightarrow 00:15:45.738$ could in OK be a possible target for therapy.

NOTE Confidence: 0.8358886

 $00{:}15{:}45.740 \dashrightarrow 00{:}15{:}48.719$ The way that we looked at this was to

NOTE Confidence: 0.8358886

00:15:48.719 --> 00:15:51.906 study the number of two thirty 5235 cells,

NOTE Confidence: 0.8358886

 $00:15:51.910 \longrightarrow 00:15:53.358$ which is a reflection

 $00:15:53.358 \longrightarrow 00:15:54.444$ of birthright expansion.

NOTE Confidence: 0.8358886

00:15:54.450 --> 00:15:57.717 We see that in our PS 19 knockdown cells,

NOTE Confidence: 0.8358886

 $00:15:57.720 \longrightarrow 00:15:59.898$ if we treat with sin LK.

NOTE Confidence: 0.8358886

00:15:59.900 --> 00:16:01.252 To knock down MLK,

NOTE Confidence: 0.8358886

 $00{:}16{:}01.252 \dashrightarrow 00{:}16{:}04.116$ we see increase in the numbers of these

NOTE Confidence: 0.8358886

 $00:16:04.116 \longrightarrow 00:16:06.797$ are three projector cells or three cells,

NOTE Confidence: 0.8358886

 $00:16:06.800 \longrightarrow 00:16:09.816$ in contrast to absence of in OK or.

NOTE Confidence: 0.8358886

 $00{:}16{:}09.820 \dashrightarrow 00{:}16{:}12.480$ Design OK or a control escape in

NOTE Confidence: 0.8358886

 $00{:}16{:}12.480 \dashrightarrow 00{:}16{:}15.443$ OK and the point here is that

NOTE Confidence: 0.8358886

 $00:16:15.443 \longrightarrow 00:16:17.603$ in the patients with DPA,

NOTE Confidence: 0.8358886

 $00{:}16{:}17.610 \dashrightarrow 00{:}16{:}19.815$ generally speaking they are treated

NOTE Confidence: 0.8358886

 $00{:}16{:}19.815 \dashrightarrow 00{:}16{:}22.020$ with steroids or transfused in

NOTE Confidence: 0.8358886

 $00:16:22.095 \longrightarrow 00:16:24.165$ when the hemoglobin is below 8.

NOTE Confidence: 0.8358886

 $00:16:24.170 \longrightarrow 00:16:27.506$ So our goal is not to necessarily improve

NOTE Confidence: 0.8358886

 $00:16:27.506 \longrightarrow 00:16:29.907$ the hemoglobin up to normal levels,

 $00:16:29.910 \longrightarrow 00:16:32.286$ but rather to increase or decreases

NOTE Confidence: 0.8358886

 $00:16:32.286 \longrightarrow 00:16:35.189$ sufficiently so that they will no longer

NOTE Confidence: 0.8358886

 $00:16:35.189 \longrightarrow 00:16:37.697$ need story therapy or wrestle transfusions.

NOTE Confidence: 0.8358886

 $00:16:37.700 \longrightarrow 00:16:39.750$ So that's really our goal.

NOTE Confidence: 0.81782615

 $00:16:42.210 \longrightarrow 00:16:44.508$ You have a hand raise. Pat Gallagher

NOTE Confidence: 0.81782615

 $00:16:44.508 \longrightarrow 00:16:46.470$ is raising his hand. OK, yes?

NOTE Confidence: 0.8330304

 $00:16:50.340 \longrightarrow 00:16:52.255$ Let me think we have to figure

NOTE Confidence: 0.8330304

 $00:16:52.255 \longrightarrow 00:16:53.923$ out how to let him talk.

NOTE Confidence: 0.8330304

00:16:53.923 --> 00:16:56.369 Hold on one second so I can do it.

NOTE Confidence: 0.8330304

 $00{:}16{:}56.370 --> 00{:}16{:}58.008$ I can do it. Go ahead.

NOTE Confidence: 0.8330304

 $00{:}16{:}58.010 \dashrightarrow 00{:}17{:}01.868$ I'm good guys are the question I got OK?

NOTE Confidence: 0.8330304

00:17:01.870 --> 00:17:05.314 Do you put my hand down there?

NOTE Confidence: 0.8330304

 $00:17:05.320 \dashrightarrow 00:17:09.139$ I don't know if I can do know how to do that.

NOTE Confidence: 0.8330304

00:17:09.140 --> 00:17:11.016 I had a question though on the

NOTE Confidence: 0.8330304

 $00:17:11.016 \longrightarrow 00:17:12.286$ previous slide you normalized

NOTE Confidence: 0.8330304

 $00:17:12.286 \longrightarrow 00:17:13.906$ the healthy control cells.

 $00:17:13.910 \longrightarrow 00:17:16.129$ Yes to 100% on the previous slide.

NOTE Confidence: 0.8330304

 $00{:}17{:}16.130 \dashrightarrow 00{:}17{:}18.731$ Yeah, and had to go back and if you

NOTE Confidence: 0.8330304

00:17:18.731 --> 00:17:21.353 knock down you had the sin LK and you

NOTE Confidence: 0.8330304

 $00:17:21.353 \longrightarrow 00:17:23.758$ gave the absolute number of cells.

NOTE Confidence: 0.8330304

 $00:17:23.760 \longrightarrow 00:17:25.986$ Is that affected in the healthy controls?

NOTE Confidence: 0.8330304

 $00:17:25.990 \longrightarrow 00:17:27.258$ Well certainly with DPA

NOTE Confidence: 0.8322626

 $00:17:27.260 \longrightarrow 00:17:29.168$ patient samples there are lower numbers

NOTE Confidence: 0.8322626

 $00:17:29.168 \longrightarrow 00:17:32.036$ of cells, but in spite of this we see

NOTE Confidence: 0.8322626

 $00{:}17{:}32.036 \dashrightarrow 00{:}17{:}33.938$ hyperactivation of energy. So I'm talking

NOTE Confidence: 0.8322626

00:17:33.940 --> 00:17:36.530 bout in the controls on the left. In

NOTE Confidence: 0.86921525

 $00:17:36.530 \longrightarrow 00:17:39.418$ the controls, if we knock if we if

NOTE Confidence: 0.86921525

 $00:17:39.418 \longrightarrow 00:17:42.287$ we look at the numbers of cells.

NOTE Confidence: 0.86921525

 $00{:}17{:}42.290 \dashrightarrow 00{:}17{:}45.332$ I don't know if we actually have done that.

NOTE Confidence: 0.86921525

00:17:45.340 --> 00:17:48.364 I mean normally we would try to

NOTE Confidence: 0.86921525

 $00:17:48.364 \longrightarrow 00:17:50.739$ normalize the cell's cell number.

 $00:17:50.740 \longrightarrow 00:17:53.484$ To make it equal on both sides,

NOTE Confidence: 0.86921525

00:17:53.490 --> 00:17:56.634 but yet it's still we see hyperactivation of.

NOTE Confidence: 0.86921525

00:17:56.640 --> 00:18:00.168 I mean we see lower levels of an OK

NOTE Confidence: 0.87146217

00:18:00.170 --> 00:18:01.738 activation. Yeah thanks yeah.

NOTE Confidence: 0.7750263

00:18:03.020 --> 00:18:04.900 So in collaboration with Johan,

NOTE Confidence: 0.7750263

 $00:18:04.900 \longrightarrow 00:18:07.150$ we were able to look at.

NOTE Confidence: 0.7750263

 $00:18:07.150 \longrightarrow 00:18:11.929$ He gave us 8 compounds that he had screened.

NOTE Confidence: 0.7750263

00:18:11.930 --> 00:18:15.106 Using a 14,000 compound library in his DBA

NOTE Confidence: 0.7750263

00:18:15.106 --> 00:18:18.666 RPS 19 knockdown mice trying to go back,

NOTE Confidence: 0.7750263

00:18:18.670 --> 00:18:23.340 but I can't. And what we saw was

NOTE Confidence: 0.7750263

 $00{:}18{:}23.340 \dashrightarrow 00{:}18{:}25.230$ these eight compounds and compounds.

NOTE Confidence: 0.7750263

 $00{:}18{:}25.230 \to 00{:}18{:}27.533$ Six and eight were the most effective

NOTE Confidence: 0.7750263

 $00:18:27.533 \longrightarrow 00:18:30.139$ as far as their third expansion.

NOTE Confidence: 0.7750263

 $00{:}18{:}30.140 \dashrightarrow 00{:}18{:}33.542$ And then if we looked at the Enoch activity,

NOTE Confidence: 0.7750263

 $00:18:33.550 \longrightarrow 00:18:35.058$ the typical substrates that

NOTE Confidence: 0.7750263

 $00:18:35.058 \longrightarrow 00:18:36.943$ we've examined in the past,

 $00:18:36.950 \longrightarrow 00:18:39.704$ we see that in these two compounds we were

NOTE Confidence: 0.7750263

 $00{:}18{:}39.704 \dashrightarrow 00{:}18{:}42.618$ able to see significantly decrease activity.

NOTE Confidence: 0.7750263

 $00:18:42.620 \longrightarrow 00:18:44.888$ Now here we have our Pierce

NOTE Confidence: 0.7750263

00:18:44.888 --> 00:18:46.400 19 knockout cells again,

NOTE Confidence: 0.7750263

 $00:18:46.400 \longrightarrow 00:18:49.793$ and we see with this is with sin LK.

NOTE Confidence: 0.7750263

 $00:18:49.800 \longrightarrow 00:18:52.474$ So when we add this compound is

NOTE Confidence: 0.7750263

00:18:52.474 --> 00:18:55.337 Scituate which is a TGF beta inhibitor.

NOTE Confidence: 0.7750263

 $00:18:55.340 \longrightarrow 00:18:58.836$ Which has an OK as an off target.

NOTE Confidence: 0.7750263

 $00:18:58.840 \longrightarrow 00:19:00.595$ We see that there's improvement

NOTE Confidence: 0.7750263

 $00:19:00.595 \longrightarrow 00:19:01.648$ in other places.

NOTE Confidence: 0.7750263

00:19:01.650 --> 00:19:04.107 However, if we knock down in OK,

NOTE Confidence: 0.7750263

 $00:19:04.110 \longrightarrow 00:19:05.860$ we still see an improvement.

NOTE Confidence: 0.7750263

 $00:19:05.860 \dashrightarrow 00:19:08.317$ So just by knocking down and OK,

NOTE Confidence: 0.7750263

 $00:19:08.320 \longrightarrow 00:19:11.128$ we see an improvement as we would expect,

NOTE Confidence: 0.7750263

 $00:19:11.130 \longrightarrow 00:19:13.531$ and then the treatment also in combination

 $00:19:13.531 \longrightarrow 00:19:16.304$ with an saw an OK gave about equal

NOTE Confidence: 0.7750263

 $00:19:16.304 \longrightarrow 00:19:18.500$ amount of improvement of earth places.

NOTE Confidence: 0.7750263

 $00:19:18.500 \longrightarrow 00:19:20.678$ So what the suggested to us

NOTE Confidence: 0.7750263

 $00:19:20.678 \longrightarrow 00:19:23.059$ was that the effect of this SD,

NOTE Confidence: 0.7750263

 $00:19:23.060 \longrightarrow 00:19:24.815$ two ages most most likely

NOTE Confidence: 0.7750263

00:19:24.815 --> 00:19:26.570 due primarily to an OK.

NOTE Confidence: 0.7750263

00:19:26.570 --> 00:19:28.970 So pushing up in OK activity.

NOTE Confidence: 0.7750263

 $00:19:28.970 \longrightarrow 00:19:30.465$ There's no effect of Miley

NOTE Confidence: 0.7750263

 $00{:}19{:}30.465 \dashrightarrow 00{:}19{:}31.960$ sales as we previously seen.

NOTE Confidence: 0.75028515

 $00:19:34.370 \longrightarrow 00:19:36.990$ This presentation also within OK

NOTE Confidence: 0.75028515

00:19:36.990 --> 00:19:40.458 Nebbish and improves with pre sis of

NOTE Confidence: 0.75028515

00:19:40.458 --> 00:19:43.498 of our mouse models again which we had

NOTE Confidence: 0.75028515

00:19:43.589 --> 00:19:46.481 shown but turn 19119 increased with

NOTE Confidence: 0.75028515

 $00:19:46.481 \longrightarrow 00:19:49.736$ treatment of our this TGF beta inhibitor.

NOTE Confidence: 0.75028515

00:19:49.736 --> 00:19:53.365 Same with RPL 11 in the fold was

NOTE Confidence: 0.75028515

00:19:53.365 --> 00:19:56.901 different a little bit but two to three

 $00:19:56.901 \longrightarrow 00:20:00.197$ fold two year 424 fold stimulation.

NOTE Confidence: 0.75028515

00:20:00.200 --> 00:20:04.778 A production of rich white cells.

NOTE Confidence: 0.75028515

 $00:20:04.780 \longrightarrow 00:20:07.090$ We we also looked at human.

NOTE Confidence: 0.75028515

00:20:07.090 --> 00:20:09.400 Our cell models, including our PSAT,

NOTE Confidence: 0.75028515

 $00:20:09.400 \longrightarrow 00:20:11.710$ knocked down and 11 knock down.

NOTE Confidence: 0.75028515

 $00{:}20{:}11.710 \dashrightarrow 00{:}20{:}14.790$ We see again Approvement at our DBA cells.

NOTE Confidence: 0.75028515

00:20:14.790 --> 00:20:17.100 No ST208 with SD too late,

NOTE Confidence: 0.75028515

00:20:17.100 --> 00:20:19.530 so we see an improvement anywhere

NOTE Confidence: 0.75028515

 $00:20:19.530 \longrightarrow 00:20:22.798$ from four and a half to about 7

NOTE Confidence: 0.75028515

00:20:22.798 --> 00:20:25.132 fold and then finally a patient

NOTE Confidence: 0.75028515

00:20:25.219 --> 00:20:27.865 sample we because he 235 levels.

NOTE Confidence: 0.75028515

 $00:20:27.870 \longrightarrow 00:20:30.362$ And found where would CSD 208 that

NOTE Confidence: 0.75028515

 $00{:}20{:}30.362 \dashrightarrow 00{:}20{:}32.535$ we see increase in our ourselves

NOTE Confidence: 0.75028515

 $00:20:32.535 \longrightarrow 00:20:34.929$ with red cells and above 2 fold

NOTE Confidence: 0.75028515

 $00:20:35.006 \longrightarrow 00:20:36.788$ increase by quantitating.

 $00:20:36.790 \longrightarrow 00:20:38.730$ So how does this occur?

NOTE Confidence: 0.75028515

 $00:20:38.730 \longrightarrow 00:20:41.458$ Well one of the reasons why we think

NOTE Confidence: 0.75028515

 $00{:}20{:}41.458 \dashrightarrow 00{:}20{:}44.317$ MLK may be important for treatment such

NOTE Confidence: 0.75028515

00:20:44.317 --> 00:20:47.654 as losing is that Lucy and may require

NOTE Confidence: 0.75028515

 $00:20:47.654 \longrightarrow 00:20:50.370$ which is a mino acid as I mentioned

NOTE Confidence: 0.75028515

 $00{:}20{:}50.370 \dashrightarrow 00{:}20{:}52.788$ before in clinical trials that it

NOTE Confidence: 0.75028515

 $00:20:52.788 \longrightarrow 00:20:55.322$ may require active mtor complex and

NOTE Confidence: 0.75028515

 $00:20:55.322 \longrightarrow 00:20:57.866$ pathways to be activated or induced.

NOTE Confidence: 0.75028515

 $00{:}20{:}57.870 \dashrightarrow 00{:}21{:}00.880$ And Raptor me and this whole complex.

NOTE Confidence: 0.75028515

 $00:21:00.880 \longrightarrow 00:21:04.618$ We believe this is attached to lysosome.

NOTE Confidence: 0.75028515

00:21:04.620 --> 00:21:07.956 In the case of an OK phosphorylating Raptor,

NOTE Confidence: 0.75028515

 $00:21:07.960 \longrightarrow 00:21:10.060$ we hypothesize that this released

NOTE Confidence: 0.75028515

00:21:10.060 --> 00:21:12.560 this complex from the license zone,

NOTE Confidence: 0.75028515

 $00:21:12.560 \longrightarrow 00:21:15.250$ thereby enabling translation to occur.

NOTE Confidence: 0.75028515

00:21:15.250 --> 00:21:17.390 So Mark perform evening up

NOTE Confidence: 0.75028515

 $00:21:17.390 \longrightarrow 00:21:19.102$ fluorescent experiments where he

00:21:19.102 --> 00:21:21.067 labeled Raptor with green life,

NOTE Confidence: 0.75028515

 $00{:}21{:}21.070 \dashrightarrow 00{:}21{:}24.024$ some in red and then looked at

NOTE Confidence: 0.75028515

 $00:21:24.024 \longrightarrow 00:21:26.172$ colocalization in the case of

NOTE Confidence: 0.75028515

00:21:26.172 --> 00:21:28.560 RPS Anki knockdown cells where he

NOTE Confidence: 0.75028515

 $00{:}21{:}28.560 \dashrightarrow 00{:}21{:}31.469$ saw was that the Raptor was cold.

NOTE Confidence: 0.75028515

 $00:21:31.470 \longrightarrow 00:21:34.470$ Localising with their lices own.

NOTE Confidence: 0.75028515

 $00:21:34.470 \longrightarrow 00:21:37.473$ And as attached seem to be attached

NOTE Confidence: 0.75028515

 $00:21:37.473 \longrightarrow 00:21:39.699$ or interacting with each other,

NOTE Confidence: 0.75028515

 $00:21:39.700 \longrightarrow 00:21:42.416$ whereas with an OK knock down we

NOTE Confidence: 0.75028515

00:21:42.416 --> 00:21:45.119 don't see wrapped or any longer

NOTE Confidence: 0.75028515

 $00{:}21{:}45.119 \dashrightarrow 00{:}21{:}47.544$ colocalizes with their license home,

NOTE Confidence: 0.75028515

 $00:21:47.550 \longrightarrow 00:21:50.710$ so this is 1 possible role model that

NOTE Confidence: 0.75028515

 $00{:}21{:}50.710 \dashrightarrow 00{:}21{:}53.768$ we propose for which YNLK inhibitors

NOTE Confidence: 0.75028515

00:21:53.768 --> 00:21:57.032 might actually work together with lysine

NOTE Confidence: 0.75028515

 $00:21:57.118 \longrightarrow 00:22:00.168$ to enhance the erythropoietic effect.

 $00:22:00.170 \longrightarrow 00:22:03.074$ And this is another experiment where

NOTE Confidence: 0.75028515

 $00{:}22{:}03.074 \dashrightarrow 00{:}22{:}06.228$ we basically took our cell model RPS

NOTE Confidence: 0.75028515

 $00:22:06.228 \longrightarrow 00:22:09.296$ 19 and looked at the number of two

NOTE Confidence: 0.75028515

 $00:22:09.296 \longrightarrow 00:22:12.112$ CD 235 cells and you can see here.

NOTE Confidence: 0.75028515

 $00:22:12.120 \longrightarrow 00:22:13.764$ So control losing increasing

NOTE Confidence: 0.75028515

00:22:13.764 --> 00:22:16.214 orbital Boyces St 208, again RTF,

NOTE Confidence: 0.75028515

00:22:16.214 --> 00:22:18.626 beta inhibitor and the combination of

NOTE Confidence: 0.75028515

 $00:22:18.626 \longrightarrow 00:22:21.586$ the two seemed to increase even more.

NOTE Confidence: 0.75028515

 $00{:}22{:}21.590 \to 00{:}22{:}23.685$ The effects through synergistic and

NOTE Confidence: 0.75028515

 $00:22:23.685 \longrightarrow 00:22:26.540$ synergism and then the we looked at.

NOTE Confidence: 0.75028515

 $00{:}22{:}26.540 \dashrightarrow 00{:}22{:}29.668$ Also for EBT one which is a downstream

NOTE Confidence: 0.75028515

 $00:22:29.668 \longrightarrow 00:22:32.089$ target of optimizing implore.

NOTE Confidence: 0.75028515

 $00:22:32.090 \longrightarrow 00:22:34.070$ And showed that the phosphorylation

NOTE Confidence: 0.75028515

00:22:34.070 --> 00:22:36.863 was also enhanced when we combine the

NOTE Confidence: 0.75028515

 $00:22:36.863 \longrightarrow 00:22:39.823$ leucine with SD in our PS90 knockdown models.

NOTE Confidence: 0.79829687

 $00:22:41.880 \longrightarrow 00:22:43.970$ So in summary of this,

 $00:22:43.970 \longrightarrow 00:22:46.546$ part of the talk we showed that in

NOTE Confidence: 0.79829687

 $00{:}22{:}46.546 \dashrightarrow 00{:}22{:}49.276$ OK is activated in every projectors

NOTE Confidence: 0.79829687

 $00:22:49.276 \longrightarrow 00:22:52.771$ from DBA patient samples as well as

NOTE Confidence: 0.79829687

 $00:22:52.771 \longrightarrow 00:22:55.255$ our other human and mouse models.

NOTE Confidence: 0.79829687

 $00{:}22{:}55.260 \dashrightarrow 00{:}22{:}57.244$ Pharmacological genetic inhibition of

NOTE Confidence: 0.79829687

 $00:22:57.244 \longrightarrow 00:22:59.724$ NLC increases with regenerx expansion

NOTE Confidence: 0.79829687

 $00:22:59.724 \longrightarrow 00:23:02.593$ in our models and then OK appears to

NOTE Confidence: 0.79829687

 $00:23:02.593 \longrightarrow 00:23:04.479$ exert influence on putting translation

NOTE Confidence: 0.79829687

 $00:23:04.479 \longrightarrow 00:23:06.957$ in DBA through the mtor pathway.

NOTE Confidence: 0.79829687

 $00{:}23{:}06.960 \dashrightarrow 00{:}23{:}09.991$ So our focus was next to begin

NOTE Confidence: 0.79829687

 $00{:}23{:}09.991 \dashrightarrow 00{:}23{:}12.210$ to identify potential the rapies.

NOTE Confidence: 0.79829687

00:23:12.210 --> 00:23:14.430 That would target in OK.

NOTE Confidence: 0.79829687

 $00{:}23{:}14.430 \dashrightarrow 00{:}23{:}16.830$ Unfortunately, the SD 208 compound is

NOTE Confidence: 0.79829687

 $00:23:16.830 \longrightarrow 00:23:19.300$ not ready for clinical application.

NOTE Confidence: 0.79829687

 $00:23:19.300 \longrightarrow 00:23:21.510$ It doesn't have the appropriate

00:23:21.510 --> 00:23:22.836 physical chemical properties,

NOTE Confidence: 0.79829687

00:23:22.840 --> 00:23:23.692 including solubility,

NOTE Confidence: 0.79829687

 $00:23:23.692 \longrightarrow 00:23:27.720$ to be able to be converted to the clinic.

NOTE Confidence: 0.79829687

00:23:27.720 --> 00:23:30.576 And So what we show is that we

NOTE Confidence: 0.79829687

 $00:23:30.576 \longrightarrow 00:23:33.978$ worked with a number of mid chemistry

NOTE Confidence: 0.79829687

00:23:33.978 --> 00:23:36.573 consultants from our Spark program,

NOTE Confidence: 0.79829687

 $00:23:36.580 \longrightarrow 00:23:40.101$ which is a program at Stanford to

NOTE Confidence: 0.79829687

 $00:23:40.101 \longrightarrow 00:23:43.480$ convert projects and lab to the clinics.

NOTE Confidence: 0.79829687

 $00:23:43.480 \longrightarrow 00:23:46.077$ And were able to examine a number

NOTE Confidence: 0.79829687

 $00:23:46.077 \longrightarrow 00:23:47.190$ of these compounds,

NOTE Confidence: 0.79829687

 $00:23:47.190 \longrightarrow 00:23:49.787$ all of which have different primary targets.

NOTE Confidence: 0.79829687

 $00:23:49.790 \longrightarrow 00:23:52.380$ But we tested a number of them.

NOTE Confidence: 0.79829687

 $00:23:52.380 \longrightarrow 00:23:54.060$ This is just a representative

NOTE Confidence: 0.79829687

 $00{:}23{:}54.060 \dashrightarrow 00{:}23{:}55.740$ experiment showing in OK in

NOTE Confidence: 0.79829687

00:23:55.807 --> 00:23:57.579 vitro kinase activity descend,

NOTE Confidence: 0.79829687

00:23:57.580 --> 00:23:59.575 everyone is familiar with which

00:23:59.575 --> 00:24:01.570 is a tyrosine kinase inhibitor

NOTE Confidence: 0.79829687

 $00{:}24{:}01.641 --> 00{:}24{:}03.507$ used to treat CML and carafe,

NOTE Confidence: 0.79829687

 $00:24:03.510 \longrightarrow 00:24:06.478$ and if it would be rough inhibitor OTS.

NOTE Confidence: 0.79829687

00:24:06.480 --> 00:24:09.042 167 is a milk inhibitor suppen

NOTE Confidence: 0.79829687

 $00:24:09.042 \longrightarrow 00:24:11.557$ assertive is an important hitter in

NOTE Confidence: 0.79829687

00:24:11.557 --> 00:24:14.105 St 208 is our TGF beta inhibitor.

NOTE Confidence: 0.79829687

 $00:24:14.110 \longrightarrow 00:24:16.240$ So just to show an example,

NOTE Confidence: 0.79829687

 $00:24:16.240 \longrightarrow 00:24:18.370$ we use low and high concentrations.

NOTE Confidence: 0.79829687

 $00{:}24{:}18.370 \dashrightarrow 00{:}24{:}21.578$ We able to show significant decrease in OK

NOTE Confidence: 0.79829687

 $00:24:21.578 \longrightarrow 00:24:24.539$ activity with this particular drug OTS 167.

NOTE Confidence: 0.79829687

 $00:24:24.540 \longrightarrow 00:24:27.988$ We examined some of these compounds for their

NOTE Confidence: 0.79829687

 $00:24:27.988 \longrightarrow 00:24:31.320$ ability to increased risk of Oasis in RPS,

NOTE Confidence: 0.79829687

 $00:24:31.320 \longrightarrow 00:24:33.440$ 19 lockdown models and here's

NOTE Confidence: 0.79829687

 $00:24:33.440 \longrightarrow 00:24:35.136$ our TGF beta inhibitor.

NOTE Confidence: 0.79829687

 $00:24:35.140 \longrightarrow 00:24:37.260$ We see various levels of

00:24:37.260 --> 00:24:38.956 increase in Rip Oasis,

NOTE Confidence: 0.79829687

 $00:24:38.960 \longrightarrow 00:24:41.498$ but mostly due to RTS 167.

NOTE Confidence: 0.79829687

 $00:24:41.500 \longrightarrow 00:24:43.615$ We also looked for phosphorylation

NOTE Confidence: 0.79829687

00:24:43.615 --> 00:24:44.884 of Raptor again,

NOTE Confidence: 0.79829687

 $00:24:44.890 \longrightarrow 00:24:47.170$ which is the target of which

NOTE Confidence: 0.79829687

 $00:24:47.170 \longrightarrow 00:24:50.001$ is in the Mentor complex and we

NOTE Confidence: 0.79829687

 $00:24:50.001 \longrightarrow 00:24:52.437$ see that there was an increase

NOTE Confidence: 0.79829687

00:24:52.437 --> 00:24:55.129 in phosphorylation of Raptor.

NOTE Confidence: 0.79829687

 $00:24:55.130 \longrightarrow 00:24:59.458$ But decrease with OTS 167 within OK activity.

NOTE Confidence: 0.82792556

 $00:25:02.000 \longrightarrow 00:25:04.382$ So our lead compound we decided

NOTE Confidence: 0.82792556

 $00:25:04.382 \longrightarrow 00:25:06.739$ to focus on was OTS 167.

NOTE Confidence: 0.82792556

 $00:25:06.740 \longrightarrow 00:25:08.720$ This is a milk inhibitor.

NOTE Confidence: 0.82792556

 $00:25:08.720 \longrightarrow 00:25:10.775$ It's currently under being studied

NOTE Confidence: 0.82792556

 $00:25:10.775 \longrightarrow 00:25:13.267$ in Phase 1/2 clinical trials and

NOTE Confidence: 0.82792556

00:25:13.267 --> 00:25:15.092 particularly in advance to acute

NOTE Confidence: 0.82792556

 $00:25:15.092 \longrightarrow 00:25:17.410$ leukemia as well as lung cancer.

00:25:17.410 --> 00:25:20.161 The drug was developed by uncle with

NOTE Confidence: 0.82792556

 $00:25:20.161 \longrightarrow 00:25:22.150$ therapy scientists and their bins.

NOTE Confidence: 0.82792556

 $00:25:22.150 \longrightarrow 00:25:25.382$ Number of studies that show that drug is

NOTE Confidence: 0.82792556

00:25:25.382 --> 00:25:28.070 still effective when milk is knocked out,

NOTE Confidence: 0.82792556

 $00{:}25{:}28.070 \longrightarrow 00{:}25{:}32.298$ so there are clearly other targets involved.

NOTE Confidence: 0.82792556

00:25:32.300 --> 00:25:35.177 This compound was shown in RP slinky

NOTE Confidence: 0.82792556

00:25:35.177 --> 00:25:37.308 knockdown cells again to increase

NOTE Confidence: 0.82792556

 $00:25:37.308 \longrightarrow 00:25:39.363$ every thread expansion and when

NOTE Confidence: 0.82792556

00:25:39.363 --> 00:25:41.775 we combine our knockdown oven OK

NOTE Confidence: 0.82792556

 $00{:}25{:}41.775 \dashrightarrow 00{:}25{:}44.351$ with this inhibitor we did not see

NOTE Confidence: 0.82792556

 $00{:}25{:}44.360 \dashrightarrow 00{:}25{:}46.348$ significant increase in irithyll.

NOTE Confidence: 0.82792556

 $00:25:46.348 \longrightarrow 00:25:49.771$ Police is suggesting to us again that

NOTE Confidence: 0.82792556

 $00{:}25{:}49.771 \dashrightarrow 00{:}25{:}52.298$ the primary target of OTS in this

NOTE Confidence: 0.82792556

 $00:25:52.298 \longrightarrow 00:25:55.089$ system to improve it or places in OK

NOTE Confidence: 0.82792556

 $00:25:55.089 \longrightarrow 00:25:57.956$ and this drug was dosed at 200 animal

 $00:25:57.956 \longrightarrow 00:26:00.840$ or every three days for one cycle.

NOTE Confidence: 0.74062574

 $00:26:04.120 \longrightarrow 00:26:06.418$ To Sir to to understand whether

NOTE Confidence: 0.74062574

00:26:06.418 --> 00:26:08.460 there there was toxicity

NOTE Confidence: 0.74062574

 $00:26:08.460 \longrightarrow 00:26:10.630$ to normal or through blast.

NOTE Confidence: 0.74062574

00:26:10.630 --> 00:26:12.665 We looked at every expansion

NOTE Confidence: 0.74062574

 $00:26:12.665 \longrightarrow 00:26:14.700$ as well as DBA cells.

NOTE Confidence: 0.74062574

00:26:14.700 --> 00:26:17.584 Knockdown cells with artist 19 we see

NOTE Confidence: 0.74062574

 $00:26:17.584 \longrightarrow 00:26:20.806$ that the maximum effect was at 300 animal,

NOTE Confidence: 0.74062574

 $00{:}26{:}20.810 \dashrightarrow 00{:}26{:}23.898$ but we begin to see an effect in

NOTE Confidence: 0.74062574

00:26:23.898 --> 00:26:26.907 North places as early as 30 nanomolar,

NOTE Confidence: 0.74062574

 $00{:}26{:}26.910 \dashrightarrow 00{:}26{:}28.538$ which shows greater than

NOTE Confidence: 0.74062574

 $00:26:28.538 \longrightarrow 00:26:29.759$ tenfold therapeutic window.

NOTE Confidence: 0.74062574

 $00:26:29.760 \longrightarrow 00:26:31.950$ Since the IC50 and normal.

NOTE Confidence: 0.74062574

 $00{:}26{:}31.950 \dashrightarrow 00{:}26{:}34.435$ Or healthy with Glass was

NOTE Confidence: 0.74062574

 $00:26:34.435 \longrightarrow 00:26:36.423$ about 480 animal are.

NOTE Confidence: 0.74062574

 $00:26:36.430 \longrightarrow 00:26:38.901$ With my lead cells we saw slightly

00:26:38.901 --> 00:26:40.939 more sensitivity of this compound.

NOTE Confidence: 0.74062574

 $00{:}26{:}40.940 --> 00{:}26{:}44.369$ Again, we can see 30 an animal or an

NOTE Confidence: 0.74062574

 $00:26:44.369 \longrightarrow 00:26:46.958$ increase in it with the police is,

NOTE Confidence: 0.74062574

 $00:26:46.960 \longrightarrow 00:26:48.832$ but then the myeloid cell we

NOTE Confidence: 0.74062574

 $00:26:48.832 \longrightarrow 00:26:50.638$ began to see decreased more

NOTE Confidence: 0.74062574

 $00:26:50.638 \longrightarrow 00:26:52.598$ significantly around 300 nanomolar,

NOTE Confidence: 0.74062574

 $00:26:52.600 \longrightarrow 00:26:54.910$ so Even so we believe the

NOTE Confidence: 0.74062574

 $00{:}26{:}54.910 \dashrightarrow 00{:}26{:}57.110$ the rapeutic window is about 10 fold.

NOTE Confidence: 0.74062574

 $00{:}26{:}57.110 \dashrightarrow 00{:}27{:}02.195$ And again, this is an in vitro assay system.

NOTE Confidence: 0.74062574

00:27:02.200 --> 00:27:03.466 So in conclusion,

NOTE Confidence: 0.74062574

 $00{:}27{:}03.466 \dashrightarrow 00{:}27{:}05.998$ Altius 167 appears to improve our

NOTE Confidence: 0.74062574

 $00{:}27{:}05.998 \dashrightarrow 00{:}27{:}08.637$ ethical thesis in our DBA models in

NOTE Confidence: 0.74062574

 $00{:}27{:}08.637 \dashrightarrow 00{:}27{:}10.790$ vitro with very little toxicity.

NOTE Confidence: 0.74062574

 $00:27:10.790 \longrightarrow 00:27:12.965$ There have been previous reports

NOTE Confidence: 0.74062574

 $00:27:12.965 \longrightarrow 00:27:15.654$ in Nora Blastoma and breast cancer

 $00:27:15.654 \longrightarrow 00:27:17.394$ Xenografted mouse models treated

NOTE Confidence: 0.74062574

 $00:27:17.394 \longrightarrow 00:27:20.671$ with OTS 167 twice a week for three

NOTE Confidence: 0.74062574

00:27:20.671 --> 00:27:23.812 weeks over the course of a month or so,

NOTE Confidence: 0.74062574

 $00:27:23.812 \longrightarrow 00:27:26.068$ and none of those mice developed

NOTE Confidence: 0.74062574

 $00:27:26.068 \longrightarrow 00:27:27.560$ bone marrow toxicity.

NOTE Confidence: 0.74062574

 $00{:}27{:}27.560 \dashrightarrow 00{:}27{:}30.050$ There are also other inhibitors to

NOTE Confidence: 0.74062574

 $00:27:30.050 \longrightarrow 00:27:31.710$ indicate that potentially they.

NOTE Confidence: 0.74062574

 $00:27:31.710 \longrightarrow 00:27:33.186$ May be effective,

NOTE Confidence: 0.74062574

 $00{:}27{:}33.186 \dashrightarrow 00{:}27{:}35.154$ including John Conyers inhibitors

NOTE Confidence: 0.74062574

 $00:27:35.154 \longrightarrow 00:27:37.534$ and we're currently testing those

NOTE Confidence: 0.74062574

 $00{:}27{:}37.534 \dashrightarrow 00{:}27{:}39.614$ and finally experiments to really

NOTE Confidence: 0.74062574

00:27:39.614 --> 00:27:41.906 test this is necessary in vivo

NOTE Confidence: 0.74062574

 $00:27:41.906 \longrightarrow 00:27:44.181$ in order for us to proceed to

NOTE Confidence: 0.74062574

 $00:27:44.190 \longrightarrow 00:27:45.438$ clinical trial stage.

NOTE Confidence: 0.79584086

 $00:27:47.720 \longrightarrow 00:27:48.956$ Another interesting observation

NOTE Confidence: 0.79584086

 $00:27:48.956 \longrightarrow 00:27:51.840$ that Mark made as far as Enoch

 $00:27:51.912 \longrightarrow 00:27:54.438$ expression is the fact that metformin,

NOTE Confidence: 0.79584086

 $00:27:54.440 \longrightarrow 00:27:56.120$ the commonly used medication

NOTE Confidence: 0.79584086

 $00:27:56.120 \longrightarrow 00:27:57.800$ for type 2 diabetes,

NOTE Confidence: 0.79584086

00:27:57.800 --> 00:28:00.320 inhibits an OK expression in small

NOTE Confidence: 0.79584086

 $00:28:00.320 \longrightarrow 00:28:02.000$ cell lung cancer cells,

NOTE Confidence: 0.79584086

 $00:28:02.000 \longrightarrow 00:28:04.478$ and this drug also improves the

NOTE Confidence: 0.79584086

00:28:04.478 --> 00:28:06.620 effect of hematopoiesis and delays

NOTE Confidence: 0.79584086

 $00:28:06.620 \longrightarrow 00:28:08.300$ tumors in Fanconi mice.

NOTE Confidence: 0.79584086

 $00:28:08.300 \longrightarrow 00:28:10.988$ This has been reported to be and metformin

NOTE Confidence: 0.79584086

00:28:10.988 --> 00:28:13.758 to be protective against aldehydes,

NOTE Confidence: 0.79584086

 $00:28:13.760 \longrightarrow 00:28:16.544$ which is one of the toxins

NOTE Confidence: 0.79584086

 $00:28:16.544 \longrightarrow 00:28:18.400$ thought to affect inhibit.

NOTE Confidence: 0.79584086

 $00:28:18.400 \longrightarrow 00:28:20.619$ He met up with stem cells and

NOTE Confidence: 0.79584086

 $00:28:20.619 \longrightarrow 00:28:22.260$ he's in this disease.

NOTE Confidence: 0.79584086

 $00:28:22.260 \longrightarrow 00:28:24.336$ There is currently a phase two

 $00:28:24.336 \longrightarrow 00:28:26.274$ trial with metformin in Fanconi

NOTE Confidence: 0.79584086

 $00{:}28{:}26.274 \dashrightarrow 00{:}28{:}28.709$ patients that's being directed by

NOTE Confidence: 0.79584086

 $00:28:28.709 \longrightarrow 00:28:30.657$ Akiko Shimamura Boston Children's.

NOTE Confidence: 0.79584086

 $00:28:30.660 \longrightarrow 00:28:33.495$ So what is the mechanism by which in Oak

NOTE Confidence: 0.79584086

 $00:28:33.495 \longrightarrow 00:28:35.488$ expression is inhibited by metformin?

NOTE Confidence: 0.79584086

 $00:28:35.490 \longrightarrow 00:28:37.602$ Well, one of the things we

NOTE Confidence: 0.79584086

 $00:28:37.602 \longrightarrow 00:28:39.629$ looked at was first of all,

NOTE Confidence: 0.79584086

 $00:28:39.630 \longrightarrow 00:28:41.615$ just metformin improve the cell

NOTE Confidence: 0.79584086

 $00:28:41.615 \longrightarrow 00:28:43.947$ numbers of C235 cells and we

NOTE Confidence: 0.79584086

 $00:28:43.947 \longrightarrow 00:28:45.837$ show that both in RPS 19 RP.

NOTE Confidence: 0.79584086

 $00:28:45.840 \longrightarrow 00:28:47.802$ 11 Knocked down models that it

NOTE Confidence: 0.79584086

 $00:28:47.802 \longrightarrow 00:28:49.110$ does increase the production

NOTE Confidence: 0.79584086

00:28:49.171 --> 00:28:50.667 of these research senators.

NOTE Confidence: 0.79584086

 $00:28:50.670 \longrightarrow 00:28:51.360$ In contrast,

NOTE Confidence: 0.79584086

 $00:28:51.360 \longrightarrow 00:28:53.430$ there's no effect of moderate cells.

NOTE Confidence: 0.79584086

 $00:28:53.430 \longrightarrow 00:28:55.160$ We don't see any phenotype,

 $00:28:55.160 \longrightarrow 00:28:57.920$ and then it would be a few economy.

NOTE Confidence: 0.79584086

 $00:28:57.920 \longrightarrow 00:28:59.640$ Or if you eat colonies,

NOTE Confidence: 0.79584086

 $00:28:59.640 \longrightarrow 00:29:01.720$ we see an increased number.

NOTE Confidence: 0.79584086

 $00:29:01.720 \longrightarrow 00:29:06.346$ With metformin. In contrast to controls.

NOTE Confidence: 0.79584086

 $00:29:06.350 \longrightarrow 00:29:09.910$ CD 235 arthritis sales.

NOTE Confidence: 0.79584086

 $00:29:09.910 \longrightarrow 00:29:11.610$ Metformin increases by five or

NOTE Confidence: 0.79584086

 $00:29:11.610 \longrightarrow 00:29:13.985$ six fold two to six fold and

NOTE Confidence: 0.79584086

 $00:29:13.985 \longrightarrow 00:29:16.078$ then be a few mirrors for it's

NOTE Confidence: 0.79584086

 $00:29:16.078 \longrightarrow 00:29:18.089$ also about two or three fold.

NOTE Confidence: 0.79584086

 $00{:}29{:}18.090 \dashrightarrow 00{:}29{:}19.725$ We've seen increasing without no

NOTE Confidence: 0.79584086

 $00:29:19.725 \longrightarrow 00:29:21.360$ effect on the mileage ourselves.

NOTE Confidence: 0.744118

 $00:29:23.780 \longrightarrow 00:29:26.315$ The metform in also improves erythropoiesis

NOTE Confidence: 0.744118

 $00{:}29{:}26.315 \dashrightarrow 00{:}29{:}28.343$ by inhibiting anoche activity.

NOTE Confidence: 0.744118

00:29:28.350 --> 00:29:31.194 Here we have again in OK

NOTE Confidence: 0.744118

 $00:29:31.194 \longrightarrow 00:29:34.449$ phosphorylation of in OK Mabel Rector.

 $00:29:34.450 \longrightarrow 00:29:38.250$ In all cases we see that the metformin

NOTE Confidence: 0.744118

 $00:29:38.250 \longrightarrow 00:29:42.659$ as well as SD 208 TGF beta inhibitor

NOTE Confidence: 0.744118

 $00:29:42.659 \longrightarrow 00:29:45.619$ decreases the activity of a van.

NOTE Confidence: 0.744118

00:29:45.620 --> 00:29:48.668 OK in our PS90 knockdown model.

NOTE Confidence: 0.744118

 $00{:}29{:}48.670 \dashrightarrow 00{:}29{:}51.240$ The RNA expression also decreases

NOTE Confidence: 0.744118

 $00:29:51.240 \longrightarrow 00:29:53.810$ which is interesting which is.

NOTE Confidence: 0.744118

 $00:29:53.810 \longrightarrow 00:29:57.030$ Primary mechanism by which we

NOTE Confidence: 0.744118

00:29:57.030 --> 00:30:00.250 believe metformin inhibits an OK.

NOTE Confidence: 0.744118

 $00{:}30{:}00.250 \dashrightarrow 00{:}30{:}02.080$ An orith Rd expansion we see

NOTE Confidence: 0.744118

 $00:30:02.080 \longrightarrow 00:30:04.180$ again with knockdown of RPS 19.

NOTE Confidence: 0.744118

00:30:04.180 --> 00:30:06.908 Any cells and knocked down again OK at

NOTE Confidence: 0.744118

 $00:30:06.908 \longrightarrow 00:30:09.745$ the same time we see that just knock down

NOTE Confidence: 0.744118

 $00:30:09.745 \longrightarrow 00:30:12.748$ of in OK and improves mini expansion.

NOTE Confidence: 0.744118

 $00:30:12.750 \longrightarrow 00:30:16.017$ There is replaces but also just say it not

NOTE Confidence: 0.744118

 $00:30:16.017 \longrightarrow 00:30:19.167$ kind of in OK here as well as metformin.

NOTE Confidence: 0.744118

 $00{:}30{:}19.170 \dashrightarrow 00{:}30{:}21.102$ So metform in not done OK in the

 $00{:}30{:}21.102 \dashrightarrow 00{:}30{:}22.915$ form it again no significant

NOTE Confidence: 0.744118

 $00{:}30{:}22.915 \dashrightarrow 00{:}30{:}25.215$ change which again suggested some

NOTE Confidence: 0.744118

00:30:25.215 --> 00:30:27.030 informants working through an OK.

NOTE Confidence: 0.69575894

 $00:30:30.090 \longrightarrow 00:30:30.860$ Sorry.

NOTE Confidence: 0.7946285

 $00:30:33.190 \longrightarrow 00:30:36.032$ This is a slide which shows our

NOTE Confidence: 0.7946285

00:30:36.032 --> 00:30:38.299 treatment of zebrafish models in

NOTE Confidence: 0.7946285

00:30:38.299 --> 00:30:40.719 collaboration with Scholin at UCLA.

NOTE Confidence: 0.7946285

 $00:30:40.720 \longrightarrow 00:30:42.935$ He created a DBA model

NOTE Confidence: 0.7946285

00:30:42.935 --> 00:30:44.707 using RPS 19 morpholinos.

NOTE Confidence: 0.7946285

 $00:30:44.710 \longrightarrow 00:30:47.804$ This is a phenotype of the fish.

NOTE Confidence: 0.7946285

 $00{:}30{:}47.810 \dashrightarrow 00{:}30{:}50.512$ After approximately 5 days you can see

NOTE Confidence: 0.7946285

 $00:30:50.512 \dashrightarrow 00:30:52.669$ that these embryos show significant

NOTE Confidence: 0.7946285

 $00{:}30{:}52.669 \dashrightarrow 00{:}30{:}54.974$ anemia compared to controls with

NOTE Confidence: 0.7946285

 $00:30:54.974 \longrightarrow 00:30:57.110$ the treatment with metformin.

NOTE Confidence: 0.7946285

 $00:30:57.110 \longrightarrow 00:30:59.994$ We see that there is again increase

 $00:30:59.994 \longrightarrow 00:31:02.292$ in risk reduction as indicated

NOTE Confidence: 0.7946285

 $00:31:02.292 \longrightarrow 00:31:04.737$ by staining with Odeon sitting.

NOTE Confidence: 0.7946285

 $00:31:04.740 \longrightarrow 00:31:06.908$ Which binds to hemoglobin.

NOTE Confidence: 0.82489973

 $00:31:09.400 \longrightarrow 00:31:11.260$ So how does metformin

NOTE Confidence: 0.82489973

00:31:11.260 --> 00:31:13.120 regulate in OK expression?

NOTE Confidence: 0.82489973

00:31:13.120 --> 00:31:16.840 One of the ideas is through micro RNAs,

NOTE Confidence: 0.82489973

 $00:31:16.840 \longrightarrow 00:31:20.032$ and so Mark created a number of

NOTE Confidence: 0.82489973

 $00{:}31{:}20.032 \dashrightarrow 00{:}31{:}22.452$ truncation mutants that would include

NOTE Confidence: 0.82489973

 $00{:}31{:}22.452 \dashrightarrow 00{:}31{:}25.338$ include a variety of micro ironies

NOTE Confidence: 0.82489973

 $00:31:25.338 \longrightarrow 00:31:29.029$ for which we can then try to identify

NOTE Confidence: 0.82489973

 $00{:}31{:}29.029 \dashrightarrow 00{:}31{:}31.446$ the specific mechanism by which

NOTE Confidence: 0.82489973

 $00:31:31.446 \longrightarrow 00:31:34.726$ metformin effects in OK expression.

NOTE Confidence: 0.82489973

 $00:31:34.730 \longrightarrow 00:31:37.915$ So this is showing the levels in

NOTE Confidence: 0.82489973

 $00:31:37.915 \longrightarrow 00:31:41.514$ humans that mirror 30 a mere 26

NOTE Confidence: 0.82489973

 $00:31:41.514 \longrightarrow 00:31:43.654$ increases with metform in treatment.

NOTE Confidence: 0.82489973

00:31:43.660 --> 00:31:47.062 So this is again in human primary

 $00:31:47.062 \longrightarrow 00:31:50.604$ cells versus mice that do not show

NOTE Confidence: 0.82489973

 $00{:}31{:}50.604 \dashrightarrow 00{:}31{:}53.580$ this increase in May 26 expression.

NOTE Confidence: 0.82489973

 $00:31:53.580 \longrightarrow 00:31:57.476$ So the idea here is metform in could be

NOTE Confidence: 0.82489973

 $00:31:57.476 \longrightarrow 00:32:00.704$ inducing near 26 which binds to the

NOTE Confidence: 0.82489973

 $00{:}32{:}00.704 \dashrightarrow 00{:}32{:}03.544$ three prime UTR event OK resulting

NOTE Confidence: 0.82489973

 $00:32:03.544 \longrightarrow 00:32:07.078$ in inhibition of by of expression.

NOTE Confidence: 0.82489973

 $00:32:07.080 \longrightarrow 00:32:09.774$ So this is a luciferase constant

NOTE Confidence: 0.82489973

 $00:32:09.774 \dashrightarrow 00:32:11.982$ Reporter construct showing activity as

NOTE Confidence: 0.82489973

 $00:32:11.982 \longrightarrow 00:32:14.404$ a reflection of transcription of in OK.

NOTE Confidence: 0.82489973

 $00{:}32{:}14.410 \dashrightarrow 00{:}32{:}17.281$ You can see that with the R in micro

NOTE Confidence: 0.82489973

 $00:32:17.281 \longrightarrow 00:32:21.028$ RNA 26 mimetic that there is we see

NOTE Confidence: 0.82489973

 $00:32:21.028 \longrightarrow 00:32:22.957$ decreased transcription also with

NOTE Confidence: 0.82489973

 $00{:}32{:}22.957 \dashrightarrow 00{:}32{:}25.806$ 181 which is in the same region

NOTE Confidence: 0.82489973

 $00:32:25.806 \longrightarrow 00:32:28.248$ the truncation mutant as mere 26

NOTE Confidence: 0.82489973

 $00:32:28.248 \longrightarrow 00:32:30.690$ we see a decrease in me.

 $00:32:30.690 \longrightarrow 00:32:31.908$ Yeah can you

NOTE Confidence: 0.77215475

 $00:32:31.910 \longrightarrow 00:32:35.158$ clarify what you mean by a mere 26

NOTE Confidence: 0.77215475

 $00:32:35.160 \longrightarrow 00:32:37.310$ mimetic? It's it's a similar.

NOTE Confidence: 0.77215475

 $00:32:37.310 \longrightarrow 00:32:39.158$ The structure, as a mere 26,

NOTE Confidence: 0.77215475

 $00:32:39.160 \longrightarrow 00:32:41.635$ so it's acting as if it were a mere

NOTE Confidence: 0.77215475

00:32:41.635 --> 00:32:44.730 26 and binding to that site, so I hope

NOTE Confidence: 0.8210358

 $00:32:44.730 \longrightarrow 00:32:46.524$ it's done with like with the

NOTE Confidence: 0.8210358

 $00:32:46.524 \longrightarrow 00:32:48.430$ retrovirus like with a hairpin or.

NOTE Confidence: 0.8210358

 $00{:}32{:}48.430 --> 00{:}32{:}51.470$ Yes, I believe so, yeah.

NOTE Confidence: 0.8210358

 $00:32:51.470 \longrightarrow 00:32:53.942$ And the same thing with an OK if

NOTE Confidence: 0.8210358

 $00{:}32{:}53.942 \dashrightarrow 00{:}32{:}56.635$ we we see that in OK expression

NOTE Confidence: 0.8210358

 $00:32:56.635 \longrightarrow 00:32:59.530$ at the protein level is decreased.

NOTE Confidence: 0.8210358

 $00:32:59.530 \longrightarrow 00:33:02.236$ But a sponge which basically is

NOTE Confidence: 0.8210358

 $00:33:02.236 \longrightarrow 00:33:04.747$ exactly what it described it in

NOTE Confidence: 0.8210358

 $00:33:04.747 \longrightarrow 00:33:07.219$ no longer allows me or 26 to bind

NOTE Confidence: 0.8210358

 $00:33:07.300 \longrightarrow 00:33:09.520$ to the three point Muti UTR.

 $00:33:09.520 \longrightarrow 00:33:11.860$ We see that there is again

NOTE Confidence: 0.8210358

 $00:33:11.860 \longrightarrow 00:33:14.483$ increase in OK at the Mr. NY.

NOTE Confidence: 0.8210358

00:33:14.483 --> 00:33:16.601 Also prefer at the Reporter assay

NOTE Confidence: 0.8210358

 $00:33:16.601 \longrightarrow 00:33:18.729$ and also the protein level.

NOTE Confidence: 0.8143553

 $00:33:19.730 \longrightarrow 00:33:22.786$ And also the Mirror 181 on the top

NOTE Confidence: 0.8143553

 $00:33:22.786 \longrightarrow 00:33:25.460$ didn't affect, but on the bottom did

NOTE Confidence: 0.8143553

 $00:33:25.460 \longrightarrow 00:33:27.370$ what's going on with that?

NOTE Confidence: 0.8143553

 $00:33:27.370 \longrightarrow 00:33:29.280$ Let me see for here.

NOTE Confidence: 0.8143553

 $00:33:29.280 \longrightarrow 00:33:32.272$ Yeah, I mean that's a good question whether

NOTE Confidence: 0.8143553

 $00:33:32.272 \longrightarrow 00:33:35.511$ the mere 181 it may not be a specific

NOTE Confidence: 0.8143553

00:33:35.511 --> 00:33:38.070 so that it doesn't sound consistent,

NOTE Confidence: 0.8143553

 $00{:}33{:}38.070 \dashrightarrow 00{:}33{:}39.985$ because you would expect me

NOTE Confidence: 0.8143553

 $00{:}33{:}39.985 \to 00{:}33{:}42.270$ or 181 to increase there too.

NOTE Confidence: 0.8143553

 $00:33:42.270 \longrightarrow 00:33:44.878$ So you know, not all the mirror when

NOTE Confidence: 0.8143553

00:33:44.878 --> 00:33:48.008 he I guess they could be inhibiting

 $00:33:48.008 \longrightarrow 00:33:50.398$ or blocking other mere sites.

NOTE Confidence: 0.8143553

 $00:33:50.400 \longrightarrow 00:33:51.575$ So, but that's something we

NOTE Confidence: 0.8143553

 $00:33:51.575 \longrightarrow 00:33:53.290$ need to look at more carefully,

NOTE Confidence: 0.8143553

 $00:33:53.290 \longrightarrow 00:33:55.900$ but thanks for noticing, yeah.

NOTE Confidence: 0.8143553

 $00:33:55.900 \longrightarrow 00:33:57.800$ An Emmy this metform in also

NOTE Confidence: 0.8143553

 $00{:}33{:}57.800 \dashrightarrow 00{:}33{:}58.940$ mediates everything voices,

NOTE Confidence: 0.8143553

 $00:33:58.940 \longrightarrow 00:34:01.523$ so again we see the C235 increase

NOTE Confidence: 0.8143553

 $00{:}34{:}01.523 \dashrightarrow 00{:}34{:}04.260$ with the mimetic that Foreman or both.

NOTE Confidence: 0.8712816

 $00{:}34{:}05.820 \dashrightarrow 00{:}34{:}06.924$ Finding interrupted right

NOTE Confidence: 0.8712816

 $00:34:06.924 \longrightarrow 00:34:08.028$ 'cause they're happening.

NOTE Confidence: 0.8712816

 $00:34:08.030 \longrightarrow 00:34:10.620$ So there's a question from the audience.

NOTE Confidence: 0.8712816

 $00:34:10.620 \longrightarrow 00:34:12.490$ Is the difference of metformin

NOTE Confidence: 0.8712816

00:34:12.490 --> 00:34:14.360 effect on human versus mouse

NOTE Confidence: 0.8712816

 $00:34:14.431 \longrightarrow 00:34:16.650$ near 26 due to changes in mere

NOTE Confidence: 0.8712816

00:34:16.650 --> 00:34:18.370 26 transcription or processing?

NOTE Confidence: 0.8712816

 $00{:}34{:}18.370 \dashrightarrow 00{:}34{:}20.578$ Yeah, that's a really good question.

 $00:34:20.580 \longrightarrow 00:34:23.156$ I don't think we really know that.

NOTE Confidence: 0.8712816

 $00:34:23.160 \longrightarrow 00:34:25.380$ We haven't really focused on the

NOTE Confidence: 0.8712816

 $00:34:25.380 \longrightarrow 00:34:28.064$ mouse system, but we do know that it's

NOTE Confidence: 0.8712816

 $00:34:28.064 \longrightarrow 00:34:30.653$ different and you know that there are

NOTE Confidence: 0.8712816

 $00:34:30.653 \longrightarrow 00:34:33.095$ clearly differences at the genomic level

NOTE Confidence: 0.8712816

 $00:34:33.095 \longrightarrow 00:34:35.707$ and that the exact reason for that.

NOTE Confidence: 0.8712816

 $00:34:35.710 \longrightarrow 00:34:38.734$ We're not sure. So it's a good.

NOTE Confidence: 0.8712816

00:34:38.740 --> 00:34:42.120 It's a great question, yeah?

NOTE Confidence: 0.8712816

 $00:34:42.120 \longrightarrow 00:34:44.922$ So both in the mouses are

NOTE Confidence: 0.8712816

 $00:34:44.922 \longrightarrow 00:34:46.790$ in the mouse system.

NOTE Confidence: 0.8712816

 $00:34:46.790 \longrightarrow 00:34:50.984$ We see an increase turn 119 and also 235.

NOTE Confidence: 0.8712816

 $00{:}34{:}50{.}990 \dashrightarrow 00{:}34{:}54{.}502$ We see increase in CD 235 R 3119

NOTE Confidence: 0.8712816

00:34:54.502 --> 00:34:57.691 the expression as you would expect

NOTE Confidence: 0.8712816

 $00{:}34{:}57.691 \dashrightarrow 00{:}35{:}00.451$ goes down with metform in treatment

NOTE Confidence: 0.8712816

00:35:00.451 --> 00:35:03.719 here in our PS90 knockdown cells.

 $00:35:03.720 \longrightarrow 00:35:04.899$ So, to summarize,

NOTE Confidence: 0.8712816

 $00:35:04.899 \longrightarrow 00:35:06.471$ metformin improves with crisis

NOTE Confidence: 0.8712816

 $00:35:06.471 \longrightarrow 00:35:08.160$ in our models of DPA.

NOTE Confidence: 0.8712816

 $00:35:08.160 \longrightarrow 00:35:10.152$ It decreases in OK expression through

NOTE Confidence: 0.8712816

 $00:35:10.152 \longrightarrow 00:35:12.760$ mere 26 A and targeting mirrors is

NOTE Confidence: 0.8712816

00:35:12.760 --> 00:35:15.190 a possible approach to DBA therapy.

NOTE Confidence: 0.8712816

 $00{:}35{:}15.190 \dashrightarrow 00{:}35{:}16.765$ There now companies that are

NOTE Confidence: 0.8712816

 $00:35:16.765 \longrightarrow 00:35:18.896$ trying to make medics or sponges

NOTE Confidence: 0.8712816

 $00{:}35{:}18.896 \dashrightarrow 00{:}35{:}20.369$ for clinical application.

NOTE Confidence: 0.8712816

 $00:35:20.370 \longrightarrow 00:35:22.590$ Although it's very in early stages.

NOTE Confidence: 0.80600363

 $00{:}35{:}25.020 \dashrightarrow 00{:}35{:}28.296$ OK, for the last part of the talk I'm

NOTE Confidence: 0.80600363

 $00:35:28.296 \longrightarrow 00:35:32.068$ going to focus on a new project that I

NOTE Confidence: 0.80600363

 $00:35:32.068 \longrightarrow 00:35:34.680$ appreciate Diane and Vanessa's input.

NOTE Confidence: 0.80600363

 $00:35:34.680 \longrightarrow 00:35:37.688$ This is a protein set B1 which is

NOTE Confidence: 0.80600363

00:35:37.688 --> 00:35:40.306 special 80 rich binding protein one

NOTE Confidence: 0.80600363

 $00{:}35{:}40.306 \dashrightarrow 00{:}35{:}43.920$ and we initially did a many years ago.

 $00:35:43.920 \longrightarrow 00:35:46.950$ Actually did a RNA seek experiment

NOTE Confidence: 0.80600363

 $00:35:46.950 \dashrightarrow 00:35:50.496$ with fetal liver human CD 34 positive

NOTE Confidence: 0.80600363

 $00:35:50.496 \longrightarrow 00:35:53.842$ cells that were transduced with RPS 19.

NOTE Confidence: 0.80600363

00:35:53.850 --> 00:35:56.130 Like if I were constructs and found 560

NOTE Confidence: 0.80600363

00:35:56.130 --> 00:35:58.469 or so genes that are differentially

NOTE Confidence: 0.80600363

 $00:35:58.469 \longrightarrow 00:36:01.464$ expressed and then we cross reference that

NOTE Confidence: 0.80600363

 $00:36:01.464 \longrightarrow 00:36:04.124$ with the list of genes identifying earlier.

NOTE Confidence: 0.80600363

 $00:36:04.130 \longrightarrow 00:36:06.643$ It'll poesis in a paper published by

NOTE Confidence: 0.80600363

 $00:36:06.643 \longrightarrow 00:36:09.436$ Mohan Orla and also Pat Gallagher and

NOTE Confidence: 0.80600363

 $00{:}36{:}09.436 {\:\dashrightarrow\:} 00{:}36{:}12.281$ found about 1700 genes over lapping were

NOTE Confidence: 0.80600363

 $00:36:12.281 \longrightarrow 00:36:14.500$ about 42 genes that we then analyze

NOTE Confidence: 0.80600363

 $00:36:14.500 \longrightarrow 00:36:17.654$ in a variety of our cell model systems

NOTE Confidence: 0.80600363

 $00{:}36{:}17.654 \dashrightarrow 00{:}36{:}20.094$ and hematopoietic stem cells both in

NOTE Confidence: 0.80600363

00:36:20.094 --> 00:36:22.164 control an RPS 19 knockdown cells

NOTE Confidence: 0.80600363

 $00:36:22.164 \longrightarrow 00:36:24.748$ and found that among those that were.

 $00:36:24.750 \longrightarrow 00:36:25.600$ Mostly regulated,

NOTE Confidence: 0.80600363

 $00{:}36{:}25.600 {\:{\circ}{\circ}{\circ}}>00{:}36{:}28.575$ the Sepy one was very interesting to

NOTE Confidence: 0.80600363

 $00:36:28.575 \longrightarrow 00:36:31.180$ us since not much had been described

NOTE Confidence: 0.80600363

 $00:36:31.180 \longrightarrow 00:36:34.688$ at all on the role of sappy one during

NOTE Confidence: 0.80600363

 $00:36:34.688 \longrightarrow 00:36:38.100$ Aritha Poesis So sappy one is a

NOTE Confidence: 0.80600363

00:36:38.100 --> 00:36:40.700 protein that basically forms chromatin

NOTE Confidence: 0.80600363

 $00:36:40.793 \longrightarrow 00:36:43.949$ loops and regulates transcription.

NOTE Confidence: 0.80600363

00:36:43.950 --> 00:36:46.155 And there's a number of their number

NOTE Confidence: 0.80600363

 $00{:}36{:}46.155 \dashrightarrow 00{:}36{:}48.729$ of papers that have very nicely

NOTE Confidence: 0.80600363

 $00:36:48.729 \longrightarrow 00:36:51.329$ described its expression and rolling

NOTE Confidence: 0.80600363

 $00:36:51.329 \longrightarrow 00:36:52.369$ hematopoietic differentiation.

NOTE Confidence: 0.80600363

 $00:36:52.370 \longrightarrow 00:36:54.375$ It's moderately expressed in HS

NOTE Confidence: 0.80600363

 $00:36:54.375 \longrightarrow 00:36:57.000$ season is required for self renewal.

NOTE Confidence: 0.80600363

 $00:36:57.000 \longrightarrow 00:36:59.105$ It's induced in lymphopoiesis and

NOTE Confidence: 0.80600363

 $00:36:59.105 \longrightarrow 00:37:01.210$ required for T cell expansion.

NOTE Confidence: 0.80600363

 $00{:}37{:}01.210 \dashrightarrow 00{:}37{:}03.340$ Knockout mouse have defects in

 $00:37:03.340 \longrightarrow 00:37:05.044$ lymphopoiesis and then downregulation

NOTE Confidence: 0.80600363

 $00:37:05.044 \longrightarrow 00:37:05.840$ in Milo.

NOTE Confidence: 0.80600363

00:37:05.840 --> 00:37:08.666 Police has been demonstrated in its

NOTE Confidence: 0.80600363

00:37:08.666 --> 00:37:11.722 requirement for PU .1 regulation in

NOTE Confidence: 0.80600363

00:37:11.722 --> 00:37:14.437 in common Milo progenitor cells.

NOTE Confidence: 0.80600363

 $00{:}37{:}14.440 \dashrightarrow 00{:}37{:}17.345$ So Mark looked at sappi 1M RNA

NOTE Confidence: 0.80600363

 $00:37:17.345 \longrightarrow 00:37:18.175$ expression anarchist.

NOTE Confidence: 0.80600363

 $00{:}37{:}18.180 --> 00{:}37{:}19.428 \ 19 \ knockdown \ cells,$

NOTE Confidence: 0.80600363

 $00:37:19.428 \longrightarrow 00:37:20.676$ day OD five.

NOTE Confidence: 0.80600363

 $00{:}37{:}20.680 \dashrightarrow 00{:}37{:}22.790$ He showed that the expression

NOTE Confidence: 0.80600363

 $00{:}37{:}22.790 \dashrightarrow 00{:}37{:}25.792$ decrease more rapidly in our PS 19

NOTE Confidence: 0.80600363

 $00{:}37{:}25.792 \dashrightarrow 00{:}37{:}27.827$ knockdown cells compared to controls

NOTE Confidence: 0.80600363

 $00{:}37{:}27.827 \dashrightarrow 00{:}37{:}30.658$ and then also at the protein level.

NOTE Confidence: 0.80600363

 $00:37:30.660 \longrightarrow 00:37:33.257$ We see that there is much more

NOTE Confidence: 0.80600363

 $00:37:33.257 \longrightarrow 00:37:35.660$ decrease in the protein levels.

 $00:37:35.660 \longrightarrow 00:37:39.548$ Is that being one day 5?

NOTE Confidence: 0.80600363

 $00{:}37{:}39.550 \dashrightarrow 00{:}37{:}42.286$ The colony essays and liquid culture

NOTE Confidence: 0.80600363

 $00{:}37{:}42.286 \dashrightarrow 00{:}37{:}44.592$ essays that we perform showed

NOTE Confidence: 0.80600363

 $00:37:44.592 \longrightarrow 00:37:46.980$ that in colony essays in RPS.

NOTE Confidence: 0.80600363

 $00:37:46.980 \longrightarrow 00:37:47.828$ 19 knockdown.

NOTE Confidence: 0.80600363

 $00:37:47.828 \longrightarrow 00:37:50.796$ HSBC's that we did not see significant

NOTE Confidence: 0.80600363

 $00:37:50.796 \longrightarrow 00:37:53.074$ increase in colony numbers per

NOTE Confidence: 0.80600363

 $00:37:53.074 \longrightarrow 00:37:54.850$ plate with three expressions.

NOTE Confidence: 0.80600363

 $00{:}37{:}54.850 \dashrightarrow 00{:}37{:}57.196$ We expressed that being one about

NOTE Confidence: 0.80600363

 $00:37:57.196 \longrightarrow 00:38:00.089$ to fold in these knockdown cells,

NOTE Confidence: 0.80600363

 $00:38:00.090 \longrightarrow 00:38:03.142$ we can see any much increase in

NOTE Confidence: 0.80600363

 $00:38:03.142 \longrightarrow 00:38:06.210$ liquid culture. However, we sell a CD.

NOTE Confidence: 0.80600363

 $00:38:06.210 \longrightarrow 00:38:09.346$ 235 sales that there was increase in.

NOTE Confidence: 0.80600363

 $00:38:09.350 \longrightarrow 00:38:12.450$ With happy one overexpression.

NOTE Confidence: 0.80600363

 $00:38:12.450 \longrightarrow 00:38:14.904$ But although there was no increase

NOTE Confidence: 0.80600363

 $00:38:14.904 \longrightarrow 00:38:16.540$ in the colony numbers,

 $00:38:16.540 \longrightarrow 00:38:19.403$ we could see by visualization that on

NOTE Confidence: 0.80600363

 $00{:}38{:}19.403 \dashrightarrow 00{:}38{:}22.267$ the colonies were huge for much larger,

NOTE Confidence: 0.80600363

 $00:38:22.270 \longrightarrow 00:38:24.310$ suggesting that there was a

NOTE Confidence: 0.80600363

 $00:38:24.310 \longrightarrow 00:38:25.942$ proliferation of these cells,

NOTE Confidence: 0.80600363

 $00{:}38{:}25.950 \dashrightarrow 00{:}38{:}29.214$ and we did see also a normal controls,

NOTE Confidence: 0.80600363

 $00:38:29.220 \longrightarrow 00:38:32.484$ but not as dramatically as in our case,

NOTE Confidence: 0.80600363

 $00:38:32.490 \longrightarrow 00:38:33.720$ 19 knockdown cells.

NOTE Confidence: 0.8144065

00:38:33.720 --> 00:38:36.990 There's also a hand up from. Yeah, it's

NOTE Confidence: 0.8144065

 $00:38:36.990 \longrightarrow 00:38:38.630$ Vince. OK, I've is.

NOTE Confidence: 0.73706543

 $00:38:41.950 \longrightarrow 00:38:44.070$ Sorry, that was sorry that

NOTE Confidence: 0.73706543

 $00:38:44.070 \longrightarrow 00:38:45.766$ was accidentally did that.

NOTE Confidence: 0.73706543

 $00:38:45.770 \longrightarrow 00:38:47.459$ Oh alright, never mind.

NOTE Confidence: 0.73706543

 $00{:}38{:}47.460 \dashrightarrow 00{:}38{:}50.225$ So colony as says we see that in

NOTE Confidence: 0.73706543

 $00:38:50.225 \longrightarrow 00:38:52.719$ knockdown of sappy when in normal

NOTE Confidence: 0.73706543

 $00:38:52.719 \longrightarrow 00:38:55.589$ cells or healthy progenitors we see a

 $00:38:55.677 \longrightarrow 00:38:58.827$ slight decrease in an doxycycline or 71

NOTE Confidence: 0.73706543

00:38:58.827 --> 00:39:01.464 knockdown was 235 in liquid culture.

NOTE Confidence: 0.73706543

 $00:39:01.464 \longrightarrow 00:39:05.112$ So we do see a mild decrease in

NOTE Confidence: 0.73706543

 $00:39:05.112 \longrightarrow 00:39:08.466$ erythroid colonies and or through glass.

NOTE Confidence: 0.73706543

 $00:39:08.470 \longrightarrow 00:39:10.070$ So sad he went on.

NOTE Confidence: 0.73706543

 $00:39{:}10.070 \dashrightarrow 00{:}39{:}11.342$ Regulation does not dramatically

NOTE Confidence: 0.73706543

00:39:11.342 --> 00:39:12.932 impact anemia phenotype in DBA.

NOTE Confidence: 0.73706543

 $00:39:12.940 \longrightarrow 00:39:15.880$ So what is the function of 71?

NOTE Confidence: 0.73706543

 $00{:}39{:}15.880 \dashrightarrow 00{:}39{:}17.588$ So healthy uncommitted Mylar

NOTE Confidence: 0.73706543

 $00:39:17.588 \longrightarrow 00:39:19.296$ projectors we see that.

NOTE Confidence: 0.73706543

 $00:39:19.300 \longrightarrow 00:39:20.294$ So again,

NOTE Confidence: 0.73706543

 $00:39:20.294 \longrightarrow 00:39:22.779$ here's our colony assay results

NOTE Confidence: 0.73706543

 $00:39:22.779 \longrightarrow 00:39:25.297$ showing mild decrease in be a few.

NOTE Confidence: 0.73706543

 $00:39:25.300 \longrightarrow 00:39:26.659$ Ease the culture.

NOTE Confidence: 0.73706543

00:39:26.659 --> 00:39:29.830 The same 235 cells but in megakaryocytes

NOTE Confidence: 0.73706543

00:39:29.912 --> 00:39:32.565 regenerators we see at least a 30%

 $00:39:32.570 \longrightarrow 00:39:35.566$ decrease in with that be one knockdown.

NOTE Confidence: 0.73706543

 $00{:}39{:}35.570 \dashrightarrow 00{:}39{:}38.276$ So our hypothesis at this point

NOTE Confidence: 0.73706543

 $00:39:38.276 \longrightarrow 00:39:41.495$ was that although in DPA we see

NOTE Confidence: 0.73706543

 $00:39:41.495 \longrightarrow 00:39:43.705$ a block here early erythroblast

NOTE Confidence: 0.73706543

 $00{:}39{:}43.705 \dashrightarrow 00{:}39{:}46.586$ stage or seek to the 235 stage.

NOTE Confidence: 0.73706543

 $00:39:46.590 \longrightarrow 00:39:49.060$ That perhaps happy one is

NOTE Confidence: 0.73706543

 $00:39:49.060 \longrightarrow 00:39:51.036$ acting upstream of this,

NOTE Confidence: 0.73706543

 $00:39:51.040 \longrightarrow 00:39:53.134$ affecting the megakaryocyte

NOTE Confidence: 0.73706543

 $00{:}39{:}53.134 \dashrightarrow 00{:}39{:}55.228$ recite progenitor stage.

NOTE Confidence: 0.73706543

 $00:39:55.230 \longrightarrow 00:39:56.856$ And so here we have again

NOTE Confidence: 0.73706543

 $00:39:56.856 \longrightarrow 00:39:58.670$ our as our PS90 knockdown.

NOTE Confidence: 0.73706543

 $00:39:58.670 \longrightarrow 00:40:00.880$ Showing calling or calling numbers.

NOTE Confidence: 0.73706543

 $00:40:00.880 \longrightarrow 00:40:03.450$ Interesting Lee with CFU GM.

NOTE Confidence: 0.73706543

 $00:40:03.450 \longrightarrow 00:40:06.528$ We see an increase in colonies.

NOTE Confidence: 0.73706543

 $00:40:06.530 \longrightarrow 00:40:10.128$ C235 expansion was noted with RPS 19.

 $00:40:10.130 \longrightarrow 00:40:13.510$ Knock down an 71.

NOTE Confidence: 0.73706543

 $00:40:13.510 \longrightarrow 00:40:16.756$ We did see also 11B expansion

NOTE Confidence: 0.73706543

 $00:40:16.756 \longrightarrow 00:40:17.838$ and overexpression.

NOTE Confidence: 0.73706543

 $00:40:17.840 \longrightarrow 00:40:21.086$ We also see CD 41 increase

NOTE Confidence: 0.73706543

 $00:40:21.086 \longrightarrow 00:40:23.250$ more even more significantly.

NOTE Confidence: 0.75487924

 $00:40:25.470 \longrightarrow 00:40:27.410$ How does this happen then?

NOTE Confidence: 0.75487924

 $00:40:27.410 \longrightarrow 00:40:30.126$ What is the mechanism downstream of that?

NOTE Confidence: 0.75487924

00:40:30.130 --> 00:40:33.082 Be? One well, one of the things one of

NOTE Confidence: 0.75487924

 $00{:}40{:}33.082 \to 00{:}40{:}36.027$ the genes that has been very important

NOTE Confidence: 0.75487924

00:40:36.027 --> 00:40:38.659 recently has been a heat shock.

NOTE Confidence: 0.75487924

 $00:40:38.660 \longrightarrow 00:40:40.600$ 70th proteins that's encoded by

NOTE Confidence: 0.75487924

 $00:40:40.600 \longrightarrow 00:40:43.322$ the gene zedex, HSP, 1A1B, and 1A,

NOTE Confidence: 0.75487924

 $00:40:43.322 \longrightarrow 00:40:45.650$ and there are three different papers.

NOTE Confidence: 0.75487924

 $00:40:45.650 \longrightarrow 00:40:47.640$ Sorry, there are three different

NOTE Confidence: 0.75487924

00:40:47.640 --> 00:40:49.630 papers that have been describing

NOTE Confidence: 0.75487924

 $00{:}40{:}49.693 \dashrightarrow 00{:}40{:}51.468$ by Hermione and Leah Dacosta.

 $00:40:51.470 \longrightarrow 00:40:55.709$ The role of HSP 70 in regulating gotta one.

NOTE Confidence: 0.75487924

00:40:55.710 --> 00:40:58.076 An agency 70 is thought to interact

NOTE Confidence: 0.75487924

 $00:40:58.076 \longrightarrow 00:41:01.107$ with God and one to prevent the caspase

NOTE Confidence: 0.75487924

 $00{:}41{:}01.107 \dashrightarrow 00{:}41{:}03.400$ dependent cleavage of God in one,

NOTE Confidence: 0.75487924

 $00:41:03.400 \longrightarrow 00:41:05.350$ so basically is stabilizing that

NOTE Confidence: 0.75487924

 $00:41:05.350 \longrightarrow 00:41:08.027$ complex or God or one to allow

NOTE Confidence: 0.75487924

 $00:41:08.027 \longrightarrow 00:41:09.977$ it to to to activate genes,

NOTE Confidence: 0.75487924

 $00{:}41{:}09.980 \longrightarrow 00{:}41{:}12.868$ and this is just a profile of chromosomes

NOTE Confidence: 0.75487924

00:41:12.868 --> 00:41:15.787 looking at various genes so they are not

NOTE Confidence: 0.75487924

 $00{:}41{:}15.787 \dashrightarrow 00{:}41{:}18.381$ affected by sappy one or influence to

NOTE Confidence: 0.75487924

 $00{:}41{:}18.381 \dashrightarrow 00{:}41{:}20.957$ rescued by setting one in the orange.

NOTE Confidence: 0.75487924

 $00:41:20.960 \longrightarrow 00:41:24.092$ So here are the two genes and if we

NOTE Confidence: 0.75487924

 $00{:}41{:}24.092 --> 00{:}41{:}27.046$ knock down step one, we see that.

NOTE Confidence: 0.75487924

 $00{:}41{:}27.046 \dashrightarrow 00{:}41{:}29.698$ This HP A1A expression goes down.

NOTE Confidence: 0.75487924

00:41:29.700 --> 00:41:30.141 Similarly,

00:41:30.141 --> 00:41:33.228 if we knock down at SABI one

NOTE Confidence: 0.75487924

 $00:41:33.228 \longrightarrow 00:41:36.127$ we see HSP one being Arnie,

NOTE Confidence: 0.75487924

 $00:41:36.130 \longrightarrow 00:41:40.618$ decreasing ansim with the protein levels.

NOTE Confidence: 0.75487924

 $00:41:40.620 \longrightarrow 00:41:43.026$ If we then look examine our

NOTE Confidence: 0.75487924

00:41:43.026 --> 00:41:44.630 peacemaking knockdown cells and

NOTE Confidence: 0.75487924

00:41:44.703 --> 00:41:46.539 then overexpress sappy one,

NOTE Confidence: 0.75487924

 $00:41:46.540 \longrightarrow 00:41:49.438$ we see that there's a rescue and

NOTE Confidence: 0.75487924

 $00:41:49.438 \longrightarrow 00:41:52.039$ the expression of these two genes

NOTE Confidence: 0.75487924

 $00{:}41{:}52.039 \dashrightarrow 00{:}41{:}55.000$ and also at the protein level here.

NOTE Confidence: 0.75313586

00:41:58.080 --> 00:42:02.176 To understand what sappy one might be doing,

NOTE Confidence: 0.75313586

 $00{:}42{:}02.180 \dashrightarrow 00{:}42{:}04.740$ and regulating house regulating it,

NOTE Confidence: 0.75313586

 $00:42:04.740 \longrightarrow 00:42:07.806$ just P1A1A and the HPA 1B.

NOTE Confidence: 0.75313586

00:42:07.810 --> 00:42:09.472 Mark performed chromatin

NOTE Confidence: 0.75313586

 $00{:}42{:}09.472 \dashrightarrow 00{:}42{:}11.688$ immunoprecipitation assays with antibodies

NOTE Confidence: 0.75313586

 $00:42:11.688 \longrightarrow 00:42:14.458$ that are specific to these H3K914,

NOTE Confidence: 0.75313586

00:42:14.460 --> 00:42:15.996 desolation HK4 trimethylation

00:42:15.996 --> 00:42:18.044 HCC 27 five metalation,

NOTE Confidence: 0.75313586

 $00:42:18.050 \longrightarrow 00:42:22.546$ and these are the sites that set the

NOTE Confidence: 0.75313586

 $00:42:22.546 \longrightarrow 00:42:26.730$ one binds and what we see is that.

NOTE Confidence: 0.75313586

00:42:26.730 --> 00:42:29.496 There's also commented peaks which that

NOTE Confidence: 0.75313586

 $00:42:29.496 \longrightarrow 00:42:32.323$ with the H3K29 acetylation and HGK

NOTE Confidence: 0.75313586

00:42:32.323 --> 00:42:35.065 4 trimethylation but not with H3K27

NOTE Confidence: 0.75313586

00:42:35.065 --> 00:42:37.382 Trimethylation. So here's our model.

NOTE Confidence: 0.75313586

 $00:42:37.382 \longrightarrow 00:42:40.622$ Here is the HSP A1A gene. Here's.

NOTE Confidence: 0.75313586

 $00:42:40.622 \longrightarrow 00:42:43.856$ Here are the seven one binding sites,

NOTE Confidence: 0.75313586

 $00{:}42{:}43.860 \dashrightarrow 00{:}42{:}47.129$ and here are the predicted enhancer and

NOTE Confidence: 0.75313586

 $00:42:47.129 \longrightarrow 00:42:49.879$ promoter sites that we would expect,

NOTE Confidence: 0.75313586

 $00:42:49.880 \longrightarrow 00:42:53.058$ and so sappy one promotes the looping

NOTE Confidence: 0.75313586

 $00{:}42{:}53.058 \dashrightarrow 00{:}42{:}56.928$ so that all of these will be enclosed.

NOTE Confidence: 0.75313586

 $00:42:56.930 \longrightarrow 00:42:58.678$ Doximity to facilitate transcription

NOTE Confidence: 0.75313586

 $00:42:58.678 \longrightarrow 00:43:00.426$ of these two genes,

 $00:43:00.430 \longrightarrow 00:43:02.730$ another approach that Mark Hughes

NOTE Confidence: 0.75313586

 $00:43:02.730 \longrightarrow 00:43:05.030$ was as chromatin confirmation capture

NOTE Confidence: 0.75313586

 $00:43:05.099 \longrightarrow 00:43:07.114$ essay where you basically treat

NOTE Confidence: 0.75313586

 $00:43:07.114 \longrightarrow 00:43:08.726$ the cells with formalin,

NOTE Confidence: 0.75313586

 $00:43:08.730 \longrightarrow 00:43:11.365$ then digest using restriction enzymes

NOTE Confidence: 0.75313586

 $00{:}43{:}11.365 \dashrightarrow 00{:}43{:}14.927$ and then perform PCR and what he

NOTE Confidence: 0.75313586

00:43:14.927 --> 00:43:17.846 showed was that with our with our

NOTE Confidence: 0.75313586

 $00:43:17.846 \longrightarrow 00:43:20.793$ control we see two nice peaks at the

NOTE Confidence: 0.75313586

 $00:43:20.793 \longrightarrow 00:43:22.714$ sites of the Enhancement Promoter

NOTE Confidence: 0.75313586

00:43:22.714 --> 00:43:25.336 regions with RPS 19 knock down,

NOTE Confidence: 0.75313586

 $00:43:25.340 \longrightarrow 00:43:26.258$ this is blunted.

NOTE Confidence: 0.75313586

 $00:43:26.258 \longrightarrow 00:43:28.400$ With knock down as happy when we

NOTE Confidence: 0.75313586

00:43:28.471 --> 00:43:30.685 basically don't see these peaks and

NOTE Confidence: 0.75313586

 $00{:}43{:}30.685 \dashrightarrow 00{:}43{:}32.815$ then with re expression of happy

NOTE Confidence: 0.75313586

 $00:43:32.815 \longrightarrow 00:43:34.873$ when we see rescue and again the

NOTE Confidence: 0.75313586

 $00:43:34.873 \longrightarrow 00:43:37.250$ peaks appear again.

00:43:37.250 --> 00:43:39.542 Another way to validate these data

NOTE Confidence: 0.75313586

 $00:43:39.542 \longrightarrow 00:43:42.510$ and look at it functionally is to

NOTE Confidence: 0.75313586

00:43:42.510 --> 00:43:46.150 perform a new technology known as Cloud Nine,

NOTE Confidence: 0.75313586

 $00:43:46.150 \longrightarrow 00:43:48.580$ and this is a approach that

NOTE Confidence: 0.75313586

 $00:43:48.580 \longrightarrow 00:43:51.160$ was developed by Kevin Wang at

NOTE Confidence: 0.75313586

 $00{:}43{:}51.160 \dashrightarrow 00{:}43{:}53.355$ Stanford where you use Casper,

NOTE Confidence: 0.75313586

00:43:53.360 --> 00:43:55.904 Chris crisper CAS 9 to basically

NOTE Confidence: 0.75313586

 $00:43:55.904 \longrightarrow 00:43:58.450$ target certain sequences in the genome,

NOTE Confidence: 0.75313586

 $00:43:58.450 \longrightarrow 00:44:00.066$ treat with abscisic acid,

NOTE Confidence: 0.75313586

 $00:44:00.066 \longrightarrow 00:44:02.086$ which then forms and reversibly

NOTE Confidence: 0.75313586

 $00:44:02.086 \longrightarrow 00:44:03.539$ loop that's induced,

NOTE Confidence: 0.75313586

 $00:44:03.540 \longrightarrow 00:44:07.516$ and then you can see the effects downstream.

NOTE Confidence: 0.75313586

 $00:44:07.520 \longrightarrow 00:44:09.896$ So this is what Mark did and this

NOTE Confidence: 0.75313586

00:44:09.896 --> 00:44:12.130 is again showing our sappy one,

NOTE Confidence: 0.75313586

 $00:44:12.130 \longrightarrow 00:44:13.442$ and the various promoters

00:44:13.442 --> 00:44:14.426 and enhancer region,

NOTE Confidence: 0.75313586

00:44:14.430 --> 00:44:16.726 and again using G Arnie dimer pairs.

NOTE Confidence: 0.75313586

00:44:16.730 --> 00:44:19.242 If you knock down S isep or sappy

NOTE Confidence: 0.75313586

00:44:19.242 --> 00:44:22.016 one treat with this, I said the one,

NOTE Confidence: 0.75313586

 $00:44:22.016 \longrightarrow 00:44:24.020$ you see that there's no expression

NOTE Confidence: 0.75313586

 $00:44:24.091 \longrightarrow 00:44:24.958$ of these two.

NOTE Confidence: 0.75313586

 $00:44:24.960 \longrightarrow 00:44:26.944$ He drop protein genes.

NOTE Confidence: 0.75313586

00:44:26.944 --> 00:44:29.920 If we then induce looping using

NOTE Confidence: 0.75313586

00:44:30.020 --> 00:44:32.318 Cloud nine at at the EP1,

NOTE Confidence: 0.75313586

 $00:44:32.320 \longrightarrow 00:44:34.918$ which is here to transducer to

NOTE Confidence: 0.75313586

 $00{:}44{:}34.918 \dashrightarrow 00{:}44{:}37.020$ facilitate expression of his HSP.

NOTE Confidence: 0.75313586

 $00:44:37.020 \longrightarrow 00:44:40.050$ A1A we see that there's expression

NOTE Confidence: 0.75313586

 $00:44:40.050 \longrightarrow 00:44:43.459$ but not with the other JPA 1D.

NOTE Confidence: 0.75313586

 $00:44:43.460 \longrightarrow 00:44:46.070$ If you then loop at EP2,

NOTE Confidence: 0.75313586

 $00:44:46.070 \longrightarrow 00:44:49.353$ which is here so that the enhancer

NOTE Confidence: 0.75313586

 $00:44:49.353 \longrightarrow 00:44:53.468$ is next to the P2 for HSV one a B1B,

 $00:44:53.470 \longrightarrow 00:44:56.122$ you see that there's expression of

NOTE Confidence: 0.75313586

 $00:44:56.122 \longrightarrow 00:44:59.559$ this gene and also slightly of HPA 1A.

NOTE Confidence: 0.75313586

 $00:44:59.560 \longrightarrow 00:45:02.892$ And then finally this is just a

NOTE Confidence: 0.75313586

 $00:45:02.892 \longrightarrow 00:45:06.139$ control showing there's no induction Eugene.

NOTE Confidence: 0.75313586

 $00{:}45{:}06.140 \dashrightarrow 00{:}45{:}08.822$ So correlating with that is also

NOTE Confidence: 0.75313586

 $00:45:08.822 \longrightarrow 00:45:10.163$ me P expansion.

NOTE Confidence: 0.75313586

 $00:45:10.170 \longrightarrow 00:45:12.858$ We we knocked down SFB one,

NOTE Confidence: 0.75313586

 $00:45:12.860 \longrightarrow 00:45:15.100$ but then we expressed it.

NOTE Confidence: 0.75313586

 $00:45:15.100 \longrightarrow 00:45:16.324$ We see increase.

NOTE Confidence: 0.75313586

00:45:16.324 --> 00:45:19.740 We also see it by looping of P1

NOTE Confidence: 0.75313586

 $00{:}45{:}19.740 \dashrightarrow 00{:}45{:}22.055$ and P2 increase in comparison

NOTE Confidence: 0.75313586

 $00:45:22.055 \longrightarrow 00:45:25.361$ to controls and we don't see any

NOTE Confidence: 0.75313586

 $00{:}45{:}25.361 \dashrightarrow 00{:}45{:}28.539$ effect on HCS or the CMP population.

NOTE Confidence: 0.75313586

 $00:45:28.540 \longrightarrow 00:45:29.440$ So can

NOTE Confidence: 0.7754213

00:45:29.440 --> 00:45:32.122 I just clarify what's the experimental

 $00:45:32.122 \longrightarrow 00:45:34.476$ design here for this expansion

NOTE Confidence: 0.7754213

00:45:34.476 --> 00:45:37.440 and the CMP right? Starting with

NOTE Confidence: 0.7978872

 $00:45:37.440 \longrightarrow 00:45:40.037$ this is looking at 34 positive CD,

NOTE Confidence: 0.7978872

 $00:45:40.040 \longrightarrow 00:45:41.890$ 71 low population of cells,

NOTE Confidence: 0.7978872

00:45:41.890 --> 00:45:44.116 but they've been cultured in vitro

NOTE Confidence: 0.7978872

00:45:44.120 --> 00:45:47.460 plus and minus for how yes, all of these

NOTE Confidence: 0.7978872

 $00:45:47.460 \longrightarrow 00:45:50.428$ yeah and then treated with the cloud 9,

NOTE Confidence: 0.7978872

 $00:45:50.430 \longrightarrow 00:45:52.536$ so you're then treating with the

NOTE Confidence: 0.7978872

 $00{:}45{:}52.536 \dashrightarrow 00{:}45{:}54.787$ dimer peers and then with abscisic

NOTE Confidence: 0.7978872

00:45:54.787 --> 00:45:57.110 acid to induce the looping. But

NOTE Confidence: 0.7978872

00:45:57.110 --> 00:45:59.330 what you're starting with me P.

NOTE Confidence: 0.7991827

 $00:46:00.770 \longrightarrow 00:46:03.002$ We're starting at the beginning and

NOTE Confidence: 0.7991827

 $00:46:03.002 \longrightarrow 00:46:05.268$ then waiting until the stage where

NOTE Confidence: 0.7991827

 $00:46:05.268 \longrightarrow 00:46:07.386$ we would see these cells appear.

NOTE Confidence: 0.7991827

 $00:46:07.390 \longrightarrow 00:46:12.268$ OK, yeah. Yeah, and then we also look.

NOTE Confidence: 0.7991827

 $00:46:12.270 \longrightarrow 00:46:15.573$ We also examine the Seppi 1S I or any

00:46:15.573 --> 00:46:18.020 induced Rizzo Blastic Buster Apolysis

NOTE Confidence: 0.7991827

 $00{:}46{:}18.020 \dashrightarrow 00{:}46{:}21.509$ showed that with the looping at P1

NOTE Confidence: 0.7991827

 $00:46:21.509 \longrightarrow 00:46:24.421$ and P2 we see increasing with robust

NOTE Confidence: 0.7991827

00:46:24.421 --> 00:46:26.958 increase in CD41A positive cells

NOTE Confidence: 0.7991827

 $00:46:26.958 \longrightarrow 00:46:31.730$ but not in Salem be my light cells.

NOTE Confidence: 0.7991827

00:46:31.730 --> 00:46:33.645 The similar findings were also

NOTE Confidence: 0.7991827

00:46:33.645 --> 00:46:36.240 observed in cells that had SHRPS 19.

NOTE Confidence: 0.7991827

 $00:46:36.240 \longrightarrow 00:46:38.496$ So in our PS90 knockdown cells

NOTE Confidence: 0.7991827

 $00:46:38.496 \longrightarrow 00:46:40.000$ the same thing happened.

NOTE Confidence: 0.7991827

 $00:46:40.000 \longrightarrow 00:46:42.639$ If we loop it P1 we see

NOTE Confidence: 0.7991827

 $00:46:42.639 \longrightarrow 00:46:44.510$ the expression of HSP 1A.

NOTE Confidence: 0.7991827

 $00:46:44.510 \longrightarrow 00:46:46.890$ If we loop at P2P CHSP A1B

NOTE Confidence: 0.7991827

 $00{:}46{:}46.890 \dashrightarrow 00{:}46{:}48.650$ expression and then controls,

NOTE Confidence: 0.7991827

 $00:46:48.650 \longrightarrow 00:46:52.082$ we don't see and then this is just

NOTE Confidence: 0.7991827

 $00:46:52.082 \longrightarrow 00:46:54.462$ Western blot showing the expression

 $00:46:54.462 \longrightarrow 00:46:57.787$ a day five for both of these.

NOTE Confidence: 0.7991827

00:46:57.790 --> 00:46:58.640 Experiments.

NOTE Confidence: 0.7247406

 $00:47:00.680 \longrightarrow 00:47:02.060$ So looping appears.

NOTE Confidence: 0.7247406

 $00:47:02.060 \longrightarrow 00:47:04.360$ He also rescues Happy one,

NOTE Confidence: 0.7247406

 $00:47:04.360 \longrightarrow 00:47:06.752$ defects and healthy megakaryocytes.

NOTE Confidence: 0.7247406

 $00{:}47{:}06.752 \dashrightarrow 00{:}47{:}09.144$ Here's colony essays showing

NOTE Confidence: 0.7247406

 $00:47:09.144 \longrightarrow 00:47:11.293$ our looping experiments and

NOTE Confidence: 0.7247406

00:47:11.293 --> 00:47:13.891 knockdown of 71 so by looping

NOTE Confidence: 0.7247406

 $00{:}47{:}13.891 \dashrightarrow 00{:}47{:}16.727$ itself we can see increasing BFUE.

NOTE Confidence: 0.7247406

00:47:16.730 --> 00:47:18.580 Nothing with the CFU GM.

NOTE Confidence: 0.7247406

 $00{:}47{:}18.580 \dashrightarrow 00{:}47{:}21.052$ Also, in liquid culture we see

NOTE Confidence: 0.7247406

00:47:21.052 --> 00:47:23.736 just with looping alone we can see

NOTE Confidence: 0.7247406

 $00:47:23.736 \longrightarrow 00:47:25.976$ improvement and C 235 and 41 a,

NOTE Confidence: 0.7247406

 $00:47:25.980 \longrightarrow 00:47:27.460$ but not in CD11B.

NOTE Confidence: 0.7247406

 $00:47:27.460 \longrightarrow 00:47:29.788$ So again, that tells us that

NOTE Confidence: 0.7247406

 $00:47:29.788 \longrightarrow 00:47:32.170$ the looping is critical for the

 $00:47:32.249 \longrightarrow 00:47:34.429$ transcription of these genes to

NOTE Confidence: 0.7247406

 $00:47:34.429 \longrightarrow 00:47:37.670$ take place and also for the effects

NOTE Confidence: 0.7247406

 $00:47:37.670 \longrightarrow 00:47:40.105$ on CD235 and C41 proliferation.

NOTE Confidence: 0.7247406

00:47:40.110 --> 00:47:43.110 Finally, to link this all back to God.

NOTE Confidence: 0.7247406

00:47:43.110 --> 00:47:45.360 Oh one. As I mentioned earlier,

NOTE Confidence: 0.7247406

 $00:47:45.360 \longrightarrow 00:47:48.104$ the God one is regulated HSP 70 being

NOTE Confidence: 0.7247406

 $00:47:48.104 \longrightarrow 00:47:50.987$ available to prevent the degradation of God.

NOTE Confidence: 0.7247406

 $00:47:50.990 \longrightarrow 00:47:51.664$ Oh one.

NOTE Confidence: 0.7247406

00:47:51.664 --> 00:47:53.686 So here we see 235 positive

NOTE Confidence: 0.7247406

00:47:53.686 --> 00:47:55.859 cells E 41 positive cells,

NOTE Confidence: 0.7247406

 $00{:}47{:}55.860 \dashrightarrow 00{:}47{:}57.051$ gotta one expression.

NOTE Confidence: 0.7247406

 $00:47:57.051 \longrightarrow 00:47:59.830$ And if we then take our knockdown

NOTE Confidence: 0.7247406

 $00:47:59.902 \longrightarrow 00:48:01.487$ of essay of sappy one,

NOTE Confidence: 0.7247406

 $00:48:01.490 \longrightarrow 00:48:04.038$ we see that the looping can restore

NOTE Confidence: 0.7247406

 $00:48:04.038 \longrightarrow 00:48:06.718$ the expression of God in one because

00:48:06.718 --> 00:48:08.986 now you're getting HP 70 expression.

NOTE Confidence: 0.7247406

 $00:48:08.990 \longrightarrow 00:48:11.240$ So it's able to prevent

NOTE Confidence: 0.7247406

 $00:48:11.240 \longrightarrow 00:48:13.040$ degradation of God 01.

NOTE Confidence: 0.7247406

 $00:48:13.040 \longrightarrow 00:48:15.110$ Godwin expression is just shown

NOTE Confidence: 0.7247406

 $00:48:15.110 \longrightarrow 00:48:17.180$ here showing the increase in

NOTE Confidence: 0.7247406

 $00:48:17.251 \longrightarrow 00:48:18.935$ expression correlating with with

NOTE Confidence: 0.7247406

 $00:48:18.935 \longrightarrow 00:48:21.896$ the data I just talked about in

NOTE Confidence: 0.7247406

 $00:48:21.896 \longrightarrow 00:48:23.721$ an MVP expansion also correlated

NOTE Confidence: 0.7247406

 $00{:}48{:}23.721 \dashrightarrow 00{:}48{:}26.000$ with with the God one expression.

NOTE Confidence: 0.82798517

 $00:48:26.000 \longrightarrow 00:48:28.835$ So if you did a western on

NOTE Confidence: 0.82798517

 $00:48:28.835 \longrightarrow 00:48:31.667$ the top right for the HSP 70,

NOTE Confidence: 0.82798517

 $00:48:31.670 \longrightarrow 00:48:34.100$ it would parallel the gotta one.

NOTE Confidence: 0.82798517

 $00:48:34.100 \longrightarrow 00:48:36.485$ That's what we believe. Yes, yes.

NOTE Confidence: 0.82798517

 $00:48:36.485 \longrightarrow 00:48:40.219$ And so that goes back to our model where we

NOTE Confidence: 0.82798517

 $00:48:40.219 \longrightarrow 00:48:43.418$ are in normal conditions that 71 gradually

NOTE Confidence: 0.82798517

 $00:48:43.418 \longrightarrow 00:48:46.627$ decreases as cells become more committed.

00:48:46.630 --> 00:48:50.184 But we do see HP 70 increasing, which

NOTE Confidence: 0.82798517

 $00{:}48{:}50.184 \dashrightarrow 00{:}48{:}52.848$ prevents gotten one from being degraded.

NOTE Confidence: 0.82798517

 $00:48:52.850 \longrightarrow 00:48:55.508$ There's some city state level there,

NOTE Confidence: 0.82798517

 $00:48:55.510 \longrightarrow 00:48:57.286$ but obviously once committed

NOTE Confidence: 0.82798517

00:48:57.286 --> 00:48:59.506 to pull through blast stage,

NOTE Confidence: 0.82798517

00:48:59.510 --> 00:49:02.168 we see significant increase in God,

NOTE Confidence: 0.82798517

 $00:49:02.170 \longrightarrow 00:49:05.042$ one in order to transduced jeans that are

NOTE Confidence: 0.82798517

 $00:49:05.042 \longrightarrow 00:49:07.500$ required for earthway differentiation,

NOTE Confidence: 0.82798517

 $00:49:07.500 \longrightarrow 00:49:08.514$ terminal differentiation.

NOTE Confidence: 0.82798517

00:49:08.514 --> 00:49:11.556 We knocked down sappy one prematurely,

NOTE Confidence: 0.82798517

 $00:49:11.560 \longrightarrow 00:49:13.870$ or, in the case of RPS,

NOTE Confidence: 0.82798517

 $00:49:13.870 \longrightarrow 00:49:15.001$ 19, insufficiency, seven.

NOTE Confidence: 0.82798517

 $00:49:15.001 \longrightarrow 00:49:18.110$ That would be 1 levels are much lower,

NOTE Confidence: 0.82798517

 $00:49:18.110 \longrightarrow 00:49:20.798$ which does not allow HSP 70 P.

NOTE Confidence: 0.82798517

 $00:49:20.800 \longrightarrow 00:49:23.495$ It just be 70 should be expressed,

00:49:23.500 --> 00:49:26.188 which then blunts the expression of God.

NOTE Confidence: 0.82798517

00:49:26.190 --> 00:49:29.422 Oh, one so that we don't get the

NOTE Confidence: 0.82798517

 $00{:}49{:}29.422 \dashrightarrow 00{:}49{:}31.577$ normal increase and got a wine.

NOTE Confidence: 0.82798517

 $00:49:31.580 \longrightarrow 00:49:34.255$ And we believe this contributes

NOTE Confidence: 0.82798517

 $00:49:34.255 \longrightarrow 00:49:36.395$ to the anemia phenotype.

NOTE Confidence: 0.82798517

 $00:49:36.400 \longrightarrow 00:49:39.165$ It's just a schematic of what we

NOTE Confidence: 0.82798517

00:49:39.165 --> 00:49:41.638 believe is happening in CMP, ME P.

NOTE Confidence: 0.82798517

 $00:49:41.638 \longrightarrow 00:49:43.358$ The ratio relative ratios that

NOTE Confidence: 0.82798517

 $00:49:43.358 \longrightarrow 00:49:45.500$ occur at this stage normally,

NOTE Confidence: 0.82798517

 $00:49:45.500 \longrightarrow 00:49:47.882$ and then at the risible stage

NOTE Confidence: 0.82798517

 $00:49:47.882 \longrightarrow 00:49:50.039$ would receive an abundance of God.

NOTE Confidence: 0.82798517

 $00:49:50.040 \longrightarrow 00:49:51.560$ One protein that basically

NOTE Confidence: 0.82798517

 $00:49:51.560 \longrightarrow 00:49:53.080$ overwhelms the HSP 70,

NOTE Confidence: 0.82798517

 $00:49:53.080 \longrightarrow 00:49:55.624$ but that we have enough to be able

NOTE Confidence: 0.82798517

00:49:55.624 --> 00:49:57.998 to induce Raceway specific genes,

NOTE Confidence: 0.82798517

 $00:49:58.000 \longrightarrow 00:50:00.520$ so I'd like to end there.

 $00:50:00.520 \longrightarrow 00:50:01.645$ And then just to acknowledge

NOTE Confidence: 0.82798517

 $00{:}50{:}01.645 \dashrightarrow 00{:}50{:}03.030$ people who have done the work.

NOTE Confidence: 0.82798517

 $00:50:03.030 \longrightarrow 00:50:04.578$ I think I've been knowledge Mark

NOTE Confidence: 0.82798517

 $00:50:04.578 \longrightarrow 00:50:05.897$ throughout the talk as well

NOTE Confidence: 0.82798517

 $00:50:05.897 \longrightarrow 00:50:07.542$ as the other Members I lab who

NOTE Confidence: 0.82798517

 $00:50:07.542 \longrightarrow 00:50:08.669$ contributed to this project.

NOTE Confidence: 0.82798517

00:50:08.670 --> 00:50:10.178 The collaborators at Stanford

NOTE Confidence: 0.82798517

 $00:50:10.178 \longrightarrow 00:50:12.063$ and also the other collaborators

NOTE Confidence: 0.82798517

 $00:50:12.063 \longrightarrow 00:50:13.568$ that not at Stanford in,

NOTE Confidence: 0.82798517

 $00:50:13.570 \longrightarrow 00:50:14.614$ particularly Diana Vanessa,

NOTE Confidence: 0.82798517

00:50:14.614 --> 00:50:17.050 for their help in doing the experiments

NOTE Confidence: 0.82798517

 $00:50:17.106 \longrightarrow 00:50:19.134$ with happy One and Megakaryocytic with

NOTE Confidence: 0.82798517

00:50:19.134 --> 00:50:20.893 recite progenitors and the funding

NOTE Confidence: 0.82798517

 $00:50:20.893 \dashrightarrow 00:50:22.668$ sources that supported this work.

NOTE Confidence: 0.82798517

 $00:50:22.670 \longrightarrow 00:50:24.800$ So I'm happy to answer questions.