WEBVTT

- NOTE duration:"00:27:41"
- NOTE recognizability:0.901
- NOTE language:en-us
- NOTE Confidence: 0.69964416
- $00:00:00.000 \dashrightarrow 00:00:06.384$  And I want to introduce Jason Shelter.
- NOTE Confidence: 0.69964416
- 00:00:06.390 --> 00:00:08.982 J<br/>ason is an assistant professor of
- NOTE Confidence: 0.69964416
- 00:00:08.982 --> 00:00:11.496 surgery and received his PhD from MIT,
- NOTE Confidence: 0.69964416
- $00:00:11.500 \longrightarrow 00:00:13.876$  where he worked in the laboratory
- NOTE Confidence: 0.69964416
- 00:00:13.876 --> 00:00:16.788 of Doctor Angelika Amon in the Koch
- NOTE Confidence: 0.69964416
- 00:00:16.788 --> 00:00:18.456 Institute for Cancer Research.
- NOTE Confidence: 0.69964416
- 00:00:18.460 --> 00:00:20.028 After completing his PhD,
- NOTE Confidence: 0.69964416
- $00{:}00{:}20.028 \dashrightarrow 00{:}00{:}21.988$  he established his own research
- NOTE Confidence: 0.69964416
- $00{:}00{:}21.988 \dashrightarrow 00{:}00{:}24.367$  group as an independent fellow at
- NOTE Confidence: 0.69964416
- 00:00:24.367 --> 00:00:26.292 the Cold Spring Harbor Laboratory.
- NOTE Confidence: 0.69964416
- $00{:}00{:}26{.}300 \dashrightarrow 00{:}00{:}28{.}604$  The Shelter Lab is broadly interested
- NOTE Confidence: 0.69964416
- $00{:}00{:}28.604 \dashrightarrow 00{:}00{:}30.648$  in understanding the genomic changes
- NOTE Confidence: 0.69964416
- $00:00:30.648 \rightarrow 00:00:32.420$  that drive cancer progression,
- NOTE Confidence: 0.69964416

 $00:00:32.420 \longrightarrow 00:00:33.174$  particularly aneuploidy,

NOTE Confidence: 0.69964416

 $00:00:33.174 \longrightarrow 00:00:35.436$  which is found in more than

NOTE Confidence: 0.69964416

 $00{:}00{:}35{.}436 \dashrightarrow 00{:}00{:}37{.}131$ 90% of human tumors.

NOTE Confidence: 0.69964416

00:00:37.131 -> 00:00:37.568 Additionally,

NOTE Confidence: 0.69964416

 $00{:}00{:}37.568 \dashrightarrow 00{:}00{:}39.753$  they're working to identify genomic

NOTE Confidence: 0.69964416

 $00{:}00{:}39.753 \dashrightarrow 00{:}00{:}41.722$  alterations that create druggable

NOTE Confidence: 0.69964416

 $00:00:41.722 \rightarrow 00:00:43.678$  the rapeutic vulnerabilities and cancer.

NOTE Confidence: 0.69964416

00:00:43.680 - 00:00:45.632 They have recently discovered

NOTE Confidence: 0.69964416

 $00{:}00{:}45.632 \dashrightarrow 00{:}00{:}47.584$  the first ever selective

NOTE Confidence: 0.69964416

 $00:00:47.584 \rightarrow 00:00:49.950$  inhibitor of the kinase CDK 11,

NOTE Confidence: 0.69964416

 $00{:}00{:}49{.}950 \dashrightarrow 00{:}00{:}53{.}094$  and developing CDK 11 inhibition as a new

NOTE Confidence: 0.69964416

 $00{:}00{:}53.094 \dashrightarrow 00{:}00{:}56.426$  strategy to treat malignancies without.

NOTE Confidence: 0.69964416

 $00:00:56.426 \longrightarrow 00:01:00.256$  Further delay Jason all yours.

NOTE Confidence: 0.959933932857143

 $00:01:00.730 \rightarrow 00:01:03.509$  Thanks so much for the kind introduction,

NOTE Confidence: 0.959933932857143

 $00{:}01{:}03.510 \dashrightarrow 00{:}01{:}05.868$  so I'm very excited to be able to share

NOTE Confidence: 0.959933932857143

 $00:01:05.868 \rightarrow 00:01:07.950$  with you today some research my lab

- NOTE Confidence: 0.959933932857143
- $00{:}01{:}07{.}950 \dashrightarrow 00{:}01{:}10{.}505$  has done about off target activity of
- NOTE Confidence: 0.959933932857143
- $00:01:10.505 \rightarrow 00:01:12.705$  cancer drugs undergoing clinical trials.
- NOTE Confidence: 0.959933932857143
- $00:01:12.710 \longrightarrow 00:01:14.768$  These are my disclosures and this
- NOTE Confidence: 0.959933932857143
- $00:01:14.768 \rightarrow 00:01:16.723$  project really comes from a journal
- NOTE Confidence: 0.959933932857143
- $00:01:16.723 \dashrightarrow 00:01:18.811$  article that I read a few years ago
- NOTE Confidence: 0.959933932857143
- 00:01:18.877 --> 00:01:20.829 that had a statistic in it that I
- NOTE Confidence: 0.959933932857143
- $00:01:20.829 \rightarrow 00:01:22.872$  found to be just absolutely shocking.
- NOTE Confidence: 0.959933932857143
- $00:01:22.872 \longrightarrow 00:01:26.284$  If you look at all drugs that enter
- NOTE Confidence: 0.959933932857143
- 00:01:26.284 --> 00:01:28.288 clinical testing and oncology,
- NOTE Confidence: 0.959933932857143
- $00:01:28.290 \longrightarrow 00:01:30.984 97\%$  of drug indication pairs that
- NOTE Confidence: 0.959933932857143
- $00:01:30.984 \rightarrow 00:01:33.740$  enter clinical trials fail during that
- NOTE Confidence: 0.959933932857143
- 00:01:33.740 --> 00:01:36.386 testing and don't end up receiving
- NOTE Confidence: 0.959933932857143
- $00{:}01{:}36{.}386 \dashrightarrow 00{:}01{:}38{.}606$  FDA approval and this 97% failure
- NOTE Confidence: 0.959933932857143
- 00:01:38.606 --> 00:01:40.574 rate for oncology drugs is the
- NOTE Confidence: 0.959933932857143
- $00:01:40.574 \rightarrow 00:01:42.657$  highest of any field of medicine.
- NOTE Confidence: 0.959933932857143

 $00:01:42.660 \longrightarrow 00:01:45.228$  So more cancer drugs fail than

NOTE Confidence: 0.959933932857143

00:01:45.228 --> 00:01:46.940 psychiatric drugs or endocrinology,

NOTE Confidence: 0.959933932857143

00:01:46.940 --> 00:01:48.808 drugs, or infectious disease,

NOTE Confidence: 0.959933932857143

 $00:01:48.808 \rightarrow 00:01:50.676$  drugs or anything else.

NOTE Confidence: 0.959933932857143

 $00:01:50.680 \longrightarrow 00:01:53.864$  And if you look at the proximate causes

NOTE Confidence: 0.959933932857143

 $00:01:53.864 \rightarrow 00:01:56.501$  for trial failure, the most common,

NOTE Confidence: 0.959933932857143

 $00:01:56.501 \longrightarrow 00:01:58.703$  immediate causes that drugs run into

NOTE Confidence: 0.959933932857143

 $00:01:58.703 \rightarrow 00:02:01.080$  are toxicity and limited efficacy.

NOTE Confidence: 0.959933932857143

 $00:02:01.080 \longrightarrow 00:02:01.644$  That is,

NOTE Confidence: 0.959933932857143

 $00:02:01.644 \longrightarrow 00:02:03.336$  the drugs have too many side

NOTE Confidence: 0.959933932857143

 $00:02:03.336 \dashrightarrow 00:02:05.338$  effects for patients to safely take.

NOTE Confidence: 0.959933932857143

 $00:02:05.340 \rightarrow 00:02:07.276$  Or maybe the patients can safely take them,

NOTE Confidence: 0.959933932857143

 $00:02:07.280 \longrightarrow 00:02:09.513$  but they have limited efficacy and they

NOTE Confidence: 0.959933932857143

 $00:02:09.513 \rightarrow 00:02:11.578$  don't actually shrink the patient's tumor.

NOTE Confidence: 0.959933932857143

 $00:02:11.580 \longrightarrow 00:02:13.267$  And while these are kind of the.

NOTE Confidence: 0.959933932857143

00:02:13.270 --> 00:02:15.302 Proximate causes for oncology

 $00:02:15.302 \longrightarrow 00:02:16.826$  drug trial failure.

NOTE Confidence: 0.959933932857143

 $00:02:16.830 \longrightarrow 00:02:18.615$  I think the underlying reasons

NOTE Confidence: 0.959933932857143

00:02:18.615 --> 00:02:21.400 why so many drugs run into these

NOTE Confidence: 0.959933932857143

 $00:02:21.400 \rightarrow 00:02:23.710$  problems isn't very well understood,

NOTE Confidence: 0.959933932857143

00:02:23.710 --> 00:02:25.950 and today I'm going to share some

NOTE Confidence: 0.959933932857143

 $00{:}02{:}25{.}950 \dashrightarrow 00{:}02{:}28{.}210$  evidence from my lab towards one

NOTE Confidence: 0.959933932857143

 $00:02:28.210 \longrightarrow 00:02:30.260$  potential explanation for this high

NOTE Confidence: 0.959933932857143

 $00:02:30.260 \rightarrow 00:02:32.404$  failure rate and the hypothesis that

NOTE Confidence: 0.959933932857143

 $00{:}02{:}32{.}404 \dashrightarrow 00{:}02{:}35{.}068$  I'm going to argue for is that many

NOTE Confidence: 0.959933932857143

 $00:02:35.068 \rightarrow 00:02:37.402$  drugs are entering clinical testing and

NOTE Confidence: 0.959933932857143

 $00:02:37.402 \rightarrow 00:02:39.543$  oncology with an incorrect understanding

NOTE Confidence: 0.959933932857143

 $00:02:39.543 \longrightarrow 00:02:41.703$  of their mechanism of action,

NOTE Confidence: 0.959933932857143

 $00{:}02{:}41.710 \dashrightarrow 00{:}02{:}43.625$  and I think this mischaracterization

NOTE Confidence: 0.959933932857143

 $00{:}02{:}43.625 \dashrightarrow 00{:}02{:}44.774$  of cancer drugs.

NOTE Confidence: 0.959933932857143

 $00{:}02{:}44.780 \dashrightarrow 00{:}02{:}48.119$  Maybe one factor by no means the only factor,

 $00:02:48.120 \dashrightarrow 00:02:50.440$  but one factor that contributes

NOTE Confidence: 0.959933932857143

 $00{:}02{:}50{.}440 \dashrightarrow 00{:}02{:}52{.}760$  to this extremely high failure

NOTE Confidence: 0.959933932857143

 $00:02:52.835 \longrightarrow 00:02:54.927$  rate for oncology the rapies.

NOTE Confidence: 0.959933932857143

00:02:54.930 --> 00:02:56.246 So in my lab,

NOTE Confidence: 0.959933932857143

 $00:02:56.246 \rightarrow 00:02:57.891$  we've been interested in using

NOTE Confidence: 0.959933932857143

 $00{:}02{:}57{.}891 \dashrightarrow 00{:}02{:}59{.}731$  genetic approaches to investigate

NOTE Confidence: 0.959933932857143

 $00{:}02{:}59{.}731 \dashrightarrow 00{:}03{:}02{.}136$  the mechanisms of action of

NOTE Confidence: 0.959933932857143

 $00:03:02.136 \rightarrow 00:03:03.990$  different experimental cancer drugs,

NOTE Confidence: 0.959933932857143

 $00{:}03{:}03{.}990 \dashrightarrow 00{:}03{:}06{.}246$  and by searching through the current

NOTE Confidence: 0.959933932857143

00:03:06.246 --> 00:03:08.116 literature and looking on clinicaltrials.gov,

NOTE Confidence: 0.959933932857143

 $00:03:08.116 \longrightarrow 00:03:10.678$  we put together a list of 12

NOTE Confidence: 0.959933932857143

 $00:03:10.678 \rightarrow 00:03:12.658$  different drugs targeting 7 different

NOTE Confidence: 0.959933932857143

 $00{:}03{:}12.658 \dashrightarrow 00{:}03{:}14.633$  cancer related proteins that we

NOTE Confidence: 0.959933932857143

00:03:14.633 - > 00:03:16.169 were interested in studying.

NOTE Confidence: 0.959933932857143

 $00:03:16.170 \longrightarrow 00:03:18.516$  These drugs have been used in

NOTE Confidence: 0.959933932857143

 $00{:}03{:}18{.}516 \dashrightarrow 00{:}03{:}20{.}672$  about 30 different clinical trials

- NOTE Confidence: 0.959933932857143
- 00:03:20.672 --> 00:03:23.247 targeting several 100 cancer patients.
- NOTE Confidence: 0.959933932857143
- $00{:}03{:}23{.}250 \dashrightarrow 00{:}03{:}25{.}980$  So six of these proteins are reported
- NOTE Confidence: 0.959933932857143
- $00{:}03{:}25{.}980 \dashrightarrow 00{:}03{:}28{.}869$  to be cancer genetic dependencies.
- NOTE Confidence: 0.959933932857143
- $00:03:28.870 \longrightarrow 00:03:31.005$  That is the function of these proteins
- NOTE Confidence: 0.959933932857143
- $00:03:31.005 \dashrightarrow 00:03:33.408$  is reported to be essential for the
- NOTE Confidence: 0.959933932857143
- $00:03:33.408 \rightarrow 00:03:35.550$  growth and proliferation of cancer cells.
- NOTE Confidence: 0.959933932857143
- $00:03:35.550 \longrightarrow 00:03:36.384$  For instance,
- NOTE Confidence: 0.959933932857143
- $00:03:36.384 \longrightarrow 00:03:38.469$  pack four is a kinase.
- NOTE Confidence: 0.959933932857143
- $00{:}03{:}38{.}470 \dashrightarrow 00{:}03{:}40{.}708$  It's been reported that Pack 4
- NOTE Confidence: 0.959933932857143
- 00:03:40.708 00:03:42.595 kinase activity is essential for
- NOTE Confidence: 0.959933932857143
- $00:03:42.595 \rightarrow 00:03:44.230$  the growth of colon cancer.
- NOTE Confidence: 0.959933932857143
- $00:03:44.230 \longrightarrow 00:03:45.718$  Lung cancer, breast cancer,
- NOTE Confidence: 0.959933932857143
- $00{:}03{:}45{.}718$  -->  $00{:}03{:}47{.}950$  and a few other cancer types.
- NOTE Confidence: 0.959933932857143
- $00{:}03{:}47{.}950 \dashrightarrow 00{:}03{:}50{.}896$  And because of that genetic data
- NOTE Confidence: 0.959933932857143
- $00:03:50.896 \dashrightarrow 00:03:53.800$  concerning pack four that motivated.
- NOTE Confidence: 0.959933932857143

 $00:03:53.800 \rightarrow 00:03:55.924$  Wiser to develop a small molecule

NOTE Confidence: 0.959933932857143

 $00:03:55.924 \longrightarrow 00:03:56.986$  pack for inhibitor

NOTE Confidence: 0.95828214444444

 $00:03:59.020 \longrightarrow 00:04:00.995$  PF 3758309 which they then

NOTE Confidence: 0.95828214444444

 $00:04:00.995 \longrightarrow 00:04:02.575$  entered into clinical testing.

NOTE Confidence: 0.95828214444444

 $00:04:02.580 \dashrightarrow 00:04:04.274$  Caspase 3 is a little bit different.

NOTE Confidence: 0.95828214444444

00:04:04.280 --> 00:04:07.160 I'm going to talk about Caspase 3 separately,

NOTE Confidence: 0.958282144444444

 $00:04:07.160 \dashrightarrow 00:04:09.274$  so we were interested in testing the

NOTE Confidence: 0.95828214444444

 $00:04:09.274 \rightarrow 00:04:11.176$  mechanism of action of these drugs

NOTE Confidence: 0.95828214444444

 $00{:}04{:}11{.}176 \dashrightarrow 00{:}04{:}13{.}054$  and seeing whether they killed cancer

NOTE Confidence: 0.95828214444444

 $00:04:13.054 \rightarrow 00:04:14.901$  cells through the inhibition of these

NOTE Confidence: 0.95828214444444

 $00{:}04{:}14{.}901 \dashrightarrow 00{:}04{:}16{.}998$  proteins and as a first step towards

NOTE Confidence: 0.95828214444444

 $00:04:16.998 \longrightarrow 00:04:19.126$  this process we wanted to confirm that

NOTE Confidence: 0.958282144444444

00:04:19.126 --> 00:04:21.564 the proteins these drugs were targeting

NOTE Confidence: 0.958282144444444

 $00:04:21.564 \rightarrow 00:04:23.639$  were truly cancer genetic dependencies.

NOTE Confidence: 0.958282144444444

 $00:04:23.640 \longrightarrow 00:04:24.770$  That is, they were essential.

NOTE Confidence: 0.95828214444444

 $00:04:24.770 \rightarrow 00:04:27.570$  For cancer growth and to investigate this,

 $00:04:27.570 \longrightarrow 00:04:29.394$  we set up a crisper competition

NOTE Confidence: 0.95828214444444

 $00:04:29.394 \rightarrow 00:04:31.400$  assay to see what happened when

NOTE Confidence: 0.95828214444444

 $00:04:31.400 \longrightarrow 00:04:33.215$  we knocked these jeans out.

NOTE Confidence: 0.95828214444444

00:04:33.220 --> 00:04:34.910 To do this CRISPR assay,

NOTE Confidence: 0.958282144444444

 $00{:}04{:}34{.}910 \dashrightarrow 00{:}04{:}36{.}940$  we transduced cancer cell lines

NOTE Confidence: 0.95828214444444

 $00{:}04{:}36{.}940 \dashrightarrow 00{:}04{:}39{.}812$  with cast 9 and then we transduced

NOTE Confidence: 0.95828214444444

 $00{:}04{:}39{.}812 \dashrightarrow 00{:}04{:}42{.}486$  them a second time with a guide

NOTE Confidence: 0.95828214444444

00:04:42.486 --> 00:04:45.060 RNA coexpressed along with GFP.

NOTE Confidence: 0.95828214444444

 $00{:}04{:}45{.}060 \dashrightarrow 00{:}04{:}47{.}148$  This would then create a mixed

NOTE Confidence: 0.95828214444444

00:04:47.148 --> 00:04:48.925 population of GFP positive cells

NOTE Confidence: 0.95828214444444

00:04:48.925 --> 00:04:51.151 that had the guide RNA and caused

NOTE Confidence: 0.95828214444444

 $00{:}04{:}51{.}151 \dashrightarrow 00{:}04{:}53{.}714$  mutations in the target gene and then

NOTE Confidence: 0.95828214444444

 $00{:}04{:}53{.}714 \dashrightarrow 00{:}04{:}55{.}579$  UN transduced non fluorescent cells.

NOTE Confidence: 0.95828214444444

 $00{:}04{:}55{.}580 \dashrightarrow 00{:}04{:}57{.}700$  We then measure the percentage

NOTE Confidence: 0.95828214444444

 $00:04:57.700 \longrightarrow 00:04:59.820$  of GFP cells over time.

 $00:04:59.820 \longrightarrow 00:05:02.190$  If the percent of GFP positive

NOTE Confidence: 0.95828214444444

00:05:02.190 --> 00:05:03.375 cells decreases overtime,

NOTE Confidence: 0.95828214444444

 $00{:}05{:}03{.}380 \dashrightarrow 00{:}05{:}05{.}096$  that tells us that what ever gene

NOTE Confidence: 0.95828214444444

00:05:05.096 --> 00:05:07.059 the guide RNA is knocking out,

NOTE Confidence: 0.95828214444444

 $00:05:07.060 \longrightarrow 00:05:09.328$  it must be required for cancer growth

NOTE Confidence: 0.95828214444444

 $00:05:09.328 \rightarrow 00:05:11.900$  because the GFP positive cells are dying.

NOTE Confidence: 0.958282144444444

00:05:11.900 --> 00:05:12.584 In contrast,

NOTE Confidence: 0.95828214444444

 $00:05:12.584 \longrightarrow 00:05:14.636$  if the percent of GFP positive

NOTE Confidence: 0.95828214444444

00:05:14.636 - 00:05:16.319 cells stays about the same,

NOTE Confidence: 0.95828214444444

 $00{:}05{:}16.320 \dashrightarrow 00{:}05{:}18.070$  then that's evidence that whatever

NOTE Confidence: 0.95828214444444

00:05:18.070 --> 00:05:19.820 this guide RNA is targeting,

NOTE Confidence: 0.95828214444444

 $00:05:19.820 \longrightarrow 00:05:21.730$  it isn't important for cancer

NOTE Confidence: 0.958282144444444

 $00:05:21.730 \longrightarrow 00:05:23.640$  growth because these GFP positive

NOTE Confidence: 0.958282144444444

 $00:05:23.703 \longrightarrow 00:05:25.218$  cells can grow just fine.

NOTE Confidence: 0.958282144444444

 $00{:}05{:}25{.}220 \dashrightarrow 00{:}05{:}27{.}788$  So that's what the assay looked like we

NOTE Confidence: 0.95828214444444

 $00:05:27.788 \rightarrow 00:05:30.230$  designed and cloned multiple guide RNA's.

 $00:05:30.230 \longrightarrow 00:05:32.180$  Targeting each of the putative

NOTE Confidence: 0.95828214444444

 $00:05:32.180 \longrightarrow 00:05:34.130$  cancer genetic dependencies we were

NOTE Confidence: 0.95828214444444

 $00:05:34.190 \rightarrow 00:05:36.321$  interested in studying and then we

NOTE Confidence: 0.95828214444444

 $00:05:36.321 \rightarrow 00:05:37.983$  did a bunch of competition assays

NOTE Confidence: 0.95828214444444

 $00{:}05{:}37{.}990 \dashrightarrow 00{:}05{:}40{.}209$  and this is what one of these

NOTE Confidence: 0.95828214444444

 $00:05:40.209 \rightarrow 00:05:41.520$  competition assays looks like.

NOTE Confidence: 0.95828214444444

 $00:05:41.520 \longrightarrow 00:05:44.608$  So here we're in MD AMB 231 sells

NOTE Confidence: 0.95828214444444

 $00:05:44.608 \dashrightarrow 00:05:46.153$  a triple negative breast cancer

NOTE Confidence: 0.95828214444444

 $00:05:46.153 \dashrightarrow 00:05:48.179$  cell line as negative controls.

NOTE Confidence: 0.95828214444444

 $00:05:48.180 \dashrightarrow 00:05:50.120$  We have guide RNA's targeting

NOTE Confidence: 0.95828214444444

 $00:05:50.120 \longrightarrow 00:05:52.628$  nonessential loci. Rosa 26 and eight.

NOTE Confidence: 0.958282144444444

 $00{:}05{:}52.628 \dashrightarrow 00{:}05{:}54.788$  The S1 guide RNA's targeting.

NOTE Confidence: 0.95828214444444

 $00:05:54.790 \dashrightarrow 00:05:57.526$  These genes exhibit no drop out.

NOTE Confidence: 0.95828214444444

 $00{:}05{:}57{.}530 \dashrightarrow 00{:}06{:}00{.}050$  As positive controls we have guide RNA's,

NOTE Confidence: 0.95828214444444

 $00{:}06{:}00{.}050 \dashrightarrow 00{:}06{:}02{.}710$  targeting the essential replication genes,

- 00:06:02.710 --> 00:06:04.202 or PA3 and PC,
- NOTE Confidence: 0.95828214444444
- $00:06:04.202 \rightarrow 00:06:06.990$  and a guide RNA's targeting these genes,
- NOTE Confidence: 0.95828214444444
- $00:06:06.990 \rightarrow 00:06:09.306$  which are required for DNA replication.
- NOTE Confidence: 0.95828214444444
- 00:06:09.310 --> 00:06:11.998 Drop out between 50 fold and 200
- NOTE Confidence: 0.95828214444444
- $00:06:11.998 \longrightarrow 00:06:14.988$  fold over 5 passages in culture.
- NOTE Confidence: 0.95828214444444
- $00:06:14.990 \rightarrow 00:06:16.970$  We then looked at the effects of guide RNA's,
- NOTE Confidence: 0.958282144444444
- $00:06:16.970 \longrightarrow 00:06:18.720$  targeting each of the putative
- NOTE Confidence: 0.95828214444444
- $00:06:18.720 \rightarrow 00:06:20.120$  cancer genetic dependencies that
- NOTE Confidence: 0.95828214444444
- 00:06:20.120 --> 00:06:22.009 we were interested in studying,
- NOTE Confidence: 0.95828214444444
- $00:06:22.010 \rightarrow 00:06:24.775$  and we were really astounded when the
- NOTE Confidence: 0.95828214444444
- $00{:}06{:}24.775 \dashrightarrow 00{:}06{:}27.550$  guide RNA's targeting these cancer drug
- NOTE Confidence: 0.95828214444444
- $00:06:27.550 \rightarrow 00:06:29.930$  targets exhibited no dropout whatsoever.
- NOTE Confidence: 0.95828214444444
- 00:06:29.930 --> 00:06:31.690 These guide RNAs behaved exactly
- NOTE Confidence: 0.958282144444444
- $00:06:31.690 \longrightarrow 00:06:33.450$  the same as guide RNA's,
- NOTE Confidence: 0.95828214444444
- $00:06:33.450 \longrightarrow 00:06:35.382$  targeting known non essential
- NOTE Confidence: 0.95828214444444
- $00:06:35.382 \rightarrow 00:06:38.730$  genes like Rosa 26 and a VS1.

 $00:06:38.730 \rightarrow 00:06:40.776$  This was incredibly surprising to us

NOTE Confidence: 0.95828214444444

 $00:06:40.776 \rightarrow 00:06:43.145$  because right now there are patients who

NOTE Confidence: 0.95828214444444

 $00:06:43.145 \longrightarrow 00:06:45.336$  are receiving anti htac 6 therapy and.

NOTE Confidence: 0.95828214444444

00:06:45.340 --> 00:06:47.338 Anti milk therapy and anti Kim.

NOTE Confidence: 0.95828214444444

 $00:06:47.340 \rightarrow 00:06:50.308$  One therapy based on the belief that these

NOTE Confidence: 0.95828214444444

 $00:06:50.308 \rightarrow 00:06:52.938$  proteins are required for cancer growth,

NOTE Confidence: 0.95828214444444

 $00:06:52.940 \longrightarrow 00:06:55.784$  but this experiment suggests that in

NOTE Confidence: 0.95828214444444

 $00{:}06{:}55{.}784 \dashrightarrow 00{:}06{:}58{.}126$  these experimental conditions in this

NOTE Confidence: 0.95828214444444

 $00:06:58.126 \rightarrow 00:07:00.765$  cell line we can eliminate these genes

NOTE Confidence: 0.95828214444444

 $00:07:00.765 \rightarrow 00:07:03.520$  without any effect on cancer whatsoever.

NOTE Confidence: 0.965021968181818

 $00:07:03.520 \longrightarrow 00:07:04.864$  So this is what it looked

NOTE Confidence: 0.965021968181818

 $00:07:04.864 \longrightarrow 00:07:06.080$  like in one cell line.

NOTE Confidence: 0.965021968181818

 $00{:}07{:}06.080 \dashrightarrow 00{:}07{:}08.814$  We ended up repeating this faceing 32

NOTE Confidence: 0.965021968181818

00:07:08.814 --> 00:07:10.338 different cancer cell lines from more

NOTE Confidence: 0.965021968181818

 $00:07:10.338 \longrightarrow 00:07:12.259$  than a dozen different cancer types,

 $00:07:12.260 \longrightarrow 00:07:13.568$  and in each of these experiments

NOTE Confidence: 0.965021968181818

 $00{:}07{:}13.568 \dashrightarrow 00{:}07{:}14.810$  we got the same result.

NOTE Confidence: 0.965021968181818

 $00{:}07{:}14.810 \dashrightarrow 00{:}07{:}17.411$  There is no drop out of the guide RNA's

NOTE Confidence: 0.965021968181818

 $00:07:17.411 \longrightarrow 00:07:19.329$  targeting these drug targets and there

NOTE Confidence: 0.965021968181818

 $00:07:19.329 \longrightarrow 00:07:21.998$  was no evidence that any of these genes

NOTE Confidence: 0.965021968181818

 $00:07:21.998 \rightarrow 00:07:24.434$  were actually dependency in any cancer type.

NOTE Confidence: 0.965021968181818

 $00:07:24.440 \rightarrow 00:07:26.610$  So this made us take a step back and think,

NOTE Confidence: 0.965021968181818

00:07:26.610 - 00:07:27.780 well, is there something that

NOTE Confidence: 0.965021968181818

 $00:07:27.780 \longrightarrow 00:07:29.480$  could be going wrong in this assay?

NOTE Confidence: 0.965021968181818

00:07:29.480 --> 00:07:31.370 Could we be you know doing

NOTE Confidence: 0.965021968181818

 $00:07:31.370 \longrightarrow 00:07:32.315$  something incorrect here?

NOTE Confidence: 0.965021968181818

 $00:07:32.320 \longrightarrow 00:07:33.624$  And so we thought.

NOTE Confidence: 0.965021968181818

00:07:33.624 --> 00:07:35.609 Well, maybe with CRISPR.

NOTE Confidence: 0.965021968181818

00:07:35.609 --> 00:07:37.268 We're generating heterozygous

NOTE Confidence: 0.965021968181818

 $00:07:37.268 \rightarrow 00:07:40.270$  mutations but not homozygous mutations.

NOTE Confidence: 0.965021968181818

00:07:40.270 - 00:07:41.785 You know, maybe we're we're

 $00:07:41.785 \rightarrow 00:07:43.300$  introducing mutations into these genes,

NOTE Confidence: 0.965021968181818

00:07:43.300 --> 00:07:45.120 but we're not really knocking

NOTE Confidence: 0.965021968181818

 $00{:}07{:}45{.}120 \dashrightarrow 00{:}07{:}46{.}576$  out the total protein.

NOTE Confidence: 0.965021968181818

 $00:07:46.580 \rightarrow 00:07:47.976$  So we thought OK,

NOTE Confidence: 0.965021968181818

 $00{:}07{:}47.976 \dashrightarrow 00{:}07{:}49.721$  instead of doing this population

NOTE Confidence: 0.965021968181818

 $00:07:49.721 \longrightarrow 00:07:50.700$  based approach,

NOTE Confidence: 0.965021968181818

00:07:50.700 --> 00:07:52.722 let's make single cell Dr Knockout

NOTE Confidence: 0.965021968181818

 $00:07:52.722 \longrightarrow 00:07:55.371$  clones and be as sure as humanly

NOTE Confidence: 0.965021968181818

00:07:55.371 - 00:07:57.426 possible that we were really

NOTE Confidence: 0.965021968181818

00:07:57.426 --> 00:08:00.160 eliminating 100% of the target protein.

NOTE Confidence: 0.965021968181818

 $00{:}08{:}00{.}160 \dashrightarrow 00{:}08{:}01{.}196$  So we did that.

NOTE Confidence: 0.965021968181818

00:08:01.196 --> 00:08:03.178 We used A2 CRISPR guide RNA strategy

NOTE Confidence: 0.965021968181818

 $00{:}08{:}03{.}178 \dashrightarrow 00{:}08{:}05{.}747$  where we designed to guide RNA targeting

NOTE Confidence: 0.965021968181818

00:08:05.747 --> 00:08:07.575 an upstream exon into downstream

NOTE Confidence: 0.965021968181818

 $00:08:07.575 \rightarrow 00:08:09.633$  exon so that we could physically

 $00:08:09.633 \longrightarrow 00:08:12.068$  cut the gene out of the genome.

NOTE Confidence: 0.965021968181818

00:08:12.070 -> 00:08:14.009 And there would be no protein left.

NOTE Confidence: 0.965021968181818

 $00:08:14.010 \longrightarrow 00:08:16.278$  So we sorted a single cells

NOTE Confidence: 0.965021968181818

 $00:08:16.278 \rightarrow 00:08:17.790$  that were double positive.

NOTE Confidence: 0.965021968181818

 $00{:}08{:}17.790 \dashrightarrow 00{:}08{:}20.070$  That picked up both guide RNA's

NOTE Confidence: 0.965021968181818

 $00{:}08{:}20{.}070 \dashrightarrow 00{:}08{:}22{.}942$  that we transduced in and then we

NOTE Confidence: 0.965021968181818

 $00:08:22.942 \longrightarrow 00:08:24.958$  verified target knockout using

NOTE Confidence: 0.965021968181818

 $00{:}08{:}24.958 \dashrightarrow 00{:}08{:}26.470$  two independent antibodies.

NOTE Confidence: 0.965021968181818

 $00:08:26.470 \longrightarrow 00:08:29.039$  So for instance 1 gene we were

NOTE Confidence: 0.965021968181818

 $00:08:29.039 \longrightarrow 00:08:31.048$  interested in studying as math K14.

NOTE Confidence: 0.965021968181818

 $00{:}08{:}31{.}050 \dashrightarrow 00{:}08{:}32{.}736$  This is the gene that encodes

NOTE Confidence: 0.965021968181818

 $00:08:32.736 \longrightarrow 00:08:34.112$  the kinase P38 alpha.

NOTE Confidence: 0.965021968181818

 $00{:}08{:}34{.}112 \dashrightarrow 00{:}08{:}36{.}548$  We generated knock out clones and we

NOTE Confidence: 0.965021968181818

 $00{:}08{:}36{.}548 \dashrightarrow 00{:}08{:}38{.}198$  verified complete target knock out

NOTE Confidence: 0.965021968181818

 $00:08:38.198 \rightarrow 00:08:40.520$  using one antibody and then verified

NOTE Confidence: 0.965021968181818

 $00:08:40.520 \rightarrow 00:08:42.877$  it again using a second antibody.

- NOTE Confidence: 0.965021968181818
- $00:08:42.880 \longrightarrow 00:08:44.120$  So that we could be,
- NOTE Confidence: 0.965021968181818
- 00:08:44.120 --> 00:08:44.830 you know,
- NOTE Confidence: 0.965021968181818
- $00:08:44.830 \longrightarrow 00:08:46.960$  as sure as physically possible that
- NOTE Confidence: 0.965021968181818
- $00:08:46.960 \rightarrow 00:08:49.918$  we had truly eliminated all trace of
- NOTE Confidence: 0.965021968181818
- $00:08:49.918 \rightarrow 00:08:52.790$  these putative cancer drivers from the cell.
- NOTE Confidence: 0.965021968181818
- 00:08:52.790 --> 00:08:53.120 However,
- NOTE Confidence: 0.965021968181818
- $00{:}08{:}53.120 \dashrightarrow 00{:}08{:}55.100$  when we tested the fitness effects
- NOTE Confidence: 0.965021968181818
- $00:08:55.100 \rightarrow 00:08:56.530$  of these knockout clones,
- NOTE Confidence: 0.965021968181818
- $00:08:56.530 \longrightarrow 00:08:58.354$  we got exactly the same result
- NOTE Confidence: 0.965021968181818
- $00:08:58.354 \rightarrow 00:09:00.643$  that we got from the competition
- NOTE Confidence: 0.965021968181818
- $00:09:00.643 \rightarrow 00:09:03.163$  assays knocking out these putative
- NOTE Confidence: 0.965021968181818
- $00:09:03.163 \dashrightarrow 00:09:05.079$  cancer genetic dependencies had
- NOTE Confidence: 0.965021968181818
- $00{:}09{:}05{.}079 \dashrightarrow 00{:}09{:}06{.}749$  no effect on cancer growth.
- NOTE Confidence: 0.965021968181818
- $00:09:06.750 \longrightarrow 00:09:08.054$  So here, for instance,
- NOTE Confidence: 0.965021968181818
- $00:09:08.054 \rightarrow 00:09:09.684$  is a proliferation assay in
- NOTE Confidence: 0.965021968181818

 $00{:}09{:}09{.}684 \dashrightarrow 00{:}09{:}11{.}268$ a Melanoma cell line.

NOTE Confidence: 0.965021968181818

00:09:11.270 --> 00:09:13.508 We have three map K14 knock out

NOTE Confidence: 0.965021968181818

 $00{:}09{:}13.508 \dashrightarrow 00{:}09{:}15.859$  clones and then two control rows

NOTE Confidence: 0.965021968181818

 $00:09:15.859 \longrightarrow 00:09:18.575$  of 26 clones and these map K14

NOTE Confidence: 0.965021968181818

 $00:09:18.575 \dashrightarrow 00:09:20.390$  knock out clones grow exactly as

NOTE Confidence: 0.965021968181818

 $00{:}09{:}20{.}390 \dashrightarrow 00{:}09{:}22{.}743$  well as the rows of 26 control.

NOTE Confidence: 0.965021968181818

00:09:22.743 --> 00:09:23.076 Jones,

NOTE Confidence: 0.965021968181818

 $00{:}09{:}23.076 \dashrightarrow 00{:}09{:}25.740$  we could also put these cells in soft

NOTE Confidence: 0.965021968181818

 $00:09:25.811 \rightarrow 00:09:28.326$  Agar challenge their clonogenic ability.

NOTE Confidence: 0.965021968181818

 $00{:}09{:}28.330 \dashrightarrow 00{:}09{:}30.565$  We saw no difference in

NOTE Confidence: 0.965021968181818

00:09:30.565 --> 00:09:31.906 Clonogenic ability either.

NOTE Confidence: 0.965021968181818

 $00{:}09{:}31{.}910 \dashrightarrow 00{:}09{:}34{.}898$  These knock out cells grew just fine.

NOTE Confidence: 0.965021968181818

00:09:34.900 --> 00:09:35.736 So to sum up,

NOTE Confidence: 0.965021968181818

 $00:09:35.736 \longrightarrow 00:09:37.272$  a whole bunch of data that I

NOTE Confidence: 0.965021968181818

 $00:09:37.272 \longrightarrow 00:09:38.616$  don't have time to show you.

NOTE Confidence: 0.965021968181818

 $00:09:38.620 \longrightarrow 00:09:40.804$  We ended up eliminating all six

- NOTE Confidence: 0.965021968181818
- 00:09:40.804 --> 00:09:42.260 different cancer driver genes
- NOTE Confidence: 0.965021968181818
- $00:09:42.321 \longrightarrow 00:09:44.659$  that we were studying in at least
- NOTE Confidence: 0.965021968181818
- $00:09:44.659 \rightarrow 00:09:46.379$  three different cancer types each,
- NOTE Confidence: 0.965021968181818
- $00{:}09{:}46{.}380 \dashrightarrow 00{:}09{:}48{.}840$  and there was no fitness effect
- NOTE Confidence: 0.965021968181818
- $00{:}09{:}48.840 \dashrightarrow 00{:}09{:}51.140$  what soever that we could discuss.
- NOTE Confidence: 0.965021968181818
- $00{:}09{:}51{.}140 \dashrightarrow 00{:}09{:}52{.}568$  So this was a really strange
- NOTE Confidence: 0.965021968181818
- $00:09:52.568 \rightarrow 00:09:54.472$  finding to us and it made us try
- NOTE Confidence: 0.965021968181818
- $00:09:54.472 \longrightarrow 00:09:56.020$  to figure out what was going on.
- NOTE Confidence: 0.965021968181818
- $00:09:56.020 \longrightarrow 00:09:57.478$  So we were looking at the
- NOTE Confidence: 0.965021968181818
- $00:09:57.478 \rightarrow 00:09:58.450$  targets of 12 different
- NOTE Confidence: 0.908104099333333
- $00{:}09{:}58{.}503 \dashrightarrow 00{:}10{:}00{.}375$  anti cancer drugs in various stages
- NOTE Confidence: 0.908104099333333
- $00{:}10{:}00{.}375 \dashrightarrow 00{:}10{:}01{.}988$  of clinical development and we
- NOTE Confidence: 0.908104099333333
- $00:10:01.988 \rightarrow 00:10:03.830$  looked at these drug targets with
- NOTE Confidence: 0.908104099333333
- $00{:}10{:}03.830 \dashrightarrow 00{:}10{:}05.240$  multiple different CRISPR techniques.
- NOTE Confidence: 0.908104099333333
- $00:10:05.240 \longrightarrow 00:10:07.640$  We did CRISPR competition assays.
- NOTE Confidence: 0.908104099333333

00:10:07.640 --> 00:10:09.584 We made CRISPR knockouts,

NOTE Confidence: 0.908104099333333

00:10:09.584 --> 00:10:10.556 but concordantly.

NOTE Confidence: 0.908104099333333

 $00{:}10{:}10{.}560 \dashrightarrow 00{:}10{:}13.262$  They both showed that we could eliminate

NOTE Confidence: 0.908104099333333

 $00:10:13.262 \rightarrow 00:10:15.456$  these jeans without a detrimental

NOTE Confidence: 0.908104099333333

 $00{:}10{:}15{.}456 \dashrightarrow 00{:}10{:}17{.}388$  effect on cancer proliferation.

NOTE Confidence: 0.908104099333333

 $00:10:17.390 \dashrightarrow 00:10:19.346$  This then raised the question well,

NOTE Confidence: 0.908104099333333

00:10:19.350 - 00:10:21.821 why were these genes believed to be

NOTE Confidence: 0.908104099333333

 $00:10:21.821 \rightarrow 00:10:23.850$  cancer essential in the 1st place?

NOTE Confidence: 0.908104099333333

 $00:10:23.850 \longrightarrow 00:10:25.698$  And when we looked into the

NOTE Confidence: 0.908104099333333

 $00:10:25.698 \rightarrow 00:10:26.930$  literature on these genes,

NOTE Confidence: 0.908104099333333

 $00{:}10{:}26{.}930 \dashrightarrow 00{:}10{:}29{.}346$  we found the two main lines of evidence

NOTE Confidence: 0.908104099333333

 $00:10:29.346 \longrightarrow 00:10:31.237$  had identified these genes as cancer,

NOTE Confidence: 0.908104099333333

 $00:10:31.240 \longrightarrow 00:10:32.130$  essential initially.

NOTE Confidence: 0.908104099333333

 $00:10:32.130 \longrightarrow 00:10:34.800$  The first line of evidence identifying

NOTE Confidence: 0.908104099333333

 $00{:}10{:}34{.}800 \dashrightarrow 00{:}10{:}37{.}562$  these genes as cancer essential were

NOTE Confidence: 0.908104099333333

 $00:10:37.562 \rightarrow 00:10:39.822$  experiments done using RNA interference.

- NOTE Confidence: 0.908104099333333
- $00:10:39.830 \longrightarrow 00:10:42.092$  The second line of evidence were
- NOTE Confidence: 0.908104099333333
- $00{:}10{:}42.092 \dashrightarrow 00{:}10{:}44.589$  experiments done using small molecule drugs,
- NOTE Confidence: 0.908104099333333
- $00:10:44.590 \longrightarrow 00:10:46.606$  many of which had then gone
- NOTE Confidence: 0.908104099333333
- $00:10:46.606 \rightarrow 00:10:48.590$  on to enter clinical testing.
- NOTE Confidence: 0.908104099333333
- $00:10:48.590 \longrightarrow 00:10:50.590$  So we wanted to see if we could
- NOTE Confidence: 0.908104099333333
- $00{:}10{:}50{.}590 \dashrightarrow 00{:}10{:}52{.}253$  backtrack a little and understand why
- NOTE Confidence: 0.908104099333333
- $00{:}10{:}52.253 \dashrightarrow 00{:}10{:}54.889$  we had come to such a different result
- NOTE Confidence: 0.908104099333333
- $00:10:54.890 \longrightarrow 00:10:57.230$  than these previous experiments done.
- NOTE Confidence: 0.908104099333333
- $00{:}10{:}57{.}230 \dashrightarrow 00{:}11{:}00{.}527$  Using RNA I and small molecule drugs.
- NOTE Confidence: 0.908104099333333
- $00:11:00.530 \longrightarrow 00:11:02.364$  So I'll first show you what we
- NOTE Confidence: 0.908104099333333
- $00:11:02.364 \rightarrow 00:11:04.087$  learned when we looked at some
- NOTE Confidence: 0.908104099333333
- 00:11:04.087 --> 00:11:05.833 of the prior RNA I experiments.
- NOTE Confidence: 0.908104099333333
- 00:11:05.840 --> 00:11:07.751 So this is an RNA I experiment
- NOTE Confidence: 0.908104099333333
- $00{:}11{:}07{.}751 \dashrightarrow 00{:}11{:}09{.}622$  published in the literature a few
- NOTE Confidence: 0.908104099333333
- $00:11:09.622 \dashrightarrow 00:11:11.608$  years ago that had identified the
- NOTE Confidence: 0.908104099333333

 $00:11:11.608 \rightarrow 00:11:13.509$  kinase pack for as essential for

NOTE Confidence: 0.908104099333333

 $00{:}11{:}13.509 \dashrightarrow 00{:}11{:}15.528$  the growth of colon cancer cells.

NOTE Confidence: 0.908104099333333

 $00{:}11{:}15{.}528 \dashrightarrow 00{:}11{:}17{.}052$  In this experiment,

NOTE Confidence: 0.908104099333333

 $00:11:17.052 \rightarrow 00:11:18.968$  the investigators took SI

NOTE Confidence: 0.908104099333333

 $00:11:18.968 \longrightarrow 00:11:20.520$  RNA's targeting pack four.

NOTE Confidence: 0.908104099333333

00:11:20.520 --> 00:11:23.478 They introduced them into HCT 116,

NOTE Confidence: 0.908104099333333

00:11:23.480 --> 00:11:24.590 colon cancer cells,

NOTE Confidence: 0.908104099333333

 $00{:}11{:}24{.}590 \dashrightarrow 00{:}11{:}27{.}785$  and they found that the SI RNAs decreased

NOTE Confidence: 0.908104099333333

00:11:27.785 --> 00:11:30.598 colon cancer cell survival data like

NOTE Confidence: 0.908104099333333

 $00{:}11{:}30{.}598 \dashrightarrow 00{:}11{:}33{.}111$  this motivated Pfizer to enter a pack

NOTE Confidence: 0.908104099333333

 $00{:}11{:}33{.}111 \dashrightarrow 00{:}11{:}35{.}729$  for inhibitor into clinical trials.

NOTE Confidence: 0.908104099333333

 $00{:}11{:}35{.}730 \dashrightarrow 00{:}11{:}37{.}930$  We had found no fitness effect when we

NOTE Confidence: 0.908104099333333

00:11:37.930 --> 00:11:39.968 had knocked out packed 4 using CRISPR,

NOTE Confidence: 0.908104099333333

 $00:11:39.970 \longrightarrow 00:11:42.322$  so we wanted to see if we could

NOTE Confidence: 0.908104099333333

 $00:11:42.322 \rightarrow 00:11:43.614$  recapitulate this result that

NOTE Confidence: 0.908104099333333

 $00:11:43.614 \rightarrow 00:11:45.546$  had been published using RNA I.

- NOTE Confidence: 0.908104099333333
- $00{:}11{:}45{.}550 \dashrightarrow 00{:}11{:}48{.}175$  Two of these SI RNA constructs were
- NOTE Confidence: 0.908104099333333
- $00{:}11{:}48.175 \dashrightarrow 00{:}11{:}49.729$  commercially available and we had
- NOTE Confidence: 0.908104099333333
- $00{:}11{:}49{.}730 \dashrightarrow 00{:}11{:}51{.}865$  HCT 116 cells growing in my lab,
- NOTE Confidence: 0.908104099333333
- $00:11:51.870 \longrightarrow 00:11:53.970$  so we purchased these siren's
- NOTE Confidence: 0.908104099333333
- $00{:}11{:}53{.}970 \dashrightarrow 00{:}11{:}56{.}070$  from this prior publication and
- NOTE Confidence: 0.908104099333333
- $00:11:56.143 \longrightarrow 00:11:58.207$  then tested them in our cells.
- NOTE Confidence: 0.908104099333333
- 00:11:58.210 --> 00:11:59.918 We transfected these siren's,
- NOTE Confidence: 0.908104099333333
- $00:11:59.918 \rightarrow 00:12:02.480$  the same from the prior publication
- NOTE Confidence: 0.908104099333333
- 00:12:02.547 --> 00:12:03.800 into HCT 116 cells,
- NOTE Confidence: 0.908104099333333
- $00{:}12{:}03{.}800 \dashrightarrow 00{:}12{:}06{.}110$  and we could confirm by Western blot.
- NOTE Confidence: 0.908104099333333
- $00{:}12{:}06{.}110 \dashrightarrow 00{:}12{:}09{.}160$  These SI RNAs decrease protein
- NOTE Confidence: 0.908104099333333
- $00{:}12{:}09{.}160 \dashrightarrow 00{:}12{:}11.845$  expression as expected and we did
- NOTE Confidence: 0.908104099333333
- $00{:}12{:}11.845 \dashrightarrow 00{:}12{:}13.840$  a self survival as say and we could
- NOTE Confidence: 0.908104099333333
- 00:12:13.908 --> 00:12:16.258 confirm that they killed HCT 116.
- NOTE Confidence: 0.908104099333333
- 00:12:16.258 --> 00:12:17.914 Colon cancer cells exactly
- NOTE Confidence: 0.908104099333333

 $00:12:17.914 \longrightarrow 00:12:19.570$  as had been reported.

NOTE Confidence: 0.908104099333333

 $00{:}12{:}19{.}570 \dashrightarrow 00{:}12{:}20{.}743$  However, using CRISPR,

NOTE Confidence: 0.908104099333333

 $00:12:20.743 \longrightarrow 00:12:23.480$  we were also able to generate a

NOTE Confidence: 0.908104099333333

 $00:12:23.554 \rightarrow 00:12:25.900$  pack for knockout clone in this

NOTE Confidence: 0.908104099333333

00:12:25.900 --> 00:12:27.900 exact same cancer cell line.

NOTE Confidence: 0.908104099333333

 $00{:}12{:}27{.}900 \dashrightarrow 00{:}12{:}30{.}105$  So here we had a pack for knockout clone.

NOTE Confidence: 0.908104099333333

 $00:12:30.110 \longrightarrow 00:12:32.406$  You can see there's no pack for

NOTE Confidence: 0.908104099333333

 $00:12:32.406 \rightarrow 00:12:34.277$  expression in either the control

NOTE Confidence: 0.908104099333333

 $00{:}12{:}34{.}277 \dashrightarrow 00{:}12{:}35{.}917$  or the knockdown condition.

NOTE Confidence: 0.908104099333333

 $00:12:35.920 \longrightarrow 00:12:37.999$  And then when we did a self

NOTE Confidence: 0.908104099333333

 $00:12:37.999 \longrightarrow 00:12:39.760$  survival assay on these cells,

NOTE Confidence: 0.908104099333333

 $00:12:39.760 \longrightarrow 00:12:41.920$  we found that transfecting the

NOTE Confidence: 0.908104099333333

00:12:41.920 --> 00:12:44.853 pack 4 knockout cells with pack 4

NOTE Confidence: 0.908104099333333

 $00:12:44.853 \rightarrow 00:12:47.872$  targeting SI RNA had exactly the same

NOTE Confidence: 0.908104099333333

 $00{:}12{:}47.872 \dashrightarrow 00{:}12{:}50.337$  detrimental impact on colon cancer

NOTE Confidence: 0.908104099333333

 $00:12:50.337 \rightarrow 00:12:53.284$  survival as it did in the pack for

- NOTE Confidence: 0.908104099333333
- $00:12:53.284 \dashrightarrow 00:12:55.459$  expressing Rosa 26 control cells.
- NOTE Confidence: 0.908104099333333
- $00{:}12{:}55{.}460 \dashrightarrow 00{:}12{:}57{.}959$  So these packed 4 targeting SI RNAs
- NOTE Confidence: 0.908104099333333
- 00:12:57.959 --> 00:12:59.979 are killing colon cancer cells,
- NOTE Confidence: 0.908104099333333
- $00:12:59.980 \longrightarrow 00:13:01.816$  but their ability to kill colon
- NOTE Confidence: 0.908104099333333
- 00:13:01.816 --> 00:13:03.040 cancer cells is entirely
- NOTE Confidence: 0.929160932631579
- $00{:}13{:}03.095 \dashrightarrow 00{:}13{:}04.527$  independent of the expression
- NOTE Confidence: 0.929160932631579
- 00:13:04.527 --> 00:13:06.317 of pack four because they're.
- NOTE Confidence: 0.929160932631579
- $00:13:06.320 \rightarrow 00:13:09.134$  Exactly as lethal in the control cells
- NOTE Confidence: 0.929160932631579
- $00{:}13{:}09{.}134 \dashrightarrow 00{:}13{:}12.045$  expressing pack four as they are in the pack.
- NOTE Confidence: 0.929160932631579
- $00:13:12.050 \longrightarrow 00:13:13.890$  4 knockout clones that we
- NOTE Confidence: 0.929160932631579
- $00:13:13.890 \longrightarrow 00:13:14.994$  generated using crisper.
- NOTE Confidence: 0.929160932631579
- $00{:}13{:}15{.}000 \dashrightarrow 00{:}13{:}16{.}960$  So this prior experiment was
- NOTE Confidence: 0.929160932631579
- 00:13:16.960 --> 00:13:18.136 was totally reproducible.
- NOTE Confidence: 0.929160932631579
- $00{:}13{:}18{.}140 \dashrightarrow 00{:}13{:}21{.}338$  These sirens killed colon cancer cells,
- NOTE Confidence: 0.929160932631579
- $00{:}13{:}21{.}340 \dashrightarrow 00{:}13{:}23{.}704$  but just the interpretation was wrong
- NOTE Confidence: 0.929160932631579

 $00:13:23.704 \rightarrow 00:13:26.319$  because the toxicity of these Sir nase,

NOTE Confidence: 0.929160932631579

 $00:13:26.320 \longrightarrow 00:13:28.120$  is just entirely independent

NOTE Confidence: 0.929160932631579

00:13:28.120 --> 00:13:29.920 of pack for expression,

NOTE Confidence: 0.929160932631579

 $00:13:29.920 \longrightarrow 00:13:32.230$  and this seems to be commonly

NOTE Confidence: 0.929160932631579

 $00{:}13{:}32{.}230 \dashrightarrow 00{:}13{:}34{.}776$  the case where we test SIRN as

NOTE Confidence: 0.929160932631579

 $00{:}13{:}34{.}776$  -->  $00{:}13{:}36{.}534$  and SH RNA's in the literature.

NOTE Confidence: 0.929160932631579

 $00{:}13{:}36{.}540 \dashrightarrow 00{:}13{:}39{.}070$  Over CRISPR derived knockout clones.

NOTE Confidence: 0.929160932631579

00:13:39.070 --> 00:13:42.310 The SI and SH RNA's may kill cancer cells,

NOTE Confidence: 0.929160932631579

 $00:13:42.310 \longrightarrow 00:13:44.676$  but it's just independent of the expression

NOTE Confidence: 0.929160932631579

 $00:13:44.676 \rightarrow 00:13:47.718$  of the gene that they were designed against.

NOTE Confidence: 0.929160932631579

 $00{:}13{:}47.720 \dashrightarrow 00{:}13{:}49.176$  The next thing that we wanted to

NOTE Confidence: 0.929160932631579

 $00{:}13{:}49{.}176 \dashrightarrow 00{:}13{:}50{.}794$  figure out was what was going on

NOTE Confidence: 0.929160932631579

 $00:13:50.794 \rightarrow 00:13:51.974$  with the small molecule drugs,

NOTE Confidence: 0.929160932631579

 $00:13:51.980 \rightarrow 00:13:54.563$  many of which had then gone on to enter

NOTE Confidence: 0.929160932631579

 $00{:}13{:}54{.}563 \dashrightarrow 00{:}13{:}56{.}441$  clinical testing and I'll show you

NOTE Confidence: 0.929160932631579

 $00:13:56.441 \rightarrow 00:13:58.749$  what happened with one of those drugs.

 $00:13:58.750 \longrightarrow 00:14:00.982$  So pack one is a drug that was

NOTE Confidence: 0.929160932631579

 $00{:}14{:}00{.}982 \dashrightarrow 00{:}14{:}03{.}214$  described with few years ago in a

NOTE Confidence: 0.929160932631579

 $00:14:03.214 \rightarrow 00:14:04.824$  paper in nature chemical biology.

NOTE Confidence: 0.929160932631579

 $00:14:04.830 \longrightarrow 00:14:07.595$  It was developed as a Caspase 3

NOTE Confidence: 0.929160932631579

 $00{:}14{:}07{.}595 \dashrightarrow 00{:}14{:}09{.}677$  activator compound so the apoptosis

NOTE Confidence: 0.929160932631579

00:14:09.677 --> 00:14:12.251 enzyme caspase 3 is normally present

NOTE Confidence: 0.929160932631579

00:14:12.251 --> 00:14:14.952 in an inactive proca<br/>spase state in

NOTE Confidence: 0.929160932631579

 $00{:}14{:}14{.}952 \dashrightarrow 00{:}14{:}17{.}785$  the cell and pack one was developed

NOTE Confidence: 0.929160932631579

 $00{:}14{:}17.785 \dashrightarrow 00{:}14{:}20.221$  to catalyze the conversion of caspase

NOTE Confidence: 0.929160932631579

 $00:14:20.221 \longrightarrow 00:14:22.246$  3 from its inactive procaspase

NOTE Confidence: 0.929160932631579

00:14:22.246 --> 00:14:25.060 state to its active caspase 3 state,

NOTE Confidence: 0.929160932631579

 $00{:}14{:}25{.}060 \dashrightarrow 00{:}14{:}27{.}076$  at which point it would then

NOTE Confidence: 0.929160932631579

 $00:14:27.076 \longrightarrow 00:14:29.116$  kill cancer cells in this drug.

NOTE Confidence: 0.929160932631579

 $00{:}14{:}29{.}116 \dashrightarrow 00{:}14{:}30{.}826$  Has been entered into three

NOTE Confidence: 0.929160932631579

 $00{:}14{:}30{.}826$  -->  $00{:}14{:}31{.}970$  different clinical trials.

00:14:31.970 --> 00:14:32.368 However,

NOTE Confidence: 0.929160932631579

 $00{:}14{:}32{.}368 \dashrightarrow 00{:}14{:}34{.}756$  this mechanism of action was worked

NOTE Confidence: 0.929160932631579

 $00{:}14{:}34.756 \dashrightarrow 00{:}14{:}37.221$  out based on in vitro biochemistry

NOTE Confidence: 0.929160932631579

 $00:14:37.221 \rightarrow 00:14:40.007$  and no one had described a mutation

NOTE Confidence: 0.929160932631579

 $00{:}14{:}40.083 \dashrightarrow 00{:}14{:}42.219$  in Caspase 3 that conferred resistance

NOTE Confidence: 0.929160932631579

 $00{:}14{:}42.219 \dashrightarrow 00{:}14{:}44.982$  to it or had assessed the effects of

NOTE Confidence: 0.929160932631579

 $00{:}14{:}44{.}982 \dashrightarrow 00{:}14{:}48{.}029$  this drug in a Caspase 3 knockout cell.

NOTE Confidence: 0.929160932631579

 $00:14:48.030 \rightarrow 00:14:50.664$  So using CRISPR we generated multiple

NOTE Confidence: 0.929160932631579

 $00:14:50.664 \longrightarrow 00:14:53.159$  Caspase 3 knockout clones and then

NOTE Confidence: 0.929160932631579

 $00:14:53.159 \longrightarrow 00:14:55.145$  we did a dose response curve.

NOTE Confidence: 0.929160932631579

 $00:14:55.150 \rightarrow 00:14:57.420$  Examining the viability of wildtype

NOTE Confidence: 0.929160932631579

 $00{:}14{:}57{.}420 \dashrightarrow 00{:}15{:}00{.}230$  and Caspase 3 knockout clones in

NOTE Confidence: 0.929160932631579

 $00:15:00.230 \longrightarrow 00:15:02.560$  different concentrations of pack one.

NOTE Confidence: 0.929160932631579

 $00:15:02.560 \longrightarrow 00:15:04.170$  So this is what it looked like

NOTE Confidence: 0.929160932631579

 $00:15:04.170 \longrightarrow 00:15:05.260$  for two control clones,

NOTE Confidence: 0.929160932631579

00:15:05.260 --> 00:15:08.880 2 clones expressing Arosa 26 guide RNA pack,

 $00:15:08.880 \rightarrow 00:15:11.680$  one is a potent anti cancer agent.

NOTE Confidence: 0.929160932631579

00:15:11.680 --> 00:15:13.976 You can see it has an IC50 value

NOTE Confidence: 0.929160932631579

 $00{:}15{:}13{.}976$  -->  $00{:}15{:}16{.}128$  of around one or two micromolar.

NOTE Confidence: 0.929160932631579

 $00:15:16.130 \longrightarrow 00:15:16.467$  However,

NOTE Confidence: 0.929160932631579

 $00{:}15{:}16{.}467{\:}-{>}00{:}15{:}19{.}500$  when we did the same as say in the Caspase

NOTE Confidence: 0.929160932631579

 $00{:}15{:}19{.}577 \dashrightarrow 00{:}15{:}22{.}007$  3 knockout clones that we generated,

NOTE Confidence: 0.929160932631579

 $00:15:22.010 \longrightarrow 00:15:23.775$  we ended up getting exactly

NOTE Confidence: 0.929160932631579

 $00:15:23.775 \longrightarrow 00:15:25.187$  the same drug curve.

NOTE Confidence: 0.929160932631579

 $00{:}15{:}25{.}190 \dashrightarrow 00{:}15{:}28{.}390$  This drug is exactly as potent in caspase

NOTE Confidence: 0.929160932631579

00:15:28.390 --> 00:15:31.447 3 knockout clones as it is in caspase

NOTE Confidence: 0.929160932631579

 $00:15:31.447 \longrightarrow 00:15:34.130$  3 expressing Rosa 26 control clones.

NOTE Confidence: 0.929160932631579

 $00{:}15{:}34{.}130 \dashrightarrow 00{:}15{:}37{.}570$  It has an IC50 value of 1 to 2 micromolar,

NOTE Confidence: 0.929160932631579

 $00{:}15{:}37{.}570 \dashrightarrow 00{:}15{:}39{.}370$  regardless of whether these

NOTE Confidence: 0.929160932631579

 $00:15:39.370 \longrightarrow 00:15:41.170$  cells express caspase 3,

NOTE Confidence: 0.929160932631579

 $00:15:41.170 \longrightarrow 00:15:42.229$  so this drug,

 $00:15:42.229 \rightarrow 00:15:43.994$  which entered clinical trials as

NOTE Confidence: 0.929160932631579

 $00:15:43.994 \rightarrow 00:15:46.149$  a caspase 3 activating compound.

NOTE Confidence: 0.929160932631579

00:15:46.150 --> 00:15:48.850 Its anti cancer activity actually

NOTE Confidence: 0.929160932631579

 $00:15:48.850 \rightarrow 00:15:51.010$  comes from something entirely

NOTE Confidence: 0.929160932631579

 $00:15:51.010 \longrightarrow 00:15:52.974$  independent of caspase 3 and this

NOTE Confidence: 0.929160932631579

 $00{:}15{:}52{.}974 \dashrightarrow 00{:}15{:}54{.}426$  is actually the case for many

NOTE Confidence: 0.929160932631579

 $00:15:54.426 \longrightarrow 00:15:56.157$  of the drugs that we studied.

NOTE Confidence: 0.929160932631579

 $00:15:56.160 \rightarrow 00:15:58.040$  So to show you a few more examples,

NOTE Confidence: 0.929160932631579

00:15:58.040 --> 00:16:00.590 HDK 6 is a histone deacetylase

NOTE Confidence: 0.929160932631579

 $00:16:00.590 \rightarrow 00:16:02.670$  Celgene has developed each DAC.

NOTE Confidence: 0.929160932631579

 $00{:}16{:}02.670 \dashrightarrow 00{:}16{:}05.250$ 6 inhibitors sitter in a statin

NOTE Confidence: 0.929160932631579

00:16:05.250 --> 00:16:06.110 richelain ISTAT.

NOTE Confidence: 0.929160932631579

00:16:06.110 --> 00:16:09.150 We knocked out HDK 6 but we saw no change

NOTE Confidence: 0.820004865555556

 $00:16:09.228 \rightarrow 00:16:11.084$  in sensitivity to these

NOTE Confidence: 0.820004865555556

00:16:11.084 --> 00:16:12.940 putative HDK 6 inhibitors.

NOTE Confidence: 0.820004865555556

00:16:12.940 --> 00:16:15.090 Milk is a cancer related

- NOTE Confidence: 0.820004865555556
- 00:16:15.090 --> 00:16:16.380 kinase uncle therapy.
- NOTE Confidence: 0.820004865555556
- 00:16:16.380 --> 00:16:17.940 Science is developed this drug,
- NOTE Confidence: 0.820004865555556
- 00:16:17.940 --> 00:16:19.732 Novartis, developed this drug.
- NOTE Confidence: 0.820004865555556
- $00:16:19.732 \rightarrow 00:16:22.420$  We use CRISPR to knockout milk.
- NOTE Confidence: 0.820004865555556
- $00:16:22.420 \rightarrow 00:16:25.458$  We saw no change in sensitivity to
- NOTE Confidence: 0.820004865555556
- $00:16:25.458 \rightarrow 00:16:27.620$  these milk inhibitory compounds.
- NOTE Confidence: 0.820004865555556
- $00:16:27.620 \longrightarrow 00:16:29.330$  So to sum up a whole bunch of data
- NOTE Confidence: 0.820004865555556
- 00:16:29.330 --> 00:16:31.037 that I don't have time to show you,
- NOTE Confidence: 0.820004865555556
- $00{:}16{:}31.040 \dashrightarrow 00{:}16{:}33.056$  we found that target knock outs conferred
- NOTE Confidence: 0.820004865555556
- $00:16:33.056 \longrightarrow 00:16:34.813$  no resistance for 12 different
- NOTE Confidence: 0.820004865555556
- $00:16:34.813 \rightarrow 00:16:36.937$  cancer drugs that we were studying.
- NOTE Confidence: 0.820004865555556
- 00:16:36.940 --> 00:16:38.692 We made these knockouts and did
- NOTE Confidence: 0.820004865555556
- 00:16:38.692 --> 00:16:40.725 these tests in at least three
- NOTE Confidence: 0.820004865555556
- 00:16:40.725 --> 00:16:42.317 different cancer types each,
- NOTE Confidence: 0.820004865555556
- $00{:}16{:}42{.}320 \dashrightarrow 00{:}16{:}46{.}020$  so this kind of leaves us in an odd position.
- NOTE Confidence: 0.820004865555556

 $00:16:46.020 \rightarrow 00:16:48.300$  We were studying 12 different preclinical

NOTE Confidence: 0.820004865555556

 $00{:}16{:}48{.}300 \dashrightarrow 00{:}16{:}50{.}419$  or clinical anti cancer drugs and

NOTE Confidence: 0.820004865555556

 $00{:}16{:}50{.}419 \dashrightarrow 00{:}16{:}52{.}571$  in each of these cases we found that

NOTE Confidence: 0.820004865555556

 $00:16:52.630 \rightarrow 00:16:54.560$  the reported mechanism of action.

NOTE Confidence: 0.820004865555556

 $00{:}16{:}54{.}560 \dashrightarrow 00{:}16{:}55{.}745$  Was actually incorrect.

NOTE Confidence: 0.820004865555556

00:16:55.745 --> 00:16:58.510 This then raised the question well if

NOTE Confidence: 0.820004865555556

 $00{:}16{:}58{.}577 \dashrightarrow 00{:}17{:}01{.}111$  these drugs are killing cancer cells at

NOTE Confidence: 0.820004865555556

 $00:17:01.111 \rightarrow 00:17:03.499$  nanomolar or low micromolar potency,

NOTE Confidence: 0.820004865555556

 $00:17:03.500 \rightarrow 00:17:05.000$  how is it they actually work?

NOTE Confidence: 0.820004865555556

 $00:17:05.000 \rightarrow 00:17:06.536$  What is it they're actually targeting?

NOTE Confidence: 0.820004865555556

 $00{:}17{:}06{.}540 \dashrightarrow 00{:}17{:}08{.}404$  We wanted to see if we could figure

NOTE Confidence: 0.820004865555556

 $00{:}17{:}08{.}404 \dashrightarrow 00{:}17{:}10{.}608$  out how they were actually functioning.

NOTE Confidence: 0.820004865555556

 $00{:}17{:}10.610 \dashrightarrow 00{:}17{:}12.884$  We've had the best success so

NOTE Confidence: 0.820004865555556

 $00:17:12.884 \rightarrow 00:17:15.598$  far with one drug called O TS964.

NOTE Confidence: 0.820004865555556

 $00:17:15.598 \rightarrow 00:17:17.796$  This is what the drug looks like.

NOTE Confidence: 0.820004865555556

 $00{:}17{:}17{.}800 \dashrightarrow 00{:}17{:}20{.}448$  It was described in a paper in science

- NOTE Confidence: 0.820004865555556
- $00{:}17{:}20{.}448 \dashrightarrow 00{:}17{:}21{.}864$  Translational Medicine a few years
- NOTE Confidence: 0.820004865555556
- 00:17:21.864 --> 00:17:24.149 ago as an inhibitor of a kinase called PBK,
- NOTE Confidence: 0.820004865555556
- $00:17:24.150 \rightarrow 00:17:26.787$  which is also called Pop K in the literature,
- NOTE Confidence: 0.820004865555556
- 00:17:26.790 --> 00:17:27.765 but using CRISPR.
- NOTE Confidence: 0.820004865555556
- $00{:}17{:}27.765 \dashrightarrow 00{:}17{:}29.715$  We knocked out PVK and we
- NOTE Confidence: 0.820004865555556
- $00:17:29.715 \longrightarrow 00:17:31.570$  saw no effect whatsoever.
- NOTE Confidence: 0.820004865555556
- 00:17:31.570 --> 00:17:33.710 On sensitivity to this compound
- NOTE Confidence: 0.820004865555556
- $00:17:33.710 \longrightarrow 00:17:36.784$  telling us that this drug O TS964
- NOTE Confidence: 0.820004865555556
- $00{:}17{:}36{.}784 \dashrightarrow 00{:}17{:}39{.}586$  must have some other cellular target.
- NOTE Confidence: 0.820004865555556
- $00:17:39.590 \longrightarrow 00:17:41.326$  To see if we could figure out
- NOTE Confidence: 0.820004865555556
- $00:17:41.326 \rightarrow 00:17:43.199$  what this drug was actually doing,
- NOTE Confidence: 0.820004865555556
- $00{:}17{:}43.200 \dashrightarrow 00{:}17{:}46.170$  we used a genetic based approach
- NOTE Confidence: 0.820004865555556
- $00{:}17{:}46{.}170 \dashrightarrow 00{:}17{:}47{.}430$  for this approach.
- NOTE Confidence: 0.820004865555556
- $00:17:47.430 \longrightarrow 00:17:49.110$  We took highly mutagenized
- NOTE Confidence: 0.820004865555556
- $00{:}17{:}49{.}110 \dashrightarrow 00{:}17{:}50{.}994$  colon cancer cells, HCT 116.
- NOTE Confidence: 0.820004865555556

 $00:17:50.994 \rightarrow 00:17:53.178$  They have a very high mutation rate

NOTE Confidence: 0.820004865555556

 $00:17:53.178 \rightarrow 00:17:54.727$  because they're microsatellite unstable

NOTE Confidence: 0.820004865555556

 $00{:}17{:}54{.}727 \dashrightarrow 00{:}17{:}57{.}457$  and then we expose these drugs to

NOTE Confidence: 0.820004865555556

 $00:17:57.526 \longrightarrow 00:17:59.446$  a nearly lethal concentration of

NOTE Confidence: 0.820004865555556

00:17:59.450 --> 00:18:03.046 O TS96 four such that about 99.9%

NOTE Confidence: 0.820004865555556

 $00:18:03.046 \longrightarrow 00:18:05.538$  of cells on the plate were killed.

NOTE Confidence: 0.820004865555556

 $00:18:05.540 \longrightarrow 00:18:05.858$  However,

NOTE Confidence: 0.820004865555556

 $00:18:05.858 \rightarrow 00:18:07.766$  there were a few stragglers that

NOTE Confidence: 0.820004865555556

 $00{:}18{:}07.766 \dashrightarrow 00{:}18{:}09.487$  remained when we cut these cells

NOTE Confidence: 0.820004865555556

 $00:18:09.487 \rightarrow 00:18:11.469$  in the drug for a period of weeks

NOTE Confidence: 0.820004865555556

 $00:18:11.469 \rightarrow 00:18:13.387$  until these cells were able to grow

NOTE Confidence: 0.820004865555556

 $00:18:13.387 \longrightarrow 00:18:15.077$  and form little micro colonies.

NOTE Confidence: 0.820004865555556

 $00:18:15.077 \rightarrow 00:18:17.351$  We then subjected these cells to

NOTE Confidence: 0.820004865555556

 $00:18:17.351 \rightarrow 00:18:19.654$  whole exome sequencing and when we

NOTE Confidence: 0.820004865555556

 $00:18:19.654 \rightarrow 00:18:22.020$  did sequencing on the resistant clones,

NOTE Confidence: 0.820004865555556

 $00:18:22.020 \longrightarrow 00:18:24.444$  what we were hoping to see was a

- NOTE Confidence: 0.820004865555556
- $00{:}18{:}24{.}444 \dashrightarrow 00{:}18{:}26{.}709$  mutation that blocked whatever it was.
- NOTE Confidence: 0.820004865555556
- 00:18:26.710 --> 00:18:28.600 This drug was actually targeting.
- NOTE Confidence: 0.820004865555556
- $00{:}18{:}28{.}600 \dashrightarrow 00{:}18{:}31{.}090$  Maybe these cells could survive a
- NOTE Confidence: 0.820004865555556
- $00:18:31.090 \rightarrow 00:18:33.214$  lethal treatment because they had
- NOTE Confidence: 0.820004865555556
- $00:18:33.214 \rightarrow 00:18:35.189$  some mutation preventing drug binding
- NOTE Confidence: 0.820004865555556
- 00:18:35.189 --> 00:18:38.288 to whatever O TS96 or was actually doing.
- NOTE Confidence: 0.820004865555556
- $00:18:38.290 \rightarrow 00:18:40.208$  So when we did whole exome sequencing
- NOTE Confidence: 0.820004865555556
- $00:18:40.208 \longrightarrow 00:18:41.030$  on these clones,
- NOTE Confidence: 0.820004865555556
- $00:18:41.030 \longrightarrow 00:18:43.333$  we were really excited to see that
- NOTE Confidence: 0.820004865555556
- $00:18:43.333 \longrightarrow 00:18:45.566$  every clone that we looked at had
- NOTE Confidence: 0.820004865555556
- $00:18:45.566 \longrightarrow 00:18:47.036$  the same mutation in it.
- NOTE Confidence: 0.820004865555556
- 00:18:47.040 --> 00:18:48.726 Every drug resistant clone had a
- NOTE Confidence: 0.820004865555556
- $00{:}18{:}48.726 \dashrightarrow 00{:}18{:}50.779$  mutation in the cyclin dependent kinase,
- NOTE Confidence: 0.820004865555556
- 00:18:50.780 --> 00:18:51.610 CDK 11.
- NOTE Confidence: 0.820004865555556
- $00{:}18{:}51{.}610$  -->  $00{:}18{:}54{.}515$  They had a glycine to serine substitution,
- NOTE Confidence: 0.820004865555556

 $00:18:54.520 \longrightarrow 00:18:56.886$  right smack dab in the middle of

NOTE Confidence: 0.820004865555556

 $00{:}18{:}56{.}886 \dashrightarrow 00{:}18{:}59{.}079$  the CDK 11 kinase domain.

NOTE Confidence: 0.820004865555556

 $00:18:59.080 \rightarrow 00:19:00.740$  So this immediately suggested to

NOTE Confidence: 0.820004865555556

 $00:19:00.740 \longrightarrow 00:19:02.400$  us that maybe this drug,

NOTE Confidence: 0.820004865555556

 $00:19:02.400 \longrightarrow 00:19:05.456$  which had been developed as a PDK inhibitor,

NOTE Confidence: 0.820004865555556

 $00:19:05.460 \rightarrow 00:19:07.208$  was actually functioning through

NOTE Confidence: 0.820004865555556

 $00{:}19{:}07{.}208 \dashrightarrow 00{:}19{:}08{.}956$  inhibition of CDK 11.

NOTE Confidence: 0.89901102625

00:19:08.960 --> 00:19:10.620 Instead, one potential limitation

NOTE Confidence: 0.89901102625

 $00:19:10.620 \longrightarrow 00:19:12.552$  to this is that, well,

NOTE Confidence: 0.89901102625

 $00:19:12.552 \rightarrow 00:19:14.456$  there actually isn't a precedent for this.

NOTE Confidence: 0.89901102625

 $00{:}19{:}14.460 \dashrightarrow 00{:}19{:}17.718$  CDK 11 hasn't been previously dropped,

NOTE Confidence: 0.89901102625

 $00{:}19{:}17{.}720 \dashrightarrow 00{:}19{:}20{.}312$  so we wanted to see if this mutation

NOTE Confidence: 0.89901102625

 $00{:}19{:}20{.}312 \dashrightarrow 00{:}19{:}22{.}463$  actually had anything to do with

NOTE Confidence: 0.89901102625

00:19:22.463 --> 00:19:25.940 sensitivity to OTS 964 in order to do that,

NOTE Confidence: 0.89901102625

 $00{:}19{:}25{.}940 \dashrightarrow 00{:}19{:}27{.}704$  we wanted to see whether this mutation

NOTE Confidence: 0.89901102625

 $00{:}19{:}27.704 \dashrightarrow 00{:}19{:}29.310$  that we discovered in the resistance.

- NOTE Confidence: 0.89901102625
- $00{:}19{:}29{.}310 \dashrightarrow 00{:}19{:}31{.}715$  Jones was actually sufficient to
- NOTE Confidence: 0.89901102625
- 00:19:31.715 -> 00:19:33.988 confer resistance to OTS 964.
- NOTE Confidence: 0.89901102625
- $00:19:33.988 \longrightarrow 00:19:35.032$  To test this,
- NOTE Confidence: 0.89901102625
- 00:19:35.032 --> 00:19:37.120 we used a CRISPR knockin strategy
- NOTE Confidence: 0.89901102625
- $00:19:37.189 \rightarrow 00:19:39.294$  where we introduced this glycine
- NOTE Confidence: 0.89901102625
- $00{:}19{:}39{.}294 \dashrightarrow 00{:}19{:}41{.}399$  to serine substitution that we
- NOTE Confidence: 0.89901102625
- $00:19:41.474 \rightarrow 00:19:43.749$  recovered in drug resistant cells.
- NOTE Confidence: 0.89901102625
- $00:19:43.750 \longrightarrow 00:19:45.904$  We knocked it into drug naive
- NOTE Confidence: 0.89901102625
- $00{:}19{:}45{.}904 \dashrightarrow 00{:}19{:}48{.}093$  cancer cells and then tested its
- NOTE Confidence: 0.89901102625
- 00:19:48.093 --> 00:19:50.241 effects on on O TS964 sensitivity.
- NOTE Confidence: 0.89901102625
- $00:19:50.241 \rightarrow 00:19:52.503$  This is what it looked like.
- NOTE Confidence: 0.89901102625
- 00:19:52.510 --> 00:19:54.175 Here we have four different
- NOTE Confidence: 0.89901102625
- $00{:}19{:}54{.}175 \dashrightarrow 00{:}19{:}56{.}261$  cancer cell lines treated with a
- NOTE Confidence: 0.89901102625
- 00:19:56.261 --> 00:19:58.104 lethal concentration of O TS964,
- NOTE Confidence: 0.89901102625
- $00{:}19{:}58.104 \dashrightarrow 00{:}20{:}00{.}396$  with a negative control guide RNA.
- NOTE Confidence: 0.89901102625

 $00:20:00.400 \longrightarrow 00:20:02.870$  Or if we just cut in the CDK 11 gene,

NOTE Confidence: 0.89901102625

 $00{:}20{:}02{.}870 \dashrightarrow 00{:}20{:}05{.}258$  we have no cancer cell viability.

NOTE Confidence: 0.89901102625

 $00{:}20{:}05{.}260 \dashrightarrow 00{:}20{:}07{.}956$  But if we introduce a repair template that

NOTE Confidence: 0.89901102625

 $00:20:07.956 \rightarrow 00:20:10.597$  includes the glycine to serine substitution,

NOTE Confidence: 0.89901102625

 $00:20:10.600 \longrightarrow 00:20:12.225$  then we can restore viability

NOTE Confidence: 0.89901102625

 $00{:}20{:}12.225 \dashrightarrow 00{:}20{:}14.442$  in the presence of an otherwise

NOTE Confidence: 0.89901102625

 $00:20:14.442 \longrightarrow 00:20:16.634$  lethal concentration of O TS964.

NOTE Confidence: 0.89901102625

 $00{:}20{:}16.634 \dashrightarrow 00{:}20{:}19.090$  So this tells us that this mutation is

NOTE Confidence: 0.89901102625

 $00{:}20{:}19.156 \dashrightarrow 00{:}20{:}21.508$  in fact both necessary and sufficient

NOTE Confidence: 0.89901102625

 $00:20:21.508 \rightarrow 00:20:23.730$  for resistance to this compound.

NOTE Confidence: 0.89901102625

 $00{:}20{:}23.730 \dashrightarrow 00{:}20{:}25.878$  We then followed this up with

NOTE Confidence: 0.89901102625

 $00:20:25.878 \longrightarrow 00:20:26.952$  some biochemical assays.

NOTE Confidence: 0.89901102625

00:20:26.960 --> 00:20:30.888 We confirmed that O TS964 inhibits CDK 11.

NOTE Confidence: 0.89901102625

 $00{:}20{:}30{.}890 \dashrightarrow 00{:}20{:}33{.}557$  With an IC50 value of around 40

NOTE Confidence: 0.89901102625

00:20:33.557 --> 00:20:35.828 to 50 animal or in vitro,

NOTE Confidence: 0.89901102625

 $00:20:35.830 \rightarrow 00:20:38.798$  and we did a cell based target engagement

- NOTE Confidence: 0.89901102625
- 00:20:38.798 --> 00:20:40.870 assay using mass spectrometry,

 $00{:}20{:}40.870 \dashrightarrow 00{:}20{:}43.132$  we found that 100 animal or

NOTE Confidence: 0.89901102625

 $00:20:43.132 \longrightarrow 00:20:44.724$  treatment with O TS964.

NOTE Confidence: 0.89901102625

00:20:44.724 --> 00:20:47.643 It didn't bind to hundreds of other

NOTE Confidence: 0.89901102625

 $00:20:47.643 \longrightarrow 00:20:50.099$  cellular kinases, but it bound.

NOTE Confidence: 0.89901102625

 $00:20:50.099 \rightarrow 00:20:53.250$  It caused about 70% of binding site

NOTE Confidence: 0.89901102625

00:20:53.250 --> 00:20:56.362 occlusion for CDK 11, and only CDK 11.

NOTE Confidence: 0.89901102625

 $00{:}20{:}56{.}362 \dashrightarrow 00{:}20{:}58{.}799$  So from this work we concluded that

NOTE Confidence: 0.89901102625

 $00:20:58.799 \rightarrow 00:21:00.503$  by profiling a mischaracterized

NOTE Confidence: 0.89901102625

 $00:21:00.503 \rightarrow 00:21:03.217$  anti cancer agent we were actually

NOTE Confidence: 0.89901102625

 $00:21:03.217 \rightarrow 00:21:05.762$  able to serendipitously discover the

NOTE Confidence: 0.89901102625

00:21:05.762 --> 00:21:10.030 first selective inhibitor of CDK 11.

NOTE Confidence: 0.89901102625

 $00{:}21{:}10.030 \dashrightarrow 00{:}21{:}12.590$  So to sum up what I told you so far,

NOTE Confidence: 0.89901102625

00:21:12.590 --> 00:21:14.564 we're kind of operating in a space

NOTE Confidence: 0.89901102625

 $00{:}21{:}14.564 \dashrightarrow 00{:}21{:}16.790$  in which the vast majority of new

 $00:21:16.790 \rightarrow 00:21:18.770$  therapies that get tested in human

NOTE Confidence: 0.89901102625

00:21:18.830 --> 00:21:21.350 patients in oncology don't end up working,

NOTE Confidence: 0.89901102625

 $00{:}21{:}21{.}350 \dashrightarrow 00{:}21{:}22.856$  and we put together a collection

NOTE Confidence: 0.89901102625

 $00:21:22.856 \longrightarrow 00:21:24.220$  of these drugs to study.

NOTE Confidence: 0.89901102625

 $00{:}21{:}24{.}220 \dashrightarrow 00{:}21{:}26{.}299$  And one thing that we found while

NOTE Confidence: 0.89901102625

 $00{:}21{:}26{.}299 \dashrightarrow 00{:}21{:}28{.}564$  studying them is that many of these

NOTE Confidence: 0.89901102625

 $00{:}21{:}28.564 \dashrightarrow 00{:}21{:}30.199$  drugs have actually been designed

NOTE Confidence: 0.89901102625

 $00:21:30.199 \longrightarrow 00:21:32.434$  to target proteins that have no

NOTE Confidence: 0.89901102625

 $00{:}21{:}32{.}434 \dashrightarrow 00{:}21{:}34{.}279$  detectable role in cancer growth.

NOTE Confidence: 0.89901102625

 $00:21:34.280 \longrightarrow 00:21:34.685$  Furthermore,

NOTE Confidence: 0.89901102625

 $00{:}21{:}34.685 \dashrightarrow 00{:}21{:}37.520$  while these drugs do kill cancer cells,

NOTE Confidence: 0.89901102625

00:21:37.520 --> 00:21:39.265 they largely kill cancer cells

NOTE Confidence: 0.89901102625

 $00{:}21{:}39{.}265 \dashrightarrow 00{:}21{:}41{.}010$  through off target effects rather

NOTE Confidence: 0.89901102625

00:21:41.073 - > 00:21:43.233 than through the target that they

NOTE Confidence: 0.89901102625

 $00:21:43.233 \rightarrow 00:21:44.673$  were initially designed against,

NOTE Confidence: 0.89901102625

 $00{:}21{:}44.680 \dashrightarrow 00{:}21{:}46.423$  and I think that this can increase

- NOTE Confidence: 0.89901102625
- $00{:}21{:}46{.}423 \dashrightarrow 00{:}21{:}48{.}372$  the burden of side effects and the
- NOTE Confidence: 0.89901102625
- $00:21:48.372 \rightarrow 00:21:49.802$  decrease the efficacy when some
- NOTE Confidence: 0.89901102625
- $00:21:49.802 \longrightarrow 00:21:51.700$  of these drugs are actually used.
- NOTE Confidence: 0.89901102625
- $00:21:51.700 \longrightarrow 00:21:53.575$  We don't truly understand how
- NOTE Confidence: 0.89901102625
- $00:21:53.575 \rightarrow 00:21:55.917$  they're working or where their anti
- NOTE Confidence: 0.89901102625
- $00:21:55.917 \dashrightarrow 00:21:57.517$  cancer activity comes from.
- NOTE Confidence: 0.89901102625
- $00{:}21{:}57{.}520 \dashrightarrow 00{:}22{:}00{.}061$  Think this conclusion has a number of
- NOTE Confidence: 0.89901102625
- $00:22:00.061 \rightarrow 00:22:02.039$  important considerations and caveats though.
- NOTE Confidence: 0.89901102625
- $00:22:02.040 \longrightarrow 00:22:04.480$  For instance, there could be
- NOTE Confidence: 0.89901102625
- $00:22:04.480 \rightarrow 00:22:06.432$  unrecognized cell type specificity.
- NOTE Confidence: 0.89901102625
- $00:22:06.440 \longrightarrow 00:22:07.912$  We did these competitions
- NOTE Confidence: 0.89901102625
- $00{:}22{:}07{.}912 \dashrightarrow 00{:}22{:}09{.}752$  in 32 cancer cell lines.
- NOTE Confidence: 0.89901102625
- $00{:}22{:}09.760 \dashrightarrow 00{:}22{:}11.124$  We generated knockout clones
- NOTE Confidence: 0.89901102625
- $00{:}22{:}11.124 \dashrightarrow 00{:}22{:}12.829$  in three cancer types each,
- NOTE Confidence: 0.89901102625
- 00:22:12.830 --> 00:22:13.328 but it was,
- NOTE Confidence: 0.89901102625

00:22:13.328 --> 00:22:13.660 you know,

NOTE Confidence: 0.829353936153846

 $00:22:13.660 \rightarrow 00:22:16.408$  physically, impossible for us to test

NOTE Confidence: 0.829353936153846

 $00:22:16.408 \rightarrow 00:22:19.558$  every subtype of leukemia or every subtype.

NOTE Confidence: 0.829353936153846

00:22:19.560 --> 00:22:20.788 Kidney cancer in existence,

NOTE Confidence: 0.829353936153846

 $00:22:20.788 \dashrightarrow 00:22:22.630$  and so we can't fully recognize

NOTE Confidence: 0.829353936153846

 $00:22:22.686 \rightarrow 00:22:24.376$  rule out some unrecognized cell

NOTE Confidence: 0.829353936153846

 $00:22:24.376 \rightarrow 00:22:26.066$  type specificity that hasn't been

NOTE Confidence: 0.829353936153846

 $00:22:26.124 \longrightarrow 00:22:27.996$  reported in the literature on these.

NOTE Confidence: 0.829353936153846

00:22:28.000 --> 00:22:30.267 Targets. Secondly,

NOTE Confidence: 0.829353936153846

 $00:22:30.267 \rightarrow 00:22:32.652$  we specifically tested the hypothesis

NOTE Confidence: 0.829353936153846

00:22:32.652 -> 00:22:35.118 that these proteins are required

NOTE Confidence: 0.829353936153846

 $00{:}22{:}35{.}118 \dashrightarrow 00{:}22{:}37{.}528$  for cell autonomous cancer growth,

NOTE Confidence: 0.829353936153846

 $00:22:37.530 \rightarrow 00:22:39.007$  that is, cells going from you know,

NOTE Confidence: 0.829353936153846

 $00:22:39.010 \longrightarrow 00:22:40.846$  one cancer cell to 2:00 to 4:00 to 8:00,

NOTE Confidence: 0.829353936153846

 $00{:}22{:}40.850 \dashrightarrow 00{:}22{:}43.235$  and so on, and this had been reported for

NOTE Confidence: 0.829353936153846

 $00:22:43.235 \longrightarrow 00:22:45.468$  each of the drugs that we had studied.

00:22:45.470 --> 00:22:47.126 However, if it turned out that,

NOTE Confidence: 0.829353936153846

 $00:22:47.130 \longrightarrow 00:22:49.979$  say, pack four had some role in

NOTE Confidence: 0.829353936153846

 $00{:}22{:}49{.}979 \dashrightarrow 00{:}22{:}52{.}509$  angiogenesis or in immune evasion,

NOTE Confidence: 0.829353936153846

 $00:22:52.510 \rightarrow 00:22:54.890$  or some other non cell autonomous process,

NOTE Confidence: 0.829353936153846

 $00:22:54.890 \longrightarrow 00:22:57.260$  that wouldn't be ruled out for

NOTE Confidence: 0.829353936153846

 $00:22:57.260 \longrightarrow 00:22:58.840$  the cell autonomous proliferation

NOTE Confidence: 0.829353936153846

 $00{:}22{:}58{.}905 \dashrightarrow 00{:}23{:}00{.}820$  focused as says that we've done.

NOTE Confidence: 0.829353936153846

 $00:23:00.820 \longrightarrow 00:23:02.686$  I think a third important consideration

NOTE Confidence: 0.829353936153846

 $00{:}23{:}02.686 \dashrightarrow 00{:}23{:}04.679$  is while our data suggests that

NOTE Confidence: 0.829353936153846

 $00:23:04.679 \rightarrow 00:23:06.399$  these drugs are promiscuous and

NOTE Confidence: 0.829353936153846

00:23:06.399 - 00:23:08.292 may have multiple targets in the

NOTE Confidence: 0.829353936153846

 $00{:}23{:}08{.}292 \dashrightarrow 00{:}23{:}10{.}161$  cell just because a cancer drug is

NOTE Confidence: 0.829353936153846

 $00{:}23{:}10.170 \dashrightarrow 00{:}23{:}11.466$  promiscuous doesn't necessarily mean

NOTE Confidence: 0.829353936153846

 $00{:}23{:}11{.}466 \dashrightarrow 00{:}23{:}13{.}840$  that it will fail in the clinic.

NOTE Confidence: 0.829353936153846

 $00{:}23{:}13.840 \dashrightarrow 00{:}23{:}15.944$  There are a number of drugs like sunitinib,

00:23:15.950 --> 00:23:18.920 Serafin, IB which do have multiple

NOTE Confidence: 0.829353936153846

 $00{:}23{:}18{.}920 \dashrightarrow 00{:}23{:}20{.}900$  targets in the cell.

NOTE Confidence: 0.829353936153846

00:23:20.900 --> 00:23:21.610 And so,

NOTE Confidence: 0.829353936153846

 $00:23:21.610 \rightarrow 00:23:23.385$  just because something is promiscuous

NOTE Confidence: 0.829353936153846

 $00:23:23.385 \rightarrow 00:23:25.418$  doesn't necessarily mean that it will fail.

NOTE Confidence: 0.829353936153846

00:23:25.420 --> 00:23:25.763 However,

NOTE Confidence: 0.829353936153846

 $00:23:25.763 \longrightarrow 00:23:28.507$  I think that if our goal in cancer

NOTE Confidence: 0.829353936153846

 $00:23:28.507 \rightarrow 00:23:31.324$  biology is to kind of reach a plateau

NOTE Confidence: 0.829353936153846

 $00:23:31.324 \rightarrow 00:23:33.206$  of targeted precision medicine where

NOTE Confidence: 0.829353936153846

 $00:23:33.206 \rightarrow 00:23:35.272$  you sequence a patient's tumor,

NOTE Confidence: 0.829353936153846

 $00{:}23{:}35{.}272 \dashrightarrow 00{:}23{:}37{.}502$  you identify the mutations and

NOTE Confidence: 0.829353936153846

 $00:23:37.502 \rightarrow 00:23:39.220$  amplifications and alterations and

NOTE Confidence: 0.829353936153846

 $00{:}23{:}39{.}220 \dashrightarrow 00{:}23{:}41{.}140$  then design a drug cocktail based

NOTE Confidence: 0.829353936153846

 $00:23:41.140 \longrightarrow 00:23:43.020$  on that particular genetic profile

NOTE Confidence: 0.829353936153846

 $00:23:43.020 \rightarrow 00:23:44.658$  in order to get to that level.

NOTE Confidence: 0.829353936153846

 $00:23:44.660 \rightarrow 00:23:47.486$  I think we need to have a really good

- NOTE Confidence: 0.829353936153846
- $00{:}23{:}47{.}486 \dashrightarrow 00{:}23{:}49{.}746$  understanding of what drugs do and how
- NOTE Confidence: 0.829353936153846
- $00:23:49.746 \rightarrow 00:23:52.049$  their anti cancer activity actually arises.
- NOTE Confidence: 0.829353936153846
- $00:23:52.050 \longrightarrow 00:23:54.390$  And what we'd suggest is that
- NOTE Confidence: 0.829353936153846
- $00:23:54.390 \rightarrow 00:23:55.950$  pre clinical genetic validation,
- NOTE Confidence: 0.829353936153846
- 00:23:55.950 --> 00:23:58.488 particularly using CRISPR instead of RNA.
- NOTE Confidence: 0.829353936153846
- $00{:}23{:}58{.}490 \dashrightarrow 00{:}24{:}00{.}682$  I may help us get genetic insight into
- NOTE Confidence: 0.829353936153846
- 00:24:00.682 --> 00:24:03.156 how anti cancer drugs work and may
- NOTE Confidence: 0.829353936153846
- $00{:}24{:}03.156 \dashrightarrow 00{:}24{:}05.021$  decrease the number of investigational
- NOTE Confidence: 0.829353936153846
- $00:24:05.084 \rightarrow 00:24:07.009$  drugs that enter clinical trials,
- NOTE Confidence: 0.829353936153846
- $00:24:07.010 \rightarrow 00:24:10.328$  but end up failing during clinical testing.
- NOTE Confidence: 0.829353936153846
- $00:24:10.330 \rightarrow 00:24:12.290$  So this is work that was done by my group.
- NOTE Confidence: 0.829353936153846
- 00:24:12.290 --> 00:24:13.008 In particular,
- NOTE Confidence: 0.829353936153846
- $00{:}24{:}13.008 \dashrightarrow 00{:}24{:}14.444$  two really talented students
- NOTE Confidence: 0.829353936153846
- $00{:}24{:}14{.}444 \dashrightarrow 00{:}24{:}16{.}470$  and Lynn and Chris Giuliano.
- NOTE Confidence: 0.829353936153846
- 00:24:16.470 --> 00:24:17.595 I'd like to acknowledge the
- NOTE Confidence: 0.829353936153846

00:24:17.595 --> 00:24:18.990 funding and thank you so much,

NOTE Confidence: 0.829353936153846

00:24:18.990 --> 00:24:20.268 I'd be happy to answer any

NOTE Confidence: 0.829353936153846

 $00:24:20.268 \longrightarrow 00:24:21.120$  questions that you have.

NOTE Confidence: 0.951348497777778

 $00:24:24.680 \rightarrow 00:24:25.892$  Thanks very much.

NOTE Confidence: 0.951348497777778

 $00{:}24{:}25{.}892 \dashrightarrow 00{:}24{:}28{.}316$  I thought that was really great.

NOTE Confidence: 0.951348497777778

00:24:28.320 --> 00:24:32.100 I think you know one of the one of the

NOTE Confidence: 0.951348497777778

 $00{:}24{:}32{.}100 \dashrightarrow 00{:}24{:}34{.}871$  things we're all aware of is that when

NOTE Confidence: 0.951348497777778

00:24:34.871 - 00:24:37.163 we combine drugs that the toxicity

NOTE Confidence: 0.951348497777778

 $00{:}24{:}37{.}163 \dashrightarrow 00{:}24{:}40{.}118$  goes way up and you know of course,

NOTE Confidence: 0.951348497777778

 $00:24:40.120 \longrightarrow 00:24:42.143$  much of the reason for that is

NOTE Confidence: 0.951348497777778

 $00:24:42.143 \longrightarrow 00:24:44.208$  that many of these drugs are

NOTE Confidence: 0.951348497777778

 $00{:}24{:}44{.}208 \dashrightarrow 00{:}24{:}46{.}088$  promiscuous and are doing much

NOTE Confidence: 0.951348497777778

 $00{:}24{:}46.088 \dashrightarrow 00{:}24{:}48.529$  more than what we need them to do.

NOTE Confidence: 0.951348497777778

 $00:24:48.530 \rightarrow 00:24:52.976$  There's a there was a question a minute ago.

NOTE Confidence: 0.951348497777778

 $00:24:52.980 \rightarrow 00:24:58.460$  Uh oh, so the from from Jeffrey Townsend.

NOTE Confidence: 0.951348497777778

00:24:58.460 -> 00:25:00.868 How were the original 12 drugs selected

- NOTE Confidence: 0.951348497777778
- $00:25:00.868 \rightarrow 00:25:02.810$  and assembled for investigation?
- NOTE Confidence: 0.885461179230769
- 00:25:03.060 --> 00:25:06.436 Yep, so I didn't have time to discuss
- NOTE Confidence: 0.885461179230769
- $00:25:06.436 \longrightarrow 00:25:08.380$  that extensively in this talk,
- NOTE Confidence: 0.885461179230769
- $00:25:08.380 \longrightarrow 00:25:10.678$  but what we were interested in
- NOTE Confidence: 0.885461179230769
- $00:25:10.680 \rightarrow 00:25:12.510$  our underlying hypothesis is that
- NOTE Confidence: 0.885461179230769
- $00{:}25{:}12.510 \dashrightarrow 00{:}25{:}14.747$  the gold standard for knowing a
- NOTE Confidence: 0.885461179230769
- $00{:}25{:}14.747 \dashrightarrow 00{:}25{:}16.805$  cancer drugs mechanism of action is
- NOTE Confidence: 0.885461179230769
- $00{:}25{:}16.805 \dashrightarrow 00{:}25{:}18.575$  the identification of a mutation
- NOTE Confidence: 0.885461179230769
- $00{:}25{:}18.575 \dashrightarrow 00{:}25{:}20.275$  that confers resistance to it.
- NOTE Confidence: 0.885461179230769
- $00{:}25{:}20{.}280 \dashrightarrow 00{:}25{:}22{.}290$  The classic example here is Gleevec
- NOTE Confidence: 0.885461179230769
- $00:25:22.290 \longrightarrow 00:25:24.169$  and the mutations in BCR ABL.
- NOTE Confidence: 0.885461179230769
- 00:25:24.170 --> 00:25:26.234 Set block, Liebeck activity and our
- NOTE Confidence: 0.885461179230769
- $00{:}25{:}26{.}234 \dashrightarrow 00{:}25{:}28{.}388$  thinking was that drugs that lacked
- NOTE Confidence: 0.885461179230769
- $00{:}25{:}28{.}388 \dashrightarrow 00{:}25{:}30{.}168$  that level of genetic validation
- NOTE Confidence: 0.885461179230769
- $00{:}25{:}30{.}168 \dashrightarrow 00{:}25{:}32{.}262$  were less likely to be acting
- NOTE Confidence: 0.885461179230769

 $00:25:32.262 \rightarrow 00:25:33.867$  through an on target mechanism.

NOTE Confidence: 0.885461179230769

 $00{:}25{:}33{.}870 \dashrightarrow 00{:}25{:}35{.}400$  So we selected drugs that

NOTE Confidence: 0.885461179230769

 $00:25:35.400 \longrightarrow 00:25:37.336$  specifically did not have that level

NOTE Confidence: 0.885461179230769

 $00:25:37.336 \rightarrow 00:25:39.066$  of genetic evidence behind them.

NOTE Confidence: 0.600856868

 $00:25:40.760 \longrightarrow 00:25:45.120$  And from from Mike Hurwitz.

NOTE Confidence: 0.600856868

 $00:25:45.120 \longrightarrow 00:25:46.260$  Sort of along that line.

NOTE Confidence: 0.600856868

00:25:46.260 --> 00:25:47.814 Do you find it striking that every

NOTE Confidence: 0.600856868

 $00:25:47.814 \rightarrow 00:25:49.590$  single one of your targets was wrong?

NOTE Confidence: 0.87856613

 $00{:}25{:}49{.}920 \dashrightarrow 00{:}25{:}53{.}140$  Yeah, so for the sake of time,

NOTE Confidence: 0.87856613

 $00:25:53.140 \rightarrow 00:25:54.538$  yeah, for the sake of time,

NOTE Confidence: 0.87856613

 $00{:}25{:}54{.}540 \dashrightarrow 00{:}25{:}57{.}172$  I focused on the ones that were

NOTE Confidence: 0.87856613

 $00:25:57.172 \longrightarrow 00:25:59.289$  where we discovered that the

NOTE Confidence: 0.87856613

 $00{:}25{:}59{.}289 \dashrightarrow 00{:}26{:}01{.}554$  mechanism of action was incorrect.

NOTE Confidence: 0.87856613

00:26:01.560 --> 00:26:03.968 However, we did have a few examples

NOTE Confidence: 0.87856613

 $00{:}26{:}03.968 \dashrightarrow 00{:}26{:}05.899$  where we could validate it,

NOTE Confidence: 0.87856613

 $00:26:05.900 \rightarrow 00:26:07.700$  and I'm just trying to here.

- NOTE Confidence: 0.87856613
- 00:26:07.700 --> 00:26:09.685 I'm going to show just
- NOTE Confidence: 0.87856613
- $00{:}26{:}09{.}685 \dashrightarrow 00{:}26{:}11{.}670$  one example of that now.
- NOTE Confidence: 0.87856613
- $00:26:11.670 \longrightarrow 00:26:14.731$  So this is not Lynn 3A.
- NOTE Confidence: 0.87856613
- $00{:}26{:}14.731 \dashrightarrow 00{:}26{:}17.619$  This is a drug that's been reported to
- NOTE Confidence: 0.87856613
- $00:26:17.619 \rightarrow 00:26:19.921$  function through P53 activation blocks.
- NOTE Confidence: 0.87856613
- 00:26:19.921 --> 00:26:22.206 The interaction between MDM two
- NOTE Confidence: 0.87856613
- $00:26:22.206 \longrightarrow 00:26:25.555$  and P53 we generated P53 knockout
- NOTE Confidence: 0.87856613
- 00:26:25.555 --> 00:26:28.445 clones using crisper and when we
- NOTE Confidence: 0.87856613
- $00{:}26{:}28{.}445 \dashrightarrow 00{:}26{:}30{.}370$  did this drug sensitivity curve
- NOTE Confidence: 0.87856613
- $00:26:30.370 \longrightarrow 00:26:32.794$  we found that a nutlin has no
- NOTE Confidence: 0.87856613
- 00:26:32.794 --> 00:26:34.781 effect on the P53 knockout clones,
- NOTE Confidence: 0.87856613
- $00{:}26{:}34{.}781 \dashrightarrow 00{:}26{:}37{.}192$  while it kills the P53 expressing
- NOTE Confidence: 0.87856613
- $00{:}26{:}37{.}192 \dashrightarrow 00{:}26{:}39{.}236$  Rosa 26 control phones.
- NOTE Confidence: 0.87856613
- $00{:}26{:}39{.}240 \dashrightarrow 00{:}26{:}40{.}878$  So in general so this is.
- NOTE Confidence: 0.87856613
- $00{:}26{:}40.880 \dashrightarrow 00{:}26{:}42.944$  What we would expect for a drug that
- NOTE Confidence: 0.87856613

00:26:42.944 --> 00:26:44.778 acts for an on target activity.

NOTE Confidence: 0.87856613

 $00{:}26{:}44.780 \dashrightarrow 00{:}26{:}46.796$  You know a huge delta between the

NOTE Confidence: 0.87856613

 $00:26:46.796 \rightarrow 00:26:48.299$  target knockouts and the target,

NOTE Confidence: 0.87856613

 $00:26:48.300 \rightarrow 00:26:49.710$  expressing control clones,

NOTE Confidence: 0.87856613

 $00{:}26{:}49{.}710 \dashrightarrow 00{:}26{:}53{.}500$  and we found a few examples of this.

NOTE Confidence: 0.87856613

00:26:53.500 --> 00:26:53.860 OK,

NOTE Confidence: 0.9160527775

 $00{:}26{:}54{.}300 \dashrightarrow 00{:}26{:}57{.}282$  and I think this is the last

NOTE Confidence: 0.9160527775

 $00{:}26{:}57{.}282 \dashrightarrow 00{:}27{:}00{.}020$  question from from Karen Anderson.

NOTE Confidence: 0.9160527775

 $00{:}27{:}00{.}020 \dashrightarrow 00{:}27{:}01{.}940$  Did you make the searing mutant of CDK

NOTE Confidence: 0.9160527775

00:27:01.940 --> 00:27:04.469 11 and show that the inhibitor was no

NOTE Confidence: 0.9160527775

 $00:27:04.469 \rightarrow 00:27:06.230$  longer effective in biochemical assays?

NOTE Confidence: 0.861997609

 $00:27:06.920 \longrightarrow 00:27:08.852$  So we have been doing the

NOTE Confidence: 0.861997609

00:27:08.852 --> 00:27:10.422 biochemical assays through ACR, oh,

NOTE Confidence: 0.861997609

 $00:27:10.422 \longrightarrow 00:27:12.396$  at the moment, we are not skilled

NOTE Confidence: 0.861997609

 $00:27:12.396 \rightarrow 00:27:14.938$  in in vitro biochemistry ourselves,

NOTE Confidence: 0.861997609

 $00:27:14.940 \longrightarrow 00:27:17.544$  and so we've just done it with

- NOTE Confidence: 0.861997609
- $00:27:17.544 \longrightarrow 00:27:19.833$  the the through the CR out and

 $00:27:19.833 \longrightarrow 00:27:21.744$  we'd be glad to to launch the

NOTE Confidence: 0.861997609

 $00:27:21.744 \rightarrow 00:27:23.280$  collaboration to investigate that,

NOTE Confidence: 0.861997609

 $00:27:23.280 \longrightarrow 00:27:24.760$  because I think that would be very powerful.

NOTE Confidence: 0.880617993333333

00:27:26.060 --> 00:27:29.894 Well, I want to thank both Jason and Kurt.

NOTE Confidence: 0.880617993333333

 $00{:}27{:}29{.}900 \dashrightarrow 00{:}27{:}33{.}020$  It makes me proud to have these kinds

NOTE Confidence: 0.880617993333333

 $00:27:33.020 \rightarrow 00:27:35.400$  of presentations on my first day here.

NOTE Confidence: 0.880617993333333

 $00:27:35.400 \longrightarrow 00:27:37.932$  So thank you very, very much

NOTE Confidence: 0.880617993333333

 $00:27:37.932 \longrightarrow 00:27:40.999$  and we'll see you all next week.