



**AB  
GRAD  
CON 23**

1  
00:00:11,770 --> 00:00:09,910

[Music]

2  
00:00:14,990 --> 00:00:11,780

hello everyone

3  
00:00:16,970 --> 00:00:15,000

my name is jono Abshire I am a second

4  
00:00:19,250 --> 00:00:16,980

year PhD student at Portland State

5  
00:00:21,650 --> 00:00:19,260

University working in the center for

6  
00:00:23,150 --> 00:00:21,660

life in extreme environments more

7  
00:00:25,070 --> 00:00:23,160

specifically out of the extreme virus

8  
00:00:27,130 --> 00:00:25,080

lab and today I'll be giving you a

9  
00:00:29,570 --> 00:00:27,140

little bit of insight into Life In Hell

10  
00:00:31,910 --> 00:00:29,580

understanding the role of toxin

11  
00:00:33,709 --> 00:00:31,920

antitoxin systems in prokaryotic genomes

12  
00:00:37,610 --> 00:00:33,719

and their potential for virus host

13  
00:00:39,170 --> 00:00:37,620

co-evolution tldr extreme organisms the

14

00:00:40,549 --> 00:00:39,180

viruses that infect them and their

15

00:00:42,709 --> 00:00:40,559

interactions in these extreme

16

00:00:44,630 --> 00:00:42,719

environments and our journey kind of

17

00:00:46,810 --> 00:00:44,640

begins right here so this is a personal

18

00:00:49,130 --> 00:00:46,820

image of Lassen Volcanic National Park

19

00:00:51,350 --> 00:00:49,140

it is one of the many areas on our

20

00:00:53,750 --> 00:00:51,360

beautiful planet that is home to these

21

00:00:56,330 --> 00:00:53,760

volcanic Hot Springs this is one of the

22

00:00:57,709 --> 00:00:56,340

main sampling sites of our lab and while

23

00:00:59,810 --> 00:00:57,719

most people wouldn't believe that there

24

00:01:02,209 --> 00:00:59,820

are things living in this environment

25

00:01:03,889 --> 00:01:02,219

there certainly are

26  
00:01:05,390 --> 00:01:03,899  
um specifically these microbes from hell

27  
00:01:07,490 --> 00:01:05,400  
and I'll get back to these guys in just

28  
00:01:09,770 --> 00:01:07,500  
a second but a little bit of background

29  
00:01:11,690 --> 00:01:09,780  
about why we why we study these

30  
00:01:13,250 --> 00:01:11,700  
particular environments so these

31  
00:01:15,170 --> 00:01:13,260  
volcanic Hot Springs are often

32  
00:01:17,210 --> 00:01:15,180  
considered and understood to be

33  
00:01:19,210 --> 00:01:17,220  
analogues for both for planets both

34  
00:01:21,950 --> 00:01:19,220  
outside and inside of our solar system

35  
00:01:25,010 --> 00:01:21,960  
referring back to some of the ancient

36  
00:01:26,570 --> 00:01:25,020  
Hot Springs found recently on Mars which

37  
00:01:29,149 --> 00:01:26,580  
may or may not have looked something

38  
00:01:31,609 --> 00:01:29,159

like this boiling Mud Pot that we find

39

00:01:32,690 --> 00:01:31,619

in Devil's Kitchen at Lassen Volcanic

40

00:01:34,789 --> 00:01:32,700

National Park

41

00:01:38,810 --> 00:01:34,799

getting back to the microbes one such

42

00:01:41,210 --> 00:01:38,820

microbe is this um micro right here uh

43

00:01:43,069 --> 00:01:41,220

spherical cell known as sacrolobus so

44

00:01:44,870 --> 00:01:43,079

factoricus this is a scanning electron

45

00:01:46,370 --> 00:01:44,880

microscope showing you the topology of

46

00:01:49,670 --> 00:01:46,380

the cells

47

00:01:52,490 --> 00:01:49,680

um coined s441 by our lab really thrives

48

00:01:54,469 --> 00:01:52,500

and uh lives in these hot acidic

49

00:01:58,730 --> 00:01:54,479

conditions with temperatures from around

50

00:02:01,190 --> 00:01:58,740

70 to 75 to 80 degrees C and ph's of

51  
00:02:04,310 --> 00:02:01,200  
around 3 to as low as one so these

52  
00:02:06,109 --> 00:02:04,320  
really extreme environments and when we

53  
00:02:09,290 --> 00:02:06,119  
think about these environments and the

54  
00:02:12,530 --> 00:02:09,300  
reach ecosystems that uh and habitat

55  
00:02:15,650 --> 00:02:12,540  
them we can also think about one of the

56  
00:02:17,570 --> 00:02:15,660  
most abundant molecules on the planet so

57  
00:02:19,790 --> 00:02:17,580  
that of viruses and yes there are these

58  
00:02:22,130 --> 00:02:19,800  
extreme viruses infecting these

59  
00:02:24,229 --> 00:02:22,140  
extremophiles uh the one which I work

60  
00:02:26,510 --> 00:02:24,239  
with in lab is sopholobus spindle shape

61  
00:02:29,809 --> 00:02:26,520  
virus one sacrilebus is a natural host

62  
00:02:32,089 --> 00:02:29,819  
of this particular virus this ssv1 this

63  
00:02:34,670 --> 00:02:32,099

is Stanley uses they them pronouns a

64

00:02:36,589 --> 00:02:34,680

really character by his lemon shape so

65

00:02:39,110 --> 00:02:36,599

these shapes are really unique to both

66

00:02:41,570 --> 00:02:39,120

archaeal organisms extremophiles and

67

00:02:43,430 --> 00:02:41,580

some of these extreme viruses

68

00:02:45,110 --> 00:02:43,440

and when we think about looking at these

69

00:02:47,270 --> 00:02:45,120

organisms and the interactions between

70

00:02:49,009 --> 00:02:47,280

them in these extreme environments we

71

00:02:52,070 --> 00:02:49,019

can look at some systems currently

72

00:02:55,550 --> 00:02:52,080

prevalent today in bacterial cells so

73

00:02:58,130 --> 00:02:55,560

these toxin antitoxin systems usually

74

00:03:00,650 --> 00:02:58,140

known as addiction modules found on

75

00:03:05,030 --> 00:03:00,660

plasmids again really prevalent in

76

00:03:07,550 --> 00:03:05,040

bacterial genomes only recently being uh

77

00:03:09,229 --> 00:03:07,560

being discovered to have some viral

78

00:03:12,470 --> 00:03:09,239

encoding

79

00:03:15,410 --> 00:03:12,480

and what this particular system confers

80

00:03:17,330 --> 00:03:15,420

is when you have a plasmid that has the

81

00:03:19,910 --> 00:03:17,340

positive uh or is positive for that

82

00:03:21,830 --> 00:03:19,920

particular addiction module that cell

83

00:03:23,690 --> 00:03:21,840

will needs that plasmid in order to

84

00:03:26,089 --> 00:03:23,700

continue its life in that particular

85

00:03:29,330 --> 00:03:26,099

environment so here we have the presence

86

00:03:31,309 --> 00:03:29,340

of the TA should that cell or progeny

87

00:03:33,110 --> 00:03:31,319

continue with the presence of that

88

00:03:35,210 --> 00:03:33,120

plasmid you would expect normal growth

89

00:03:37,009 --> 00:03:35,220

in the environment whereas if it were to

90

00:03:38,930 --> 00:03:37,019

lose that plasmid it would have

91

00:03:41,149 --> 00:03:38,940

been addicted to that particular genome

92

00:03:43,070 --> 00:03:41,159

it would die off whereas in a negative

93

00:03:45,410 --> 00:03:43,080

system you would get growth either way

94

00:03:47,330 --> 00:03:45,420

whether or not there's plasmid loss and

95

00:03:49,130 --> 00:03:47,340

we can further visualize this particular

96

00:03:51,949 --> 00:03:49,140

mechanism by looking at what might

97

00:03:53,869 --> 00:03:51,959

happen to some uncolonized cells some

98

00:03:55,789 --> 00:03:53,879

sort of event happens in which the

99

00:03:57,050 --> 00:03:55,799

addiction module is introduced to that

100

00:03:59,030 --> 00:03:57,060

population

101  
00:04:01,190 --> 00:03:59,040  
and you have your addicted Survivor

102  
00:04:02,809 --> 00:04:01,200  
going on to make new progeny and then

103  
00:04:05,270 --> 00:04:02,819  
conferring that group protection and

104  
00:04:08,149 --> 00:04:05,280  
persistence so these systems are really

105  
00:04:11,449 --> 00:04:08,159  
understood in bacteria to confer a

106  
00:04:14,270 --> 00:04:11,459  
microbial persistence phenotype some

107  
00:04:17,509 --> 00:04:14,280  
have been referred to as fate defense

108  
00:04:21,229 --> 00:04:17,519  
mechanisms and largely considered uh as

109  
00:04:23,330 --> 00:04:21,239  
plasmid stabilizers on the plasm

110  
00:04:25,010 --> 00:04:23,340  
um and we can also uh assume that

111  
00:04:26,270 --> 00:04:25,020  
another group of uncolonized cells were

112  
00:04:28,189 --> 00:04:26,280  
to come in here perhaps that group

113  
00:04:30,950 --> 00:04:28,199

encounter you would still get that toxic

114

00:04:33,230 --> 00:04:30,960

culling uh from that toxin antitoxin

115

00:04:35,030 --> 00:04:33,240

system uh one important note they are

116

00:04:37,490 --> 00:04:35,040

characterized as two genes that are

117

00:04:39,770 --> 00:04:37,500

typically right next to each other

118

00:04:42,290 --> 00:04:39,780

um the toxin being just Downstream of

119

00:04:44,990 --> 00:04:42,300

the antitoxin uh pretty stable toxin

120

00:04:47,570 --> 00:04:45,000

pretty unstable antitoxin

121

00:04:49,310 --> 00:04:47,580

and looking at this system with ssv1 a

122

00:04:54,469 --> 00:04:49,320

previous student did quite a bit of work

123

00:04:56,510 --> 00:04:54,479

in uh mutagenesis of the ssv1 genome and

124

00:04:59,150 --> 00:04:56,520

we use this at least these mutants to

125

00:05:00,650 --> 00:04:59,160

test whether or not some of these uh

126

00:05:03,170 --> 00:05:00,660

genes or open reading frames are

127

00:05:05,870 --> 00:05:03,180

essential to the virus and two genes in

128

00:05:08,090 --> 00:05:05,880

particular this T3 and TX transcript

129

00:05:11,090 --> 00:05:08,100

these particular two genes we can make

130

00:05:14,090 --> 00:05:11,100

changes to what I think is the antitoxin

131

00:05:15,710 --> 00:05:14,100

and we don't see too much

132

00:05:18,050 --> 00:05:15,720

um difference in terms of viral

133

00:05:20,390 --> 00:05:18,060

infection viral function whereas if we

134

00:05:21,830 --> 00:05:20,400

were to delete this uh or insert a

135

00:05:23,749 --> 00:05:21,840

sequence into this particular open

136

00:05:26,390 --> 00:05:23,759

reading frame what I think is the toxin

137

00:05:28,070 --> 00:05:26,400

we do see differences in infection

138

00:05:32,450 --> 00:05:28,080

mechanisms

139

00:05:34,430 --> 00:05:32,460

these extreme viruses and just virus

140

00:05:36,950 --> 00:05:34,440

reproduction overall viruses can

141

00:05:39,469 --> 00:05:36,960

typically go through two Pathways or a

142

00:05:42,409 --> 00:05:39,479

combination of both so a lyric cycle in

143

00:05:44,090 --> 00:05:42,419

which the viral genome or the viral DNA

144

00:05:46,310 --> 00:05:44,100

will insert itself into the cell

145

00:05:49,010 --> 00:05:46,320

you have some replication going on

146

00:05:51,170 --> 00:05:49,020

eventually assembly of those virions and

147

00:05:53,090 --> 00:05:51,180

the cell will then burst die and those

148

00:05:55,969 --> 00:05:53,100

viruses will go on to infect other cells

149

00:05:58,249 --> 00:05:55,979

whereas in a lysogenic cycle the viral

150

00:06:00,350 --> 00:05:58,259

genome is incorporated into the host

151

00:06:02,629 --> 00:06:00,360

genome and there's usually a latent

152

00:06:05,529 --> 00:06:02,639

phase in which it kind of just stays

153

00:06:09,230 --> 00:06:05,539

there maybe some induction event happens

154

00:06:11,810 --> 00:06:09,240

in ssv's case it actually just buds from

155

00:06:13,550 --> 00:06:11,820

the cell without killing the cell and

156

00:06:16,249 --> 00:06:13,560

one of the one of the two ways that we

157

00:06:18,650 --> 00:06:16,259

use to test for this viral reproduction

158

00:06:20,990 --> 00:06:18,660

is of course PCR so amplifying a

159

00:06:23,590 --> 00:06:21,000

sequence that's specific to viral DNA

160

00:06:27,230 --> 00:06:23,600

looking for its presence in cell-free

161

00:06:29,210 --> 00:06:27,240

supernatants another way is through Halo

162

00:06:30,710 --> 00:06:29,220

assay so very similar to a plaque assay

163

00:06:31,670 --> 00:06:30,720

in which you look where cells are

164

00:06:35,150 --> 00:06:31,680

bursting

165

00:06:37,249 --> 00:06:35,160

we cultivate a lot of cells uninfected

166

00:06:39,950 --> 00:06:37,259

sacraloba cells like you saw in the

167

00:06:41,990 --> 00:06:39,960

slide a while back and then we spot cell

168

00:06:43,490 --> 00:06:42,000

free supernatants which contain these

169

00:06:46,129 --> 00:06:43,500

mutant viruses

170

00:06:48,950 --> 00:06:46,139

onto the plate and look for clearings

171

00:06:51,950 --> 00:06:48,960

and this Halo assay is made possible

172

00:06:55,189 --> 00:06:51,960

because of the fact that ssv1 buds from

173

00:06:57,590 --> 00:06:55,199

its cell without killing the cell

174

00:06:59,689 --> 00:06:57,600

um and the onion the infected cells grow

175

00:07:02,029 --> 00:06:59,699

quite slower so it really converts this

176

00:07:03,950 --> 00:07:02,039

growth stunting phenotype in the

177

00:07:06,469 --> 00:07:03,960

sacrolobus cell

178

00:07:08,150 --> 00:07:06,479

and looking at the toxin protein and

179

00:07:09,590 --> 00:07:08,160

large a large majority of my work has

180

00:07:13,490 --> 00:07:09,600

been looking at mutants that we've made

181

00:07:15,409 --> 00:07:13,500

in this toxin protein and several uh

182

00:07:17,029 --> 00:07:15,419

machine learning softwares have

183

00:07:20,450 --> 00:07:17,039

indicated that there's a quite a high

184

00:07:22,969 --> 00:07:20,460

probability of a cleavage site over here

185

00:07:25,730 --> 00:07:22,979

behind this very hydrophobic

186

00:07:27,589 --> 00:07:25,740

uh Helix you right here and so I sought

187

00:07:30,830 --> 00:07:27,599

to characterize at least this particular

188

00:07:33,050 --> 00:07:30,840

mechanism in ssv1 through using some of

189

00:07:36,650 --> 00:07:33,060

these mutants that we already have

190

00:07:38,330 --> 00:07:36,660

um and while largely these over here are

191

00:07:40,550 --> 00:07:38,340

our wild type so really shouldn't expect

192

00:07:43,430 --> 00:07:40,560

any kind of change in viral function you

193

00:07:46,309 --> 00:07:43,440

can see some really nice Halos and we

194

00:07:48,290 --> 00:07:46,319

can confer that you know the cells are

195

00:07:49,969 --> 00:07:48,300

not dying but they're sick

196

00:07:53,749 --> 00:07:49,979

and in our mutants where we have

197

00:07:55,909 --> 00:07:53,759

insertions in in that specific toxin uh

198

00:07:57,469 --> 00:07:55,919

open reading frame we don't see any Halo

199

00:08:00,589 --> 00:07:57,479

formation

200

00:08:02,930 --> 00:08:00,599

um likewise I sought to substitute those

201  
00:08:05,689 --> 00:08:02,940  
two uh residues at the cleavage site

202  
00:08:08,629 --> 00:08:05,699  
since it's largely important for protein

203  
00:08:10,490 --> 00:08:08,639  
maturation protein function and we still

204  
00:08:13,430 --> 00:08:10,500  
don't see a Halo just switching those

205  
00:08:15,170 --> 00:08:13,440  
two residues but the main takeaway from

206  
00:08:16,969 --> 00:08:15,180  
this is that we still get virus

207  
00:08:19,550 --> 00:08:16,979  
replication

208  
00:08:21,230 --> 00:08:19,560  
um we still get quite a bit of at least

209  
00:08:23,869 --> 00:08:21,240  
high levels of viral replication from

210  
00:08:25,610 --> 00:08:23,879  
some of these mutants and one of these

211  
00:08:28,249 --> 00:08:25,620  
mutants particular we didn't see

212  
00:08:29,869 --> 00:08:28,259  
anything so maybe that one's detrimental

213  
00:08:32,630 --> 00:08:29,879

to that particular open reading frame

214

00:08:34,550 --> 00:08:32,640

but we do see that there is virus

215

00:08:38,389 --> 00:08:34,560

replication happening in the cell

216

00:08:40,250 --> 00:08:38,399

supernatant when we screen for DNA

217

00:08:42,829 --> 00:08:40,260

um and it's important to kind of note

218

00:08:44,329 --> 00:08:42,839

that this this protein isn't just unique

219

00:08:46,190 --> 00:08:44,339

to ssv1

220

00:08:48,650 --> 00:08:46,200

um in in looking at these systems in

221

00:08:50,690 --> 00:08:48,660

these extreme organisms uh this one in

222

00:08:53,210 --> 00:08:50,700

particular up here is another SSV so

223

00:08:55,190 --> 00:08:53,220

another spindle shaped virus in which we

224

00:08:58,370 --> 00:08:55,200

see really low sequence similarity

225

00:09:01,490 --> 00:08:58,380

across these particular genes but quite

226

00:09:03,769 --> 00:09:01,500

a hot quite a high bit of um structural

227

00:09:05,410 --> 00:09:03,779

similarity between these genes and a lot

228

00:09:08,210 --> 00:09:05,420

of these are

229

00:09:11,150 --> 00:09:08,220

genomes of extremophiles while some are

230

00:09:13,430 --> 00:09:11,160

virally encoded

231

00:09:15,410 --> 00:09:13,440

and uh one of the main takeaways really

232

00:09:17,210 --> 00:09:15,420

just wrapping uh wrapping it all back

233

00:09:19,070 --> 00:09:17,220

around looking for these particular

234

00:09:20,750 --> 00:09:19,080

mechanisms and how they confer

235

00:09:23,570 --> 00:09:20,760

persistence with both their

236

00:09:26,210 --> 00:09:23,580

extremophilic organism and their viruses

237

00:09:28,970 --> 00:09:26,220

would be a really nice first start in

238

00:09:32,630 --> 00:09:28,980

looking for these biosignatures perhaps

239

00:09:34,790 --> 00:09:32,640

um and ways that we can look into how

240

00:09:36,829 --> 00:09:34,800

the microbes might be interacting

241

00:09:39,050 --> 00:09:36,839

in these particular extreme environments

242

00:09:41,389 --> 00:09:39,060

this is another sampling video

243

00:09:43,910 --> 00:09:41,399

um just of Lassen Volcanic Park pretty

244

00:09:45,290 --> 00:09:43,920

pretty close to where s441 was isolated

245

00:09:47,329 --> 00:09:45,300

from

246

00:09:49,130 --> 00:09:47,339

uh looks pretty similar to maybe

247

00:09:50,870 --> 00:09:49,140

something we'd see on another planet

248

00:09:54,410 --> 00:09:50,880

that we could sample at

249

00:09:56,389 --> 00:09:54,420

and then a Shameless plug but looking

250

00:09:58,009 --> 00:09:56,399

for viruses in space I think is the

251

00:10:00,050 --> 00:09:58,019

obvious next step we know that they're

252

00:10:03,350 --> 00:10:00,060

the most abundant molecule on the planet

253

00:10:04,970 --> 00:10:03,360

and just understanding that these

254

00:10:07,490 --> 00:10:04,980

mechanisms and that these systems are

255

00:10:10,190 --> 00:10:07,500

out there allowing these cells to thrive

256

00:10:13,370 --> 00:10:10,200

in this particular environment would be

257

00:10:16,009 --> 00:10:13,380

a great uh in starter conversation for

258

00:10:17,269 --> 00:10:16,019

looking for these small biomolecules in

259

00:10:19,490 --> 00:10:17,279

space

260

00:10:20,870 --> 00:10:19,500

and with that I just want to thank my

261

00:10:23,449 --> 00:10:20,880

lab

262

00:10:26,570 --> 00:10:23,459

um my group over here at Boiling Springs

263

00:10:31,770 --> 00:10:26,580

Lake when we collected samples and my Pi

264

00:10:31,780 --> 00:10:35,570

[Music]

265

00:10:43,250 --> 00:10:38,210

thank you General we have a time for

266

00:10:46,850 --> 00:10:45,350

hey my name is Pia and I'm really

267

00:10:48,650 --> 00:10:46,860

interested in like Christopher Cross

268

00:10:50,150 --> 00:10:48,660

Community of microbes

269

00:10:52,009 --> 00:10:50,160

um have you looked at all into the

270

00:10:53,210 --> 00:10:52,019

interactions of other phage defense

271

00:10:57,170 --> 00:10:53,220

systems

272

00:10:59,449 --> 00:10:57,180

um yeah so at least in terms of uh this

273

00:11:01,670 --> 00:10:59,459

toxin antitoxin system they're pretty

274

00:11:03,170 --> 00:11:01,680

understood in bacterial cells to like

275

00:11:06,650 --> 00:11:03,180

incorporate themselves into the genomic

276

00:11:09,110 --> 00:11:06,660

crispr cas9 there are some proteins that

277

00:11:10,970 --> 00:11:09,120

um that look pretty similar to this a291

278

00:11:14,990 --> 00:11:10,980

and some of these other virally encoded

279

00:11:18,530 --> 00:11:15,000

genes that do have some kind of crispr

280

00:11:20,449 --> 00:11:18,540

casts editing mechanism so yeah I think

281

00:11:21,829 --> 00:11:20,459

they it might be leaning towards that

282

00:11:24,050 --> 00:11:21,839

way but they're pretty well

283

00:11:26,750 --> 00:11:24,060

characterized in terms of like how they

284

00:11:28,910 --> 00:11:26,760

work with proteins and mRNA so um yeah

285

00:11:30,530 --> 00:11:28,920

not too far there yet but

286

00:11:44,090 --> 00:11:30,540

um definitely definitely somewhere in

287

00:11:49,790 --> 00:11:46,310

uh thank you for the talk um Marshall

288

00:11:51,470 --> 00:11:49,800

Seton JPL and I I was curious so um

289

00:11:53,569 --> 00:11:51,480

you're talking about looking for viruses

290

00:11:55,190 --> 00:11:53,579

and extraterrestrial environments um do

291

00:11:57,230 --> 00:11:55,200

you think there would be or I'm curious

292

00:11:59,990 --> 00:11:57,240

I'm completely ignoring yeah so no no I

293

00:12:02,210 --> 00:12:00,000

dropped out of biotube so for this

294

00:12:04,370 --> 00:12:02,220

um for viruses specifically like it

295

00:12:07,370 --> 00:12:04,380

would small molecule biosignature

296

00:12:09,650 --> 00:12:07,380

classes vary from viruses to like what

297

00:12:12,769 --> 00:12:09,660

you'd be looking for to cells yeah yeah

298

00:12:14,210 --> 00:12:12,779

I I for sure and that kind of brings me

299

00:12:15,769 --> 00:12:14,220

at least back to a point like of looking

300

00:12:18,170 --> 00:12:15,779

for fossils and stuff like that so

301

00:12:20,150 --> 00:12:18,180

viruses and sediment

302

00:12:22,490 --> 00:12:20,160

um and at least our lab there's been

303

00:12:24,470 --> 00:12:22,500

some work in like silica and coding

304

00:12:26,990 --> 00:12:24,480

viruses and how you know the stability

305

00:12:28,910 --> 00:12:27,000

of them so maybe there's a pretty silica

306

00:12:30,710 --> 00:12:28,920

Rich environment in which we might be

307

00:12:32,870 --> 00:12:30,720

able to look for any kind of these

308

00:12:35,509 --> 00:12:32,880

signatures so like proteins

309

00:12:37,190 --> 00:12:35,519

um or or yeah just specific things that

310

00:12:38,870 --> 00:12:37,200

might might have been preserved in those

311

00:12:41,150 --> 00:12:38,880

particular environments

312

00:12:43,790 --> 00:12:41,160

um considering that these viruses live

313

00:12:45,590 --> 00:12:43,800

and thrive in in pretty extremes

314

00:12:47,269 --> 00:12:45,600

um you know they're they're they got

315

00:12:50,210 --> 00:12:47,279

they got to be out there at least in my

316

00:12:52,310 --> 00:12:50,220

mind so you mentioned um looking at like

317

00:12:54,590 --> 00:12:52,320

morphology and fossils and things like

318

00:12:57,350 --> 00:12:54,600

that I know that at least for

319

00:13:00,110 --> 00:12:57,360

um looking at cells and things they've

320

00:13:02,629 --> 00:13:00,120

been shown to to mimic abiotic systems

321

00:13:06,170 --> 00:13:02,639

uh pretty well because humans are very

322

00:13:08,949 --> 00:13:06,180

good pattern recognition and so um has

323

00:13:13,190 --> 00:13:08,959

anything been seen like that for viruses

324

00:13:14,750 --> 00:13:13,200

yeah not the not that I'm familiar with

325

00:13:16,670 --> 00:13:14,760

um I know it's at least a brand new

326

00:13:18,829 --> 00:13:16,680

conversation of starting to look for

327

00:13:21,170 --> 00:13:18,839

like viruses in space Astro virology

328

00:13:23,269 --> 00:13:21,180

things like that so uh yeah we're just

329

00:13:24,590 --> 00:13:23,279

kind of getting up and running oh yeah

330

00:13:26,389 --> 00:13:24,600

yeah no sorry I don't mean to like play

331

00:13:32,590 --> 00:13:26,399

20 Questions oh no I just thought it was

332

00:13:37,610 --> 00:13:35,870

yeah there's a there's a little cure lab

333

00:13:40,009 --> 00:13:37,620

that like goes on right now that I teach

334

00:13:44,030 --> 00:13:40,019

so I'm obviously not there but they're

335

00:13:44,040 --> 00:13:57,520

any other questions