



1
00:00:00,790 --> 00:00:07,320

[Music]

2
00:00:11,860 --> 00:00:09,299

[Applause]

3
00:00:13,629 --> 00:00:11,870

what I'm gonna be talking about today is

4
00:00:17,080 --> 00:00:13,639

some work that we've been trying to do

5
00:00:19,750 --> 00:00:17,090

to look at how we might be able to

6
00:00:24,370 --> 00:00:19,760

distinguish the identity of different

7
00:00:26,620 --> 00:00:24,380

carbon fixation pathways in deep time so

8
00:00:28,660 --> 00:00:26,630

I mean we care about this right because

9
00:00:30,849 --> 00:00:28,670

not only are we taking carbon dioxide

10
00:00:34,120 --> 00:00:30,859

and converting it to biomass but it has

11
00:00:36,370 --> 00:00:34,130

to do with energetics material transfer

12
00:00:39,340 --> 00:00:36,380

and perhaps more importantly the

13
00:00:41,350 --> 00:00:39,350

Lindemann style trophic dynamics of the

14

00:00:43,720 --> 00:00:41,360

global biosphere at the time and one of

15

00:00:46,300 --> 00:00:43,730

the things that you can do to identify

16

00:00:48,850 --> 00:00:46,310

which autotrophic pathway may actually

17

00:00:52,420 --> 00:00:48,860

be fixing carbon is by looking at the

18

00:00:54,610 --> 00:00:52,430

ratios of carbon-13 to carbon-12 it

19

00:00:57,730 --> 00:00:54,620

turns out that due to the tension

20

00:00:59,890 --> 00:00:57,740

between equilibrium isotope effects and

21

00:01:03,070 --> 00:00:59,900

enzymatically controlled kinetic isotope

22

00:01:05,649 --> 00:01:03,080

effects you will end up with at the end

23

00:01:08,550 --> 00:01:05,659

of this process biomass that's enriched

24

00:01:11,980 --> 00:01:08,560

in the light isotope of carbon and

25

00:01:13,960 --> 00:01:11,990

leaving behind carbon dioxide that is

26
00:01:18,510 --> 00:01:13,970
enriched in the heavy isotope of carbon

27
00:01:21,570 --> 00:01:18,520
okay so how do we actually get at this

28
00:01:24,790 --> 00:01:21,580
effect and how do we use it to identify

29
00:01:27,100 --> 00:01:24,800
different carbon fixation pathways well

30
00:01:29,200 --> 00:01:27,110
one of the things that gave me a great

31
00:01:33,040 --> 00:01:29,210
pleasure to here is that I'm gonna be

32
00:01:35,820 --> 00:01:33,050
talking before Jody who has one of what

33
00:01:39,070 --> 00:01:35,830
I think the most crystal-clear

34
00:01:41,830 --> 00:01:39,080
presentations on how you actually can go

35
00:01:44,350 --> 00:01:41,840
out into the environment and figure out

36
00:01:46,690 --> 00:01:44,360
from the carbon isotopic composition of

37
00:01:48,460 --> 00:01:46,700
dissolve CO₂ or its counterpart

38
00:01:51,010 --> 00:01:48,470

dissolved bicarbonate and then the

39

00:01:53,560 --> 00:01:51,020

carbon isotopic composition of primary

40

00:01:55,899 --> 00:01:53,570

biomass in here a particulate organic

41

00:01:58,660 --> 00:01:55,909

carbon and figure out from the isotopic

42

00:02:00,880 --> 00:01:58,670

difference there the isotope

43

00:02:02,950 --> 00:02:00,890

fractionation associated with primary

44

00:02:04,390 --> 00:02:02,960

production and if you go and read this

45

00:02:06,580 --> 00:02:04,400

paper one of the amazing things about

46

00:02:08,410 --> 00:02:06,590

this data set is that if you correct for

47

00:02:10,539 --> 00:02:08,420

nitrate and you correct for temperature

48

00:02:12,570 --> 00:02:10,549

you're left with the temporal signal

49

00:02:17,199 --> 00:02:12,580

that may have something to do with

50

00:02:18,940 --> 00:02:17,209

enhanced co2 access as time has

51
00:02:20,680 --> 00:02:18,950
increased okay

52
00:02:23,770 --> 00:02:20,690
so one of the things though that we have

53
00:02:27,460 --> 00:02:23,780
to worry about is that rocks are not

54
00:02:31,270 --> 00:02:27,470
biomass and dissolved co2 so how do we

55
00:02:33,550 --> 00:02:31,280
go from the dissolve co2 carbon isotopic

56
00:02:35,620 --> 00:02:33,560
composition the composition of biomass

57
00:02:37,630 --> 00:02:35,630
and figure out something from the

58
00:02:39,550 --> 00:02:37,640
geologic record well the way we do we do

59
00:02:41,949 --> 00:02:39,560
is we approach it in this sort of haze

60
00:02:44,020 --> 00:02:41,959
diagram since we've got the carbon

61
00:02:46,240 --> 00:02:44,030
isotopic composition of these different

62
00:02:48,400 --> 00:02:46,250
compounds on the Left axis here is what

63
00:02:51,009 --> 00:02:48,410

we want this difference between dissolve

64

00:02:52,990 --> 00:02:51,019

co2 and biomass but we have to be

65

00:02:55,960 --> 00:02:53,000

concerned with the temperature dependent

66

00:02:58,420 --> 00:02:55,970

isotopic fractionation associated with

67

00:03:02,130 --> 00:02:58,430

carbonate speciation we have to be

68

00:03:05,590 --> 00:03:02,140

concerned with the mineral logic

69

00:03:07,780 --> 00:03:05,600

associated fractionation associated with

70

00:03:09,580 --> 00:03:07,790

the different precipitation of carbonate

71

00:03:11,289 --> 00:03:09,590

minerals and then of course there's

72

00:03:13,059 --> 00:03:11,299

going to be some small but important

73

00:03:15,280 --> 00:03:13,069

fractionation associated with the

74

00:03:19,120 --> 00:03:15,290

reworking of primary biomass into

75

00:03:21,729 --> 00:03:19,130

sedimentary total organic carbon okay so

76
00:03:24,940 --> 00:03:21,739
one of the things is that what does this

77
00:03:27,940 --> 00:03:24,950
record of mineral carbon isotopic

78
00:03:30,940 --> 00:03:27,950
compositions and total organic carbon

79
00:03:33,220 --> 00:03:30,950
isotopic compositions look like in deep

80
00:03:35,680 --> 00:03:33,230
time one of the things that were very

81
00:03:38,830 --> 00:03:35,690
fortunate for is that there was recently

82
00:03:41,110 --> 00:03:38,840
an incredible statistical reanalysis of

83
00:03:43,660 --> 00:03:41,120
the carbon isotope record in deep time

84
00:03:45,940 --> 00:03:43,670
and the carbon isotope record in deep

85
00:03:47,470 --> 00:03:45,950
time it turns out has two distinct types

86
00:03:48,819 --> 00:03:47,480
of States there's going to be the

87
00:03:51,690 --> 00:03:48,829
intervals that we're going to ignore

88
00:03:54,099 --> 00:03:51,700

here so called transitional intervals

89

00:03:56,199 --> 00:03:54,109

associated with climatic upheavals and

90

00:03:58,210 --> 00:03:56,209

non steady-state behavior in the carbon

91

00:04:00,460 --> 00:03:58,220

cycle but we're going to be focusing

92

00:04:03,039 --> 00:04:00,470

instead on these non transitional

93

00:04:05,949 --> 00:04:03,049

intervals largely the phanerozoic middle

94

00:04:08,349 --> 00:04:05,959

Proterozoic and back in time so what we

95

00:04:11,710 --> 00:04:08,359

want to do is we want to invert this

96

00:04:14,440 --> 00:04:11,720

haze diagram then and go from the carbon

97

00:04:16,870 --> 00:04:14,450

isotopic composition of sedimentary

98

00:04:18,670 --> 00:04:16,880

carbonate minerals the carbon isotopic

99

00:04:21,279 --> 00:04:18,680

composition of sedimentary organic

100

00:04:24,879 --> 00:04:21,289

carbon and go through a procedure that

101
00:04:27,430 --> 00:04:24,889
allows us to extract to the best of our

102
00:04:30,850 --> 00:04:27,440
ability the isotopic difference between

103
00:04:31,510 --> 00:04:30,860
dissolve co2 in primary biomass in deep

104
00:04:35,920 --> 00:04:31,520
time

105
00:04:39,999 --> 00:04:35,930
so it turns out that this technique was

106
00:04:41,650 --> 00:04:40,009
developed by Sarah Hurley a postdoc I'm

107
00:04:43,600 --> 00:04:41,660
fortunate enough to be working with and

108
00:04:45,309 --> 00:04:43,610
what it happens as we go like this let's

109
00:04:48,520 --> 00:04:45,319
look at a histogram for a certain time

110
00:04:50,830 --> 00:04:48,530
period of the isotopic composition of

111
00:04:52,620 --> 00:04:50,840
carbon eight carbon and what Sara

112
00:04:55,629 --> 00:04:52,630
realized is that if you do an agnostic

113
00:04:58,089 --> 00:04:55,639

resampling of the isotopic effects

114

00:05:00,420 --> 00:04:58,099

associated with carbonate mineral

115

00:05:02,670 --> 00:05:00,430

precipitation the temperature derived

116

00:05:05,649 --> 00:05:02,680

isotopic effects associated with

117

00:05:08,469 --> 00:05:05,659

carbonate speciation what you can end up

118

00:05:11,469 --> 00:05:08,479

with is you can end up with a frequency

119

00:05:13,930 --> 00:05:11,479

distribution of the Delta 13c carbon

120

00:05:15,490 --> 00:05:13,940

isotopic composition of dissolved co2

121

00:05:18,270 --> 00:05:15,500

for the time period that you're

122

00:05:21,249 --> 00:05:18,280

interested similarly if you take the

123

00:05:23,830 --> 00:05:21,259

isotopic distribution associated with

124

00:05:26,700 --> 00:05:23,840

sedimentary organic carbon and run it

125

00:05:28,870 --> 00:05:26,710

through a random resampling looking at a

126

00:05:32,920 --> 00:05:28,880

distribution that might have something

127

00:05:36,610 --> 00:05:32,930

to do with how or primary biomass is

128

00:05:39,339 --> 00:05:36,620

fractionated during microbial reworking

129

00:05:41,589 --> 00:05:39,349

you can end up with a frequency

130

00:05:44,260 --> 00:05:41,599

distribution of the carbon isotopic

131

00:05:46,959 --> 00:05:44,270

composition of biomass in deep time

132

00:05:48,850 --> 00:05:46,969

combining these two effectively just

133

00:05:51,370 --> 00:05:48,860

taking their difference gives you a

134

00:05:54,010 --> 00:05:51,380

frequency distribution of the so called

135

00:05:56,469 --> 00:05:54,020

epsilon P for primary productivity

136

00:05:59,800 --> 00:05:56,479

epsilon for a difference in Delta values

137

00:06:02,409 --> 00:05:59,810

that then may tell you something about

138

00:06:07,480 --> 00:06:02,419

what carbon fixation pathways were

139

00:06:11,680 --> 00:06:07,490

operating in deep time okay so when

140

00:06:15,339 --> 00:06:11,690

Sarah did this with the non transitional

141

00:06:18,370 --> 00:06:15,349

intervals in the carbon isotope record

142

00:06:21,010 --> 00:06:18,380

what she returned was the following type

143

00:06:22,930 --> 00:06:21,020

of distributions if we start over here

144

00:06:24,939 --> 00:06:22,940

something centered at about 50 million

145

00:06:27,580 --> 00:06:24,949

years ago going back to something about

146

00:06:30,670 --> 00:06:27,590

250 million years ago what you can see

147

00:06:33,700 --> 00:06:30,680

is a slight increase in the most

148

00:06:35,709 --> 00:06:33,710

frequent value of carbon isotope

149

00:06:38,350 --> 00:06:35,719

fractionation by the dominant primary

150

00:06:40,390 --> 00:06:38,360

producers at that time though there are

151
00:06:42,310 --> 00:06:40,400
a number of evolutionary events that may

152
00:06:44,620 --> 00:06:42,320
be associated with this it's also

153
00:06:44,890 --> 00:06:44,630
possible that what you're looking at is

154
00:06:47,680 --> 00:06:44,900
CIN

155
00:06:50,590 --> 00:06:47,690
we the effect of the gross long-term

156
00:06:53,620 --> 00:06:50,600
increase in co2 as we go back through

157
00:06:56,439 --> 00:06:53,630
the phanerozoic and as co2 increases

158
00:06:59,590 --> 00:06:56,449
what you end up with is you end up with

159
00:07:02,140 --> 00:06:59,600
less of a more efficient operation of

160
00:07:03,820 --> 00:07:02,150
the carbon co2 concentrating mechanisms

161
00:07:06,700 --> 00:07:03,830
that are associated with these organisms

162
00:07:10,029 --> 00:07:06,710
leading to a larger fractionation we're

163
00:07:11,920 --> 00:07:10,039

going to focus next on two intervals for

164

00:07:14,680 --> 00:07:11,930

a case study the case study one is this

165

00:07:16,480 --> 00:07:14,690

mid Proterozoic interval and what we

166

00:07:19,120 --> 00:07:16,490

want to do is we want to ask the

167

00:07:21,790 --> 00:07:19,130

following question was the middle

168

00:07:24,760 --> 00:07:21,800

Proterozoic actually the age of

169

00:07:26,939 --> 00:07:24,770

cyanobacteria what I'm showing here is a

170

00:07:30,850 --> 00:07:26,949

diagram that has a bunch of different

171

00:07:35,320 --> 00:07:30,860

compilations of oxygen levels from today

172

00:07:37,000 --> 00:07:35,330

back to the early archaea and shown here

173

00:07:40,120 --> 00:07:37,010

is of course the great oxidation

174

00:07:43,450 --> 00:07:40,130

oxidation interval but also shown down

175

00:07:46,540 --> 00:07:43,460

here in green are some indications both

176
00:07:49,420 --> 00:07:46,550
from geochemistry and also from micro

177
00:07:52,000 --> 00:07:49,430
fossils of cyanobacterial oxygenic

178
00:07:56,320 --> 00:07:52,010
photosynthesis and then on this side of

179
00:08:00,520 --> 00:07:56,330
the diagram some information from

180
00:08:02,890 --> 00:08:00,530
molecular clocks from micro fossil

181
00:08:06,219 --> 00:08:02,900
evidence and from biomarker evidence for

182
00:08:08,950 --> 00:08:06,229
a primary production global primary

183
00:08:11,560 --> 00:08:08,960
biospheric primary production dominated

184
00:08:14,290 --> 00:08:11,570
by algae and what we're asking is in

185
00:08:16,570 --> 00:08:14,300
this mid Proterozoic interval wasn't

186
00:08:19,659 --> 00:08:16,580
really an age of cyanobacteria can we

187
00:08:22,000 --> 00:08:19,669
use this technique to say okay the

188
00:08:24,490 --> 00:08:22,010

primary producer at this time which it

189

00:08:26,529 --> 00:08:24,500

seems like it was a sign of

190

00:08:31,719 --> 00:08:26,539

bacterial population or was it something

191

00:08:34,120 --> 00:08:31,729

else so one of the things that is a

192

00:08:37,350 --> 00:08:34,130

complicating factor in this is that all

193

00:08:39,969 --> 00:08:37,360

extant cyanobacteria have a CO_2

194

00:08:42,610 --> 00:08:39,979

concentrating mechanism probably the

195

00:08:45,100 --> 00:08:42,620

most famous bacterial organelle is known

196

00:08:48,010 --> 00:08:45,110

as the carboxy zone it's where cyanosis

197

00:08:51,160 --> 00:08:48,020

today have their Rubisco have all their

198

00:08:53,860 --> 00:08:51,170

carbonic anhydrase they bring in bicarb

199

00:08:55,990 --> 00:08:53,870

it gets converted through the activity

200

00:08:58,690 --> 00:08:56,000

of carbonic anhydrase to carbon dioxide

201
00:09:00,280 --> 00:08:58,700
right at the site of carbon fixation

202
00:09:02,920 --> 00:09:00,290
and what you can do is you can overcome

203
00:09:04,810 --> 00:09:02,930
some of the catalytic inefficiency of

204
00:09:08,350 --> 00:09:04,820
Rubisco in that way and there have been

205
00:09:10,710 --> 00:09:08,360
a whole bunch of different estimates for

206
00:09:13,000 --> 00:09:10,720
when the cyanobacterial CO_2

207
00:09:15,280 --> 00:09:13,010
concentrating mechanism may have come

208
00:09:19,540 --> 00:09:15,290
into play and what it is usually

209
00:09:21,970 --> 00:09:19,550
associated with is drops in carbon

210
00:09:24,340 --> 00:09:21,980
dioxide levels it makes sense if carbon

211
00:09:27,220 --> 00:09:24,350
dioxide levels fall you would expect

212
00:09:31,510 --> 00:09:27,230
that a CO_2 concentrating mechanism would

213
00:09:35,740 --> 00:09:31,520

be developed so we decided to test this

214

00:09:39,610 --> 00:09:35,750

and Sarah in the lab was able to create

215

00:09:42,240 --> 00:09:39,620

a carboxy zone this mutant of the

216

00:09:51,400 --> 00:09:42,250

wild-type cyanobacterium Senate caucus

217

00:09:53,920 --> 00:09:51,410

7002 and these green fluorescence images

218

00:09:58,150 --> 00:09:53,930

show the distribution of Rubisco in

219

00:10:01,870 --> 00:09:58,160

these mutant cyanobacteria she grew them

220

00:10:04,330 --> 00:10:01,880

under different co2 levels and it turns

221

00:10:06,370 --> 00:10:04,340

out that they're a high carbon requiring

222

00:10:08,580 --> 00:10:06,380

mutant which means that growing them at

223

00:10:10,840 --> 00:10:08,590

very low co2 levels is very difficult

224

00:10:12,910 --> 00:10:10,850

fortunately though it's a relatively

225

00:10:14,950 --> 00:10:12,920

simple system to model where you just

226

00:10:16,990 --> 00:10:14,960

have co2 diffusing into the site of

227

00:10:19,150 --> 00:10:17,000

carbon fixation so we were able to model

228

00:10:23,380 --> 00:10:19,160

what might happen at lower co2 levels

229

00:10:26,590 --> 00:10:23,390

even at relatively low co2 levels what

230

00:10:30,300 --> 00:10:26,600

you can see is that the range of carbon

231

00:10:33,990 --> 00:10:30,310

isotopes that are produced by these

232

00:10:37,690 --> 00:10:34,000

potentially physiologically ancestral

233

00:10:41,380 --> 00:10:37,700

cyanobacteria does not cover more than

234

00:10:44,050 --> 00:10:41,390

about 30% of the epsilon primary

235

00:10:47,080 --> 00:10:44,060

productivity range that we see in the

236

00:10:49,450 --> 00:10:47,090

middle Proterozoic so what happens when

237

00:10:52,120 --> 00:10:49,460

you just take a wild-type cyanobacterium

238

00:10:54,100 --> 00:10:52,130

and grow it again under different co2

239

00:10:56,650 --> 00:10:54,110

levels well what happens is you produce

240

00:10:59,950 --> 00:10:56,660

a range of carbon isotope values that

241

00:11:01,780 --> 00:10:59,960

covers 95% of the middle proterozoic

242

00:11:05,320 --> 00:11:01,790

epsilon primary productivity

243

00:11:10,180 --> 00:11:05,330

distribution suggesting then that one of

244

00:11:11,950 --> 00:11:10,190

the simplest interpretations of the

245

00:11:14,260 --> 00:11:11,960

middle Proterozoic

246

00:11:17,320 --> 00:11:14,270

primary productivity record is that

247

00:11:19,269 --> 00:11:17,330

cyanobacteria with a carbon can't see

248

00:11:22,950 --> 00:11:19,279

you too concentrating mechanism for the

249

00:11:25,630 --> 00:11:22,960

dominant primary producer at this time

250

00:11:28,389 --> 00:11:25,640

great so what I'm gonna do now is I'm

251
00:11:31,510 --> 00:11:28,399
gonna speed into the second case study

252
00:11:34,300 --> 00:11:31,520
which is looking in very deep time and

253
00:11:36,519 --> 00:11:34,310
saying okay what might this carbon

254
00:11:39,850 --> 00:11:36,529
isotope fractionation distribution mean

255
00:11:41,260 --> 00:11:39,860
there's some similarity here between

256
00:11:43,750 --> 00:11:41,270
what you see in the middle Proterozoic

257
00:11:46,930 --> 00:11:43,760
and what you see in the early Archaean

258
00:11:49,449 --> 00:11:46,940
and in fact that may be used to say okay

259
00:11:51,460 --> 00:11:49,459
the same carbon fixation pathways were

260
00:11:52,990 --> 00:11:51,470
an operation at this time what we

261
00:11:56,260 --> 00:11:53,000
decided to do is a thought experiment

262
00:11:59,380 --> 00:11:56,270
and working with some colleagues at BU

263
00:12:02,560 --> 00:11:59,390

Daniel se gray and Josh Gould furred we

264

00:12:08,220 --> 00:12:02,570

decided to look at what might

265

00:12:10,660 --> 00:12:08,230

potentially happen if you created a

266

00:12:12,340 --> 00:12:10,670

endergonic and biomass capable

267

00:12:15,519 --> 00:12:12,350

metabolism from a simple set of

268

00:12:18,570 --> 00:12:15,529

potential prebiotic seed compounds shown

269

00:12:21,820 --> 00:12:18,580

down here acetate carbon dioxide 4 made

270

00:12:24,280 --> 00:12:21,830

hydrogen sulfide and some others and so

271

00:12:26,320 --> 00:12:24,290

essentially what Daniel and Josh did is

272

00:12:33,430 --> 00:12:26,330

they went ahead and they led to a

273

00:12:35,440 --> 00:12:33,440

network expansion associated with this

274

00:12:37,420 --> 00:12:35,450

metabolism and they were able to show

275

00:12:39,579 --> 00:12:37,430

that not only was it endergonic it was

276

00:12:40,870 --> 00:12:39,589

capable of producing biomass and what we

277

00:12:44,970 --> 00:12:40,880

wanted to know is what's the carbon

278

00:12:47,590 --> 00:12:44,980

isotopic consequences of this expansion

279

00:12:50,220 --> 00:12:47,600

so it turns out the carbon isotope

280

00:12:54,220 --> 00:12:50,230

fractionation is both incredibly

281

00:12:59,680 --> 00:12:54,230

confusing but at some level relatively

282

00:13:02,560 --> 00:12:59,690

straightforward so it depends on two

283

00:13:04,600 --> 00:13:02,570

things primarily one the nearest

284

00:13:07,240 --> 00:13:04,610

neighbors that the carbon atom is bonded

285

00:13:09,340 --> 00:13:07,250

to and then the nearest neighbors to

286

00:13:11,650 --> 00:13:09,350

those nearest neighbors and so what a

287

00:13:14,470 --> 00:13:11,660

colleague Chris butch was able to do was

288

00:13:17,530 --> 00:13:14,480

take this full set hundreds of different

289

00:13:22,620 --> 00:13:17,540

compounds and actually calculate what

290

00:13:25,060 --> 00:13:22,630

the isotopic fractionation would be as

291

00:13:32,210 --> 00:13:25,070

this explained

292

00:13:35,660 --> 00:13:32,220

happens okay and here shown on the left

293

00:13:39,439 --> 00:13:35,670

is the iterations here are two images of

294

00:13:41,509 --> 00:13:39,449

that epsilon primary productivity on the

295

00:13:43,730 --> 00:13:41,519

top the number of compounds in the

296

00:13:45,470 --> 00:13:43,740

network is shown linearly on the bottom

297

00:13:47,720 --> 00:13:45,480

we've got it plotted versus the log

298

00:13:49,400 --> 00:13:47,730

number of networks allowed number of

299

00:13:51,650 --> 00:13:49,410

compounds in the network this one is

300

00:13:54,800 --> 00:13:51,660

easier to see the differences the early

301
00:13:58,490 --> 00:13:54,810
networks are shown here in blue and in

302
00:14:00,759 --> 00:13:58,500
red but as the network increases in

303
00:14:03,620 --> 00:14:00,769
diversity and potentially in complexity

304
00:14:08,199 --> 00:14:03,630
isotope fractionation actually goes the

305
00:14:12,110 --> 00:14:08,209
other way it is narrowing up and

306
00:14:14,120 --> 00:14:12,120
approaching a single mode at about 18

307
00:14:21,610 --> 00:14:14,130
per mil for the difference between

308
00:14:24,139 --> 00:14:21,620
dissolve co₂ and biomass so what's the

309
00:14:27,050 --> 00:14:24,149
reason behind this it turns out that

310
00:14:28,519 --> 00:14:27,060
when you count the bonds and the bonding

311
00:14:30,470 --> 00:14:28,529
partners that are involved in this as

312
00:14:35,240 --> 00:14:30,480
Chris has done here what you see is that

313
00:14:37,220 --> 00:14:35,250

as the network expands your overall

314

00:14:39,009 --> 00:14:37,230

bonding environment is getting more

315

00:14:42,170 --> 00:14:39,019

reduced you're seeing a lot more

316

00:14:44,689 --> 00:14:42,180

carbon-carbon bonds and carbon hydrogen

317

00:14:46,819 --> 00:14:44,699

bonds and here what I've shown is the

318

00:14:49,160 --> 00:14:46,829

distribution of Chris's chemo

319

00:14:51,740 --> 00:14:49,170

informatics prediction along with the

320

00:14:53,210 --> 00:14:51,750

early Archaean distribution and though

321

00:14:55,759 --> 00:14:53,220

they're not exactly the same there may

322

00:14:57,949 --> 00:14:55,769

be some kinetic effects here and less

323

00:15:00,230 --> 00:14:57,959

equilibrium effects here one thing

324

00:15:02,780 --> 00:15:00,240

that's very clear is that an epsilon

325

00:15:05,030 --> 00:15:02,790

primary productivity of around 18 per

326

00:15:07,639 --> 00:15:05,040

mil or 20 per mil a'somethin that would

327

00:15:10,400 --> 00:15:07,649

have been in the past associated with

328

00:15:13,460 --> 00:15:10,410

say the carbon fixation by the Calvin

329

00:15:15,620 --> 00:15:13,470

cycle may simply reflect the development

330

00:15:18,439 --> 00:15:15,630

in these networks of aliphatic

331

00:15:20,420 --> 00:15:18,449

carbon-carbon bond okay I'll leave you

332

00:15:22,699 --> 00:15:20,430

now with what we know what we think we

333

00:15:26,009 --> 00:15:22,709

know and what we want to know and thanks

334

00:15:27,280 --> 00:15:26,280

you