

2 ASTROBIOLOGY
0 GRADUATE
1 CONFERENCE
7



CHARLOTTESVILLE, VA

1
00:00:00,790 --> 00:00:07,419

[Music]

2
00:00:12,530 --> 00:00:09,830

so hello everyone my name is Nicolas

3
00:00:14,209 --> 00:00:12,540

Kovich my advisor is dr. Lauren Williams

4
00:00:16,670 --> 00:00:14,219

and we are at the Georgia Tech at

5
00:00:18,500 --> 00:00:16,680

Georgia Tech in Atlanta Georgia on the

6
00:00:20,269 --> 00:00:18,510

bioinformatics PhD candidate the school

7
00:00:22,640 --> 00:00:20,279

of chemistry and biochemistry and my

8
00:00:25,130 --> 00:00:22,650

talk today is titled frozen in time the

9
00:00:26,900 --> 00:00:25,140

history of proteins so I'll be talking

10
00:00:29,599 --> 00:00:26,910

about a really large molecular structure

11
00:00:32,150 --> 00:00:29,609

but I swear it pertains to the origin of

12
00:00:35,569 --> 00:00:32,160

life so just bear with the introduction

13
00:00:37,520 --> 00:00:35,579

I'll get to it eventually um so just a

14

00:00:40,430 --> 00:00:37,530

brief introduction about the central

15

00:00:41,780 --> 00:00:40,440

dogma of molecular biology I'm sure all

16

00:00:43,610 --> 00:00:41,790

of you guys are familiar with this but

17

00:00:50,360 --> 00:00:43,620

just to touch base with it for some of

18

00:00:53,090 --> 00:00:50,370

those geologists or some other people so

19

00:00:55,549 --> 00:00:53,100

here on the Left we have DNA and it's a

20

00:00:58,430 --> 00:00:55,559

nucleic acid polymer and on it

21

00:01:00,410 --> 00:00:58,440

we have encoded genes amongst other

22

00:01:02,510 --> 00:01:00,420

things but genes where we can talk about

23

00:01:05,630 --> 00:01:02,520

and over here in the blue we have an

24

00:01:08,060 --> 00:01:05,640

enzyme RNA polymerase which reads the

25

00:01:09,980 --> 00:01:08,070

DNA and looks at just the genes and

26

00:01:14,000 --> 00:01:09,990

copies those into RNA

27

00:01:16,249 --> 00:01:14,010

another nucleic acid polymer and so it's

28

00:01:19,130 --> 00:01:16,259

RNA then just has just the one gene on

29

00:01:22,070 --> 00:01:19,140

it and this gene on the mRNA goes to the

30

00:01:25,940 --> 00:01:22,080

ribosome and the ribosome then reads the

31

00:01:29,060 --> 00:01:25,950

mRNA it makes a protein and amino acid

32

00:01:31,910 --> 00:01:29,070

polymer and so I will be talking about

33

00:01:36,399 --> 00:01:31,920

the ribosome right here the key in it

34

00:01:39,170 --> 00:01:36,409

which reads RNA and makes protein so

35

00:01:41,210 --> 00:01:39,180

looking at the ribosome it has two

36

00:01:44,090 --> 00:01:41,220

subunits the small subunit and the large

37

00:01:45,620 --> 00:01:44,100

subunit the log the small subunit is all

38

00:01:49,280 --> 00:01:45,630

left and the large subunit is on the

39

00:01:52,130 --> 00:01:49,290

right RNA I have colored in pink and

40

00:01:54,560 --> 00:01:52,140

yellow and in blue are ribosomal

41

00:01:56,660 --> 00:01:54,570

proteins and then right there in the

42

00:01:59,480 --> 00:01:56,670

green at the very center of the large

43

00:02:02,030 --> 00:01:59,490

subunit is where peptide bond catalysis

44

00:02:04,399 --> 00:02:02,040

happens that is called the peptidyle

45

00:02:08,630 --> 00:02:04,409

transferase center and this particular

46

00:02:11,820 --> 00:02:08,640

transferase center is exclusively RNA so

47

00:02:15,520 --> 00:02:11,830

the ribosome is a ratchet design

48

00:02:18,580 --> 00:02:15,530

okay so like I said the ribosome reads

49

00:02:20,650 --> 00:02:18,590

RNA it makes protein or it reads nucleic

50

00:02:22,990 --> 00:02:20,660

acid polymers and it makes an amino acid

51
00:02:25,360 --> 00:02:23,000
polymer or another way of saying that is

52
00:02:28,960 --> 00:02:25,370
its input as information and its output

53
00:02:30,610 --> 00:02:28,970
is a function so in 2009 the Nobel Prize

54
00:02:32,830 --> 00:02:30,620
in Chemistry was awarded to thanking

55
00:02:34,720 --> 00:02:32,840
ramakrishnan commerce sites in a tea

56
00:02:37,240 --> 00:02:34,730
oneth for the studies of the structure

57
00:02:39,520 --> 00:02:37,250
and function of the ribosome and they

58
00:02:41,590 --> 00:02:39,530
made this really cool animation and it's

59
00:02:42,910 --> 00:02:41,600
very complex but is this is the

60
00:02:45,610 --> 00:02:42,920
introduction of the ribosome so I just

61
00:02:47,410 --> 00:02:45,620
want to show you that on the bottom here

62
00:02:50,440 --> 00:02:47,420
we have the small subunit that's in

63
00:02:52,480 --> 00:02:50,450

transparent pink on the top we have and

64

00:02:55,270 --> 00:02:52,490

this transparent yellow that would be

65

00:02:56,950 --> 00:02:55,280

the large subunit the mRNA is in blue

66

00:02:59,950 --> 00:02:56,960

and it's being read by the small subunit

67

00:03:01,750 --> 00:02:59,960

and these structures going in and out of

68

00:03:04,900 --> 00:03:01,760

here are T RNAs and they're bringing

69

00:03:06,400 --> 00:03:04,910

amino acids and these amino acids are

70

00:03:08,020 --> 00:03:06,410

brought to the peptidyl transferase

71

00:03:10,510 --> 00:03:08,030

center right in the middle of the large

72

00:03:12,700 --> 00:03:10,520

subunit and they get strung together to

73

00:03:19,570 --> 00:03:12,710

make a peptide that then emerges out the

74

00:03:23,080 --> 00:03:19,580

back of the ribosome so now to look at

75

00:03:24,030 --> 00:03:23,090

the sequence of the ribosome so I'm sure

76

00:03:26,530 --> 00:03:24,040

many of you are familiar with

77

00:03:28,240 --> 00:03:26,540

phylogenetic trees of life where we have

78

00:03:31,150 --> 00:03:28,250

the three domains of life bacteria

79

00:03:32,800 --> 00:03:31,160

archaea and Eukarya over here we have

80

00:03:35,680 --> 00:03:32,810

animals and their distant relatives a

81

00:03:37,600 --> 00:03:35,690

fungi and plants then we have our cans

82

00:03:39,760 --> 00:03:37,610

which are then you can barley distantly

83

00:03:40,960 --> 00:03:39,770

related to eukaryotes and then bacteria

84

00:03:44,260 --> 00:03:40,970

are all over here

85

00:03:47,170 --> 00:03:44,270

and so these phylogenetic trees are made

86

00:03:50,560 --> 00:03:47,180

by looking at the sequence of ribosomal

87

00:03:52,900 --> 00:03:50,570

genes so the ribosomal RNA and ribosomal

88

00:03:57,690 --> 00:03:52,910

proteins are encoded on the genome of

89

00:04:01,570 --> 00:03:57,700

all of every organism and so just the

90

00:04:04,320 --> 00:04:01,580

ribosomal RNA genes are just not made

91

00:04:06,370 --> 00:04:04,330

into protein they just stay as RNA but

92

00:04:07,840 --> 00:04:06,380

we can just be looking at the sequence

93

00:04:12,330 --> 00:04:07,850

and that's how we construct these

94

00:04:14,710 --> 00:04:12,340

phylogenetic trees so the ribosome has

95

00:04:18,340 --> 00:04:14,720

within it like the interrelatedness of

96

00:04:20,560 --> 00:04:18,350

all of life so looking at the ribosome

97

00:04:22,650 --> 00:04:20,570

from across the Tree of Life we can see

98

00:04:24,560 --> 00:04:22,660

that bacteria the large subunit has two

99

00:04:26,720 --> 00:04:24,570

ribosomal RNAs and

100

00:04:29,540 --> 00:04:26,730

33 ribosomal proteins where it's a small

101
00:04:32,330 --> 00:04:29,550
subunit has one ribosomal RNA and 22

102
00:04:34,520 --> 00:04:32,340
ribosomal proteins are kaons then have a

103
00:04:37,130 --> 00:04:34,530
few more ribosomal proteins and both

104
00:04:40,760 --> 00:04:37,140
both its large subunit and small subunit

105
00:04:42,530 --> 00:04:40,770
and then eukaryotes have one more one or

106
00:04:44,510 --> 00:04:42,540
two more ribosomal RNAs in its large

107
00:04:47,510 --> 00:04:44,520
subunit and a few more ribosomal

108
00:04:49,310 --> 00:04:47,520
proteins in both its l-s-u large living

109
00:04:53,060 --> 00:04:49,320
it and SS you small so you

110
00:04:55,070 --> 00:04:53,070
and so bacterias ribosomes are pretty

111
00:04:57,350 --> 00:04:55,080
small whereas our cans are a little bit

112
00:05:01,370 --> 00:04:57,360
bigger and eukaryotes have very large

113
00:05:03,380 --> 00:05:01,380

ribosomes so to conclude this

114

00:05:05,750 --> 00:05:03,390

introduction on the ribosome its

115

00:05:07,850 --> 00:05:05,760

sequence and structure ribosomes are

116

00:05:09,980 --> 00:05:07,860

made of both RNA and protein ribosomal

117

00:05:12,470 --> 00:05:09,990

RNA and protein sequences describe the

118

00:05:14,480 --> 00:05:12,480

interrelatedness of life and ribosomes

119

00:05:15,320 --> 00:05:14,490

size is highly correlated with species

120

00:05:18,350 --> 00:05:15,330

complexity

121

00:05:20,660 --> 00:05:18,360

here's another phylogenetic tree so on

122

00:05:23,780 --> 00:05:20,670

the top we have homo sapiens and below

123

00:05:26,260 --> 00:05:23,790

that shrimps so those are mammals and

124

00:05:27,850 --> 00:05:26,270

then farther down we have chicken and

125

00:05:31,790 --> 00:05:27,860

[Music]

126

00:05:35,840 --> 00:05:31,800

let's see mosquito right there so those

127

00:05:37,790 --> 00:05:35,850

are animals and then further down here

128

00:05:43,610 --> 00:05:37,800

we have single-celled eukaryotes like

129

00:05:44,060 --> 00:05:43,620

ester Vissi a fungus and what else we

130

00:05:46,550 --> 00:05:44,070

have here

131

00:05:49,490 --> 00:05:46,560

I think Plasmodium falciparum which is

132

00:05:51,020 --> 00:05:49,500

the parasite that causes malaria and

133

00:05:52,640 --> 00:05:51,030

then down here I just have archaea and

134

00:05:55,700 --> 00:05:52,650

bacteria because what I'm trying to show

135

00:05:58,190 --> 00:05:55,710

in this phylogenetic tree is the size of

136

00:06:00,560 --> 00:05:58,200

the ribosome so bacteria the ribosome is

137

00:06:02,750 --> 00:06:00,570

all about the same size but the sequence

138

00:06:04,190 --> 00:06:02,760

is very different and the same with our

139

00:06:06,140 --> 00:06:04,200

kaons their ribosomes a little bit

140

00:06:08,360 --> 00:06:06,150

larger than bacteria but their sequence

141

00:06:10,670 --> 00:06:08,370

is very different within the whole

142

00:06:12,530 --> 00:06:10,680

entire domain whereas eukaryotes there's

143

00:06:15,050 --> 00:06:12,540

like a really big difference in their

144

00:06:17,450 --> 00:06:15,060

size of the ribosomes and so that's what

145

00:06:20,060 --> 00:06:17,460

I have over here and these circles so

146

00:06:22,640 --> 00:06:20,070

bacteria the ribosome is small and

147

00:06:24,890 --> 00:06:22,650

Arcadians is also fairly small then the

148

00:06:26,990 --> 00:06:24,900

single cellular eukaryotes have you know

149

00:06:28,460 --> 00:06:27,000

medium sized ribosomes these animals

150

00:06:30,650 --> 00:06:28,470

have even larger ribosomes

151

00:06:32,870 --> 00:06:30,660

then mammals have the largest ribosomes

152

00:06:34,880 --> 00:06:32,880

that we see this so this is very highly

153

00:06:37,640 --> 00:06:34,890

correlated to how complex a species

154

00:06:38,430 --> 00:06:37,650

species is people used to think oh well

155

00:06:41,610 --> 00:06:38,440

maybe like the length

156

00:06:43,140 --> 00:06:41,620

you know the DNA of an organ organism

157

00:06:44,640 --> 00:06:43,150

would tell you how complex that species

158

00:06:46,290 --> 00:06:44,650

is but there is some really big

159

00:06:49,170 --> 00:06:46,300

anomalies that really just throw it off

160

00:06:52,320 --> 00:06:49,180

for instance the marble bunk fish has

161

00:06:52,620 --> 00:06:52,330

its genome was 120 billion nucleotides

162

00:06:56,250 --> 00:06:52,630

long

163

00:06:57,510 --> 00:06:56,260

whereas humans it's 6 billion so really

164

00:07:01,710 --> 00:06:57,520

that's not like a really good indication

165

00:07:03,960 --> 00:07:01,720

of how complex this species is okay so

166

00:07:05,130 --> 00:07:03,970

what am i doing in my project so I'm

167

00:07:08,190 --> 00:07:05,140

looking at the ribosome and I'm

168

00:07:09,990 --> 00:07:08,200

constructing a data set of its structure

169

00:07:12,060 --> 00:07:10,000

and the reason I'm doing that is because

170

00:07:14,190 --> 00:07:12,070

the ribosome is the origin of life it

171

00:07:17,220 --> 00:07:14,200

contains within its structuring sequence

172

00:07:19,350 --> 00:07:17,230

the history of protein folding and I

173

00:07:21,150 --> 00:07:19,360

believe that by mining the structure of

174

00:07:25,320 --> 00:07:21,160

it and the sequence set I can unravel

175

00:07:27,210 --> 00:07:25,330

how life has evolved so like I said I'm

176

00:07:30,180 --> 00:07:27,220

looking at structures of ribosomes and

177

00:07:32,070 --> 00:07:30,190

in 2009 the Nobel Prize was awarded for

178

00:07:34,050 --> 00:07:32,080

the structure for crystallize in

179

00:07:37,680 --> 00:07:34,060

ribosome and since then we have quite a

180

00:07:40,110 --> 00:07:37,690

few structures so we have bacteria E

181

00:07:42,390 --> 00:07:40,120

coli and thermus thermophilus on the

182

00:07:44,610 --> 00:07:42,400

Left we have the large subunit and on

183

00:07:46,050 --> 00:07:44,620

the right we have the small subunit we

184

00:07:48,390 --> 00:07:46,060

have the Arcadians halo rocky lamar

185

00:07:50,850 --> 00:07:48,400

smart i and pirate caucus furiosa s'

186

00:07:52,409 --> 00:07:50,860

halo arkla we only have a large subunit

187

00:07:55,260 --> 00:07:52,419

not the small subunit we can get

188

00:07:57,780 --> 00:07:55,270

crystallized we also have the protozoan

189

00:08:01,260 --> 00:07:57,790

a single cellular eukaryote again we

190

00:08:03,450 --> 00:08:01,270

only have the large subunit we have the

191

00:08:06,690 --> 00:08:03,460

single cellular fungi *Saccharomyces*

192

00:08:09,150 --> 00:08:06,700

cerevisiae the animals *Drosophila* like

193

00:08:11,400 --> 00:08:09,160

gaster and *Homo sapiens* fruit fly in

194

00:08:13,650 --> 00:08:11,410

humans and we also have the human

195

00:08:17,730 --> 00:08:13,660

protozoan parasites that cause malaria

196

00:08:20,370 --> 00:08:17,740

and African sleeping sickness and so

197

00:08:23,840 --> 00:08:20,380

what happens when we align all these on

198

00:08:26,030 --> 00:08:23,850

to each other well as you can see

199

00:08:27,960 --> 00:08:26,040

bacteria and archaea have pretty small

200

00:08:29,730 --> 00:08:27,970

ribosomes but then when you start

201
00:08:34,110 --> 00:08:29,740
putting on eukaryotes it gets a little

202
00:08:35,880 --> 00:08:34,120
bigger and larger and larger and then I

203
00:08:40,649 --> 00:08:35,890
have these human protozoan parasites

204
00:08:42,690 --> 00:08:40,659
over here for a discussion later on that

205
00:08:44,400 --> 00:08:42,700
maybe you have but for the focus of this

206
00:08:47,130 --> 00:08:44,410
talk I'm not going to talk too much

207
00:08:48,840 --> 00:08:47,140
about them but as you can see these

208
00:08:52,110 --> 00:08:48,850
structures are quite a bit larger than

209
00:08:54,629 --> 00:08:52,120
the E coli and other bacterial species

210
00:08:56,819 --> 00:08:54,639
and when we get is we get this big glob

211
00:08:59,009 --> 00:08:56,829
of like hundreds of millions of atoms

212
00:09:00,900 --> 00:08:59,019
and we really can't make sense of it

213
00:09:02,489 --> 00:09:00,910

when you look at that but when you look

214

00:09:05,730 --> 00:09:02,499

deep within its structure you can see

215

00:09:07,379 --> 00:09:05,740

that the structures of the RNA and the

216

00:09:11,400 --> 00:09:07,389

protein that make up the ribosome and

217

00:09:13,739 --> 00:09:11,410

the center of it are miraculously just

218

00:09:16,259 --> 00:09:13,749

similar their sequence is different but

219

00:09:18,989 --> 00:09:16,269

their structure is just crazy the same

220

00:09:20,879 --> 00:09:18,999

going from the center all the way out to

221

00:09:23,009 --> 00:09:20,889

near the surface of it it's only the

222

00:09:26,429 --> 00:09:23,019

surface of the ribosomal ribosomal

223

00:09:28,860 --> 00:09:26,439

structures that we see any variation so

224

00:09:30,929 --> 00:09:28,870

the center of it is just a common core

225

00:09:33,540 --> 00:09:30,939

of structure that just doesn't divulge

226

00:09:35,819 --> 00:09:33,550

from any species that we know of and

227

00:09:42,739 --> 00:09:35,829

it's when we look at the surface then

228

00:09:48,629 --> 00:09:46,499

so looking at that one helix from the

229

00:09:50,160 --> 00:09:48,639

last slide this is helix 25 and I'll

230

00:09:51,299 --> 00:09:50,170

just use this as an example to show you

231

00:09:55,319 --> 00:09:51,309

how ripe Izumo

232

00:09:58,410 --> 00:09:55,329

RNA has evolved so this helix 25 of

233

00:10:00,869 --> 00:09:58,420

bacteria we have our k ends protozoans

234

00:10:03,449 --> 00:10:00,879

fungi animals and our parasitic

235

00:10:06,210 --> 00:10:03,459

protozoan x' and then when we align

236

00:10:08,249 --> 00:10:06,220

these onto each other and this is

237

00:10:09,840 --> 00:10:08,259

aligning the ribosomes I'm just I'm just

238

00:10:11,460 --> 00:10:09,850

cutting away the rest of the ribosome

239

00:10:13,799 --> 00:10:11,470

and only looking at this helix that's on

240

00:10:16,889 --> 00:10:13,809

the surface we can see that the

241

00:10:18,720 --> 00:10:16,899

structure is also very conserved at that

242

00:10:22,049 --> 00:10:18,730

very first helix but there's HeLa C's

243

00:10:23,549 --> 00:10:22,059

budding off of that and as you can see

244

00:10:25,980 --> 00:10:23,559

going through more and more complex

245

00:10:28,319 --> 00:10:25,990

species you can see that this helix is

246

00:10:30,780 --> 00:10:28,329

just has more and more HeLa C's just

247

00:10:33,600 --> 00:10:30,790

budding off of it and so we believe that

248

00:10:35,429 --> 00:10:33,610

this is how the ribosome has a ribosome

249

00:10:38,100 --> 00:10:35,439

while RNA has evolved is that just HeLa

250

00:10:43,009 --> 00:10:38,110

sees just butted off from other HeLa C's

251

00:10:45,629 --> 00:10:43,019

and so this kind of so to wrap that up

252

00:10:49,470 --> 00:10:45,639

the ribosome has a common core and it's

253

00:10:51,119 --> 00:10:49,480

made of RNA and protein and the RNA is

254

00:10:53,610 --> 00:10:51,129

really conserved in structure but not

255

00:10:55,259 --> 00:10:53,620

sequence and these sites of ribosomal

256

00:10:57,329 --> 00:10:55,269

RNA evolution are referred to as

257

00:10:59,579 --> 00:10:57,339

expansion segments and they but offer

258

00:11:01,499 --> 00:10:59,589

the common core of the ribosome and they

259

00:11:05,879 --> 00:11:01,509

do not perturb the common core structure

260

00:11:07,559 --> 00:11:05,889

and the sounds to make an analogy

261

00:11:09,509 --> 00:11:07,569

it's kind of like a tree when you look

262

00:11:11,280 --> 00:11:09,519

at a tree you can tell the newest

263

00:11:14,009 --> 00:11:11,290

sections of it are the leaves whereas

264

00:11:15,629 --> 00:11:14,019

the oldest section is the trunk and so

265

00:11:18,419 --> 00:11:15,639

we're looking at these structures and

266

00:11:20,220 --> 00:11:18,429

we're seeing these eukaryotic expansion

267

00:11:22,559 --> 00:11:20,230

segments those are like the leaves and

268

00:11:24,329 --> 00:11:22,569

the outermost branches and when we take

269

00:11:26,639 --> 00:11:24,339

away these leaves and take away these

270

00:11:29,579 --> 00:11:26,649

branches we can just go deeper and

271

00:11:31,919 --> 00:11:29,589

deeper into the past the ribosome and we

272

00:11:34,650 --> 00:11:31,929

eventually get to just one helix and

273

00:11:37,139 --> 00:11:34,660

that would just be the trunk of the tree

274

00:11:40,829 --> 00:11:37,149

this one helix is the peptidyl

275

00:11:43,470 --> 00:11:40,839

transferase center so it's where peptide

276

00:11:48,389 --> 00:11:43,480

bonds are made and so we can look at how

277

00:11:50,609 --> 00:11:48,399

the ribosomal RNA has evolved okay so

278

00:11:52,559 --> 00:11:50,619

that's the RNA of the ribosome well what

279

00:11:55,199 --> 00:11:52,569

about the protein well the protein is

280

00:11:57,749 --> 00:11:55,209

also very similar in some regards so

281

00:12:01,859 --> 00:11:57,759

this is universal protein number four of

282

00:12:04,429 --> 00:12:01,869

the large subunit ul4 and so for those

283

00:12:07,229 --> 00:12:04,439

of you who are familiar with proteins or

284

00:12:09,150 --> 00:12:07,239

maybe not I guess up here we have a

285

00:12:11,069 --> 00:12:09,160

globular domain and that looks a lot

286

00:12:13,229 --> 00:12:11,079

like the proteins we see in all white

287

00:12:14,939 --> 00:12:13,239

today but these ribosomal proteins are

288

00:12:17,129 --> 00:12:14,949

very weird looking they have these

289

00:12:18,989 --> 00:12:17,139

really strange like tails that just go

290

00:12:20,489 --> 00:12:18,999

into the ribosome and they don't really

291

00:12:23,309 --> 00:12:20,499

look like other proteins that we know

292

00:12:24,509 --> 00:12:23,319

and so a lot of people just don't really

293

00:12:26,159 --> 00:12:24,519

look at them and they throw them out of

294

00:12:27,329 --> 00:12:26,169

their analysis and stuff like that but

295

00:12:30,689 --> 00:12:27,339

I'm looking at them because I think

296

00:12:33,329 --> 00:12:30,699

they're interesting and so this is ul4

297

00:12:35,039 --> 00:12:33,339

from a bacterial species and that's uo4

298

00:12:38,189 --> 00:12:35,049

from an our canyon and that's you all

299

00:12:40,079 --> 00:12:38,199

for from eukaryote and then when we look

300

00:12:42,840 --> 00:12:40,089

at them within these aligned crystal

301

00:12:47,129 --> 00:12:42,850

structures we see that they line right

302

00:12:49,739 --> 00:12:47,139

on top of each other and they all have

303

00:12:51,569 --> 00:12:49,749

this tail that goes deep into the

304

00:12:53,789 --> 00:12:51,579

ribosome and they have this globular

305

00:12:56,309 --> 00:12:53,799

domain that hasn't really changed in

306

00:12:57,929 --> 00:12:56,319

structure either but then arcane's and

307

00:13:00,239 --> 00:12:57,939

eukaryotes have this little loop here

308

00:13:01,919 --> 00:13:00,249

and then eukaryotes have a little bit

309

00:13:04,559 --> 00:13:01,929

more over here and there's large alpha

310

00:13:07,049 --> 00:13:04,569

helix that buds off there and this goes

311

00:13:09,539 --> 00:13:07,059

on to the surface of the ribosome where

312

00:13:12,269 --> 00:13:09,549

the ribosome has been evolving but all

313

00:13:14,879 --> 00:13:12,279

has this common core right here and this

314

00:13:17,340 --> 00:13:14,889

common core goes deep into the ribosome

315

00:13:18,790 --> 00:13:17,350

and it goes right next to the peptidyl

316

00:13:22,569 --> 00:13:18,800

transferase center

317

00:13:24,369 --> 00:13:22,579

right here and so it's almost like if

318

00:13:27,609 --> 00:13:24,379

you are looking make a reaction

319

00:13:29,829 --> 00:13:27,619

coordinate of this that you know here

320

00:13:32,019 --> 00:13:29,839

would be like the oldest peptides and

321

00:13:34,960 --> 00:13:32,029

here would be the newer ones like I

322

00:13:37,210 --> 00:13:34,970

showed in that last figure the very

323

00:13:39,970 --> 00:13:37,220

center of the ribosome is the oldest

324

00:13:41,859 --> 00:13:39,980

part it's the oldest ribosomal RNA

325

00:13:44,470 --> 00:13:41,869

whereas the outside is the newest and

326

00:13:48,069 --> 00:13:44,480

when you look at these proteins it's the

327

00:13:50,109 --> 00:13:48,079

same the proteins that protein the amino

328

00:13:52,269 --> 00:13:50,119

acids and peptides are closest to the

329

00:13:54,489 --> 00:13:52,279

petiole transfer a center or the oldest

330

00:13:57,309 --> 00:13:54,499

ones where the ones on the exterior of

331

00:13:59,859 --> 00:13:57,319

the ribosome are the newest ones and so

332

00:14:04,600 --> 00:13:59,869

looking at this we can see how proteins

333

00:14:06,910 --> 00:14:04,610

have evolved since the origin of life so

334

00:14:10,119 --> 00:14:06,920

I made this like protein folding peak

335

00:14:14,530 --> 00:14:10,129

and so basically what it was showing is

336

00:14:15,939 --> 00:14:14,540

that close to the center of the peptidyl

337

00:14:18,609 --> 00:14:15,949

transferase center we get these

338

00:14:20,319 --> 00:14:18,619

unstructured peptides but then as you

339

00:14:22,030 --> 00:14:20,329

move farther and farther out from it you

340

00:14:24,699 --> 00:14:22,040

start saying okay well there's like some

341

00:14:27,160 --> 00:14:24,709

you know beta sheets or beta hairpins

342

00:14:29,230 --> 00:14:27,170

and then you move farther out then you

343

00:14:30,819 --> 00:14:29,240

start seeing like beta barrels and as

344

00:14:32,699 --> 00:14:30,829

you go into the surface of the ribosome

345

00:14:35,980 --> 00:14:32,709

then you see is really complex looking

346

00:14:39,549 --> 00:14:35,990

proteins that we see in all of cellular

347

00:14:41,379 --> 00:14:39,559

life and so it it's a reaction

348

00:14:43,929 --> 00:14:41,389

coordinate of how proteins have evolved

349

00:14:46,509 --> 00:14:43,939

since the origin of life and so that's

350

00:14:47,859 --> 00:14:46,519

what I have been looking at and so to

351

00:14:50,710 --> 00:14:47,869

conclude these ribosomal protein

352

00:14:53,289 --> 00:14:50,720

evolution you can see that ribosomal

353

00:14:54,910 --> 00:14:53,299

proteins have also have expansions and

354

00:14:57,519 --> 00:14:54,920

this is very similar to the ribosomal

355

00:14:59,109 --> 00:14:57,529

RNA however these ribosomal protein

356

00:15:00,970 --> 00:14:59,119

structural differences are much more

357

00:15:02,559 --> 00:15:00,980

pronounced we just know a lot more about

358

00:15:04,749 --> 00:15:02,569

proteins proteins have been struck

359

00:15:06,639 --> 00:15:04,759

studied for a while whereas RNA the

360

00:15:08,319 --> 00:15:06,649

structures and everything haven't been

361

00:15:11,199 --> 00:15:08,329

very interesting to a lot of people for

362

00:15:13,359 --> 00:15:11,209

much time and these that these ribosomal

363

00:15:16,269 --> 00:15:13,369

proteins there's a common core to it and

364

00:15:18,850 --> 00:15:16,279

the common core just like RNA no protein

365

00:15:20,530 --> 00:15:18,860

is all exactly the same the sequence

366

00:15:23,650 --> 00:15:20,540

might differ but the structure is the

367

00:15:25,569 --> 00:15:23,660

same and so we believe that ribosomal

368

00:15:28,030 --> 00:15:25,579

evolution happens in both RNA and

369

00:15:29,769 --> 00:15:28,040

protein and the RNA and protein have

370

00:15:30,810 --> 00:15:29,779

evolved together in a mutualistic

371

00:15:34,530 --> 00:15:30,820

fashion

372

00:15:36,390 --> 00:15:34,540

and then so to conclude I just like to

373

00:15:38,480 --> 00:15:36,400

do show this one slide from a project

374

00:15:41,160 --> 00:15:38,490

I'm working on because we're all

375

00:15:42,540 --> 00:15:41,170

astrobiologists here and it's a little

376

00:15:44,340 --> 00:15:42,550

bit talking about metals and I just

377

00:15:46,560 --> 00:15:44,350

maybe want to show some people so that

378

00:15:51,090 --> 00:15:46,570

maybe they can give me some input later

379

00:15:53,610 --> 00:15:51,100

on so a lot of these ribosomal proteins

380

00:15:55,560 --> 00:15:53,620

are binding zinc and zinc is very

381

00:15:57,660 --> 00:15:55,570

important for a lot of proteins in

382

00:16:00,570 --> 00:15:57,670

looking at any organism that we see

383

00:16:03,320 --> 00:16:00,580

around today you know four to 10% of

384

00:16:05,400 --> 00:16:03,330

their proteins need zinc to function and

385

00:16:07,650 --> 00:16:05,410

zinc is found in many of these ribosomal

386

00:16:09,420 --> 00:16:07,660

proteins and these proteins that it's in

387

00:16:12,150 --> 00:16:09,430

are usually very simple-looking

388

00:16:13,500 --> 00:16:12,160

so here's one of them and so there's the

389

00:16:16,250 --> 00:16:13,510

zinc that's coordinating with four

390

00:16:19,560 --> 00:16:16,260

cystines and it's just made of like to

391

00:16:23,160 --> 00:16:19,570

beta hairpins and so and I'm looking at

392

00:16:24,690 --> 00:16:23,170

this it's like it's pretty these

393

00:16:27,360 --> 00:16:24,700

proteins are usually found pretty deep

394

00:16:28,760 --> 00:16:27,370

within the ribosome and I believe that

395

00:16:33,180 --> 00:16:28,770

these could be some very ancient

396

00:16:35,160 --> 00:16:33,190

peptides or proteins and I'm looking at

397

00:16:37,350 --> 00:16:35,170

the literature and you know there's all

398

00:16:39,240 --> 00:16:37,360

sorts of you know hypothesis of how like

399

00:16:41,460 --> 00:16:39,250

the world came about and everything and

400

00:16:43,110 --> 00:16:41,470

what kind of concentrations models were

401
00:16:45,600 --> 00:16:43,120
at and everything and you know there's

402
00:16:48,290 --> 00:16:45,610
some debate about zinc and whatnot but

403
00:16:50,760 --> 00:16:48,300
our lab has also shown in the past that

404
00:16:53,010 --> 00:16:50,770
you know the ribosome needs magnesium

405
00:16:55,200 --> 00:16:53,020
cations to stabilize its tertiary

406
00:16:58,290 --> 00:16:55,210
interactions between the RNA HeLa C's

407
00:16:59,640 --> 00:16:58,300
because they're very negative and there

408
00:17:02,760 --> 00:16:59,650
have been studies in our lab and other

409
00:17:04,800 --> 00:17:02,770
labs that have shown that you know RNA

410
00:17:08,430 --> 00:17:04,810
doesn't necessarily need magnesium but

411
00:17:10,200 --> 00:17:08,440
it could be using reduced iron and I

412
00:17:13,410 --> 00:17:10,210
found some literature saying that these

413
00:17:15,570 --> 00:17:13,420

peptides these zinc ribbons that these

414

00:17:20,400 --> 00:17:15,580

original proteins have that they can

415

00:17:21,870 --> 00:17:20,410

also bind iron instead of zinc and so

416

00:17:24,300 --> 00:17:21,880

hopefully tomorrow I can like look at

417

00:17:26,070 --> 00:17:24,310

some posters and maybe some talks and

418

00:17:29,160 --> 00:17:26,080

hopefully some we talking about zinc or

419

00:17:30,630 --> 00:17:29,170

iron or something like that but so I

420

00:17:32,430 --> 00:17:30,640

guess that will be conclude it um

421

00:17:35,220 --> 00:17:32,440

there's really not much to put in this

422

00:17:43,230 --> 00:17:35,230

presentation summary but this is our lab

423

00:17:50,430 --> 00:17:45,700

all right we got time for one quick

424

00:17:57,970 --> 00:17:53,770

um yeah I'm so my question if you can go

425

00:18:02,890 --> 00:17:57,980

back to slides sorry uh what are the

426

00:18:05,020 --> 00:18:02,900

arrows on the protein fold okay so it's

427

00:18:09,100 --> 00:18:05,030

just showing like so I go the way you

428

00:18:11,140 --> 00:18:09,110

like depict the protein is these arrows

429

00:18:12,880 --> 00:18:11,150

are just beta sheets and so that's just

430

00:18:15,760 --> 00:18:12,890

how it's like structured and so right

431

00:18:18,340 --> 00:18:15,770

between that like this is a simplistic

432

00:18:20,320 --> 00:18:18,350

diagram you have your amino acids and

433

00:18:22,420 --> 00:18:20,330

they're aligning and hydrogen bonding to

434

00:18:24,720 --> 00:18:22,430

each other and then the arrow is just

435

00:18:26,620 --> 00:18:24,730

pointing to like okay so this is the

436

00:18:30,370 --> 00:18:26,630

n-terminal and then this is the

437

00:18:31,810 --> 00:18:30,380

c-terminus of the protein so it's just a

438

00:18:35,320 --> 00:18:31,820

simple way of drawing a protein

439

00:18:35,860 --> 00:18:35,330

structure mm-hmm alright thank you very

440

00:18:37,180 --> 00:18:35,870

much

441

00:18:38,900 --> 00:18:37,190

Thanks