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RUTGERS

*The Environment as a Driving Force  
in the Coevolution of Ferredoxin Termini*

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Developed under the guidance of Dr. Vikas Nanda

1  
00:00:03,350 --> 00:00:02,070  
hello my name is julianna giacomo with

2  
00:00:04,630 --> 00:00:03,360  
rutgers university

3  
00:00:06,230 --> 00:00:04,640  
and i'm going to be discussing the

4  
00:00:08,470 --> 00:00:06,240  
environment as a driving force in the

5  
00:00:11,270 --> 00:00:08,480  
co-evolution of paradox and termini

6  
00:00:13,030 --> 00:00:11,280  
very briefly uh the study begins to

7  
00:00:14,709 --> 00:00:13,040  
specify the evolutionary factors of

8  
00:00:16,630 --> 00:00:14,719  
life's earliest proteins capable of

9  
00:00:17,590 --> 00:00:16,640  
redox reactions which are octa

10  
00:00:19,109 --> 00:00:17,600  
reductases

11  
00:00:20,630 --> 00:00:19,119  
these proteins are so diverse it has

12  
00:00:21,750 --> 00:00:20,640  
only reasonably been determined that

13  
00:00:23,910 --> 00:00:21,760

they are polyphyletic

14

00:00:26,310 --> 00:00:23,920  
where 50 of all metal containing

15

00:00:28,710 --> 00:00:26,320  
osteoductases have a single origin

16

00:00:30,150 --> 00:00:28,720  
from a protein fold represented by

17

00:00:31,830 --> 00:00:30,160  
accident ferridoxins

18

00:00:33,750 --> 00:00:31,840  
therefore the study of the origin of

19

00:00:36,870 --> 00:00:33,760  
ferridoxin serves as a proxy

20

00:00:38,950 --> 00:00:36,880  
for the origin of oxo reductases in

21

00:00:39,910 --> 00:00:38,960  
order to address this we split paradox

22

00:00:41,990 --> 00:00:39,920  
and sequences

23

00:00:43,830 --> 00:00:42,000  
from methanic archaea in half and

24

00:00:46,310 --> 00:00:43,840  
created phylogenetic trees overlaid with

25

00:00:47,990 --> 00:00:46,320  
optimal growth temperature and ph

26

00:00:49,590 --> 00:00:48,000

the real takeaway from this study is

27

00:00:51,189 --> 00:00:49,600

that each half of ferredoxin likely

28

00:00:53,110 --> 00:00:51,199

co-evolved independently

29

00:00:54,709 --> 00:00:53,120

where temperature was one of the driving

30

00:00:56,950 --> 00:00:54,719

factors for residue change in each

31

00:00:58,790 --> 00:00:56,960

terminus and ph was not

32

00:01:01,029 --> 00:00:58,800

and these factors can therefore be