

✕   You're being mentored by **iHiD** on **Asteroid Exploration Co.**

End session **F2**

 **Discussion** 2

 Scratchpad

Guidance

 **Iteration 12** latest 
Submitted via CLI 2 mins ago

● PASSED

ANALYSING

Compare

hamming.rb

hammingTests.rb

[readme.md](#)

 OUTDATED

```

1  # Hamming, determine the divergence of two DNA strands
2  module Hamming
3    def self.compute(strand_a, strand_b)
4      raise ArgumentError if strand_a.length != strand_b.length
5
6      strand_a.each_char.zip(strand_b.each_char).reduce(0) do |distance, pair|
7        + a, b = pair
8          distance += 1 if a != b
9          distance
10     end
11   end
12 end
13

```

Who's mentoring you

Jeremy Walker  

@iHiD

English, Spanish, French He / Him / His

+ Add to favorites

53 past discussions 

Track goals >

 **Iteration 3** was submitted  5mo ago

 WDNeumann received auto feedback

 **ErikSchierboom** 10d ago

Is there a ruby-er way to zip the two lists?
For example, this is another paragraph.

 iHiD started mentoring you

iHiD
Nope, that's a good way to do it.

ErikSchierboom 5m ago
Righty ho then. Thanks a lot!

Add a comment

H B / < > 1 2 3

Remember! Be polite and palatable. You can read our [comms guide here](#).