

Introduction to Bioinformatics course: IBT

Introduction Part 6 Vula forums vs. Vula chatroom

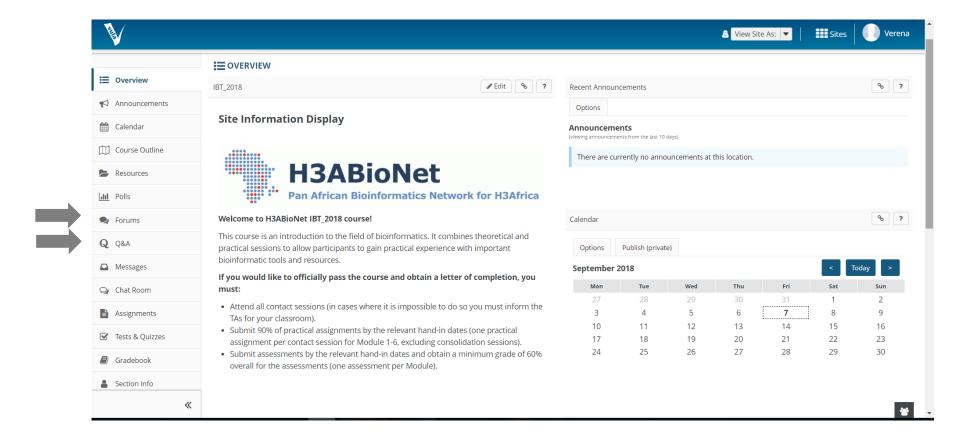






Forums vs. Chatroom









Forums vs. Chatroom



Forums

- Asking questions that you would like to easily find the answer to later
- Questions and answers or discussions that you think others would benefit from

Chatroom

for 'chat' J







Forums



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Module 2: Genomics New Topic | Forum Settings | More ▼

    session 1 0 unread of 0 messages Topic Settings | More ▼

    session 2 7 unread of 30 messages Topic Settings | More ▼
 Session 3 10 unread of 10 messages Topic Settings | More →
  Session 4 18 unread of 21 messages Topic Settings | More ▼
Module 3: Linux New Topic | Forum Settings | More ▼

    session 1 0 unread of 2 messages Topic Settings | More ▼
  Session 2 0 unread of 1 message Topic Settings | More ▼
  Session 3 0 unread of 11 messages Topic Settings | More ▼
Module 4: Sequence Alignment Theory and Application New Topic | Forum Settings | More ▼

    session 1 5 unread of 26 messages Topic Settings | More ▼

    session 2 5 unread of 9 messages Topic Settings | More ▼
  Session 3 5 unread of 7 messages Topic Settings | More ▼
Module 5: Multiple Sequence Alignment New Topic | Forum Settings | More ▼
  Session 1 7 unread of 10 messages Topic Settings | More ▼
  Session 2 0 unread of 3 messages Topic Settings | More ▼
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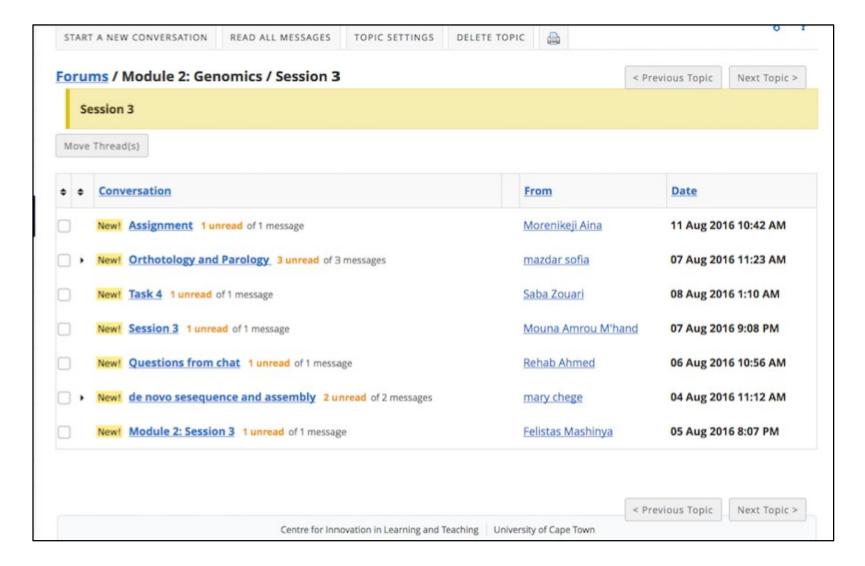






Forums





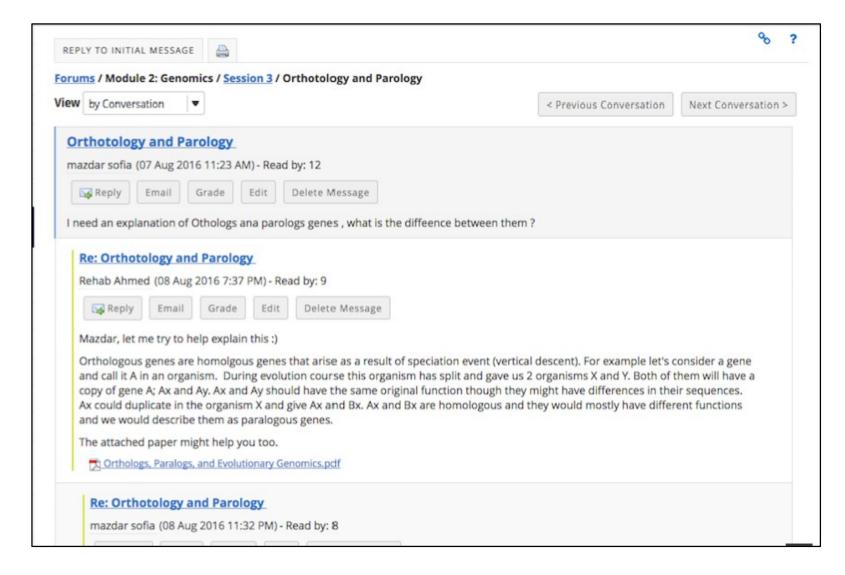






Forums











Chatroom



thanks, Rehab! :)

Meriem FASSATOUI @ (05-Oct-2016 11:20 SAST) @Dr.Jean i didn't ask about the assignment of today, i asked about the assignment of last week session 2

Jean-Baka Domelevo Entfellner 🝵 (05-Oct-2016 11:24 SAST) Yes, I figured it out, then the answer posted by Rehab just before you asked your question was relevant to you.

Meriem FASSATOUI @ (05-Oct-2016 11:26 SAST) thanks for the response

Rehab Ahmed (05-Oct-2016 13:16 SAST) Meriem, the matrix is provided for GTR and we also did not find matrix for HKY85

Adil EL HAMOUCHI 🏫 (05-Oct-2016 17:23 SAST) bonjour Jean Baka

can you explain more how we can choose the best method for the construction of phylogenetic tree ...? thank you

Dankwa Bismark (05-Oct-2016 22:44 SAST) my facebook name(Ing Bismark Bombelli)...send a request if you want to stay in touch

Donwilliams Otieno (06-Oct-2016 11:33 SAST) Hi Jean? What are the best bootstrap values to report for academic work? I am currently inferring phylogenies using maximum likelihood methods (PhyML and RAXML programs).

Donwilliams Otieno (06-Oct-2016 11:48 SAST) The is an ongoing debate on the bootstrap values, with some experts saying that values above 50% are good while others have stuck to atleast 70%. These, they say, is because of the random nature of the bootstrapping process. I don't know which values to stick to, kindly assist.

Rehab Ahmed

 (06-Oct-2016 11:52 SAST) Donwilliams, will post your interesting question in Questions and discussion

Rehab Ahmed (06-Oct-2016 11:55 SAST) You did it already:) thank you

Kim Gurwitz (06-Oct-2016 11:57 SAST) Hi All - the practical assignment and dataset for today's session (Module 7 session 4) has been updated - now under the same assignment in 'Assignments'

Adil EL HAMOUCHI 🝵 (06-Oct-2016 12:54 SAST) hello jean bakacan you please recommend us a free offline software to do the practical session

Donwilliams Otieno ∰ (06-Oct-2016 12:56 SAST) Thanks Rehab

Jean-Baka Domelevo Entfellner (06-Oct-2016 13:04 SAST) Dear Adil, I would recommend PhyML (http://www.atgc-montpellier.fr/phyml/) or RAxML (http://sco.h-its.org/exelixis/software.html), but RAxML is sometimes too customizable a tool, especially for beginners, although it's arguably the gold standard of ML software currently out there.

Cassandra Soo

(06-Oct-2016 14:34 SAST) Hi there, does anyone know how to do Task 3 from yesterday's practical (practical 3)?

Adil EL HAMOUCHI

(06-Oct-2016 14:57 SAST) big thanks JB

Kafayat Fakoya (07-Oct-2016 06:54 SAST) | submitted my module 7 session 1 assignment exactly 22.59 pm on the 5 th October, 2016 and made a resubmission 01:03 am 6th October withun he stipulated accepted period for resubmission. I was surprised to see late written against this submission. Was there anything I possibly did wrong or did not doas required to make a resubmission?

Kim Gurwitz (07-Oct-2016 10:19 SAST) Hi Kafayat, I am not sure why Vula marked your resubmitted assignment at late - as you correctly noted, resubmissions are allowed a few hours after the due date and time, just in case there was an error in the uploading. Not to worry though, your assignment submission has been captured. We do not take note of whether assignments are 'late' or not. 'Late' just indicates that the assignment was submitted after the due date and time, within a specified window (we allow a few hours 'grace period' in case participants get confused with the timezone conversions etc). If one tries to submit an assignment after this period, however, one will physically not be able to upload - the system will not allow it.

Ocung Guido 😭 (07-Oct-2016 11:01 SAST) Hi Kim. What next, will our vula accounts be pulled down after the IBT 2016 course?

Kim Gurwitz (07-Oct-2016 11:22 SAST) Hi Ocung, We are waiting for a response from the Vula team - I hope that we will be able to keep the site open. Will keep you posted. We will send out a Vula announcement soonest with 'next steps'

Natalie Smyth 🏫 (07-Oct-2016 14:34 SAST) Hi there, does anyone know how to do Task 3 from yesterday's practical (practical 3)?

Kafayat Fakoya et (08-Oct-2016 22:56 SAST) Thanks for the reassurance, Kim.







Next



20 Min. BREAK

THEN Watch video labeled: Part 7



