

**Demographic:**

1. What is your role?
2. How long have you been working your role?
3. Have you created a workflow for a bioinformatics process?
  - 3.1. If YES, briefly describe what did you do? (Language, software)

**Interaction with RE artefact:**

**(We are going to provide them with the RE template and a short explanation)**

4. Do you understand this?
  - 4.1 If NOT, Is there anything that you can't understand?
5. Could this be used? Why, why not?
  - 5.1 If YES, who and how?
6. Is this document missing anything?
  - 6.1 If YES, what is it missing?
7. Did you identify any field that is unnecessary or that you will never need or use?

**Interaction with Examples (1 and 2) and concrete syntax:**

**(We are going to provide them with both examples numbered and the concrete syntax list).**

*(Give the concrete syntax)*

8. Are these concepts useful?
9. Are any of this redundant?
10. Are we missing anything?

*(Explain the two different ways)*

11. Which ... do you like better? Why?
  - 11.1. Loop (1a or 2a)
  - 11.2. Thresholds (1a or 2a)
  - 11.3. Input/output (1b or 2b)
  - 11.4. Datastore (1b or 2b)
  - 11.5. Tools (1c or 2c)
  - 11.6. Diagrams separators/connectors (1b and 1c or 2b and 2c)

12. Did you think you can draw and use a model like this in your work? Why, why not? Would anyone else be able to use that? Who?

13. Would the template be helpful to draw the diagram or not?