

(03)

- @ I can give my input as computer engineer in personalized drug design.

current computer-Aided drug design aims to publish all the latest developments in drug design based on computational techniques. The field of computer-aided drug design has had extensive impact in the area of drug design.

computer-aided drug design could benefit from a greater understanding of how errors arise and propagate in biomolecular modeling.

Yes, I think we need a protocol to monitor transgenic animal trade.

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There some protocols are mentioned below; process to make transgenic animals

① Isolate DNA.

② Cut the DNA out of the chromosome.

③ make millions of copies of the DNA.

④ Put the DNA into a vector to carry to the organism.

⑤ Put the DNA into the host cell

and turn on it.

And we have to think about environment while releasing transgenic crops into the environment may have direct effects including: gene transfer

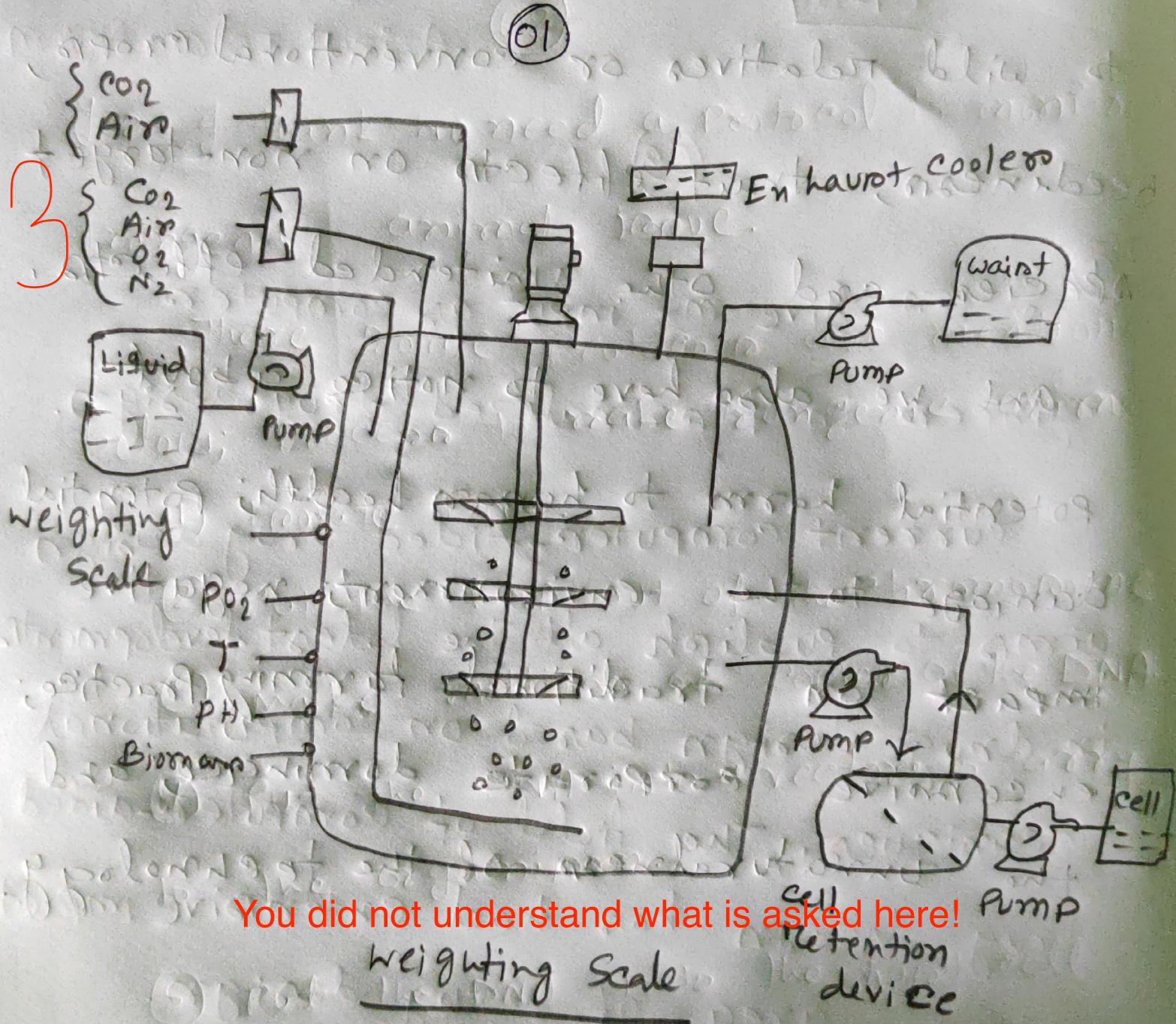
to wild relatives or conventional crops, weediness, trait effects on non-target species and other unintended effects. And we also have to notice about potential harm to human health; Potential damage to the environment; negative impact on traditional farming practice; encroaching corporate dominance; and the unnaturalness of the technology.

Some specific protocols can be mentioned;

- ① We can apply gene modification on those animal but we have to take care of their own habits, that cannot be destroyed.

② while clinical trial, some reaction
can be happen then, I think until
it's complete fully and be safe, we
have to keep them away from other
animal.

③ we have to give more observation
to them after trial. If we find
some defaults or impacts we can
take necessary action to solve
those problems.



You did not understand what is asked here!

weighting scale

cell
Retention
device

Pump

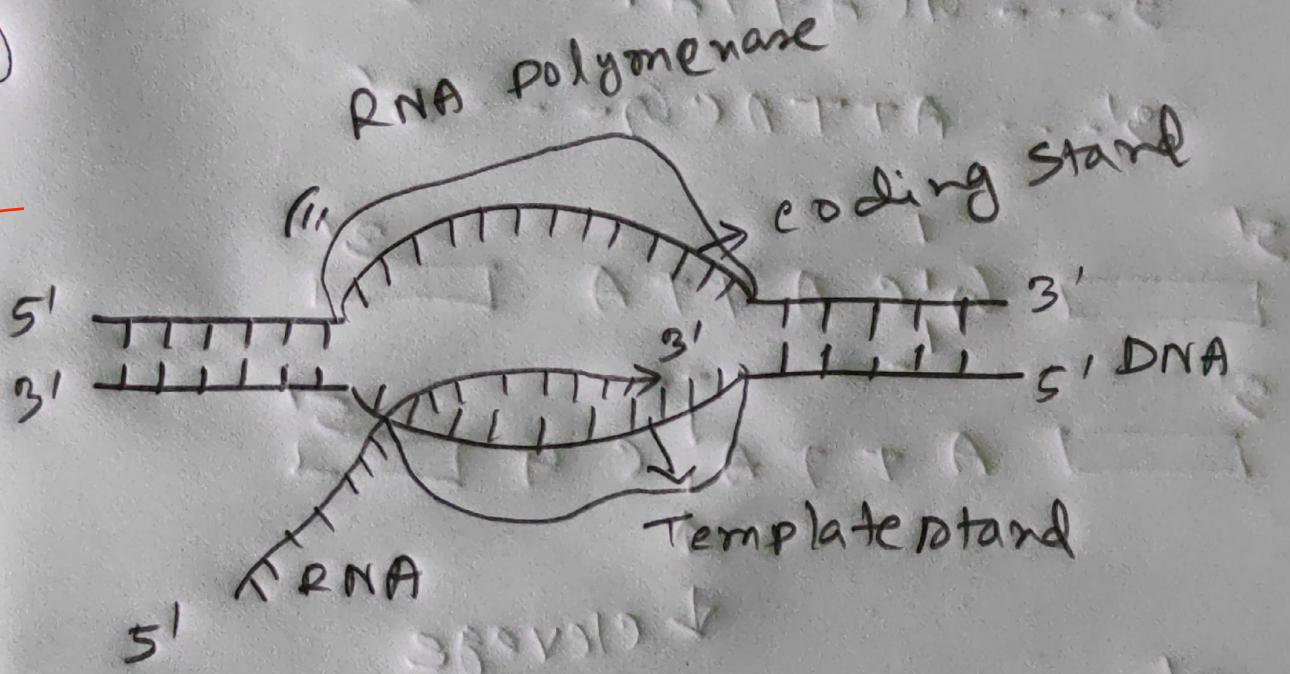
Pictorial view of the nutrients supply

and temperature control.

• Know about RNA (03) ~~about~~ address carbon

(b)

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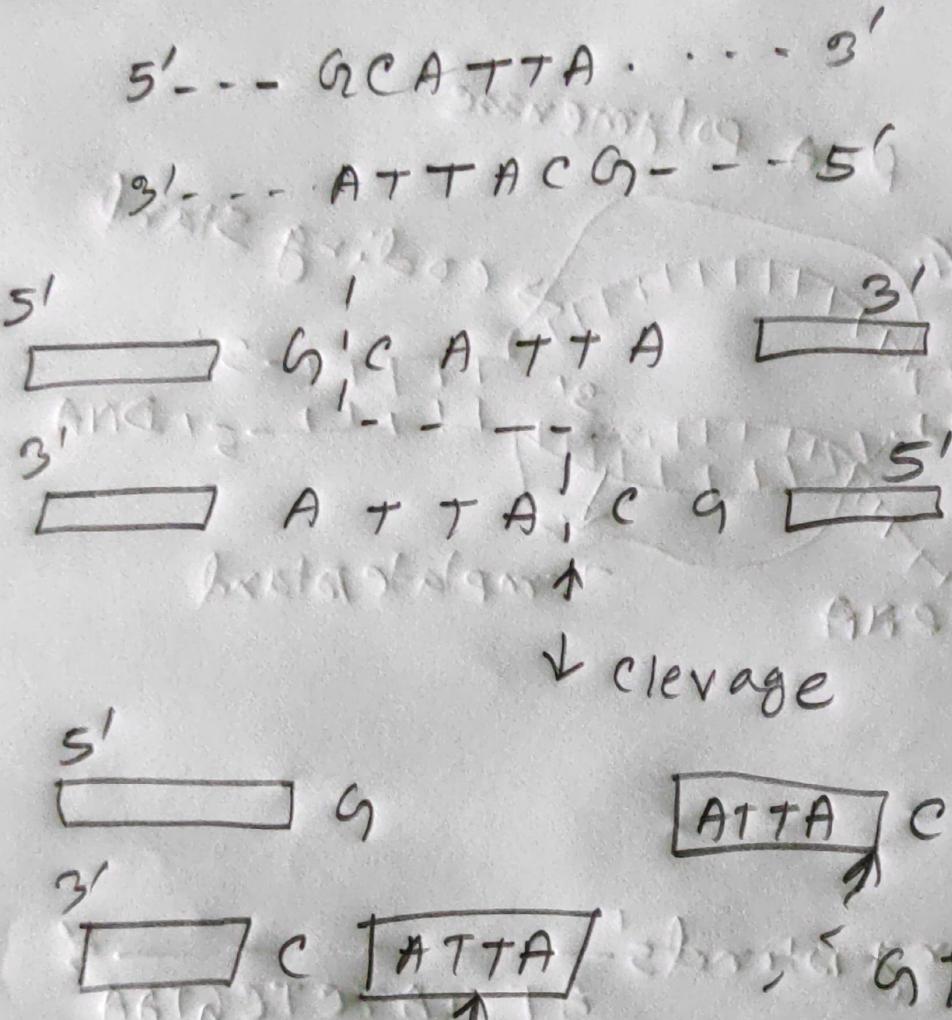


coding strand 5' ————— 3'
AT G A T C T C G T A A

RNA 5' ————— 3'
A [U] G A [U] C

Template 3' ————— 5'
T A C T A A G C A T T

process can be done by this way;



sequence GCAATTAA in PCR and have a fragment of DNA with 25 npaces and will repeat itself after every 25 npace sequence. To make it fit on right hand side, we have to work with restriction enzymes and other enzymes.