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# What to do with a YAML file
# Someone else write
get id back(CeleryScript, UUID)
delete object("regimen", "id")
unique name() # returns a name not currently used in internal storage
# Rachel writes these:
load commands(yaml file):
  self.CSV = yaml file{"CSV"}
  for yaml object in yaml file:
    make yaml object(yaml object)
make yaml object(yaml object):
  if "start time" in yaml object:
    make farm event(yaml object)
  else if "schedule" in yaml object:
    make regimen(yaml object)
  else if "actions" in yaml object:
    make sequence(yaml object)
  else:
    "Error: YAML object is not correctly formatted."
make farm event(yaml object):
  write the start of the CeleryScript
  if "repeat event" in vaml object:
    write parts of the CeleryScript a little differently so the event repeats (DONE)
  if "schedule" in yaml object:
    take the "schedule" part of the YAML object and send it to make regimen()
    use the ID make regimen() returns to finish writing the CeleryScript
  else if "actions" in yaml object:
    if type(item["actions"]) is not str: # if we need to make a sequence:
      take the "actions" part of the YAML object and send it to make sequence()
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else: # if the "actions" refer to the name of a sequence defined somewhere else in the file
      get the id and type of the object by calling obj from name()
    use the ID make sequence() returns to finish writing the CeleryScript
  Send the CeleryScript and get back the ID
  Write a Farm Event YAML object to internal storage in the corrent format, with the ID
  return
make regimen(yaml object):
  if "name" not in yaml object:
    auto = True
  else:
    auto = False
    check if the regimen is already in the internal storage, if it is, we have to update and delete a lot
of things (for later)
  write the start of the CeleryScript
  for item in yaml object["schedule"]:
    if type(item["actions"]) is not str: # if we need to make a sequence
      send the "actions", "groups", and "types" as a single YAML object to make sequence()
    else: # if the "actions" refer to the name of a sequence defined somewhere else in the file
      find the sequence in the file and send it to make sequence()
    take the returned ID of the sequence and finish writing your CeleryScript
  Send the CeleryScript and get back the ID
  Write a Regimen YAML object to internal storage in the corrent format, with the ID
  return Regimen ID
  make sequence(yaml object):
    if "name" not in yaml object:
      auto = True
    else:
      auto = False
      check if the sequence is already in the internal storage, if it is, we have to update and delete a
lot of things (for later)
    write the start of the CeleryScript
    if "group" in yaml object:
      loop over the entire CSV:
        if row is the right "group":
          CeleryScript + make actions(yaml object["actions"], x, y, z)
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else if "group" in yaml object:
      loop over the entire CSV:
        if row is the right "group":
        CeleryScript + make actions(yaml_object["actions"], x, y, z)
    else if "type" in yaml_object:
      loop over the entire CSV:
        if row is the right "type":
        CeleryScript + make actions(yaml object["actions"], x, y, z)
    Send the CeleryScript and get back the ID
    Write a Sequence YAML object to internal storage in the corrent format, with the ID
    return Sequence ID
name 1:
  kind: "farm event"
  auto: False
  ID: #
name 2:
  kind: "regimen"
  auto: True/False
  ID: #
name 3:
  kind: "sequence"
  auto: True/False
  ID: #
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