CODES WITHOUT COMMAS

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(ns commas
  "CODES WITHOUT COMMAS -- with apologies to F.H.C. Crick, et al"
  (:require [clojure.set :as s]
            [clojure.math.combinatorics :as c]))

"The problem of how a sequence of four things (nucleotides) can
determine a sequence of twenty things (amino acids) is known as
the 'coding' problem."

(comment "http://www.pnas.org/content/43/5/416" is the paper.)

(comment Now is 1957.  What do we know?)

(def nucleotides ["Adenine" "Cytosine" "Guanine" "Thymine"])
nucleotides ; => ["Adenine" "Cytosine" "Guanine" "Thymine"]

(def ACGT (map first nucleotides))
ACGT                                  ; => (\A \C \G \T)

(reverse ACGT)                        ; => (\T \G \C \A)

(def pair (zipmap ACGT (reverse ACGT)))
pair                                  ; => {\A \T \C \G \G \C \T \A}

(rand-nth ACGT)                       ; => \G
(rand-nth ACGT)                       ; => \A

(def strand (repeatedly (fn [] (rand-nth ACGT))))

(take 7 strand)                       ; => (\G \G \C \T \G \C \C)

(def dna (map (fn [b] [b (pair b)]) strand))

(take 6 dna)    ; => ([\G \C] [\C \G] [\T \A] [\T \A] [\A \T] [\A \T])
(take 7 (map first  dna))             ; => (\G \C \T \T \A \A \G)
(take 7 (map second dna))             ; => (\C \G \A \A \T \T \C)

(def essential {:F "phenylalanine"
                :H "histidine"
                :I "isoleucine"
                :K "lysine"
                :L "leucine"
                :M "methionine"
                :T "threonine"
                :V "valine"
                :W "tryptophan"})

(def conditional {:C "cysteine"
                  :G "glycine"
                  :P "proline"
                  :Q "glutamine"
                  :R "arginine"
                  :Y "tyrosine"})

(def dispensible {:A "alanine"
                  :D "aspartic acid"
                  :E "glutamic acid"
                  :N "asparagine"
                  :S "serine"})

(def amino (merge essential conditional dispensible))

(count amino)                         ; => 20

(comment Now ... What can we find out?)
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(c/selections ACGT 2)                 ; => ((\A \A)
                                      ;     (\A \C)
                                      ;     (\A \G)
                                      ;     (\A \T)
                                      ;     (\C \A)
                                      ;     (\C \C)
                                      ;     (\C \G)
                                      ;     (\C \T)
                                      ;     (\G \A)
                                      ;     (\G \C)
                                      ;     (\G \G)
                                      ;     (\G \T)
                                      ;     (\T \A)
                                      ;     (\T \C)
                                      ;     (\T \G)
                                      ;     (\T \T))

(count (c/selections ACGT 2))           ; => 16
(count (c/selections ACGT 3))           ; => 64
(> (count (c/selections ACGT 2)) (count amino)) ; => false
(> (count (c/selections ACGT 3)) (count amino)) ; => true

(def string (partial apply str))

(def triples (map string (c/selections ACGT 3)))

(take 7 triples)          ; => ("AAA" "AAC" "AAG" "AAT" "ACA" "ACC" "ACG")
(take 7 (reverse triples)) ; => ("TTT" "TTG" "TTC" "TTA" "TGT" "TGG" "TGC")
(count triples)           ; => 64

(take (count ACGT) (partition (count ACGT) 1 (cycle ACGT)))
;; => ((\A \C \G \T) (\C \G \T \A) (\G \T \A \C) (\T \A \C \G))

(def rotations
  (fn [s] (let [n (count s)]
            (map string (take n (partition n 1 (cycle s)))))))

(rotations ACGT)                      ; => ("ACGT" "CGTA" "GTAC" "TACG")
(rotations (take 3 ACGT))             ; => ("ACG" "CGA" "GAC")

(take 7 (map rotations triples))        ; => (("AAA" "AAA" "AAA")
                                        ;     ("AAC" "ACA" "CAA")
                                        ;     ("AAG" "AGA" "GAA")
                                        ;     ("AAT" "ATA" "TAA")
                                        ;     ("ACA" "CAA" "AAC")
                                        ;     ("ACC" "CCA" "CAC")
                                        ;     ("ACG" "CGA" "GAC"))

(def codons (set (map (comp set rotations) triples)))

(count codons)                          ; => 24
(take 7 codons)                         ; => (#{"ACC" "CCA" "CAC"}
                                        ;     #{"GGG"}
                                        ;     #{"TTT"}
                                        ;     #{"GCC" "CGC" "CCG"}
                                        ;     #{"CAA" "ACA" "AAC"}
                                        ;     #{"CTC" "CCT" "TCC"}
                                        ;     #{"AGC" "CAG" "GCA"})

(count (group-by count codons))         ; => 2
(map first (group-by count codons))     ; => (3 1)

(def sense-codons (filter (fn [g] (= 3 (count g))) codons))

(count sense-codons)                    ; => 20 Eureka!
```

```
(take 7 sense-codons)                   ; => (#{"ACC" "CCA" "CAC"}
                                        ;     #{"GCC" "CGC" "CCG"}
                                        ;     #{"CAA" "ACA" "AAC"}
                                        ;     #{"CTC" "CCT" "TCC"}
                                        ;     #{"AGC" "CAG" "GCA"}
                                        ;     #{"TAT" "TTA" "ATT"}
                                        ;     #{"GAG" "GGA" "AGG"})

(def sense (map (comp first sort) sense-codons))

(count sense)   ; => 20

(take 7 sense)  ; => ("ACC" "CCG" "AAC" "CCT" "AGC" "ATT" "AGG")

(def nonsense (s/difference (set triples) (set sense)))

(count nonsense)                ; => 44

(= (count triples)
   (+ (count sense)
      (count nonsense)))        ; => true

(def code (zipmap (map vec (sort sense)) (sort (keys amino))))

(sort-by second code)                   ; => ([[\A \A \C] :A]
                                        ;     [[\A \A \G] :C]
                                        ;     [[\A \A \T] :D]
                                        ;     [[\A \C \C] :E]
                                        ;     [[\A \C \G] :F]
                                        ;     [[\A \C \T] :G]
                                        ;           …
                                        ;     [[\C \G \G] :R]
                                        ;     [[\C \G \T] :S]
                                        ;     [[\C \T \G] :T]
                                        ;     [[\C \T \T] :V]
                                        ;     [[\G \G \T] :W]
                                        ;     [[\G \T \T] :Y])

(map string (take 7 (partition 3 1 strand)))
;; => ("TTC" "TCG" "CGG" "GGT" "GTG" "TGA" "GAT")

(def amino-keys (remove nil? (map code (partition 3 1 strand))))

(take 7 amino-keys)                     ; => (:R :W :M :W :G :T :D)

(def aminos (map amino amino-keys))

(take 17 aminos)                        ; => ("arginine"
                                        ;     "tryptophan"
                                        ;     "methionine"
                                        ;     "tryptophan"
                                        ;     "glycine"
                                        ;     "threonine"
                                        ;     "aspartic acid"
                                        ;     "alanine"
                                        ;     "glutamic acid"
                                        ;     "alanine"
                                        ;     "glycine"
                                        ;     "valine"
                                        ;     "asparagine"
                                        ;     "alanine"
                                        ;     "glutamic acid"
                                        ;     "glutamine"
                                        ;     "glutamine")

"Note: This is fun programming but (as of 1961) bad biology."
```

https://github.com/tbl3rd/clj-intro/blob/master/exercises/commas.clj