

I thought it over, and for a change, you are right. There is a problem. However, we are trying to solve the problem by creating more problems, so let's try something different.

What are we doing?

Nomenclature.

How should we do it?

Systematically.

What is the system?

Nucleotide → Codon → tRNA → Amino Acid

But is a sequential system.

Nucleotide → Codon → tRNA → Amino Acid

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There are two different things we should care about and so we are trying to name:

1. Pairs.
2. Relationships between pairs.

Nucleotide → Codon → tRNA → Amino Acid
↕ ↕ ↕ ↕
Nucleotide → Codon → tRNA → Amino Acid

So, I propose a system of prefix and suffix where the prefix identifies the type and the suffix identifies the thing.

Prefixes:

Nucl

Cod

Trin

Amin

Suffixes:

et

Iture

Names:

Nucllet, Nucliture

Codet, Coditure

Trinet, Triniture

Aminet, Aminiture