

Code: ACG, CGU

GCATR Tool

June 26, 2019

```
is_code = code_check_if_code(params$code)
cn_circular <- code_check_if_cn_circular(params$code)
circular <- code_check_if_circular(params$code)

comma_free <- code_check_if_comma_free(params$code)
self_comp <- code_check_if_self_complementary(params$code)
acid <- code_get_acid(params$code)
```

## 1 Prperties

- acid: RNA Tuple length  $\ell = 3$
- Circular: TRUE
- Comma-Free: TRUE
- $C_\ell$  Circular: TRUE
- Self-Complementary: TRUE

```
G <- code_factor_graph(params$code, TRUE, TRUE)

## 2 - -1

plot(G)
```

```
if(circular) {
  G <- code_factor_longest_path(params$code)
} else {
  G <- code_factor_cycle(params$code)
}
plot(G)
```

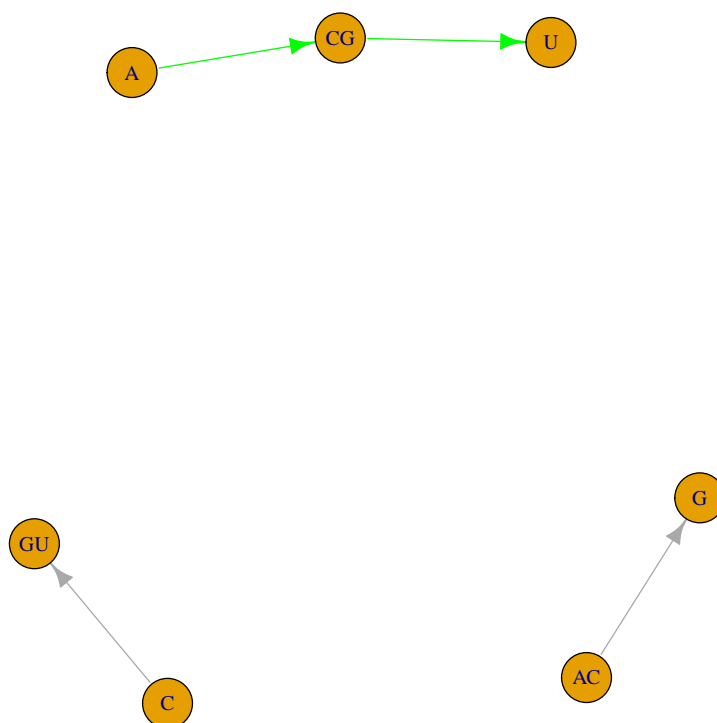


Figure 1: Representing Graph

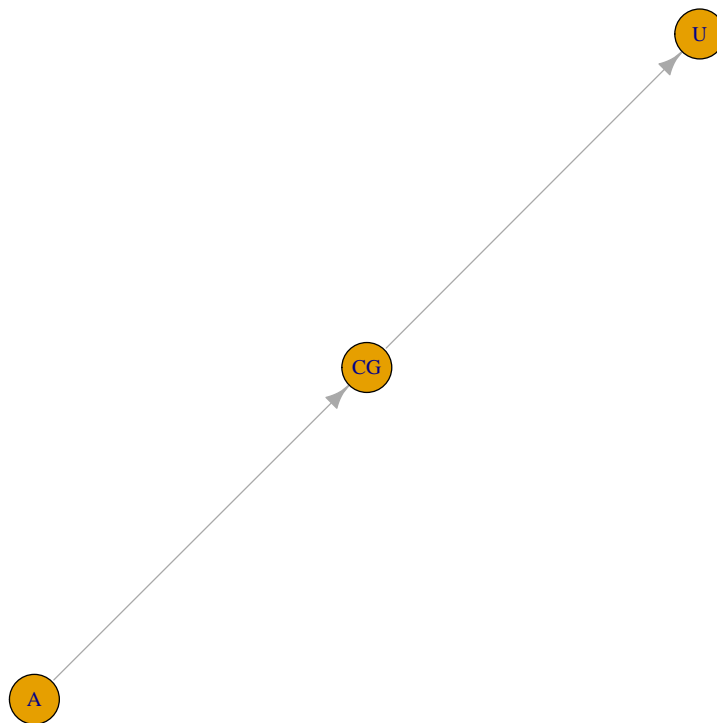


Figure 2: Representing Graph longest path or cycle

|     | X_U_      | X_C_      | X_A_       | X_G_       |
|-----|-----------|-----------|------------|------------|
| U_U | UUU - Phe | UCU - Ser | UAU - Tyr  | UGU - Cys  |
| U_C | UUC - Phe | UCC - Ser | UAC - Tyr  | UGC - Cys  |
| U_A | UUA - Leu | UCA - Ser | UAA - Stop | UGA - Stop |
| U_G | UUG - Leu | UCG - Ser | UAG - Stop | UGG - Trp  |
| C_U | CUU - Leu | CCU - Pro | CAU - His  | CGU - Arg  |
| C_C | CUC - Leu | CCC - Pro | CAC - His  | CGC - Arg  |
| C_A | CUA - Leu | CCA - Pro | CAA - Gln  | CGA - Arg  |
| C_G | CUG - Leu | CCG - Pro | CAG - Gln  | CGG - Arg  |
| A_U | AUU - Ile | ACU - Thr | AAU - Asn  | AGU - Ser  |
| A_C | AUC - Ile | ACC - Thr | AAC - Asn  | AGC - Ser  |
| A_A | AUA - Ile | ACA - Thr | AAA - Lys  | AGA - Arg  |
| A_G | AUG - Met | ACG - Thr | AAG - Lys  | AGG - Arg  |
| G_U | GUU - Val | GCU - Ala | GAU - Asp  | GGU - Gly  |
| G_C | GUC - Val | GCC - Ala | GAC - Asp  | GGC - Gly  |
| G_A | GUA - Val | GCA - Ala | GAA - Glu  | GGA - Gly  |
| G_G | GUG - Val | GCG - Ala | GAG - Glu  | GGG - Gly  |