## Code: ACG, CAG

## GCATR Tool

June 4, 2020

```
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)
tuple_l = code_tuple_length(code)</pre>
```

## 1 Prperties

• acid: DNA

• Tuple length  $\ell = 3$ 

• Circular: TRUE

• Comma-Free: TRUE

•  $C_3$  Circular: TRUE

• Self-Complementary: FALSE

```
G <- code_factor_graph(params$code, TRUE, TRUE)
plot(G)</pre>
```

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

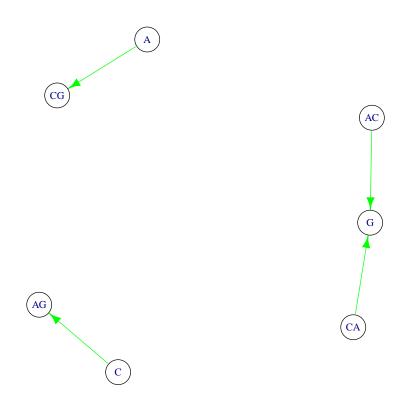


Figure 1: Representing Graph

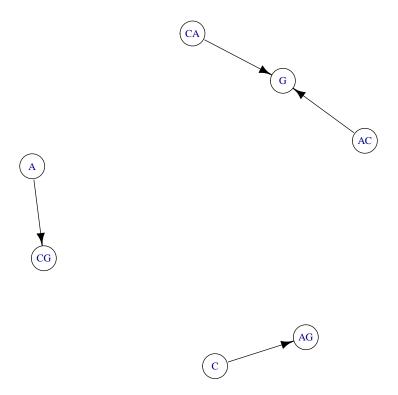


Figure 2: Representing Graph longest path or cycle

	$X_{-}U_{-}$	X_C_	XA	X_G_
U_U	UUU - Phe	UCU - Ser	UAU - Tyr	UGU - Cys
$U_{-}C$	UUC - Phe	UCC - Ser	UAC - Tyr	UGC - $Cys$
UA	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
CG	CUG - Leu	CCG - Pro	CAG - $Gln$	CGG - $Arg$
$A_{-}U$	AUU - Ile	ACU - Thr	AAU - $Asn$	AGU - Ser
$A_{-}C$	AUC - Ile	$\mathrm{ACC}$ - $\mathrm{Thr}$	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 - $Asp$	$\operatorname{GGC}$ - $\operatorname{Gly}$
$G_A$	GUA - Val	GCA - Ala	GAA - Glu	GGA - $Gly$
$G_{-}G$	GUG - Val	GCG - Ala	GAG - Glu	GGG - Gly