

Tree scale: 1

Colored ranges

- frog Ly6
- GPIHBP1
- Ly6L
- PSCA
- Ly6H
- SLURP2
- LYNX1
- SLURP1
- Ly6D
- Ly6E
- Ly6K (secreted?)
- LYPD2
- Ly6K
- Ly6 other
- Reptilian Ly6 group 1
- Reptilian Ly6 group 3
- Reptilian Ly6 group 2
- Reptilian Ly6 group 4
- Reptilian Ly6 group 1-2
- Pre-3FTx
- Non-standard
- Colubrid non-standard C #8 -> Y/L
- Plesiotypic
- Viperid non-standard C #8 -> Y
- Short-chain
- Basal viper genomic 8-cysteine
- Basal genomic 8-cysteines
- Long-chain
- Colubrid non-standard C #8 -> Y
- Dimeric
- Ly6
- class2c
- class2b
- class2d
- class1a
- class3a
- class2a
- Clade in plesiotypics
- Australian long-chain reversion
- Bungarus non-standard
- Kappa
- IIalpha
- Orphan XVII
- Orphan II
- Orphan IV
- Orphan V
- Aminergic
- Orphan IX
- Orphan VIII
- Orphan VII
- Dendroaspis non-standard
- Dendroaspis basal 8-cysteine
- class3c
- Cobra+ cytotoxin
- Laticauda long-chain reversion
- class1b
- class3b
- Basal non-standard
- Micrurus 8-cysteine in plesiotypic
- 4.0

