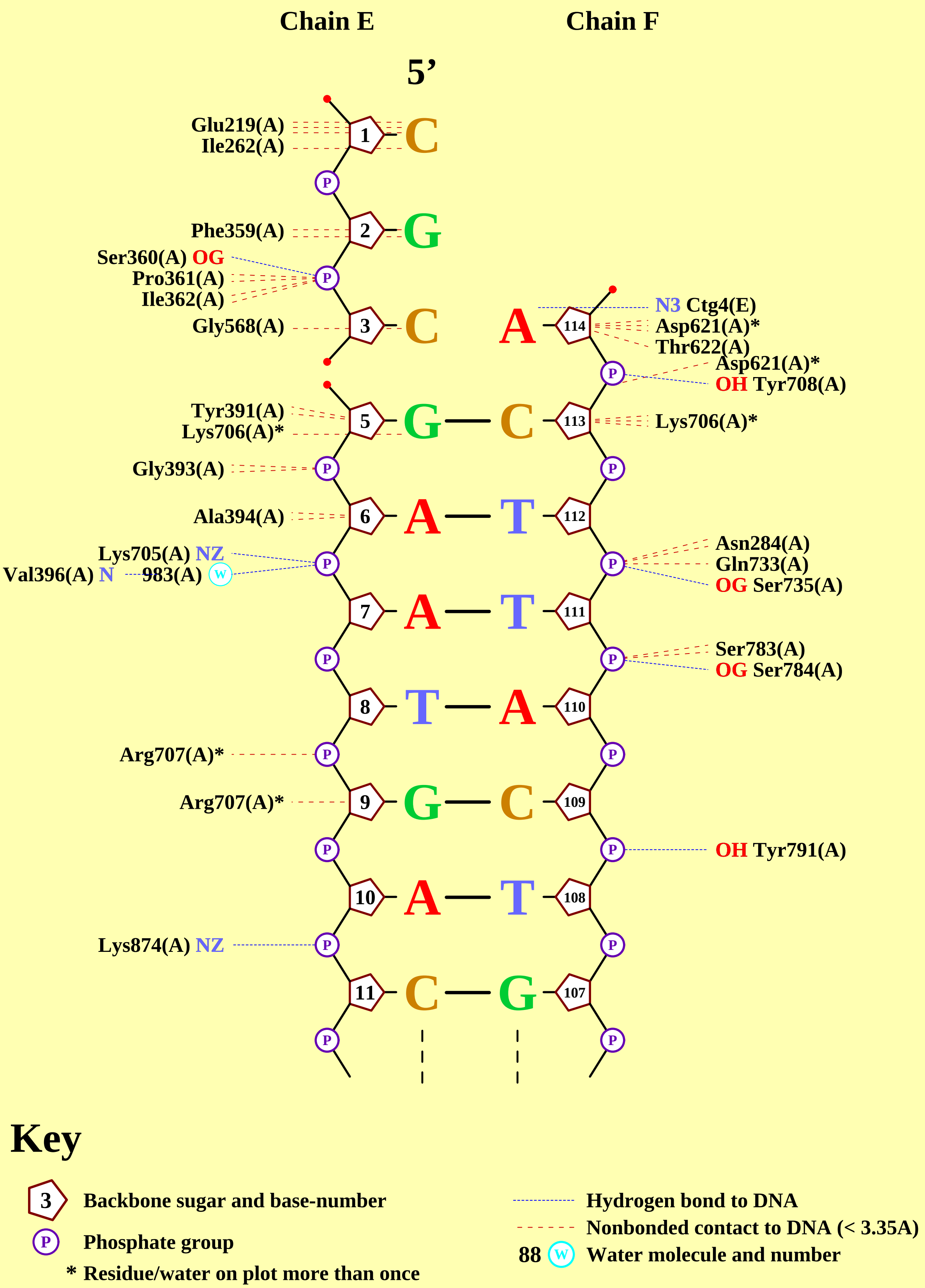
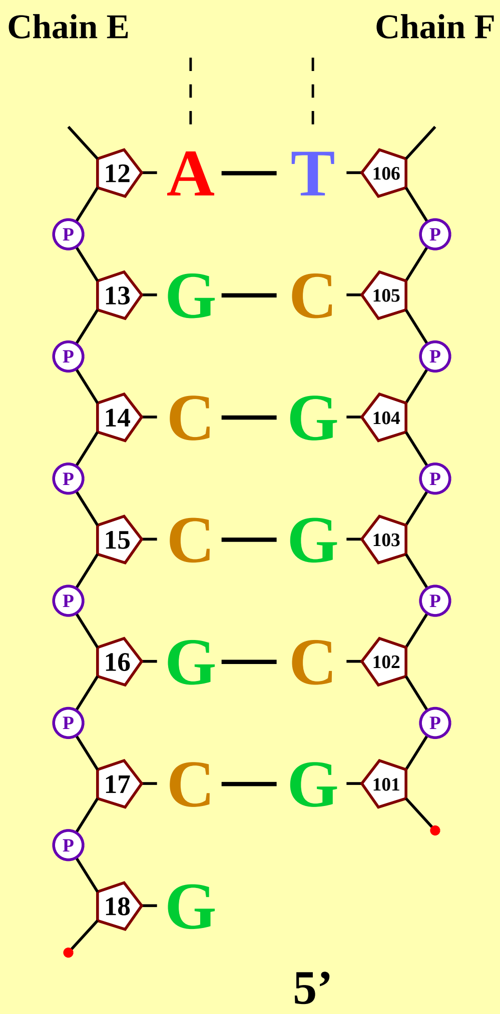

Chain E
Chain F
5’
Glu219(A)
Ile262(A)
Phe359(A)
Ser360(A) OG
Pro361(A)
Ile362(A)
Gly568(A)
Tyr391(A)
Lys706(A)*
Gly393(A)
Ala394(A)
Lys705(A) NZ
Val396(A) N
983(A)
Arg707(A)*
Arg707(A)*
Lys874(A) NZ
N3 Ctg4(E)
Asp621(A)*
Thr622(A)
Asp621(A)*
OH Tyr708(A)
Lys706(A)*
Asn284(A)
Gln733(A)
OG Ser735(A)
Ser783(A)
OG Ser784(A)
OH Tyr791(A)
C G C G A A T G A C
A C T T A C T G
Key
Backbone sugar and base-number
Phosphate group
* Residue/water on plot more than once
Hydrogen bond to DNA
Nonbonded contact to DNA (< 3.35A)
88 Water molecule and number

file



Key

3 Backbone sugar and base-number

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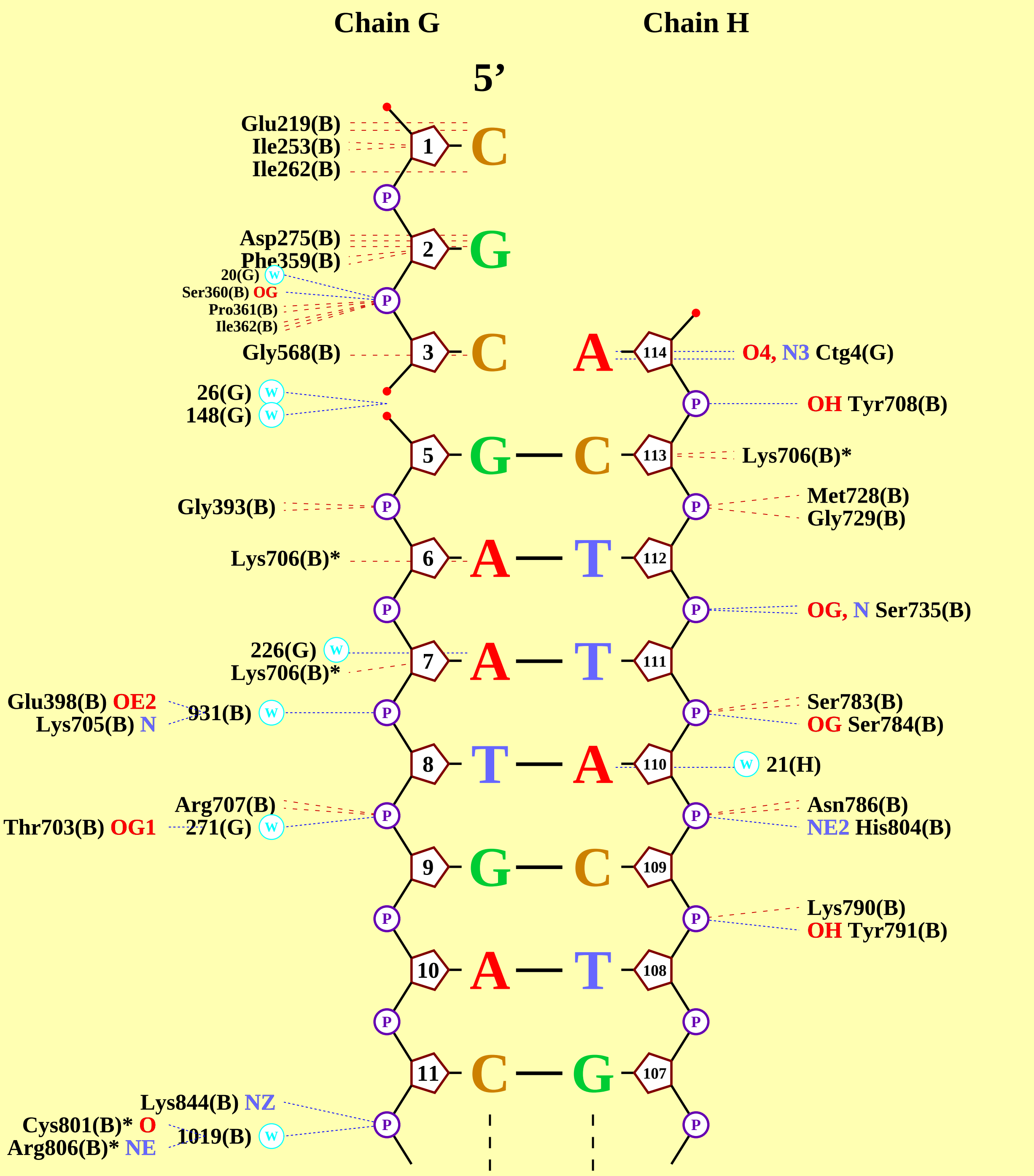
88 W Water molecule and number

file



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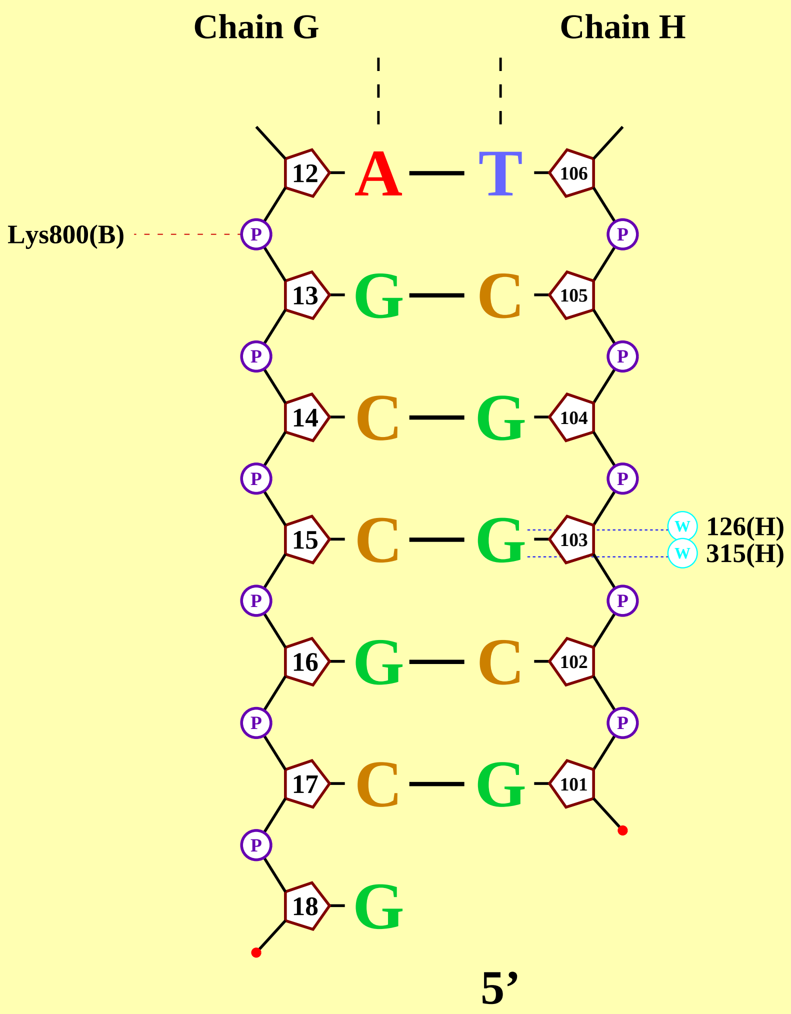
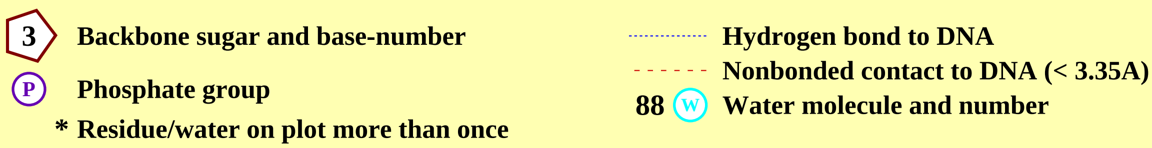
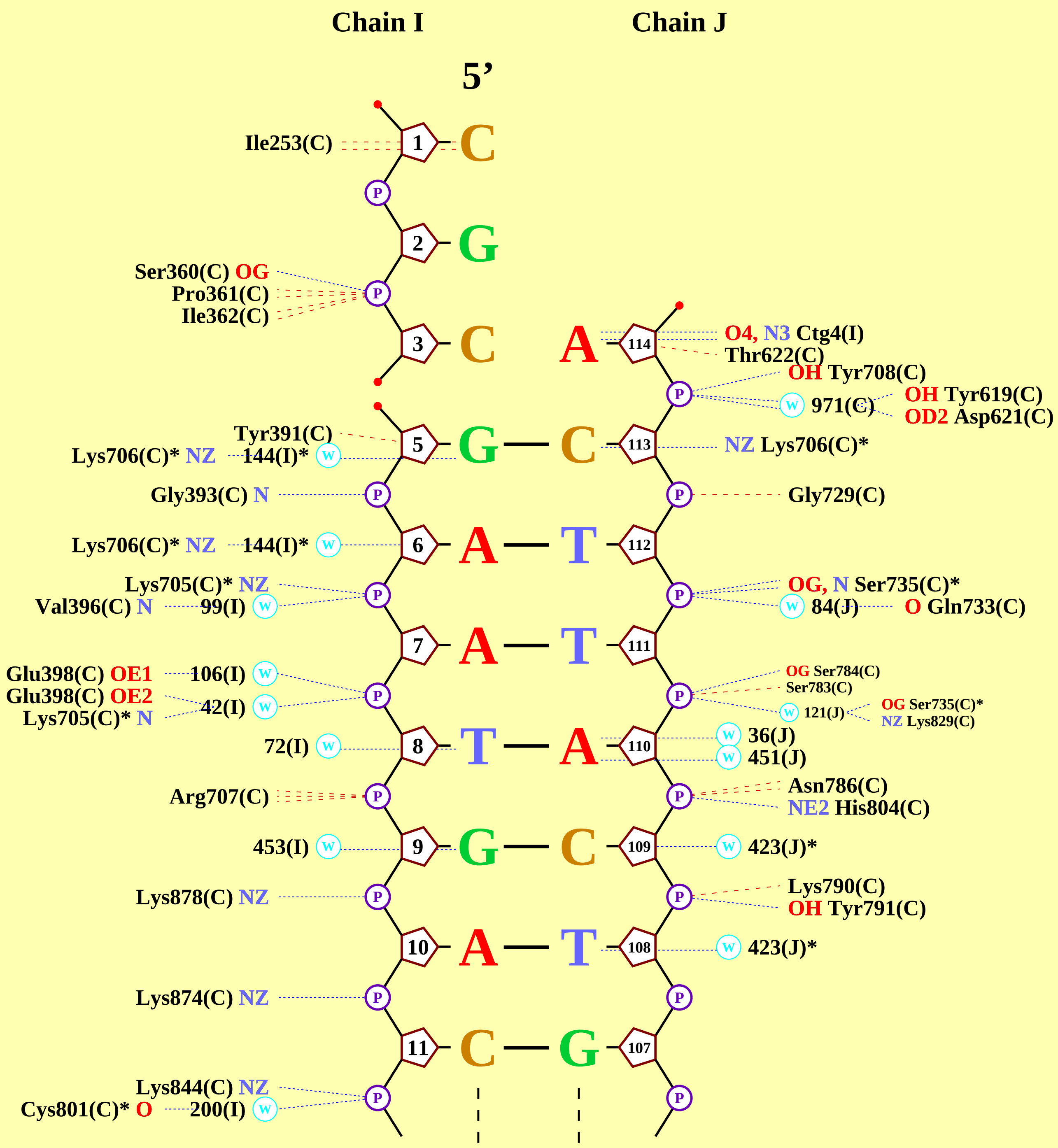
file


Chain G
Chain H
Lys800(B)
12
A
T
106
13
G
C
105
14
C
G
104
15
C
G
103
126(H)
315(H)
16
G
C
102
17
C
G
101
18
G
5’

Key


3
Backbone sugar and base-number
P
Phosphate group
* Residue/water on plot more than once
Hydrogen bond to DNA
Nonbonded contact to DNA (< 3.35A)
88
Water molecule and number

file



Key

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Phosphate group

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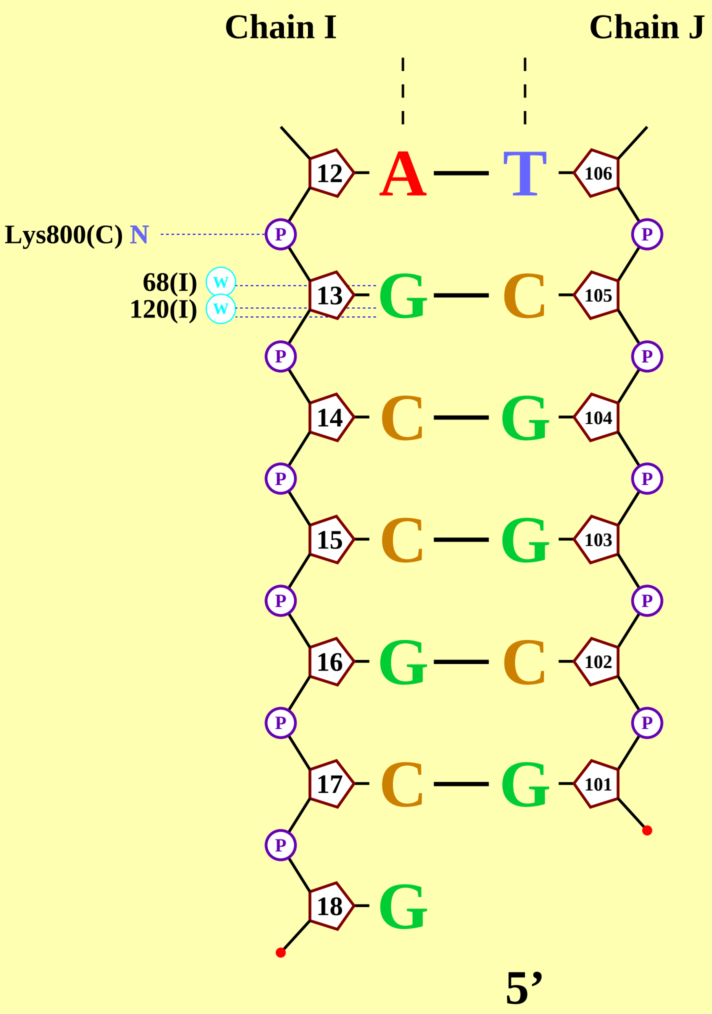

Key

- 3 Backbone sugar and base-number
- P Phosphate group
- * Residue/water on plot more than once
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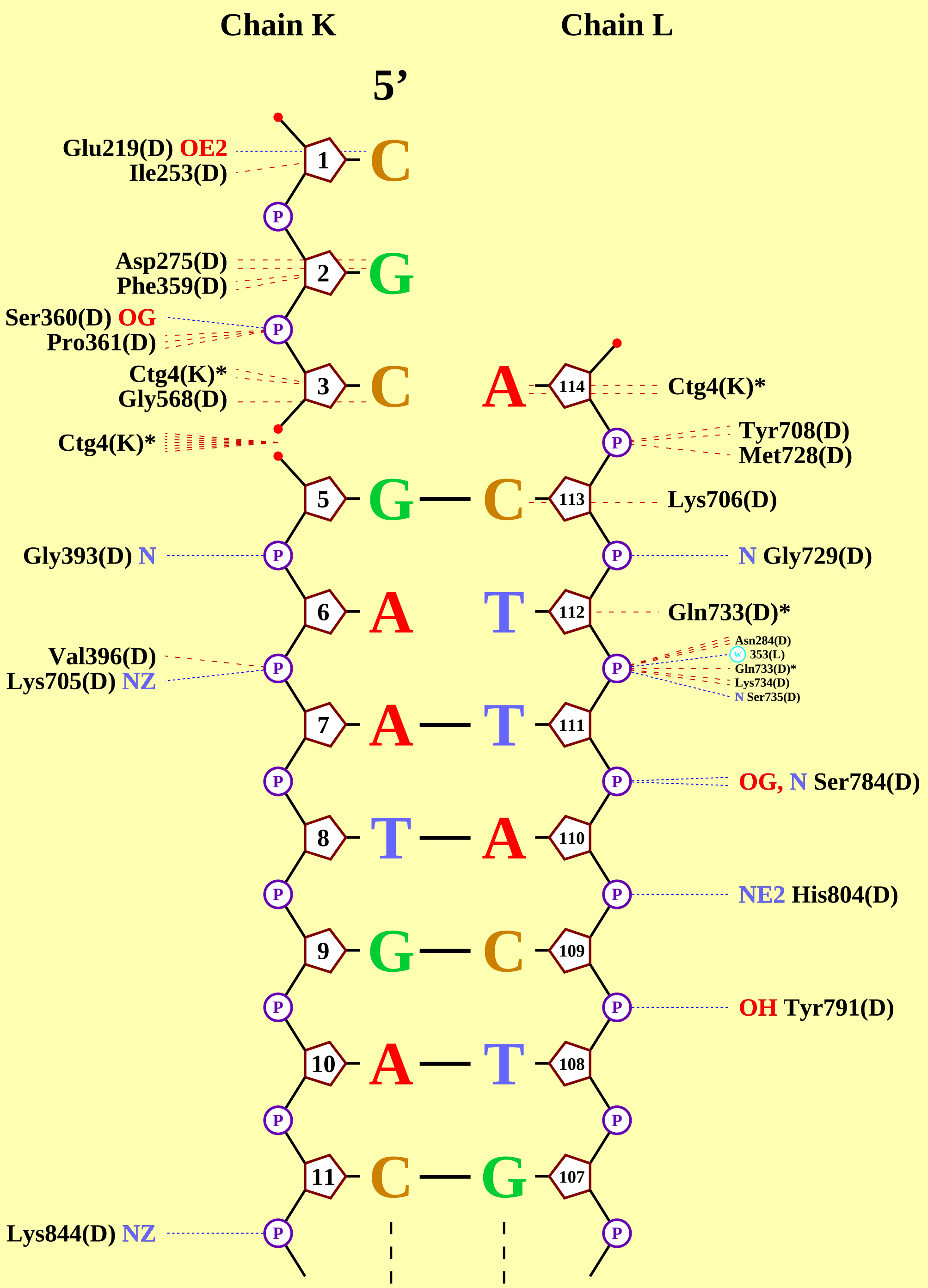
file



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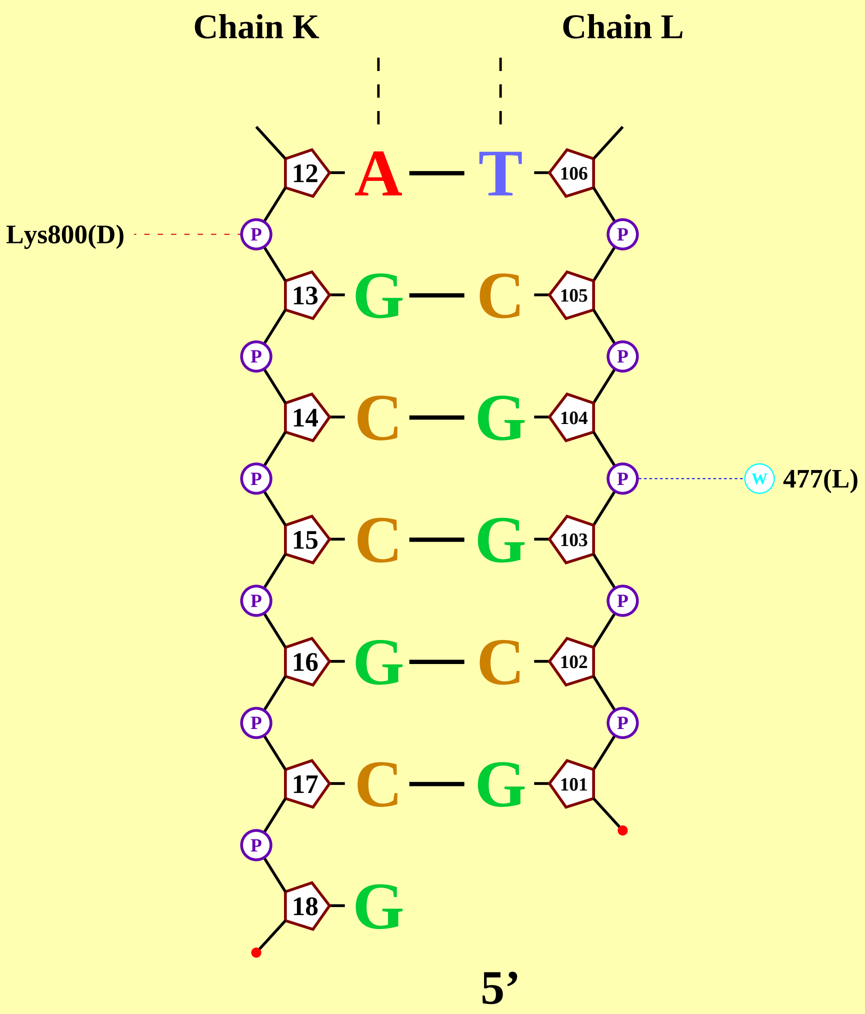

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