The Pathways in Genetics

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Abstract The positive and negative characteristic of his-pro chemistry has been revealed the molecular mechanism of genetics implicated from oxy-time to protein expansion. The molecular suppression and consequently expression has been clarified to some extent. The difference of 100 or 1000 in the structure would be motivating the influx of electro-gravitational impulses into the system towards equilibrium.

Keywords Histidine, Proline, t-RNA, Genetic suppression, Oxy-time

1. Introduction

According to positive and negative 'factors of opposite' 154 and 6 (or 6*0.0019 = 0.0114), histidine (155.1552) and proline (115.1311) may be called positive and negative amino acids or vice versa. Adding molecular weights, 155.1552 + 115.1311 = 270.2863 where 0.2863 or 2863 is an expressed form of 1063 or 163 (a point where proteolytic cleavage occurs due to electro-gravitational conflicts) where suppression values are multiple of 900 [1].

As there are so many transitions occurs into the system, the decimals have been avoided somewhere. There are 0.0001-0.0002(1-2) adjustable time differences into the system which are not specifically clarified.

There are two points of bisection found 0.3667(193) and 0.3496(184) and the molecular mechanism would be blocked under suppression and leads to expressions.

Avoiding decimals, 3667 − 2863 = 804 and 3496 − 2863 = 633 that can be expressed as 3333 = 1451 (gln ht) + 1882 (lys ht) and also 804 + 114 (factor of opposite) = 918 = 3618 (in expressed form) = 1736 (leu ht) + 1882(lys ht) thus formed a leu-lys-gln complex. Correspondingly, 146 (lys vt) + 146 (gln vt) = 292 = 900 − 608(oxy-time) and 608 − 292 = 316 (suppressed form of oxy-time) where 316 = 153 + 163 and also 131 (leu vt) + 146 (lys vt) = 277. It is seen 1605 (lunar gravity) + 277 = 1882 and 1605 − 154 = 1451 and also 1882 − 1451 = 431 = 285(15) + 146 and 285 − 154 = 131 (leu vt that suppressed form of electromagnetic values 1031) supports electro-gravitational chemistry in genetics. It is seen the integer part (vertical time) of a molecular weight possess a separate identity and the decimal part (horizontal time) are interrelated.

The t-RNA factors 66A^0 (distance of constancy) and

CCA(357) are aligned to the system in terms of values. Now, 66*0.0019 = 0.1254 = 0.0354 (in suppressed form) and 0.0003(3) time differences with 357 (CCA). Mathematically, 155 − 66 = 89 and 115 + 66 = 181 makes a tyr-ala complex where 181.1894 + 89.0935 = 270.2829 where 2863 − 2829 = 34 and where 34*0.0019 = 0.0646, a significant values in genetics, the core-values of tyr(0.1545) with 0.0001 time difference where 645 = 3345 (in expressed form) = 1894 (tyr ht) + 1451 (gln ht) and correspondingly 181 + 146 = 327 = 163*2(app.) and 181 − 146 = 35. With some differentiation, 645 = 4245 = 2124(met ht)*2 with 0.0003 time differences and 0.0645(34) − 0.0034(ht) = 0.0611 = 611(on transitions) = 617 − 6 is important.

Tyrosine (181.1894) itself an enigmatic amino acid in tyrosine kinase where 646 − 357 = 289 = 1894 − 1605 = 135(A) + 154 and 181 − 94 (i.e. values beyond 900) = 87 = 222(CC) − 135(A).

The proline (115.1311) is directly linked to genetics where 333(CCC) + 154 = 487 and 333 − 6 = 327. It is seen 333 + 357 = 690 = 69*10 and 357 − 333 = 24(or, 0.0456) are linked to his-pro chemistry and the protein expansion would be explained. The lunar time, 184 = 115 + 69 (or 0.1311) gives 250 (suppressed form) and 690, separately multiplying by 10 where 155 - 24 = 131 and 705 − 456(24) = 249 = 250 -1. It is significant that 155 + 38 = 193 and 652(0.1552) + 38 = 690 where 967(193) − 690 = 277 = 377(TTT) − 100 and 690 − 411 (or, 0.1311) = 279 = 277 + 2. The his-pro suppressions curved the system by 163 where 1031 − 967 = 64 = 164 - 100 makes a difference of 100 in structural biology would be motivating influx of electro-gravitational influxes into the system towards equilibrium. Since 1000 = 900 (values of suppression) + 100, a structural matter, 969(51) + 31 = 1000 = 1031 − 31 would be balancing the system where 51 + 63 = 114(factor of opposite).

The methionine (149.2124) shows positive-negative interactions where 155 − 6 = 149 while cysteine(121.1590) shows negative-negative interactions where 115 + 6 = 121. It is seen 2124 − 1552 = 572 = 574 -2 where 1605 − 1031 = 574

and $1590 - 1311 = 279 = 277 + 2$ where $690 + 279 = 969(51)$. The values 1590 would be expressed form of $690 = 1380/2$.

Mathematically, $1590 + 2124 = 3714$ where $3714 - 3667 = 47 = 851 - 804$ and $3714 - 3496 = 218 = 851 - 633$. This is significant that $511 + 293 = 804$ and $511 - 293 = 218$. Since $293 + 100 = 393$ (p53 protein expansion) that corresponds to $611 = 617 - 6$(factor of opposite) of JAK2 protein.

In addition, JAK2 and TP53 would be considered as positive and negative segments clarified later.

2. Discussions

The leu-lys-gln complex:

The molecular weight of leucine 131.1736 where the suppression of $1736 = 836(44) = 705$(suppression of lunar gravity) $+ 131$ (suppression of electromagnetic values) is fundamental in genetics where $836 - 277 = 559 = 405$(AAA) $+ 154$ is directly related to lysine (146.1882). The suppressed values $= 1736 + 1882 = 3618 = 918 = 900 + 18$ where $146 + 18 = 164$(point of proteolytic cleavage) and the suppressed values $= 1736 + 1451 = 3187 = 487$ where $836 + 487 = 1323$ (core values of glutamine) and $900 - 487 = 413 = 423$ (suppressed values of gln core values) $- 10$(difference of 10) in the structure. It is seen $1882 - 1451 = 431 = 267 + 164$ where $900 - 267 = 633 = 1064 - 431$. It is also seen $163*2 = 326 = 327 - 1$ and $163*3 = 489 = 487 + 2$ are significant.

The met-tyr complex:

The molecular suppression changes the scenario of values e.g., the vertical time of tyr $= 181*0.0019 = 0.3439$ or 739 where $739 + 230 = 969(51)$ and where $519 - 289 = 230$ and $519 + 289 = 808$ derived from $2124 - 1605$(lunar gravity) $= 519$ and $1894 - 1605 = 289$. It is seen $739 + 69 = 808$ and correspondingly $230 + 181 = 411(69)$. Again, $900 - 739 = 161 = 181 - 20$, a difference of 20 (i.e. halved of 40 or $2*10$) has been found in the system related to protein expansion where 670 (i.e. $900 - 230$) $+ 20 = 690 = 1380/2$ and $460 + 20 = 480$ (suppressed form of 1380) and also 518 (suppressed form of 2318) $- 38 = 480$ where $38*10 = 380 = 1380 - 1000$. The protein expansion drawn from JAK2 G1849T V617F and TP53 G469T V157F where $1849 - 469 = 1380$ and $1849 + 469 = 2318$ are considered as positive and negative segments.

Now, $149 + 181 = 330$ and correspondingly $2124 + 1894 = 4018 = 418$ (under suppression) where 645 (tyr core values) $+ 418 = 1063$ and $804 - 645 = 159$ where $159 + 171 = 330$ and $645 - 12 = 633$ where $171 - 159 = 12$.

Tyrosine is an enigmatic amino acid where $181 + 3 = 184$ and $3496(184) - 1605 = 1891 = 1894 - 3$ and also $1894 + 969(51) = 2863$. It is seen $969 - 739 = 230$ and correspondingly $969 + 739 = 1708$ what is suppressed form of 808 and also 324 (suppressed form of 2124) $+ 739 = 1063$ and $739 - 324 = 415 = 515 - 100$. I have shown a bisection of

149 in the system where $149*0.0019 = 0.2831 = 2831$ (on transition) $= 1415*2$ (app.) where $1605 - 190 = 1415$ [2] and correspondingly $804 + 190 = 994$ (suppressed form of tyr ht) where $515 + 289 = 804 = 705 + 100$ (app.) and $94 + 69 = 163$ in the structure.

About cancer:

The mutations JAK2 G1849T V617F and TP53 G469T V157F would be considered as segregation of positive and negative segment where $1849 - 469 = 1380 = 480$ (under suppression) $= 690*2 = 460*3$ and correspondingly $1849 + 469 = 2318 = 518$ (under suppression) where $808 - 518 = 290 = 155*2 - 20$ and $460/2 = 230$ in met-tyr complex.

The mutational values for both mutations are 0.0754 (core values of val) $- 0.1235 = -0.0481$ and 475 (i.e., genetic mutation $= 151 - 126 = 25$ where $25*0.0019 = 0.0475$) $+ 6 = 481$ that should be added to corresponding molecular point gives 1098 (or 198) and 638 where $1098 + 638 = 1736 = 836$ (suppressed form) and shows electro-gravitational chemistry in cancer.

It is seen $836 = 774 + 62$ where $617 + 157 = 774$ and conversely $617 - 157 = 460 = 574 - 114(6)$ and also $638 = 475 + 163$ and correspondingly $198 + 163 + 114(6) = 475$.

In opposite direction, $1235 + 754 = 1989 = 189$ (under suppression) $= 378(TTT)/2$ where $189 + 44(836) = 233$ (2033 in expressed form) and $189 - 44 = 145 = 573 - 428$ and where $2033 - 1605 = 428$ and $1605 - 1031 = 574$ makes a complicated mechanism towards equilibrium in the system. It is also seen $428 - 51 = 377 = 184 + 193$ under suppression.

The values 193 or 0.3667 is significant where $193 = 131 + 62$ and $1031 - 62 = 969(51)$. A time difference of 0.0002 is found here where $969 - 967 = 2$ and $62 + 2 = 64 = 32*2$ (oxy-time).

3. Conclusions

The histidine-proline suppression is otherwise suppression of points of bisection or replication 193 and 184 gives $193 + 184 = 377(TTT)$ making a difference of 100 after grabbing 64 beyond 967(193) in the system that would be motivating influx of electro-gravitational impulses and leads to expressions. The efficacy of oxy-time to suppress time directionally leads to lifeline. The cancer disease has been shown in light of electro-gravitational chemistry.

REFERENCES

[1]   American Journal of Medicine and Medical Sciences 2023, 13(4): 496-498.

[2]   American Journal of Biochemistry 2022, 12(1): 4-7.