A Comparative Study on JAK2 G1849T V617F and TP53 G469T V157F Mutations

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Abstract The methionine-tyrosine chemistry gives a clear concept of structural genetics. The protein expansions are result of electro-gravitational chemistry. Cysteine is a significant residue found in these structural mutations. Lunar gravity (0.1605) possesses a significant role in biophysics. The influx of electro-gravitational impulses into cell occurs following electro-gravitational structural imbalance towards equilibrium through cell processes.

Keywords Point mutation, Cysteine, Methylation, Phosphorylation, Carboxylation

1. Introduction

These cancer developing mutations v157f and v617f can be explained from methionine(AUG)- tyrosine (UAC) structural interactions. The protein mutations is followed by genetic mutations G1849T and G469T where 1849 and 469 are molecular or genetic points are structural factors. These can be related as 1849 + 469 = 2318(122) =0.2831(149, met gravitational values) - 0.0513 (electronic)time values) and conversely 1849 - 469 = 1380 = 0.2831(149)-0.1451 (electromagnetic values) in the structure. The lunar time 183*0.0019 = 0.3477 (diameter of our moon with transitional decimal factor) = 0.1159(61)*3 where 0.1159*2= 0.2318(122) and 460*3 = 1380 where 9(0.0171)*3 =27(0.0513) shows molecular point is a chemistry of gravitational and electromagnetic values. The genetic point 1849 shows an electronic values apart where 1849 + 27(0.0513) = 1876 = 469*4.

It is seen 0.0460-0.0171(9)=0.0289=0.1894 (tyr horizontal time) - 0.1605 (lunar gravity) and conversely 0.1840-0.0171(9)=0.1669=0.1150+0.0519 where 0.2124 (met horizontal time) - 0.1605=0.0519 that related to 0.1159(61). In addition, 0.0460 (expansion unit) + 0.0171=0.0631=0.1150-0.0519 and 0.2124-0.1894=0.0230=230=460/2 suggests protein expansion is due to electro-gravitational chemistry.

Mathematically, 519 + 289 = 808 = 1616/2 where 0.1616 = 0.1605 + 0.0011 and where 11 = 9 + 2(or 0.0038) are structural factors. Again, 1380 / 2 = 690 where 1690 is symmetrical to 221(i.e. 183 + 38) since 121.1590 (cys) is structurally associated.

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The core values of val = 117*0.0019 - 0.1469 = 0.0754 while pre-transitional values 0.1469 - 0.0117 = 0.1352 gives 0.1352 + 0.0754 = 0.2106 (curvature of tyr) where 0.2000 - 0.0106 = 0.1894 (tyr ht). Accordingly, 0.1735 (phe pre-transitional values) + 0.1235(phe core values) = 0.2970 where 0.2000 - 0.0970 = 0.1030 = 0.1605(lunar gravity) - 0.0575 where 0.3477(183) - 0.1451*2 = 0.0575 would actuates influx of electro-gravitational impulses into cell and leads to cancer development.

It is seen 181*0.0019 = 0.3439 (tyr vertical time) = 0.2970 + 0.0469 where 0.2970 - 0.1849 = 0.1121 that symmetrical to 0.2590 according to cysteine (121.1590) where 0.2970 - 0.2590 = 0.0380 = 0.1380 - 0.1000 (a structural factor) where 1000 - 575 = 425 (electromagnetic values in opposite direction).

The methionine and tyrosine are structurally symmetrical and molecular points 157 or 617 would be specified for valine and after mutation phenylalanine would hamper structural symmetry that leads to cancer development.

Since transitions are common factor in structural genetics, decimals have been avoided somewhere. A time difference 0f 0.0001-0.0002 has been found in the system.

2. Discussions

Met(149.2124)-Tyr(181.1894) structural symmetry and cancerous mutations:

The core values or hidden time is associated with the genetic structural system. The addition of core values = 0.0707(met) + 0.1545(tyr) = 0.2252 = 0.3667(193) - 0.1415 where 0.2831(149)/2 = 0.1415(app.) and where 0.1605 - 0.1415 = 0.0190(10) in the structure [1]. Conversely, 0.1545 - 0.0707 = 0.0838 = 0.0418*2 + 0.0002 where 0.3477(183) - 0.3059 = 0.0418 and 0.3667 - 0.3059 = 0.0608 (oxy-time)

according to fundamental molecular structure. The lunar time (0.3477) is associated with earth-moon curvature of time (0.0368), a point of bisection while 0.3667(193) would be associated with 0.0378 (TTT) would be also a point of bisection.

It is seen 1834 (i.e. 3668/2) -289 = 1545 (core values of tyrosine) while 1834 + 289 = 2123 (met ht with 0.0001 time difference) and also 1849 - 1545 = 304(16), the existence of oxy-time in the structure.

Mathematically (avoiding decimals), 3439(181) - 469 = 2970 = 1235 (phe C_v) + 1735 (phe pre-transitional values), a significant values after mutation and 3439 - 1849 = 1590 (cys ht) where 2318(122) - 1590 = 728 and where 728 - 19(1) = 709 (cys C_v) and 709 - 19 = 690 = 1380/2 in the structure. In terms of lunar gravity, 1605 - 60 = 1545 where 183/3 = 61 = 60 + 1 and 181 - 121 = 60.

From genetics point of view, 1628 (i.e. 814*2) – 469 = 1159(61) and 1849 - 1628 = 221 that dimensional to 1590 + 100 = 1690 = 1000 + 690 (i.e. 460*3/2) in cysteine (121.1590) where 814 = 487 (deoxyribonucleotide triphosphates avg. MW) + 327 (deoxyribonucleotide monophosphates avg. MW) in the structure.

The doubling of cysteine = 121.1590*2 = 242.3180 since cys-cys pairing is significant in the structure. It is seen 3180 = 1605 + 1575 = 1451 + 1729 where 3477 - 2902 = 575 (electro-gravitational factor) and 575 + 154 (factor of opposite) = 729 and 729 + 154 = 883 = 460 (expansion unit) + 423 (electro-magnet values in opposite direction with 0.0002 time difference) where 423 - 193 = 230 = 2124 - 1894. Conversely, 1605 - 1575 = 30 = 36 - 6 (factor of opposite) and 1729 - 1451 = 278 = 272 (i.e. 136*2) + 6 where 1605 (lunar gravity) - 1469 (cys pre-transitional values) = 136 = 1849 - 1713 (tyr pre-transitional values).

The tyr-cys complex shows 136 + 153 (factor of opposite) = 289 = 1894 - 1605 and 3439(181) - 2106 = 1333 = 1713 - 380 that related to protein expansion.

Since the system is structurally compatible, the mutations would leads to cancer. There would have been inclination between positive and negative side under mutations. The tyr-cys complex is associated with stem cell biochemistry [2] where gravitational side (181) and anti-gravitational side (0.1894) in tyrosine would possesses an individual identity and co-related that would be helpful for investigation of JAK2 and TP53 and its consequences under mutations.

3. About Molecular Bio-Markers

The biological processes like phosphorylation, carboxylation, methylation etc. show electro-gravitational chemistry. The phosphorylation of tyrosine= $181.1894 + 94.9714 (PO_4^{-3}) = 276.1608$ where 0.1608 + 0.0276 = 0.1884 and where 0.0884 = 0.0460 + 0.0424 = 0.0729 + 0.0154 (factor of opposite) with 0.0001 time difference and also where 0.0575 + 0.0154 = 0.0729 would be causing influx of electro-gravitational impulses into cell towards equilibrium and raises bioactivity.

The carboxylation of glutamic acid = 147.1299 + 28.01 = 175.1399 where 0.1399 + 0.0175 = 0.1574. There is an analogy between glutamic acid and tryptophan while the pre-transitional values of amino acids are measured from lunar gravity exists in upper and lower level.

The methylation of arginine and lysine gives molecular weights 188.2300 g/mol and 160.2100 g/mol respectively. It is seen 2300-188=2112=1605+507 where 3477-2970=507 and 2300+188=2488=1605+883 in the structure.

Again, 2100 - 160 = 1940 = 2970 - 1030 where 1605 - 1030 = 575 = 3477 (lunar time) - 2902(1451*2, electromagnetic values).

Molecular biomarkers would be key values to combat with cancer but needed highly experimentation.

4. Conclusions

The structural symmetry of methionine-tyrosine is associated with cancer development while mutation occurs in molecular point a structural component intrinsically aligned to met-tyr symmetry. The protein expansion is due to electro-gravitational chemistry. The electro-gravitational structural imbalance leads to influx of electro-gravitational impulses into cell towards equilibrium through cell processes and sometimes goes detrimental.

REFERENCES

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