

Research and Reviews: Journal of Medical and Health Sciences

Genes and Gene Expression

Sravya. I*

Department of Biotechnology, JNTU, Hyderabad, Telangana

Commentary Article

Received: 23/03/2015
Revised: 22/04/2015
Accepted: 30/04/2015

*For Correspondence

Sravya.I, Department of Biotechnology, JNTU, Hyderabad, Telangana, Tel: 7794804883; Email: sravyai@gmail.com

Gene Expression

Gene expression is the procedure by which hereditary guidelines^[1] are utilized to orchestrate quality items. These items are normally proteins, which go ahead to perform vital capacities as compounds, hormones^[2] and receptors^[3], for instance. Genes^[4] that don't code for proteins, for example, ribosomal RNA^[5] or exchange RNA^[6] code for utilitarian RNA items^[7].

Genes

Genes are subunits of DNA^[8-10], the data database of a cell that is contained inside the cell core. This DNA conveys the hereditary outline that is utilized to make all the proteins the cell needs. Each quality^[11-15] contains a specific arrangement of guidelines that code for a particular protein^[16].

DNA exists as two^[17] since a long time ago, matched strands that frame a twofold helix. Each of these strands is comprised of individual building squares called nucleotide bases. These bases incorporate adenine, thymine, cytosine, and guanine (A, T, C and G),^[18-20] which are masterminded in triplets, with every triplet speaking to a particular amino corrosive^[21].

DNA^[22-25] is found in all cells exhibit in the body beside those that don't contain a core, for example, developed red platelets or the cornified cells of nails and skin. Every human cell is comprised of 46 chromosomes^[26-28], each of which contains profoundly dense and wound DNA comprising of a huge number of quality groupings. In every cell, 23 chromosomes^[29] are acquired from the father and 23 are acquired from the mother. Twenty-two of the chromosomes [30] from every guardian are autosomes and the remaining chromosome^[31,32] is a X or Y sex chromosome^[33].

Quality Expression Mechanism

Quality expression^[34-36] is the procedure by which DNA is utilized to make proteins, which then go ahead to perform different imperative capacities in the body. The protein^[37] could be a chemical, hormone or receptor, for instance.

The procedure of quality expression incorporates the accompanying steps:

- **Transcription** - Transcription is the procedure by which a section of DNA^[38-40] is utilized to produce a RNA layout. The DNA section is "read" by a catalyst called RNA polymerase, which delivers a strand of RNA that is complimentary to the DNA. In this correlative RNA strand, all thymine bases are supplanted by uracil.

- **Processing** - This essential RNA transcript^[41,42] is then adjusted to change over it into adult delegate RNA (mRNA) that can be utilized as a part of interpretation. The mRNA experiences joining to uproot the non-coding parts of the transcript (introns) so that just the coding areas (exons) remain.
- **Non-coding RNA development** - Non-coding locales of RNA^[43] (ncRNA) are interpreted as forerunners which are then handled further. For instance, these locales may be translated as preribosomal RNA (pre rRNA) which then experiences cleavage to wind up ribosomal RNA (rRNA)^[44].
- **RNA trade** - The dominant part of experienced RNA^[45] is then transported from the core to the cytoplasm. Albeit some RNAs work in the core, most are helped through pores in the core into the cytosol, incorporating all RNAs included in protein blend.
- **Translation** - The last mRNA^[46] conveys the data expected to code for proteins. Each three base combines on the mRNA relates to a coupling site for an exchange RNA (tRNA) which conveys an amino corrosive. The amino acids are then connected together in a tie by a ribosome to make a simple protein chain.
- **Protein collapsing** - The long chain of amino acids^[47] folds to frame a three-dimensional structures utilizing catalysts called chaperones. This three-dimensional structure is the last, practical type of the protein.

Quality Expression Measurement

Quality expression^[48] is the procedure by which hereditary directions are utilized to incorporate quality items. Measuring this quality expression is a key component in the investigation of life sciences. Case in point, understanding the level of quality expression in a cell, tissue or organic entity can give important data as far as distinguishing viral contamination, deciding malignancy powerlessness or testing whether a microscopic organisms is impervious to penicillin^[49].

Regulation of Gene Expression

Quality expression^[50] is a standout amongst the most firmly controlled procedures in the body. This procedure needs to be entirely managed to guarantee that cells create the right measure of proteins when they require them. Any disturbance to this regulation^[51-53] can prompt genuine outcomes, including tumor.

Quality expression is directed by requirements of the cells. In the event that the cell is presented to a situation where specific quality items are required, the declaration of that quality item will be expanded. Cells likewise^[54] create particular quality items in light of outer signs or cell harm.

Two illustrations of quality expression regulation^[55] incorporate the control of insulin expression to guarantee blood glucose levels are controlled and the control of cyclin expression to guarantee ordinary movement of the cell cycle. Along these lines, quality regulation decides the cell's general structure and capacity, representing cell separation, cell morphology and the cell's flexibility to its surroundings.

Quality expression^[56] is controlled at diverse stages. Case in point, the translation step may be avoided to stop DNA being changed over to RNA or the post-translational change of a protein^[57-60] may be ceased.

Transcriptional regulation

Translation may be directed in three principle routes, as takes after:

- Genetic – Where a control variable collaborates with the quality^[61-63]
- Modulation – Where a control variable collaborates^[63-65] with the interpretation apparatus
- Epigenetic – Where non-grouping changes in DNA structure^[66-69] influence interpretation

Post-transcriptional regulation

Regulation at the post-transcriptional stage is controlled by importin^[70] and exportin proteins^[71] that impact the vehicle of RNA all through the core.

Translational regulation

Regulation at the interpretation stage^[72] is less basic than at different stages. This type of regulation is utilized by anti-toxins^[73-75] and poisons, for instance, which restrain protein interpretation with a specific end goal to disturb the cell's typical quality expression^[76,77], hence prompting cell death^[78].

Conclusion

To live, cells must have the capacity^[79] to react to changes in their surroundings. Regulation of the two principle ventures of protein creation – interpretation and interpretation – is basic to this flexibility. Cells can control which qualities get interpreted and which transcripts^[80] get deciphered; further, they can biochemically prepare transcripts and proteins to influence their movement. Regulation of translation and interpretation happens in both prokaryotes^[81] and eukaryotes^[82], yet it is significantly more mind boggling in eukaryotes.

References

1. Karthikeyan S, Hoti SL, Prasad NR. Resveratrol Modulates Expression of ABC Transporters in Non-Small Lung Cancer Cells: Molecular Docking and Gene Expression Studies. *J Cancer Sci Ther.* (2014) 6:497-504.
2. Yoshimitsu M, Higuchi K, Arima N, Medin JA, Takenaka T. Successful Induction of Pluripotent Stem Cells From a Fabry Disease Mouse Model: Toward the Development of Safe Lentiviral Gene Therapy. *J Stem Cell Res Ther.* (2015) 5:259.
3. Wang G, Cao F, Wang G, Yousry A, El-Kassaby. Role of Temperature and Soil Moisture Conditions on Flavonoid Production and Biosynthesis-Related Genes in Ginkgo (*Ginkgo biloba* L.) Leaves. *Nat Prod Chem Res.* (2015)3:162.
4. Hackeny AC, Lane AR . Exercise Endocrinology: Guidance for Future Research Direction and Focus. *J Steroids Hormon Sci.* (2015) 6:e114.
5. Saber HB, Elloumi M. Efficiently Mining Gene Expression Data via Novel Binary Biclustering Algorithms. *J Proteomics Bioinform.* (2015) S9: 008.
6. Kitamura N, Akazawa K, Yoshihara K . Statistical Properties and Power Analysis of Cox's Proportional Hazards Model Regularized by Various Penalties for DNA Microarray Gene Expression Survival Data. *J Health Med Informat* (2015) 6:180.
7. Heikal TM, Mossa ATH, Khalil WKB . Protective Effects of Vitamin C against Methomyl-Induced Injuries on the Testicular Antioxidant Status and Apoptosis-Related Gene Expression in Rat. *J Environ Anal Toxicol.* (2015) 5:255.
8. Hessmann E, Ellenrieder V, Koenig A. NFAT in Pancreatic Carcinogenesis. *J Carcinog Mutagen.* (2014) 5:203.

9. Uchiumi F, Larsen S and Tanuma S . Alteration in Transcriptional State, as a First Step in Cancer Development. *Pharm Anal Acta*. (2014) 5:e166.
10. Tulu A, Ishaque A, Egiebor E, Chambers C, Jagu R . Separate and Joint Effects of Polycyclic Aromatic Hydrocarbons (PAH) and Polychlorinated Biphenyls (PCB) on Aromatase CYP19A Transcription Level in Atlantic Tomcod (*Microgadus tomcod*). *J Marine Sci Res Dev*. (2014) 4:151.
11. Kamiuchi S, Fukaya M, Usui T, Iwata N, Okazaki M, et al. MatrIn 3 Augments the Transcriptional Activity of an SV40 Promoter- Mediated Luciferase Gene with a Highly Repetitive DNA Component. *J Mol Genet Med*. (2014) 8:146.
12. Zhang W, Hamidouche Z, Pourcher G, Gribova V, Haghighi F, et al. A Population of Human Mesenchymal Stem Cells Specific to the Fetal Liver Development. *J Stem Cell Res Ther*. (2014) 4: 238.
13. Huang J, Wang L, Jiang M, Lin H, Qi L, et al. (2012) Parathyroid Hormone- Like Hormone (PTHrP) Feedback Mitosis to Downstream DNA Replication Coupling Postreplication Repair-Induced Cell Proliferation Network in No-Tumor Hepatitis/Cirrhotic Tissues (HBV or HCV Infection) by Systems-Theoretical Analysis. *Mol Biol* 1:106.
14. Valenzuela MS .Initiation of DNA Replication in the Human Genome. *Hereditary Genetics*. (2012) S1:003.
15. Moscarelli L, Mjeshtri A, Annunziata F, Tsalouchos A, Bertoni E .BKV-DNA Replication in Renal Transplant Recipients: Early Discontinuation of Mycophenolate Mofetil or an Early Combination Therapy with Fluoroquinolones and Activated Vitamin D. What Is the Best Strategy for Management? *J Nephrol Therapeutic*. (2012) S4:005.
16. Daoud SS . Genome Wide Identification of FGFR2 Alternative Splicing in Hepatitis C: Potential Roles in Malignant ransformation. *J Pharmacogenomics Pharmacoproteomics*. (2014) 5: e140.
17. Korenkova V, Jones A, Hoy WE, Morais C, Cooper MA, et al. Urinary Biomarkers for Detection of Early and Advanced Chronic Kidney Disease - A Pilot Study. *Med chem*. (2015) 5:096-103.
18. Giulio F, Chiara G, Paolo R.Corneal Neovascularization: A Translational Perspective. *J Clin Exp Ophthalmol*. (2015) 6:387.
19. Naeem A, Khan TA, Fazili NA.Protein Folding and Misfolding: A Perspective from Theory. *J Glycomics Lipidomics*. (2015) 5:128
20. Ahmad S, Siddiqui Z . Protein Glycation: A Firm Link to Cause Metabolic Disease and their Complications. *J Glycomics Lipidomics*. (2015) 4:127.
21. Mingming Li, Pengcheng Fan and Yu Wang . Lipidomics in Health and Diseases - Beyond the Analysis of Lipids. *J Glycomics Lipidomics*. (2015) 5:126.
22. Eman Allam, Mia Recupito, Hend Mohamed, L. Jack Windsor. In Vitro Effects of Nicotine, Cigarette Smoke Condensate, and *Porphyromonas gingivalis* on Monocyte Chemoattractant Protein-1 Expression from Cultured Human Gingival Fibroblasts. *J Interdiscipl Med Dent Sci*. (2015) 3:171.
23. Goldfeld M, Malec A, Podella C, Rulison C . Proteins as Surfactant Enhancers for Environmental and Industrial Applications. *J Pet Environ Biotechnol*. (2015) 6:211.
24. Muravlyova LE, Luchanskiy VB, Bakirova RY, Kolesnikova YA, Nurgaliyeva AS, et al. The Determination of Oxidized Proteins and Albumin in Plasma of Chronic Kidney Disease Patients. *Biol Med (Aligarh)*. (2015) 7: 230.

25. Nakai R, Naganuma T . Oligoflexia, the Newest Class of the Phylum Proteobacteria, Consisting of only One Cultured Species and Uncultured Bacterial Phylotypes from Diverse Habitats. *J Phylogen Evolution Biol.* (2015) 3:141.
26. Ramdani L, Bourboulou R, Belkouch M, Jebors S, Tauran Y, et al. Multifunctional Curcumin-Nanocarriers Based on Host-Guest Interactions for Alzheimer Disease Diagnostic. *J Nanomed Nanotechno.* (2015) 1 6:270.
27. Zhang K, Li J, Wang J, Liu T, Wang X, et al. Combined REDV Polypeptide and Heparin onto Titanium Surface for the Hemocompatibility and Selectively Endothelialization. *J Cell Sci Ther.* (2015) 6:198.
28. Bech L, Busk PK, Lange L. Cell Wall Degrading Enzymes in *Trichoderma asperellum* Grown on Wheat Bran. *Fungal Genom Biol.* (2015) 4:116.
29. Tsuda B, Kametani Y, Miyamoto A, Miyako H, Kumaki N, et al. The Effect of Peptide Treatment on the HLA-Binding and Antibody Production in Peripheral Blood Mononuclear Cells Obtained from Japanese Breast Cancer Patients. *J Vaccines Vaccin.* (2015) 6:270.
30. Weltman JK Identification of Invariant Peptide Domains within Ebola Virus Glycoprotein GP1, 2. *J Med Microb Diagn* (2015) 4:176.
31. Wang H. Pull the Trigger, it Fires: The Critical Role of Insulin-Stimulated Caveolin-1 Tyrosine 14 Phosphorylation in Regulation of Insulin Trans-Endothelial Transport. *J Metabolic Syndr.* (2015) 4:e114.
32. Yao Y, Marais TLD, Costa M. Chromatin Memory in the Development of Human Cancers. *Gene Technology.* (2015) 4:114.
33. Martins MD, Castilho RM. Histones: Controlling Tumor Signaling Circuitry. *J Carcinog Mutagen.* (2013) S5:001.
34. Chahin H, Ekong B, Fandy TE . Epigenetic Therapy in Malignant and Chronic Diseases. *J Pharmacogenom Pharmacoproteomics.* (2013) 4:118.
35. Pang ALY, Rennert OM. Protein Acetylation and Spermatogenesis. *Reproductive Sys Sexual Disord.* (2013) S1:005.
36. Su B, Xiang SL, Su J, Tang HL, Liao QJ, et al. Diallyl Disulfide Increases Histone Acetylation and P21^{WAF1} Expression in Human Gastric Cancer Cells In vivo and In vitro. *Biochem Pharmacol.* (2012) 1:106.
37. Wang Y, Zhang G, Li F, Kan L, Ben J, et al. Altered Expression and DNA Methylation Profiles of ERCC6 Gene in Lens Tissue from Age-Related Cortical Cataract. *J Clin Exp Ophthalmol.* (2015) 6:392.
38. Walawalkar YD, Tiwary K, Saha T, Nayak V . Significance of Microsatellite Instability and Gene Methylation as Prognostic Biomarkers during Gallbladder Cancer Progression: A Review. *J Cell Sci Ther.* (2015) 6:196.
39. Sonohara F, Nomoto S, Hayashi M, Hishida M, Inokawa Y, et al. STEAP4 Inactivation Correlates Poor Prognosis and might be a Possible Cause of steatotic Change in Hepatocellular Carcinoma, Detected by Triple-Combination Array Analysis. *J Carcinog Mutagen.* (2014) 5:201.
40. Wang H. Pull the Trigger, it Fires: The Critical Role of Insulin-Stimulated Caveolin-1 Tyrosine 14 Phosphorylation in Regulation of Insulin Trans-Endothelial Transport. *J Metabolic Syndr.* (2015) 4:e114
41. Ling Ho H . Functional Roles of Plant Protein Kinases in Signal Transduction Pathways during Abiotic and Biotic Stress. *JBiodivers Biopros Dev.* (2015) 2:147.
42. Nichols TW . Hyperphosphorylation of Tau Protein in Downâ€™s Dementia and Alzheimerâ€™s Disease: Methylation and Implications in Prevention and Therapy. *J Alzheimers Dis Parkinsonism.* (2014) 4: 159.

43. Ling Ho H. Functional Roles of Plant Protein Kinases in Signal Transduction Pathways during Abiotic and Biotic Stress. *JBiodivers Biopros Dev.* (2015) 2:147
44. Ling Ho H. Plant Protein Kinase and Protein-Protein Interaction. *J Biodivers Biopros Dev.* (2014) 2:142
45. Praveen Mannam, Anup Srivastava, Jaya Prakash Sugunraj, Patty J Lee, Maor Sauler. Oxidants in Acute and Chronic Lung Disease. *J Blood Lymph.* (2014) 4:128.
46. Serteyn D, Pouyade G R, Sandersen C, Salciccia A, Grulke S, et al. Muscle Mitochondrial Dysfunction in Horses Affected by Acute Laminitis. *Bioenergetics.* (2014) 3:120.
47. Nichols TW. Hyperphosphorylation of Tau Protein in Downâ€™s Dementia and Alzheimerâ€™s Disease: Methylation and Implications in Prevention and Therapy. *J Alzheimers Dis Parkinsonism.* (2014) 4: 159.
48. Palego L, Betti L, Giannaccini G. Sulfur Metabolism and Sulfur- Containing Amino Acids: I- Molecular Effectors. *Biochem Pharmacol (Los Angel).* (2015) 4:158.
49. Zhang Q, Wu H, Zheng H. Aberrantly Methylated CpG Island Detection in Colon Cancer. *J Proteomics Bioinform.* (2015) S9:007.
50. Laxton RC, Doey L, Aizpurua M, Bodi I, King A, et al. Co-deletion of 1p/19q is Strongly Correlated with a High Level of MGMT Promoter Methylation in High Grade Gliomas as Revealed by Pyrosequencing. *J Mol Genet Med.* (2015) 9:151.
51. Young W, Dâ€™Souza SL, Lemischka IR, Schaniel C . Patientspecific Induced Pluripotent Stem Cells as a Platform for Disease Modeling, Drug Discovery and Precision Personalized Medicine. *J Stem Cell Res Ther.* (2012) S10:010.
52. Das S, Jena S, Kim EM, Zavazava N, Levasseur DN, et al. Transcriptional Regulation of Human NANOG by Alternate Promoters in Embryonic Stem Cells. *J Stem Cell Res Ther.* (2012) S10:009.
53. Hosoda A, Komaba A, Kishimoto M, Tamura H . Combination of Reverse Transcription and Multienzyme Restriction Fragment Length Polymorphism Analysis for Rapid Detection of Escherichia Coli. *J Microb Biochem Technol.* (2013) 6:001-008.
54. Almasi MA, Dehabadi SH .Colorimetric Immunocapture Reverse Transcription Loop-Mediated Isothermal Amplification Assay for Rapid Detection of the Potato virus Y. *J Plant Pathol Microb.* (2013) 4:188
55. Tolmachov OE, Tolmachova T. Design and Production of mRNA-based Gene Vectors for Therapeutic Reprogramming of Cell Fate. *Gene Technology.* (2015) 4:117.
56. Kumar M, Matta A, Masui O, Srivastava G, Kaur J, et al. Interactome Analysis Reveals Novel Binding Partners of Heterogeneous Nuclear Ribonucleoproteind in Oral Cancer and its Increased Nuclear Localization is A Predictor of Poor Disease Prognosis. *J Carcinog Mutagen.* (2015) 6:211.
57. Banerjee BD, Mustafa MD, Sharma T, Tyagi V, Ahmed RS, et al. Assessment of Toxicogenomic Risk Factors in Etiology of Preterm Delivery. *Reprod Syst Sex Disord.* (2014) 3:129
58. Hafidh S, Potesil D, Zdrahal Z, Honys D . DEAD-Box RNA Helicases are among the Constituents of the Tobacco Pollen mRNA Storing Bodies. *J Plant Biochem Physiol.* (2013) 1:114.
59. Hussein KR, Waines PL, Nisir RB, Glegg G, Bradley G . Development and use of Bacteroides 16S rRNA Polymerase Chain Reaction Assay for Source Tracking Dog Faecal Pollution in Bathing Waters. *Hydrobiol Current Res.* (2014) 5:163.
60. Wang Y, Zhang G, Li F, Kan L, Ben J, et al. Altered Expression and DNA Methylation Profiles of ERCC6 Gene in Lens Tissuefrom Age-Related Cortical Cataract. *J Clin Exp Ophthalmol.* (2015) 6:392.

61. Jafar E, Shakibaie MR, Poormasoomi L. Isolation of a Novel Antibiotic Resistance Plasmid DNA from Hospital Isolates of *Pseudomonas aeruginosa*. *J Clin Exp Pathol.* (2013) 3:140.
62. Khan MS, Akhtar N, Haque ME, Barua A, Chowdhury T, et al. Isolation and Identification of non-plasmid Multidrug Resistant *E.coli* from Poultry Wastes in Chittagong Region, Bangladesh. *J Bacteriol Parasitol.* (2014) 5: 182.
63. Barseem NF, El- Samalehy MF, Kasemy ZA . Transcription Factor 7-like 2 (TCF7L2) rs7903146 Polymorphism, Association with Type 2 Diabetes Mellitus Susceptibility. *J Obes Weight Loss Ther.* (2015) 5:250.
64. Takano S, Matsuda S, Hirayama Y, Sato T, Takamura I, et al. Genome-Wide Comparative Transcriptional Analysis of Developing Seeds among Seven *Oryza sativa* L. Subsp. Japonica Cultivars Grown near the Northern Limit of Rice Cultivation. *J Rice Res.* (2015) 3: 130.
65. Rotratsirikun Y, Talmadge RJ . Rapid Changes in NFAT-directed Transcriptional Activity after Muscle Paralysis Induced by a Spinal Cord Injury. *Int J Phys Med Rehabil.* (2014) 2:241.
66. Adekunle OC, Onilude AA . Antimicrobial Resistance and Plasmid Profiles of *Campylobacter* Species from Infants Presenting with Diarrhoea in Osun State, Nigeria. *J Med Microb Diagn.* (2015) 4:172.
67. Adejo GO, Atawodi SE. Acute Toxicity and Genotoxic Effects of all Parts of *Morinda lucida* Benth on pUC18 Plasmid DNA . *Nat Prod Chem Res.* (2014) S1: 006.
68. Bhaduri S, Smith JL, Phillips JG . Evaluation of the Efficiency of a Congo-Red Uptake Technique for Detection and Isolation of Plasmid-Bearing (P_{ly}) *Yersinia pestis* KIM5 in Retail Raw Ground Beef and Pork. *J Food Process Technol.* (2014) 5:375.
69. Panelli D, Lorusso FP, Papa F, Sardanelli AM, Papa S. Alternative Splicing and Nonsense Mediated Decay in Mitochondrial Complex-I Biogenesis and its Implication in Human Diseases. *J Bioanal Biomed.* (2013) S3:006.
70. .Almasi MA, Jafary H, Moradi A, Zand N, Ojaghkandi MA, et al. Detection of Coat Protein Gene of the Potato Leafroll Virus by Reverse Transcription Loop-Mediated Isothermal Amplification. *J Plant Pathol Microb.* (2013) 4:156.
71. Takemoto S. My Views and Thoughts on Translational Medicine. *Transl Med.* (2015) 5:147.
72. Meyer C, Hill K, Hill S, Dow B . Translating Falls Prevention Knowledge for Community-Dwelling People Living With Dementia: Design Protocol for a Mixed-Method Intervention. *J Alzheimers Dis Parkinsonism.* (2015) 5:185. Adams GW .Novel Concepts for the Application of Rapid DNA Technology as a Sentinel Event Prophylactic in the Criminal Justice System. *J Forensic Res.* (2015) 6:1000278.
73. Dmitriev LF, Titov VN . DNA Replication and Telomere Shortening: Key Factors Related to the Production of C3-Aldehydes and the Interaction of One of them with DNA Guanine Residues. *J Gerontol Geriat Res.* (2014) 3:175.
74. Ouwerkerk W, Zwinderman AH. Alternative Splice Variants in Gene Expression Values in Patients with Marfanâ€™s Syndrome. *J Proteomics Bioinform.* (2015) 8:001-008.
75. Shankarling GS. Regulation of pre-mRNA Alternative Splicing by the RNA Processing Factor hnRNPL. *J Biomol Res Ther.* (2013) 2:e118.
76. Sablok G, Harikrishna JA, Min XJ . Next Generation Sequencing for Better Understanding Alternative Splicing: Way Ahead for Model and Non-Model Plants. *Transcriptomics.* (2013) 1:e103 Morris KM, Cheng Y,

- Warren WC, Papenfuss AT, Belov K . Identification and Analysis of Divergent Immune Gene Families within the Devil Genome. *Immunome Res.* (2015) 11:088.
77. Arancio W, Genovese SI, Pizzolanti G, Giordano C .Hutchinson Gilford Progeria Syndrome: A Therapeutic Approach via Adenoviral Delivery of CRISPR/cas Genome Editing System. *J Genet Syndr Gene Ther.* (2015) 6:256.
78. Takano S, Matsuda S, Hirayama Y, Sato T, Takamure I, et al. Genome-Wide Comparative Transcriptional Analysis of Developing Seeds among Seven *Oryza sativa* L. Subsp. Japonica Cultivars Grown near the Northern Limit of Rice Cultivation. *J Rice Res.* (2015) 3: 130.
79. Daoud SS. Genome Wide Identification of FGFR2 Alternative Splicing in Hepatitis C: Potential Roles in Malignant transformation. *J Pharmacogenomics Pharmacoproteomics.* (2014) 5: e140.
80. El-Sebay HM, Badr EAE, El-Ghobashi Y, Khalil MM, El-Mashad GM. Specific IgE Antibodies in Infant with Cow's Milk Protein Allergy. *J Nutr Food Sci .* (2015) 5:350.
81. Lukiw WJ, Zhao Y and Dua P. Microbial Sources of Amyloid and Relevance to Amyloidogenesis and Alzheimer's Disease (AD). *J Alzheimers Dis Parkinsonism.* (2015) 5:177.
82. Rice KM, Fannin J, Para R, Thulluri S, Triest W, et al. Age-Associated Alterations of Morphology and Protein Signaling in the Female F344xBN Rat Aorta. *J Gerontol Geriatr Res.* (2015) 4:196.