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PCR Technique with its Application

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ABSTRACT

PCR (Polymerase Chain Reaction) is a revolutionary method developed by Kary B Mullis (awarded Nobel Prize for chemistry in 1993) in the 1983. PCR is based on using the ability of DNA polymerase to synthesize new strand of DNA complementary to the template strand of DNA. This technique is having an impact on many areas of molecular cloning, genetics, recombinant DNA research molecular biology, forensic analysis, evolutionary biology, and medical diagnostics.

INTRODUCTION

The Polymerase Chain Reaction (PCR) is a method of replicating DNA, it makes numerous copies of a specific segment of DNA quickly and accurately [1]. It is capable of taking a small amount of DNA or even single molecule and amplifying a specific region exponentially such that once the reaction is finished, there may exist up to 230 copies of each starting DNA molecule. Before the development of PCR, the methods used to amplify, or generate copies of recombinant DNA fragments were time-consuming and labour-intensive. But PCR reactions can complete many rounds of replication and produce billions of copies of a DNA fragment only in few hours [2-4].

The Basics of PCR Cycling

The three major steps in a PCR cycling reactions, which are repeated upto 20 to 40 cycles. It is always done on an automated thermo cycler, which has ability to heat and cool the reaction tubes in a very short period of time [5,6]. (Figure 1)

- Denaturation (95 °C), 30 sec.
- Annealing (55–60 °C), 30 sec.
- Extension (72 °C), time depends on product size.

Denaturation (94 °C)

During this stage, the double strand melts open to form single stranded DNA, all enzymatic reactions stop [7].

Annealing (54°C)

Hydrogen bonds are constantly formed and broken between the single stranded primer and the single stranded template. If the primers exactly fit the template, the hydrogen bonds formed are so strong that the primer stays attached [8].

Extension (72°C)

The bases (complementary to the template) are coupled to the primer on the 3' side (the polymerase adds dNTP's from 5' to 3'side, reading the template from 3' to 5' side, bases are added complementary to the template) [9-11].

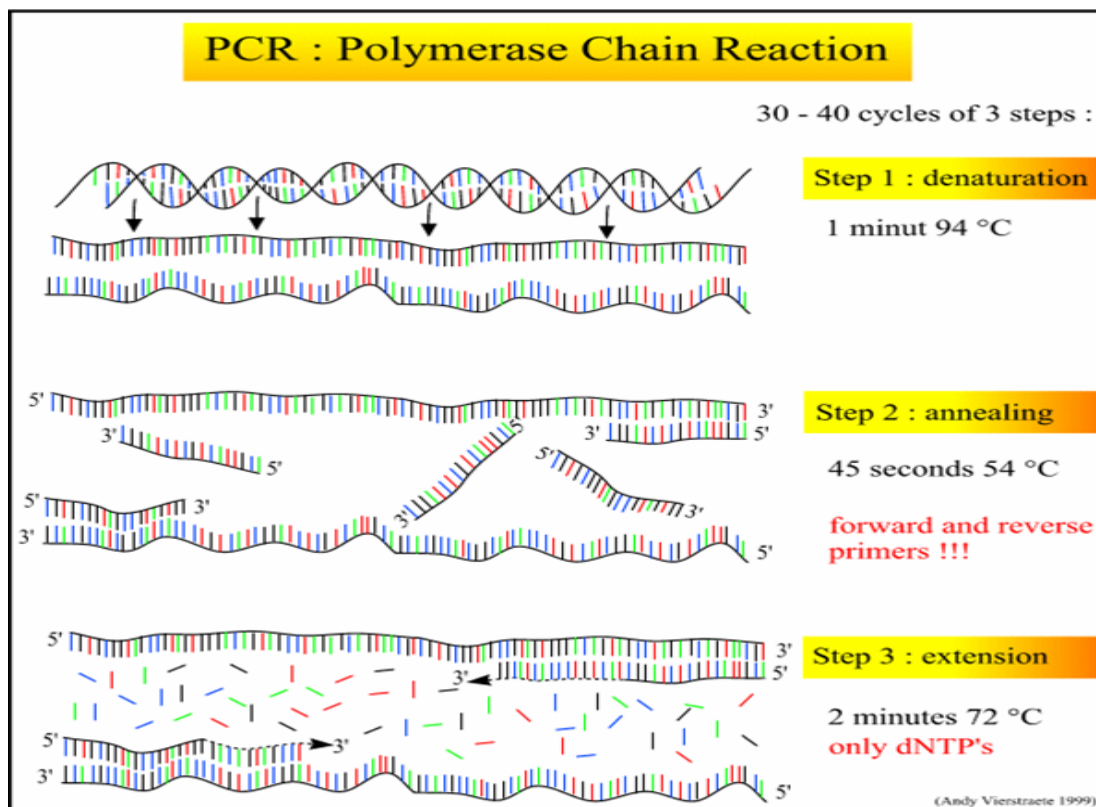


Figure 1: Basics of PCR Cycling.

The PCR technique is based on process, a cell uses to replicate a new DNA strand. The integral component is the template DNA (contains the region to be copied). Even single DNA molecule can serve as a template [12]. The fragment needed for this to be replicated is the sequence of two short regions of nucleotides at either end of the region of interest. The two short template sequences must be known hence two primers (short stretches of nucleotides) that correspond to the template sequences can be synthesized. The primers anneal to the template at their complementary sites and act as the starting point for copying [13]. DNA synthesis at one primer is directed toward the other thus resulting in replication of the desired sequence. Also needed are free nucleotides used to build the new DNA strands and a DNA polymerase does the building by sequentially adding on free nucleotides according to the instructions of the template [14-15].

Every cycle results in a doubling of the number of strands DNA present. After starting stage of few cycles, most of the product DNA strands made are the same length as the distance between the primers. The result is amplification of DNA that exists between the primers [16]. The amount of amplification is 2 raised to the n power; n represents the number of cycles that are performed. After 20 cycles, this would give approximately 1 million fold amplification. After 40 cycles the amplification would be 1×10^{12} [17,18]. (Figure 2)

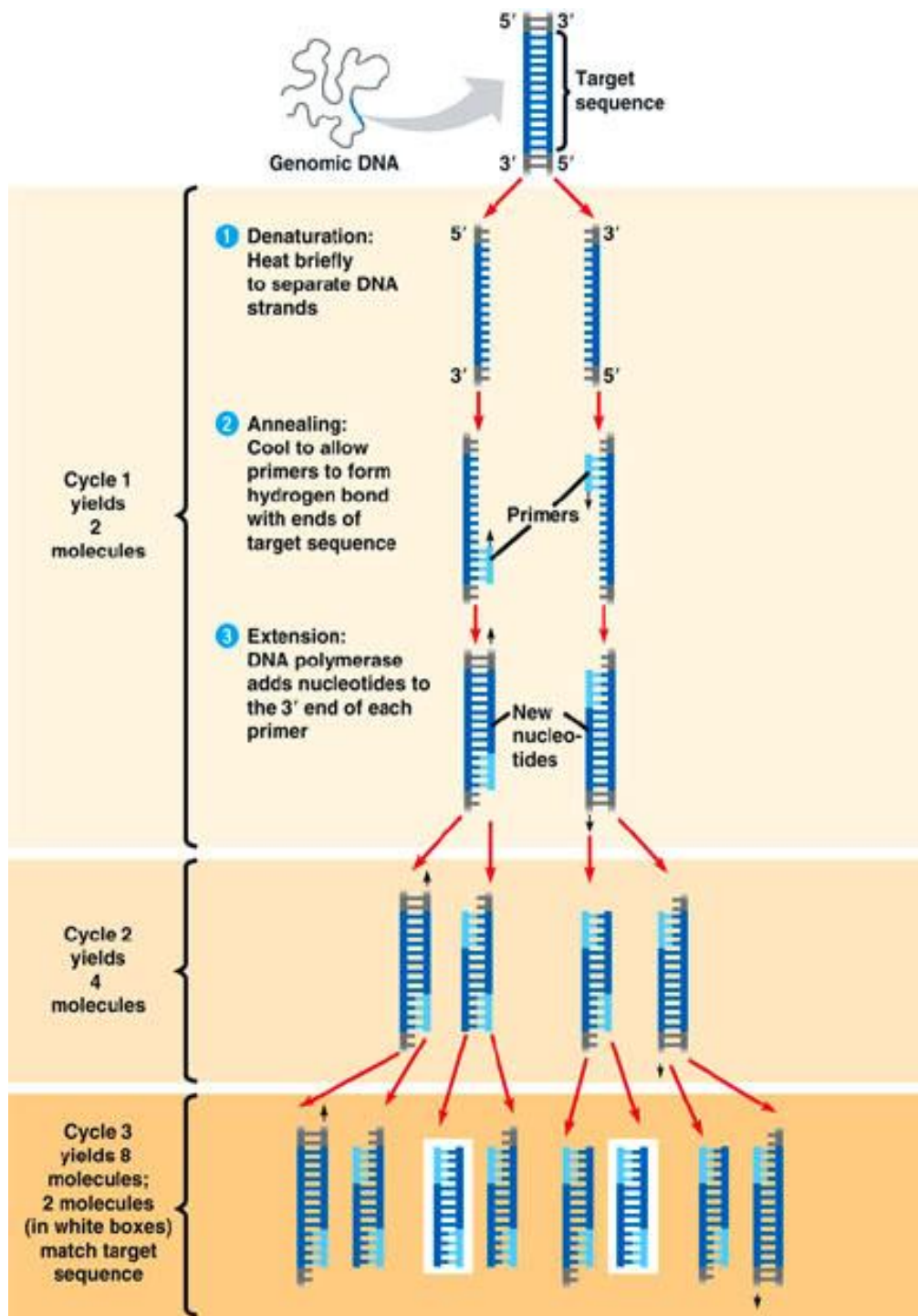


Figure 2: The amount of amplification after 3 cycles.

PREPARATION OF 18 MICRO LITRES REACTION MIXTURE

- | | |
|---------------------------|-------------------|
| • 1mM Nucleotides (dNTPs) | 2.5 micro litres |
| • Taq DNA polymerase | 0.4 micro litre |
| • 10X Taq Buffer | 2.0 micro litres |
| • Sterile water | 13.1 micro litres |

For each tube

- 18 micro litres of reaction mixture

- 1 micro litre of DNA sample
- 1 micro litre of primer

Altogether each tube contains 20 micro litres of sample, This tube is subjected to pcr procedure by following steps of pcr cycling ,the target DNA is amplified. This procedure takes 3 to 4 hrs to complete 40 cycles [19-23].

DIFFERENT TYPES OF PCR

Nested PCR
RT-PCR or Reverse Transcriptase PCR
Real Time PCR
Gradient PCR
Multiplex PCR
AFLP PCR
Allele-Specific PCR
Assembly PCR
Assymetric PCR
Colony PCR
Hot Start PCR
Inverse PCR
In Situ PCR
ISSR PCR
Late-PCR
Long PCR
Single Cell PCR
Standard PCR [24-28]

Capacity of Few Different Types of PCR

- Inverse PCR otherwise called IPCR and was initially depicted by Ochman et al. in 1988. The primary constraint of standard PCR is that 5' and 3' flanking areas of DNA part of interest must be known however in converse PCR permits you to lead PCR when just data of one interior succession is given [29].
- Reverse translation polymerase chain response (RT-PCR) is the most delicate procedure utilized for mRNA discovery and quantitation at present. Contrasted with the Northern blotch examination and RNase security measure which are fundamentally utilized for evaluating mRNA levels RT-PCR can likewise be utilized to evaluate mRNA levels from littler examples [30]. RT PCR is so touchy and effective that it can measure mRNA from a solitary cell. RT-PCR is thought to be most quick, effective and profoundly touchy to infection confinement and is very prescribed as an essential instrument for infection recognition [31].
- Multiplex RT PCR is a method utilized for concurrent enhancement of more than one objective arrangement in a solitary response tube utilizing different groundwork sets. This RT PCR is sufficiently touchy however it has numerous objective arrangements [32]. Multiplex RT PCR is effective furthermore spares the bother of doing numerous PCR responses. Multiplex RT PCR is a strategy utilized for Gene Deletion and Mutation Detection [33].
- Nested polymerase chain response is an adjustment of polymerase bind response expected to decrease the defilement in items because of the intensification of surprising groundwork tying locales [34].
- Polymerase chain response itself is the procedure used to intensify DNA tests, by means of a temperature-interceded DNA polymerase. The items can be utilized for sequencing or

examination, and this procedure is a key piece of numerous hereditary qualities research labs, alongside utilizations in DNA fingerprinting for criminology and other human hereditary cases [35]. Routine PCR obliges groundworks correlative to the ends of the objective DNA. A regularly happening issue is preliminaries tying to inaccurate districts of the DNA, giving unforeseen items [36].

- Nested polymerase chain response includes two arrangements of preliminaries, utilized as a part of two progressive keeps running of polymerase chain response, the second set expected to intensify an auxiliary focus inside of the first run item [37].

Process

The objective DNA experiences the first keep running of polymerase chain response with the first arrangement of groundworks. The choice of option and comparable groundwork tying locales gives a determination of items, stand out containing the planned succession. The item from the first response experiences a second keep running with the second arrangement of preliminaries. It is impossible that any of the undesirable PCR items contain tying destinations for both the new preliminaries, guaranteeing the item from the second PCR has little sully from undesirable results of groundwork dimers and option preliminary target groupings [38-40].

- Reverse interpretation PCR (RT-PCR) utilizes a couple of groundworks which are integral to a characterized grouping on each of the two strands of the cDNA [41]. These preliminaries are reached out by a DNA polymerase and a duplicate of the strand is made after every cycle, prompting exponential enhancement [42].
- RT-PCR incorporates three noteworthy steps. The main step is reverse interpretation (RT), in which RNA is opposite translated to cDNA utilizing converse transcriptase. This stride is imperative keeping in mind the end goal to perform PCR since DNA polymerase can act just on DNA layouts [43]. The RT step can be performed either in the same tube with PCR (one-stage PCR) or in a different one (two-stage PCR) utilizing a temperature somewhere around 40 °C and 50 °C, contingent upon the properties of the opposite transcriptase utilized .All the remaining strides in pcr amplication is same [44].
- Multiplex polymerase chain response (Multiplex PCR) is a change of polymerase anchor response with a specific end goal to quickly recognize cancellations or duplications in a substantial quality. This procedure increases genomic DNA tests utilizing different preliminaries and a temperature-interceded DNA polymerase in a warm cyler [45]. Multiplex-PCR was initially portrayed in 1988 as a technique to recognize erasures in the dystrophin gene.It has likewise been utilized with the steroid sulfatase gene [46].

APPLICATIONS OF PCR

- The polymerase chain response is utilized by a wide range of researchers in a continually expanding scope of experimental orders. In microbiology and atomic science, for instance, PCR is utilized as a part of exploration labs in DNA cloning techniques, Southern smudging, DNA sequencing, recombinant DNA innovation, to give some examples. In clinical microbiology labs PCR is priceless for the analysis of microbial diseases and epidemiological studies. In nourishment science PCR has turn out to be progressively essential to the horticultural and sustenance businesses as an important distinct option for customary identification strategies [47]. PCR is additionally utilized as a part of crime scene investigation labs and is particularly valuable on the grounds that just a little measure of unique DNA is needed, for instance, adequate DNA can be gotten from a bead of blood or a solitary hair [48].
- Constant PCR (or qPCR) is right now utilized as a part of all applications set up of customary, legacy PCR. Constant PCR has applications in all branches of organic science. Applications

incorporate agrarian and nourishment commercial ventures, quality expression examination, the conclusion of irresistible ailment and human hereditary testing [49]. Because of their capacity in fluorimetry the continuous machines are additionally good with option enhancement routines [50], for example, NASBA gave a fluorescence end-point is accessible. A portion of the use of PCR are as per the following

- PCR can be utilized for hereditary testing, where an example of DNA is broke down for the vicinity of hereditary illness transformations [51].
- PCR can be utilized as a major aspect of a touchy test for tissue writing, key to organ transplantation [52].
- PCR can be utilized for HIV test (vicinity of the HIV infection that causes AIDS can be resolved utilizing PCR on platelets. PCR tests have been created with the goal that it can recognize one viral genome among the DNA of more than 50,000 host cells [53].
- PCR can be utilized for Genetic fingerprinting (scientific science) can remarkably separate one individual from the whole populace of the world [54]. Moment tests of DNA can be detached from a wrongdoing scene and contrasted with that from suspects or from a DNA database of prior confirmation or convicts [55].
- PCR can be utilized for DNA fingerprinting can help in Parental testing (DNA sequencing) [56].
- PCR can be utilized for DNA cloning - It can concentrate fragments for insertion into a vector from a bigger genome, which may be accessible in little amounts [57].
- PCR can be utilized for the investigation of examples of quality expression. Tissues or individual cells can be examined at diverse stages to see which qualities have get to be dynamic or which have been exchanged off [58].
- PCR can at the same time increase a few loci from individual sperm has significantly upgraded hereditary mapping by considering chromosomal hybrids after meiosis [59].
- PCR can be utilized to build up connections among species in human studies and developmental science [60].
- PCR can be utilized to help recognize antiquated human stays in paleohistory.
- PCR can be utilized PCR to intensify DNA from terminated bugs saved in golden for 20 million years [61].
- PCR can be utilized to identify the vicinity of a quality moved into a living being (transgene) [62].
- PCR can be utilized to focus the sex of developing lives. Accordingly sex of in vitro treated steers incipient organisms could be resolved utilizing Y chromosome particular preliminaries before their implantation in the uterus [63].
- PCR innovation encourages the location of DNA or RNA of pathogenic living beings and, thusly, helps in clinical analytic tests for a scope of irresistible specialists like infections, microbes, protozoa [64] and so on. These PCR-based tests have various points of interest over routine counter acting agent based indicative systems that focus the body's resistant reaction to a pathogen [65]. Specifically, PCR-based tests are equipped to distinguish the vicinity of pathogenic operators ahead of time than serologically-based routines, as patients can take weeks to create antibodies against an infectious specialists [66]. PCR-based tests have been produced to specify the measure of infection in a man's blood ('viral burden') in this manner permitting doctors to check their patients' sickness movement and reaction to treatment. This has fantastic potential for enhancing the clinical administration of ailments created by popular disease, including AIDS [67] and hepatitis [68], appraisal of viral load all through and after treatment [69].
- PCR-based diagnostics tests are accessible for identifying and/or evaluating various pathogens, including:

1. HIV-1, which causes AIDS
 2. Hepatitis B and C infections, may prompt liver malignancy
 3. Human Papillomavirus, may bring about cervical growth
 4. Chlamydia trachomatis, may prompt fruitlessness in ladies
 5. Neisseria gonorrhoeae, may prompt pelvic provocative illness in ladies
 6. Cytomegalovirus, may bring about existence debilitating illness in transplant patients and other immunocompromised individuals, including HIV-1/AIDS patients
 7. Mycobacterium tuberculosis, which in its dynamic state causes tuberculosis and can prompt tissue harm of tainted organs ^[70-75].
- The utilization of PCR in diagnosing hereditary maladies, whether because of intrinsic hereditary changes or as a consequence of a characteristic hereditary transformations, is turning out to be more normal. Irregularity can be analyzed even preceding conception. Single-strand conformity polymorphism (SSCP), or single-strand chain polymorphism, is characterized as conformational contrast of single-stranded nucleotide groupings of indistinguishable length as actuated by contrasts in the arrangements under certain trial conditions ^[76]. Nowadays, SSCP is most pertinent as an analytic instrument in sub-atomic science. It can be utilized as a part of genotyping to recognize homozygous people of diverse allelic states, and also heterozygous people who acquire hereditary abnormalities ^[77].
 - Genetic advising is ruined the folks to check the record of hereditary malady in advance to settle on a choice on having kids ^[78]. This is obviously administered by national laws and rules. Identification of hereditary malady before implantation of an incipient organism in IVF (In vitro preparation) otherwise called preimplantation analysis should likewise be possible abusing PCR based strategy. Further to analyze acquired or an unconstrained sickness, either symptomatic or asymptomatic (as a result of family history like Duchene solid dystrophy) PCR based technique is exceptionally valuable ^[79].
 - Genetic finger impression is a standout amongst the most abused utilization of PCR (otherwise called DNA profiling). Profiles of particular extends of DNA are utilized as a part of hereditary fingerprinting (by and large 13 loci are looked at) which is contrast from individual to individual ^[80]. PCR additionally assumes a part in examination of genomic or mitochondrial DNA, in which examiners utilized examples from hair shafts and bones when different specimens are not open ^[81].
 - PCR is a fundamental method in cloning technique which permits era of a lot of unadulterated DNA from minor measure of format strand and further investigation of a specific quality. A few changes to the PCR convention can create transformations (general or site-coordinated) in a grouping either by an embedded part or base change ^[82]. PCR is utilized for sequence-labeled destinations (STSs) as a pointer that a specific section of a genome is available in a specific clone. A typical utilization of Real-time PCR is the investigation of expression examples of qualities amid diverse formative stages. PCR can likewise explore 'ON or OFF' of specific qualities at diverse stages in tissues (or even in individual cells) ^[83].
 - PCR has various applications in different fields. The Human Genome Project (HGP) for deciding the grouping of the 3 billion base combines in the human genome, depended vigorously on PCR. The qualities connected with an assortment of infections have been distinguished utilizing PCR. For instance, Duchenne strong dystrophy, which is created by the change of a quality, recognized by a PCR strategy called Multiplex PCR ^[84]. PCR can help to study for DNA from different life forms, for example, infections or microscopic organisms. PCR has been utilized to distinguish and to investigate connections among species in the field of transformative science. In human studies, it is likewise used to comprehend the antiquated human movement

designs. In paleontology, it has been utilized to recognize the antiquated human race [85,86]. PCR normally utilized by Paleontologists to open up DNA from terminated species or cryopreserved fossils of millions of years and accordingly can be further studied to elucidate on.

REFERENCES

1. Flanagan PV et al. Dynamic Changes in Aromatic Hydrocarbon Associated Catabolic Gene Profiles Linked to Aerobic and Anaerobic Microcosm Studies. *Innovative Energy Policies*. 2014; 3:111.
2. Flanagan PV et al. Dynamic Changes in Aromatic Hydrocarbon Associated Catabolic Gene Profiles Linked to Aerobic and Anaerobic Microcosm Studies. *Innovative Energy Policies*. 2014; 3:111.
3. Alfonso Y et al. Multiplex PCR to Detect *T. gondii* Infection based on B1 Gene and 529 bp Repetitive Element. *J AIDS Clin Res*. 2015; 6:435.
4. Alibi S et al. Identification of Clinical *Corynebacterium striatum* Strains by PCR Restriction Analysis Using the RNA Polymerase β subunit gene (*rpoB*). *J Bacteriol Parasitol*. 2015; 6:21-9.
5. Magd MAE et al. Effect of SNPs in Prolactin Promoter on Milk Traits in Egyptian Buffalo. *J Adv Dairy Res*. 2015; 3:128.
6. Ramesan K and Ramegowda D. Novel technique for plant irrigation using dynamic head of evaporation. *J AIDS Clin Res*. 2013; 2(8):1-8.
7. Sultana A et al. Antibiotic resistance molecular diversity verotoxin characterization of *e coli* isolated from clinical samples. *J AIDS Clin Res*. 2014; 3(2):661-5.
8. Verma S et al. Current scenario of *Escherichia coli* and its serotype O157:H7 in Indian subcontinent. *J AIDS Clin Res*. 2013; 2(7):125-9.
9. Ahangari G et al. Significant Association between Catecholamine O-Methyltransferase (COMT) Gene Expression Changes and Breast Cancer Pathogenesis. *J Carcinog Mutagen*. 2015; 6:219.
10. Ebrahimzadeh A et al. Allelic Diversity of Polymorphic AMA 1 (Apical Membrane Antigen 1) Vaccine Candidate Antigen of *Plasmodium falciparum* in Two Populations of Imported and Indigenous Cases in South East of Iran using Nested PCR and RFLP. *J Trop Dis*. 2014; 2:149.
11. Minaeva E and Ermilova E. Sequencing and Expression Analysis of the Gene Encoding PII Signal Protein in *Chlorella Variabilis* NC64A. *J Plant Biochem Physiol*. 2015; 3:142.
12. Lim HK et al. Improvement and Immobilization of a new Endo β 1,4 xylanases KRICT PX1 from *Paenibacillus* sp. HPL 001. *J Bioprocess Biotech*. 2015; 5:215.
13. Gawish G E H. The Prevalence of Inherited Thrombophilic Polymorphisms in Saudi Females with Recurrent Pregnancy Loss Confirmed using Different Screening Protocols of PCR. *J Mol Genet Med*. 2015; 9:156.
14. Goto I. Measurement of the Total Number of Bacteria in Saliva Using Quantitative Real Time PCR During Treatment for Head and Neck Malignancy: A Series of Cases. 2015; 14:85-90.
15. Zaâbi S et al. Preliminary Study of Population Genetic Diversity of *Hyalinoecia tubicola* (Polychaeta: Onuphidae) from the North East Coast of Tunisia (Western Mediterranean) using Random Amplified Polymorphic DNA Markers. *J Coast Zone Manag*. 2015; 18:397.
16. Goudarzi AM. Identification of anaerobic rumen fungi using molecular. *J Coast Zone Manag*. 2014; 717-722.
17. Hemati B et al. The study of complex vertebral malformation genetic defect. *J Mol Genet Med*. 2014; 485-487.
18. Pandey A et al. Impact of molecularly diagnose of diseases in some important fruit crops. *J Coast Zone Manag*. 2012; 213-226.
19. Sangamithra A et al. Detection and characterization based on *s* and *s* gene segments. *J Mol Genet Med*. 2014; 335-339.

20. Pandey A et al. Molecular study of the prevalence of feline leukemia virus. *J Carcinog Mutagen.* 2012; 728-733.
21. Ndiaye A et al. Genetic discrimination of two cowpea *vigna unguiculata* l walp bruchid coleoptera chrysolmelidae bruchinae. *J Carcinog Mutagen.* 2011; 196-201.
22. Jehan S and Al-Abrahim. Molecular identification of alfalfa mosaic virus isolated. *J AIDS Clin Res.* 2014; 348-352.
23. Gaur K and Nigam R. Pooled Conventional PCR to Detect *Tritrichomonas Foetus* Infected Beef Bulls. *J Biodivers Endanger Species.* 2015; 3:148.
24. Zeedan GSG et al. Isolation and Molecular Diagnosis of Orf Virus from Small Ruminants and Human in Egypt. *J Antivir Antiretrovir.* 2014; 6:154-161.
25. Elawad HE et al. Microbial Amidases and their Industrial Applications: A Review . *J Med Microb Diagn.* 2015; 4:174.
26. Selim ME et al. Evaluation and Characterization of Some Egyptian *Fusarium oxysporum* Isolates for their Virulence on Tomato and PCR Detection of (SIX) Effector Genes. *J Bioprocess Biotech.* 2015; 5:204.
27. Arkun K et al. Effect of Lewy Bodies on Mitochondrial DNA Copy Numbers and Deletion Burden in Parkinson's Disease Substantia nigra Neurons. *J Alzheimers Dis Parkinsonism.* 2015; 5:175.
28. Osborne AE et al. Palm Fruit Juice Mitigates AZT Mitochondrial Genotoxicity and Dose-Dependent Cytotoxicity. *J AIDS Clin Res.* 2014; 5:400.
29. Stone CB and Mahony JB . Molecular Detection of Bacterial and Viral Pathogens "Where Do We Go From Here?. *Clin Microbiol.* 2014; 3:175.
30. Mazumdar A et al. Correlation of nucleic acid amplification based detection and conventional methods of Identification of *Aspergillus Flavus* Species in Chronic Rhinosinusitis. *J Med Microb.* 2013; 61-66.
31. Hossain MA et al. First Genotype Characterization of *Giardia intestinalis* Assemblage E from Goat Kids in Bangladesh. *J Veterinar Sci Technol.* 2014; 5:201.
32. Mahadhy A et al. PCR-Free Ultrasensitive Capacitive Biosensor for Selective Detection and Quantification of Enterobacteriaceae DNA. *J Anal Bioanal Tech.* 2014; 5:210
33. Venkatesan C and Sahul Hameed AS. Analysis of Immune Genes and Heat Shock Protein Genes under Exposure to White Spot Syndrome Virus (WSSV) and Herbal Immune Stimulant in *Litopenaeus vannamei*. *J Bacteriol Parasitol.* 2014; 5:205.
34. Pradeep J et al. Expression Analysis of Solute Carrier (SLC2A) Genes in Milk Derived Mammary Epithelial Cells during Different Stages of Lactation in Sahiwal (*Bos indicus*) Cows. *J Bacteriol Parasitol.* 2014; 5:210.
35. Perez-Marquez J. SQPrimer: The Utility of Designing Homologous Primers for the Genetic Analysis Based on the PCR. *J Comput Sci Syst Biol.* 2014; 7:229-234.
36. Iyer K. *Tritrichomonas foetus* in Beef Bulls. *J Veterinar Sci Technol.* 2014; 5:194.
37. Zeedan GSG et al. Antimicrobial, Antiviral Activity and GC-MS Analysis of Essential Oil Extracted from *Achillea fragrantissima* Plant Growing In Sinai Peninsula, Egypt. *J Microb Biochem Technol.* 2014; S8:006.
38. Kojouri GA et al. The First Molecular Detection of Caprine Arthritis Encephalitis Virus (Caev) in Iran. *J Veterinar Sci Technol.* 2014; 5:184.
39. Veerasami M et al. Individual and Multiplex PCR Assays for the Detection of Adventitious Bovine and Porcine Viral Genome Contaminants in the Commercial Vaccines and Animal Derived Raw Materials. *J Veterinar Sci Technol.* 2014; 5:179.
40. Jordaan G et al Identification of Histone Epigenetic Modifications with Chromatin Immunoprecipitation PCR Array in Chronic Lymphocytic Leukemia Specimens. *J Cancer Sci Ther.* 2014; 6:325-332.

41. Abdel-Ghaffer H et al. Age-Dependent Gene Expression of Blow Fly *Lucilia cuprina* (Diptera: Calliphoridae) during Egg Development Improving Age Estimation in Forensic Entomology. *J Forensic Res.* 2014; 5:238.
42. Lokman N et al. Mutation Analysis of Gastrointestinal Stromal Tumors in a Pathology Laboratory with 42 Cases of Formalin-Fixed Paraffin-Embedded Specimens. *J Clin Exp Pathol.* 2014; 4:185.
43. Lokman N et al. pcr antigen detection and peripheral smear for diagnosis of malaria. *J Clin Exp Pathol.* 2014; 4:185.
44. <http://omicsonline.com/open-access/need-for-pcr-analysis-in-assessing-severe-malaria-infections-with-plasmodium-vivax.pdf?aid=29633>
45. Chen D et al. Development of Two Multiplex Real-Time PCR Assays for the Rapid Detection of RNA and DNA Viruses Associated with Gastroenteritis in Pediatric Patients. *Pediat Therapeut.* 2014; 4:208.
46. Sarajlić A. A Proposal for Advanced Flow based Charging in PCC Architecture. *J Telecommun Syst Manage.* 2014; 3:112.
47. Ahmed OB et al. Detection of Salmonella in Food Samples by Culture and Polymerase Chain Reaction Methods. *J Bacteriol Parasitol.* 2014; 5:187.
48. Hewedy OA et al. Phenotypic and Genotypic Diversity of Rhizobia Nodulating Faba Bean from Various Egyptian Locations. *J Bioprocess Biotech.* 2014; 4:170.
49. Tetrault et al. Single- Tube Mutation Scanning of the Epidermal Growth Factor Receptor Gene using Multiplex LATE-PCR and Lights-On/Lights-Off Probes. *J Mol Biomark Diagn.* 2014; 5:175.
50. Mishra RK et al. Direct Colony Nested-PCR for the Detection of *Fusarium oxysporum* f. sp. *Psidii* Causing Wilt Disease in *Psidium guajava* L.. *J Horticulture.* 2014; 1:105.
51. Hartshorne T et al. A High-throughput Real-time PCR Approach to Pharmacogenomics Studies. *J Pharmacogenomics Pharmacoproteomics.* 2014; 5:133.
52. Valenzuela-Muñoz V et al. Molecular Characterization and Transcription Analysis of P-Glycoprotein Gene from the Salmon Louse *Caligus rogercresseyi*. *J Aquac Res Development.* 2014; 5:236
53. Barnor JS et al. Establishment of In-House Quantitative Real-Time RT-PCR Assay for HIV-1 Viral Load Measurement: Application to Evaluate Efficacy of ART in Ghanaian Patients in an Urban Setting . *J AIDS Clin Res.* 2014; 5:305.
54. Poh JJ and Gan SKE. The Determination of Factors Involved in Column-Based Nucleic Acid Extraction and Purification. *J Bioprocess Biotech.* 2014; 4:157
55. Ebrahimzadeh A et al. Nested Polymerase Chain Reaction (PCR) on Fixed Stained Slides in Comparison to Whole Blood as a Source of DNA in Southeast of Iran. *J Trop Dis.* 2014; 2:136.
56. Noorali S et al. Effect of Differentially Expressed MicroRNAs 602 and 323-5p on Hepatitis C Virus Genotype 1b Viral Load in Infected Liver Cells. *J Infect Dis Ther.* 2014; 2:138.
57. Claudia V et al. Targeting Cancer Related Genes by Multiplex PCR Followed by High Throughput Parallel Sequencing. *Int J Genomic Med.* 2014; 2:115.
58. Mariani BMP et al. The Use of Real Time PCR Genotyping to Detect Activating GNAS Mutations in McCune-Albright Syndrome. *Hereditary Genet.* 2014; 3:126.
59. Bendtsen M et al. An Anti TNF-A Receptor Antagonist does not Augment the Effect of Autologous Mesenchymal Stem Cell Therapy in Experimental Intervertebral Disc Degeneration in Göttingen Minipigs. *J Stem Cell Res Ther.* 2014; 4: 187.
60. Liuzzi JP. Up-Regulation of miR-34a by Zinc Deficiency. *Vitam Miner.* 2014; 3:119.
61. Shahid M et al. Molecular Characterization of Trichodermaviride Isolated from Rhizospheric Soils of Uttar Pradesh Based on rDNA Markers and Analysis of Their PCR-ISSR Profiles. *J Mol Biomark Diagn.* 2014; 5:169.
62. Jaayid TA et al. Sex Determination of Fetus Prenatal from Maternal Plasma in Goats Using Duplex PCR. *Clon Transgen.* 2014; 3:123.

63. Hatta M and Sultan AR. Main Issues Behind our Ongoing Failure against Tuberculosis. *J Mycobac Dis.* 2013; 4:e125.
64. Park SW and Wei LN. Can GPCR, such as Kappa Opioid Receptor, be a Viable Therapeutic Target for Reducing or Preventing Neuroinflammation? *Clin Exp Pharmacol.* 2014; 4:e127.
65. Iida T et al. Calcified Granulomatous Lung Lesions Contain Abundant Mycobacterium tuberculosis Components. *J Mycobac Dis.* 2014; 4:142.
66. Zar MS et al. Forensic DNA Typing of Old Skeletal Remains Using AmpFISTR® Identifiler® PCR Amplification Kit. *J Forensic Res.* 2013; 5: 211.
67. Huws SA et al. Differential Colonization of Plant Parts by the Rumen Microbiota is likely to be due to Different Forage Chemistries. *J Microb Biochem Technol.* 2014; 6: 080-086.
68. Olusegun AF et al. Co- Endemicity of Plasmodium falciparum and HIV-Infections in Treated Patients is Uncorrelated in Benin City, Nigeria. *J Bacteriol Parasitol.* 2013; 4:176.
69. Shakal MA et al. Surveillance on Avian Influenza H5N1 and H9N2 Subtypes In Egypt 2012-2013. *Poult Fish Wildl Sci.* 2013; 2:111.
70. Paziienza M et al. Application of Real-Time PCR to Identify Residual Bio-Decontamination of Confined Environments after Hydrogen Peroxide Vapor Treatment: Preliminary Results. *J Microb Biochem Technol.* 2013; 6:024-028.
71. Heifetz A et al. From Receptors to Ligands: Fragment-assisted Drug Design for GPCRs Applied to the Discovery of H3 and H4 Receptor Antagonists. *Med chem.* 2013; 4:313-321.
72. Fernandes S et al. Y-Chromosome Detection in Turner Syndrome. *Human Genet Embryol.* 2013; 3:115.
73. Sinnathamby G et al. EDDR1 is a Potential Immunotherapeutic Antigen in Ovarian, Breast, and Prostate Cancer. *J Clin Cell Immunol.* 2011; 2:106.
74. Eroglu F et al. Identification of Causative Species in Cutaneous Leishmaniasis Patients Using PCR-RFLP. *J Bacteriol Parasitol.* 2011; 2:113.
75. Reddy AD et al. Incidence of White Spot Syndrome Virus (Wssv) in Indian Farmed Frozen Shrimp Products and Testing for Viability Through Bio-Inoculation Studies. *J Aquac Res Development.* 2010; 1:102.
76. Mohanty S et al. Genotypic and phenotypic diversity of Bacillus spp. isolated from Freshwater Ecosystems. *J Aquac Res Development.* 2011; 2:112.
77. Tewary A and Patra BC. Oral administration of baker's yeast (*Saccharomyces cerevisiae*) acts as a growth promoter and immunomodulator in *Labeo rohita* (Ham.). *J Aquac Res Development.* 2011; 2:109.
78. Nkeze J et al. Comparison of HIV-1 Viral Load between Abbott m2000 and Roche COBAS TaqMan Methods. *J Antivir Antiretrovir.* 2010; 2: 042-045.
79. Newman JP et al. Migration of Human Fetal Bone Marrow-derived Mesenchymal Stem Cells: Possible Involvement of GPCR and MMPs. *J Stem Cell Res Ther.* 2011; S2:007.
80. Bricker BJ. Past, Present and Future of Molecular Technology Applications for the Epidemiology of Bacterial Diseases. *J Anal Bioanal Tech.* 2010; S10:001.
81. Mei L et al. Malt Genotypic Screening of Polymorphism Information Content (PIC) of SSR Markers Based on Physiological Traits in Barley. *Molecular Biology.* 2012; 1:101.
82. Leino E et al. Quantitation of Megakaryocytic Progenitors in Apheresis Products by Flow Cytometry and Real Time PCR. *J Stem Cell Res Ther.* 2011; S3:001.
83. Banerjee HN et al. Molecular Diagnosis of Helicobacter Pylori Strain by 16S rDNA PCR Amplification and Direct Sequencing. *J Bioprocess Biotechniq.* 2011; 1:105e.
84. Rocco L et al. Genotoxicity in Zebrafish (*Danio rerio*) Exposed to two Pharmacological Products from an Impacted Italian River. *J Environment Analytic Toxicol.* 2011; 1:103.

85. Dave HV et al. Differential Gene Expression Pattern of Transforming Growth Factor Beta-1 in Early and Advanced Breast Cancers. *J Cancer Sci Ther.* 2011; 3: 244-249.
86. Malki Fatima EL et al. Identification and Characterization of Staphylococcus Isolates in Fes-Meknes Region in Morocco. *Pharm Anal Acta.* 2012; S15:003