



## New Molecular Methods for Detection of Bioterrorism Agents

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### Abstract

In this study we reviewed the newly and featured molecular methods which are increasingly used in bioterrorism surveillance. Some cornerstone developments in the detection of bioterrorism agent are; diversities of mainly emergent disease agent, phage engineering as an effective means to detect or eliminate biological agents, PCR and pre-PCR processing which an important tool for surveillance, biological agent detection kits, mass spectrometric detection and Metagenomics which the genetics subfield studying the combined genomes of a sample are now mainly quickly growing fields.

**Key words:** Bioterrorism, disease detection, Metagenomics, biothreat agents.

## INTRODUCTION

Bioterrorism is a term used for terrorism incident using biological agents, such as pathogen microorganisms and biological toxins such as ricin or botulinum. Bioterrorism outbreak works like a natural outbreak and we cannot foretell which disease may spread and when it will begin. Recently we have seen that *Ebola* outbreaks cases first confirmed in Sierra Leone following an outbreak in Guinea, unfortunately there is no cure or vaccine for *Ebola* which one of the world's deadliest viruses. MERS outbreaks also threatened the human beings in Middle East and The bad side of the situation there is no any treatment for these disease too. In the MERS outbreaks camels are thought to be the main carriers of the virus that has killed at least 200 people mainly in the world, but researchers now believe other animals could also be spreading the infection. We must learn to live with this indefiniteness and found ways to protect ourselves against any disease outbreaks. In the field of diagnostic microbiology, rapid molecular methods are critically important for detecting pathogens. With rapid and accurate detection, preventive measures can be put in place early, thereby preventing lives and further spread of any disease. For this necessity we have reviewed featured molecular methods developments in bioterrorism studies.

### Some Cornerstone Developments in Detection of Bioterrorism Agent

#### *Agent diversity works*

When we look at the prominent studies, it is thought that *Bacillus anthracis*, the ethiological agent of anthrax which relatively common throughout the world, can be used as an agent of bioterrorism. For naturally occurring outbreaks scenarios or criminal release of this pathogen a fast and accurate diagnosis is crucial to an effective response and threat mitigation [1]. Microbiological forensic studies and epidemiologic investigations progressively using molecular markers, such as polymorphisms in DNA sequence, to obtain reliable information regarding the identification or source of a suspicious strain. Recently significant research efforts have been undertaken to develop genotyping methods with increased power to differentiate *B. anthracis* or other emergent strains. A growing number of DNA markers have been identified and used to survey any biological agent's diversity in the world, leading to rapid advances in our understanding about the pathogens [1,2,3].

#### *Phage engineering as an effective means to detect and eradicate threats*

Microorganisms can cause widespread disease outbreaks and some of those can be used as agents of bioterrorism. From a biosecurity standpoint, the capacity to detect become key challenges [4]. scientist turned to phage engineering as a potentially highly effective means to both detect and mitigation of threats originating from emergent

bacterial strains [4]. For example they developed technologies allowing us to modify multiple regions of the sequence in a gene while conserving intact the remain, so reversibly interrupt the lytic cycle of T4 within its host. They carry out efficient insertion with homologous recombination and reactivate the lytic cycle and this lead to the production of engineered infective virulent recombinant progeny. This allows the production of very large, genetically engineered lytic phage banks. Screening of such a bank may allow the rapid isolation of recombinant T4 particles capable of detecting, infecting, and destroying hosts bacteria [4,5,6].

#### *PCR processing*

Diagnostic DNA analysis using polymerase chain reaction (PCR) has become a valuable tool for rapid detection of biothreat agents. However, analysis is often challenging because of the limited size, quality, and purity of the biological target [7]. Pre-PCR processing is an integrated concept. sampling methods must maximize target uptake and minimize uptake of extraneous substances that could impair the analysis with PCR inhibitors which must be known [7]. In sample treatment extensive purification leads to lossing of DNA. A cornerstone of pre-PCR processing is to apply DNA polymerase-buffer systems that should tolerant to specific sample impurities, so lowering the need for costly purification steps and maximizing DNA we have. Ineffective sample processing give way to costly and ambiguous results. This situation may hinder the decision process in a bioterrorism crisis. Quality control methods for pre-PCR processing to simplify the analysis of various biothreat agents is important [7-13]. Several quantitative PCR assays also have been developed for biopreparedness which targeting different agents. this methods employ new detection and identification principles. In one, the dissociation pattern of the DNA markers generated with universal primers studied through high-resolution melting and enabled the differentiation of 16S rRNA gene from different microorganisms, so identify 100 relevant bioterrorism agents [22-26)].

#### *Detection kits*

Many different methods based on both immuno assays and PCR-based detection systems have been developed to help detect toxins. Detection kits are commercially available for the detection of different toxins in suspicious samples. While reports does not evaluate the performance of commercially available tests. Such as ricin that is extracted from seeds of the *Ricinus communis* is the most encountered biothreat agent. The seeds yields a highly toxic product that has been used as threats to community. We have rapid detection assays such as lateral flow assays (LFAs). But we have complicating factors associated with LFAs. Such as the incorporation of antibodies of poor specificity that cross-react with near-neighbors or lectins which are cause nonspecific cross-linking for the antibodies [14,15].

#### *Mass spectrometric detection*

Mass spectrometric detection of protein-based toxins such as botulinum neurotoxin, anthrax toxins and ricin which are among the most toxic substances is important. They are composed of 2 polypeptide chains responsible for cell uptake and enzymatic function with the ability to destroy basic cellular functions [16]. Traditionally, large molecules like proteins have been detected using immunological techniques. Although sensitive, these

methods suffer from some drawbacks, such as the risk of false-positives due to crossreactions and detection of inactive toxin. In Recent years, instrumental methods based on mass spectrometry for the reliable detection of botulinum neurotoxins, anthrax toxins and ricin developed. Identification of a protein based toxin can be carried out by mass spectrometry based amino acid sequencing. Furthermore, in combination with antibody affinity preconcentration and biochemical tests with mass spectrometric detection demonstrating the toxin's enzymatic activity, very powerful analytical methods have been described. The advent of sensitive, easily operated mass spectrometers provides new possibilities for the detection of protein-based toxins[16]. The use of synthetic positive control templates containing small modifications outside the primer and probe regions is essential to ensure all aspects of the assay are functioning right. But, a general PCR assay is suffer from differentiating products which generated from positive controls and biological samples because the fluorescent probe signals generated from each type of amplicon are similar. To solve this problem scientists also have developed a new application of electrospray ionization mass spectrometry to rapidly differentiate qPCR amplicons generated with positive biological samples from those generated with synthetic positive controls. The method supports sure determination of the presence of a biothreat agent [16-18].

#### *Metagenomics*

It is important to quickly detect and confirm the specific infection for effective outbreak control. It is estimated that a one week delay in implementing control measures for a respiratory syndrome resulted in a 2.6 fold increase in the epidemic size and an extension of one month. The traditional culture based, immunologic and nucleic acid based methods suffer from drawbacks when dealing with new organisms. There are no culture based methods for all organisms and sometimes methods have poor specificity also. Immunological and nucleic acid based methods have a high specificity, so unable to detect all organisms. But in metagenomics the combined genomes of all organisms present in a sample are analyzed that requires no prior knowledge of the target also. This can be of crucial importance when encountering a new target, as it removes the initial delay caused by identification process. Metagenomics is a growing field in mapping interactions in a microbial community and you aren't need for culturing [19-20].

## **RESULT and DISCUSSION**

Not only anthrax, plaque, botilinum toxins or ricin which we are accustomed in the past but also newly emergent deseases such as ebola or MERS can be used for intentionally contamination of the people. In the situation of bioterror event, rapid molecular methods are critically important for detecting pathogens or toxins. With rapid and accurate detection, preventive measures can be put in place early, thereby preventing loss of life and further spread of any disease. When we look at the prominent studies Over the past decade, significant research efforts have been undertaken to develop genotyping methods to differentiate *B. anthracis* strains which was used in US Following the terrorist attacks of September 11 and the anthrax attacks in 2001. Scientist survey *B. anthracis* diversity in nature, leading to rapid advances in our understanding of the global population of this pathogen [1,2,3]. Phage

engineering as a effective means to detect and eradicate threats is a newly introduced field that we could benefit future in mitigation of bioterrorist event's potential devastating results [4,5,6]. Diagnostic DNA analysis using PCR has become a valuable tool for rapid detection of biothreat agents. We now can use this commonly used tool but when it comes to this bioterrorism topic this tool must be more portable and fast. Because they will not have time to wait a few days long laboratory process [7]. Many different methods based on both immuno assays and PCR based detection systems have been developed and evaluated to help detect toxins and diagnose toxin poisoning in patients but they should develop more. The sampling method should maximize target uptake and minimize uptake of extraneous substances that could impair the analysis [7-13]. Detection kits have also been developed and are commercially available for the detection of bioterrorist agents. Mass spectrometric detection of toxins is also important introduction. It especially useful for proteins so may be used for the bacterial toxins such as botulinum and anthrax as well as the plant toxin ricin which are all protein based toxins [14,15]. Metagenomics gaining importance in studying the combined genomes of a sample and their interaction. It does offer a rapid high throughput method for mapping interactions in a microbial community and circumvents the need for culturing of an agent in a bioterrorism event survey. The use of wholegenome sequencing techniques deserves attention, as it will be an important tool in future bioterrorism and also normal surveillance research. Whole-genome sequencing can be used to detect any microorganism in a sample with one analysis, thus making it truly a technique to detect the unexpected agents [19,20]. It can be foreseen that the introduction of whole-genome sequencing, not only in the field of bioterrorism, but also in the fields of public human health and animal health or as a diagnostic tool used by hospitals and general practitioners, will greatly advance treatment of infectious diseases and human health [21].

## REFERENCES

- [1] Turnbull, PC. 2002. Anthrax history, disease and ecology. In: Koehler TM, ed. Anthrax. Springer-Verlag 1-19.
- [2] Mock M, Mignot T. 2003. Anthrax toxins and the host: a story of intimacy. *Cell Microbiology* 5:15-23.
- [3] Derzelle S, Thierry S. 2013. Genetic Diversity of Bacillus anthracis in Europe: Genotyping Methods in Forensic and Epidemiologic Investigations. *Biosecurity and Bioterrorism* 11(1).
- [4] Pouillot F, Blois H, Iris F. 2010. Genetically Engineered Virulent Phage Banks in the Detection and Control of Emergent Pathogenic Bacteria. *Biosecurity and Bioterrorism* 8(2).
- [5] Summers WC. 2001. Bacteriophage therapy. *Annu Rev Microbiol* 55:438-450.
- [6] Uzan M. 2009. RNA processing and decay in bacteriophage T4. *Prog Mol Biol Transl Sci* 85:43-89.
- [7] Karlsson OE, Bela'k S, Granberg F. 2013. The Effect of Preprocessing by Sequence-Independent, Single-Primer Amplification (SISPA) on Metagenomic Detection of Viruses. *Biosecurity and Bioterrorism* 11(1).
- [8] Fredricks DN, Relman DA. 1999. Application of polymerase chain reaction to the diagnosis of infectious diseases. *Clin Infect Dis*;29(3):475-486
- [9] Hugenholtz P. 2002. Exploring prokaryotic diversity in the genomic era. *Genome Biol* 3(2) review 3.
- [10] Morse SA, Budowle B. 2006. Microbial forensics: application to bioterrorism preparedness and response. *Infect Dis Clin North Am* 20(2):455-473.
- [11] Eisen JA. 2007. Environmental shotgun sequencing: its potential and challenges for studying the hidden world of microbes. *PLoS Biol* 5(3)
- [12] Riesenfeld CS, Schloss PD, Handelsman J. 2004. Metagenomics: genomic analysis of microbial communities. *Annu Rev Genet* 38:525-552.
- [13] Hedman J, Knutsson R, Ansell R, Ra°dstro'm P, Rasmusson B. 2013. Pre-PCR Processing in Bioterrorism Preparedness: Improved Diagnostic Capabilities for Laboratory Response Networks. *Biosecurity and Bioterrorism* 11(1).
- [14] Ramage J G, Prentice KW, Morse SA, Carter A J, Datta S. 2014. Comprehensive Laboratory Evaluation of a Specific Lateral Flow Assay for the Presumptive Identification of Abrin in Suspicious White Powders and Environmental Samples. *Biosecurity and Bioterrorism* 12(1).
- [15] Hodge DR, Prentice KW, Ramage J G, Prezioso S, Gauthier C. 2013. Comprehensive Laboratory Evaluation of a Highly Specific Lateral Flow Assay for the Presumptive Identification of Ricin in Suspicious White Powders and Environmental Samples. *Biosecurity and Bioterrorism* 11(4).
- [16] A ° berg A T, Bjo°rnstad K, Hedeland M. 2013. Mass Spectrometric Detection of Protein-Based Toxins. *Biosecurity and Bioterrorism* 11( 1)
- [17] Demirev PA, Fenselau C. 2008. Mass spectrometry in biodefense. *J Mass Spectrom* 43:1441-1457.
- [18] Lim DV, Simpson JM, Kearns EA, Kramer MF. 2005. Current and developing technologies for monitoring agents of bioterrorism and biowarfare. *Clin Microbiol Rev* 18:583-607.
- [19] Motley ST, Redden CL, Sannes-Lowery KA, Eshoo MW, Hofstadler SA. 2013. Differentiating Microbial Forensic qPCR Target and Control Products by Electrospray Ionization Mass Spectrometry. *Biosecurity and Bioterrorism* 11(2).
- [20] Karlsson OE, Hansen T, Knutsson R, Lo°fstro°m C, Granberg F, Berg M. 2013. Metagenomic Detection Methods in Biopreparedness Outbreak Scenarios. *Biosecurity and Bioterrorism* 11(1).
- [21] Wielinga PR. 2013. Detecting Bioterrorism: How to Detect the Unexpected? *Biosecurity and Bioterrorism* 11(1).
- [22] Versage JL, Severin DDM, Chu MC, Petersen JM. 2003. Development of a multitarget real-time TaqMan PCR assay for enhanced detection of Francisella tularensis in complex specimens. *J Clin Microbiol* 41(12):5492-5499.
- [23] Bell CA, Uhl JR, Hadfield TL. 2002. Detection of Bacillus anthracis DNA by LightCycler PCR. *J Clin Microbiol* 40(8):2897-2902.
- [24] Espy MJ, Cockerill FR, Meyer RF. 2002. Detection of smallpox virus DNA by LightCycler PCR. *J Clin Microbiol* 40(6):1985-1988.
- [25] Yang S, Ramachandran P, Rothman R. 2009. Rapid identification of biothreat and other clinically relevant bacterial species by use of universal PCR coupled with highresolution melting analysis. *J Clin Microbiol* 47(7):2252-2255.
- [26] Jacob D, Sauer U, Housley R. 2012. Rapid and highthroughput detection of highly pathogenic bacteria by Ibis PLEX-ID technology. *PLoS One* 7(6).