

Nature of Host Parasite Adaptation: A Comprehensive Review Article

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ABSTRACT

Among all arthropods, mosquitoes are indubitably important to human beings to their consequence in which acting as vectors for many infective disease and also act as primary and sometimes secondary host for many viruses and protozoan parasites, many of which shows impact considerably on human and veterinary health. In the present review article represent the brief discussion on the relation with genetic diversity with the parasite with the mosquitos in vice-versa. It associates with high-risk level during the seasonal outbreaks demonstrated the potential vulnerable to human communities infected by the mosquito vectors. In present review will open debate discussion, how the protozoan parasite spread only through Anopheles species and viral parasite spread through only by Aedes Species. Let us understand some unanswered proofs to understand the genetic analysis of malaria.

INTRODUCTION

Insects are known for a transmission of various disease-causing organisms to humans and other living organisms related to the veterinary, some of them especially like mosquito borne diseases which cause most dreadful disease like Malaria, Dengue, Japanese encephalitis and many more. The distribution and abundance of various mosquito vectors is important in the determination of disease prevalence in disease-endemic areas. The vectors like mosquitoes have the shows the sensitivity and specificity to the particular disease or a particular virus. Female Anopheles mosquitoes which acts a vector for malaria which is a still life threatening about forty percent of the world's population were affected with that and well as Dengue which comes second among the mosquito borne disease which kills nearly about 10-12% percent of the population, majority of affected person were African natives, then followed by Asian people and then native of Latin American [1-5].

Study of Causative Organism

Despite of over a century attempts to eradicate or control vector borne disease, the vector borne diseases remains a still a vital and creating danger to the general prosperity and financial change of countries in the tropical and subtropical zones of the world. Around 40% of the aggregate people live in endemic zones. There are a common 300-500 million cases and up to 2.7 million infected with vector borne disease. The mortality levels of the infected people are remarkable in sub-Saharan Africa [6-9], where children under 5 years of age record for 90% of all infected with insect borne disease. Human wild fever is carried on by infected with intracellular parasites of the family Plasmodium sps that is transmitted by Anopheles mosquitoes. Of the four sorts of Plasmodium that damage the each individual of human beings, is resulting in most fatal diseases. Immunity to against malarial and insect, the decay of general structure, masses movements, political turmoil, and steady changes are adding to the spread of infection. In countries with endemic stage fever, the yearly related change rates over a 25-year time cross were 1.5% lower than in various countries [9-15]. This incites the cemented effect of the lower the yearly budgetary yield in a wild fever endemic country was a half diminishment in the per capita GDP showed up contrastingly in association with a non-

malarious country. Late studies suggest that the measure of intestinal enduring cases may twofold in 20 years if new methods for control are not made and finished [16-25].

Based on the availability of a literature revealed only that the causative organism are specific to specific in relation: [Dengue virus](#) show specificity *Aedes* sps, [Plasmodium](#) sps show a specificity *Anopheles* sps Japanese encephalitis virus show to *Culex* sps, Babesiosis specific to Tick and also Tick-borne encephalitis virus through the Tick. However, they were many insects which act like a vector to communicable or non-communicable diseases [4,9,16,26-32].

New Target on understanding the parasite drugs

Mohammed Hassan [33] have presented his work with a title "*The protein and genomes and its implication as malaria parasites drugs*" at International Conference on Medical Parasitology and Zoology, to understand the impact of protein as malarial drugs to eradicate the malaria or insect borne diseases. The field of bio discovery and plant based pesticides steadily progresses and the development of botanical insecticides have become more rigorous in recent years with calls for more standardization. In recent years, a large number of botanical products, including plant extracts, essential oils, and pure metabolites, have been proposed for eco-friendly control of mosquito vectors and other blood-sucking arthropods. Mosquito larvae live in a wide variety of aquatic environments, which makes them attractive targets for insecticides [34-40].

The field of bio discovery and plant based pesticides steadily progresses and the development of botanical insecticides have become more advanced in recent years with calls for more standardization. In recent years, a large number of botanical products, including plant extracts, essential oils, and pure metabolites, have been proposed for eco-friendly control of mosquito vectors and other blood-sucking arthropods [41-50]. Mosquito larvae live in a wide variety of aquatic environments, which makes them attractive targets for insecticides. Topics of understanding the parasite [2nd International Conference on Parasitology](#) towards the Parasite Remedies which helps in the understanding the most widely-used mosquito control strategies at present use various formulations of chemical insecticides to kill larvae and adults [51-58]. However, these tools are not always efficient, applicable, or available. Increased resistance of mosquitoes to chemical insecticides and the cost of these interventions make them prohibitive in many settings

Considering these drawbacks, pursuing informed, efficient ways of using chemical insecticides is essential. In the formulation process it is important to choose the best compounds, including synergists in order to achieve safety, environmental and economic benefits [60-66]. Our results support the conclusion that dillapiole is useful for public health applications targeting mosquitoes when used together with pyrethroids.

Similar to the observations with *Aedes aegypti*, the association α -cypermethrin/dillapiole increased mortality of *A. albopictus*. No statistically significant difference was noted at 15 minutes. The association α -cypermethrin/dillapiole caused 100% mortality between 60 and 75 min, while alive mosquitoes were still present at 90 minutes of exposure in the assays performed with each separate compound [67-75]. The association of these two compounds shortened the time necessary to kill mosquitoes, estimated as TL50 and TL90. These results further support our conclusion that dillapiole is useful in targeting mosquitoes when used together with pyrethroids.

An essential component identified with the improvement of *Anopheles melas* is the relationship between the larval environment of this species and the saltiness of its reproducing into adult form of mosquitoes [76-79]. It is this marvel which legitimizes the diminishing of *Anopheles melas* amongst October and December while thickness of *Anopheles gambiae sensu stricto* expanded. In fact, the last quarter of the year is the time of surge and subsidence. In this manner, the tidal pond of Ouidah is loaded with a lot of new water fundamentally from the northern Benin amongst September and October that adjusts *Anopheles melas* hatchlings environment. The desalination brings about a general reduction in mosquito thickness particularly *Anopheles melas*. Amid retreat (November-December), the stores left by the withdrawal of water are good for *Anopheles gambiae sensu stricto* and clarifies the high thickness of *Anopheles gambiae sensu stricto* saw amid this period [79-83]. In February, with the expansion in saltiness of the tidal pond of Ouidah, we recorded 92.18% of *Anopheles melas*.

The seasonal and temporal variation of *Anopheles melas* density in our study area was similar to that observed by Akogbeto [84]. The outcomes of the study showed *Anopheles melas* [81] that is active during the in the dry and raining season. Similar results were observed in Eastern Ghats which predominant in rainy season respectively.

The presence of a relatively high density of *Anopheles gambiae* may also be due to the urbanization of the traditional lagoon area and the reduction of mangrove (mangrove wood exploitation for the production of salt), natural

habitat of *Anopheles melas* [85-90]. The change of the conventional environment great to biology *Anopheles melas* makes an irregularity contrasted with the underlying populaces of vectors and results in a decline of *Anopheles melas* and an expansion in *Anopheles gambiae sensu stricto* thickness. The salinization of the tidal pond of West Bengal and Andhra Pradesh is additionally dynamic, and helps *Anopheles gambiae sensu stricto* adjustment to high saltiness. A research center review could be considered to indicate the resistance furthest reaches of both species [91-94].

CONCLUSION

Whether reclamation of natural in urban streams would give reasonable rearing territories for some mosquitoes was not clear yet It was in this manner essential to decide how adjusted stream conditions influence mosquito environment. Month to month information on water quality and larval thickness were gotten to determine the impacts of waterway frameworks on the dispersion and plenitude of youthful mosquitoes in two coastal cities in Eastern and Western India. Altogether, five species inside two genera of mosquitoes were gathered and recognized in habitat with vegetation from three positive waterways. *Culex pipiens* was the most bottomless and widely distributed species [95-98]. Another species (*Culex fuscanus*) was accounted for in specific regions. Physico-chemical parameters of waterway water were vital, however not by any means the only, arrangement of impacts on juvenile mosquito breeding. Sea-going vegetation could improve the probability of mosquito reproducing while manufactured aeration might keep the approach of mosquitoes. Moderate moving water may be another potential minor habitat type for some *Culex* and *Aedes albopictus*. Variety of stream framework with environmental rebuilding might influence the plenitude and circulation of juvenile mosquitoes [99,100].

REFERENCES

1. Voravuth S, et al. Antimalarial and anti-hemolytic properties of aqueous crude extract of gynostemma pentaphyllum leaves against plasmodium berghei infection in mice. *Malaria Contr Elimination*. 2016;150
2. Cohen A, et al. Micro RNAs and Malaria – a dynamic interaction still incompletely understood. *J Neuroinfect Dis*. 2014;5:165.
3. Tulara NK. Concurrent infection with malaria, dengue and hepatitis a virus together. *J Trop Dis*. 2015 3:155
4. Mohan K, et al. Clinical presentation and management of neonatal malaria:A Review. *Malar Chemoth Cont Elimination*. 2014;3:126
5. Verma P and Sharma YD .Malaria genome project and its impact on the disease. *J Vector Borne Dis*. 2013 40:9-15.
6. Cohen A, et al. (2014) Micro RNAs and malaria – a dynamic interaction still incompletely understood. *J Neuro infect Dis*. 5:165.
7. Timothy JB et al. Therapeutic Combination of nanoliposo malsafingol and nanoliposomalceramide for Acute Myeloid Leukemia. *J Leuk*. 2013;1:110
8. Salem A. Crosstalk between the Smad and the mitogen-activated protein kinase pathways is essential for erythroid differentiation of erythroleukemia cells induced by $\text{tgf-}\beta$, activin, hydroxyurea and butyrate. *J Leuk*. 2013;1:109
9. Gonul H. High-dose glucocorticoid for the treatment of myeloid sarcoma. *J Leuk*. 2013;1:103
10. Bin Fu, et al. Detection of BRAF V600E Mutation in Langerhans Cell Histiocytosis Using High-resolution Melting Analysis in Decalcified, Paraffin-embedded Tissue. *J Leuk*. 2013;1:101
11. JR Patel et al. Plant Derived Compounds Having Activity against P388 and L1210 Leukemia Cells. *CSJ*. 2011;2:024
12. Jayaraman A and Jamil K. Clusters of CDK2, CCND1, and CMYC genes involved in cancers:Acute Lymphocytic Leukemia (ALL) as a model. *Biol Med*. 2012;4:157
13. Harold K et al. Leucopenia in acute myeloid leukemia presenting as myocardial infarction:A Case Report. *J Bone Marrow Res*. 2013;1:110
14. Guillermo J, et al. Ruxolitinib chronic myelomonocytic leukemia-associated myelofibrosis:A Case Report. *J Bone Marrow Res*. 2013;1:109
15. Tao YF, et al. CDH13 are frequently inactivated by promoter hyper methylation in pediatric acute myeloid leukemia (AML). *J Hematol Thrombo Diseases*.2013;1:111
16. Rashid M, et al. Epigenetic silencing of *dapk1* gene is associated with faster disease progression in india populations with chronic myeloid leukemia. *J Cancer Sci Ther*. 2013;5.4:144-149

17. Oliver B. the current molecular site of the myelo proliferative neoplasms - tet-àtête with the jakpot but no Ink so far to resolve complexity. *J Leuk.* 2013;1:108
18. Sara C and Andrew WB. Expression and function of the eph receptor family in leukemia and hematopoietic malignancies:prospects for targeted therapies. *J Leuk.* 2013;1:107
19. Yasser H, et al. Abl kinase domain mutations in imatinib-treated egyptian patients with chronic myeloid leukemia. *J Leuk.* 2013;1:106
20. Nassima M, et al. Dysregulation of the antigen-induced nf-? b signaling pathway in the development of human b-cells lymphomas. *J Leuk.* 2013;1:105
21. Ota F, et al. Lenalidomide therapy of myelodysplastic syndromes. *J Leuk.* 2013;1:104
22. Ana L and Graciana SL. Role of methylenetetrahydrofolatereductase (mthfr), glutathione s-transferases (gsts m1 and t1) and haptoglobin (hp) gene polymorphisms in susceptibility to chronic myeloid leukemia (cml). *J Hematol Thrombo Diseases.*2013;1:103
23. Kiran K, et al. An uncommon morphology of acute undifferentiated leukemia:report of a rare case. *J Bone Marrow Res.* 2013;1:103
24. Sehar A and Zakariya AS. Allogeneic Stem Cell Transplantation for Chronic Myeloid Leukemia in the Era of Tyrosine Kinase Inhibitors- What are the Limitations?. *J Carcinogene Mutagene*, S14- 001
25. Massimo G, et al. Serological Survey to Determine the Occurrence of Blue Tongue Virus, Bovine Leukemia Virus and Herpesvirus Infections in the Japanese Small Ruminant Population from Northern Districts. *Clin Microbiol.* 2013;2:104
26. Seloilwe ES, et al. Parent and youth communication patterns on hiv and aids, stis and sexual matters:opportunities and challenges. *J Child Adolesc Behav.* 2015;3:203.
27. Indridason H, et al. Long term nationwide analysis of HIV and aids in Iceland, 1983-2012. *J AIDS Clin Res.* 2014;5:387.
28. Tao YanFang, et al. CDH13 is frequently inactivated by promoter hyper methylation in pediatric acute myeloid leukemia (AML). *J Hematol Thrombo Diseases.*2013;1:111
29. Rashid Mir, et al. Epigenetic Silencing of DAPK1 Gene is Associated with Faster Disease Progression in India Populations with Chronic Myeloid Leukemia. *J Cancer Sci Ther.* 2013;5.4:144-149
30. Oliver Bock. The Current Molecular Site of the Myelo proliferative Neoplasms - TET-àtête with the JAK pot but no LNK so far to Resolve Complexity. *J Leuk.* 2013;1:108
31. Sara C and Andrew W. Expression and function of the eph receptor family in leukemia and hematopoietic malignancies:prospects for targeted therapies. *J Leuk.* 2013;1:107
32. Yasser H, et al. Abl Kinase Domain Mutations in Imatinib-treated Egyptian Patients with Chronic Myeloid Leukemia. *J Leuk.* 2013;1:106
33. Nassima M, et al. Dysregulation of the antigen-induced NF-?B signaling pathway in the development of human B-cells lymphomas. *J Leuk.* 2013;1:105
34. Ota F, et al. Lenalidomide therapy of myelodysplastic syndromes. *J Leuk.* 2013;1:104
35. Ana L, et al. Role of methylene tetra hydro folate reductase (MTHFR), glutathione s-transferases (GSTS m1 and T1) and haptoglobin (HP) gene polymorphisms in susceptibility to chronic myeloid leukemia (Cml). *J Hematol Thrombo Diseases.* 2013;1:103
36. Kiran K et al. An uncommon morphology of acute undifferentiated leukemia:report of a rare case. *J Bone Marrow Res.* 2013;1:103
37. Seha A and Zakariya AS. Allogeneic Stem Cell Transplantation for Chronic Myeloid Leukemia in the Era of Tyrosine Kinase Inhibitors- What is the Limitations. *J Carcinogene Mutagene*. S14- 001
38. Massimo G, et al. Serological survey to determine the occurrence of blue tongue virus, bovine leukemia virus and herpesvirus infections in the Japanese small ruminant population from northern districts. *Clin Microbiol.* 2013;2:104
39. Imtiyaz A, et al. Inactivation of riz1 gene by promoter hypermethylation is associated with disease progression and resistance to imatinib in indian chronic myelogenous leukemia patients, first study from India. *J Cancer Sci Ther* 2013;5.2:045-051
40. Wen YC. Cancer acquired resistance:a new lesson from chronic myelogenous leukemia. *J Bone Marrow Res.* 2013;1:e101
41. Borkowska M, et al. The cytotoxic effect of polyelectrolyte shells coated bacterial cells on human leukemia cells. *J Nanomed Nanotechnol.* 2012;3:152

42. Fan H and Tong Y. Potential dual-use of bacteriophage related technologies in bioterrorism and biodefense. *J Bioterr Biodef.* 2012;3:121
43. Patriarca A, et al. The impact of molecular genetic in acute myeloid leukemia's. *J Blood Disorders Transf.* 2015;6:252
44. Bhensdadia DV, et al. Isolation, Molecular characterization and insight into the genome sequence of *E. coli* bacteriophage adb-2 from poultry fecal sample. *Next Generat Sequenc& Applic.* 2014;1:101
45. Filippov AA, et al. Bacteriophages against bio threat bacteria:diagnostic, environmental and therapeutic applications. *J Bioterr Biodef.* 2013;S3:010
46. Nora BC. How to Have Dual Lives: Proteins that Bind DNA and RNA. *Mol Biol* 2014;3:e120
47. Dutt S, et al. Immobilization of P22 bacteriophage tails pike protein on si surface for optimized salmonella capture. *J Anal Bioanal Techniques.* 2013;S7:007
48. Singh PK. Vitamin E Analogs as Radiation Counter measures:Beyond the Antioxidant Activities. *Mol Biol.* 2014;3:e116
49. Patsouris D and Jeschk MG. Stress Induced Insulin Resistance in Regards to Cellular Organelles, Inflammasome and Inflammation and Lipids. *Mol Biol.* 2014;3:e114
50. Nallaseth. Is there a role for an evolutionary genetics based rational health policy in global biomedical, health and economic policies? *.Mol Biol.* 2014;3:1000e118
51. Proost J, et al. Filamentous Phages Displaying Multivalent Peptide Motives with Specific Affinity to Anodic Alumina Surfaces. *J Bioeng Biomed Sci.* 2015;6:162
52. Takis A. Celebrating 30 years since the conception of the human genome project (hgp):new concepts ahead-molecular biology tools to efficiently modify the hg and/or other species-genomes-implications for health and disease. *Mol Biol.* 2014;3:e119
53. Bhowmick S and Tripathy S. A tale of effectors;their secretory mechanisms and computational discovery in pathogenic, non-pathogenic and commensal microbes. *Mol Biol.* 2014;3:118
54. Remes AM, et al. Functional MRI in patients with the c9orf72 expansion associate frontotemporal dementia. *Mol Biol.* 2014;3:117
55. Perepechaeva M, et al. Altered m-rna expression of "ahr-nrf2 gene batteries" in the retinas of senescence-accelerated oxy rats during development of amd-like retinopathy. *J Mol Genet Med.* 2014;8:105
56. Rosenberg I and Rosenberg E. Prebiotics and probiotics within the framework of the hologenome concept. *J Microbial Biochem Technol.* 2011;S1:001
57. Turton JF, et al. Isolation of bacteriophage against currently circulating strains of *acinetobacter baumannii*. *J Med Microb Diagn.* 2012;1:109
58. Higgins PJ. PAI-1 Promoter-specific Oligonucleotide Decoys:Transcription Factor and Potential Utility as Wound Healing Therapeutics. *Cell Dev Biol.* 2014;3:143
59. Agyare C, et al. Wound Healing and Anti-Infective Properties of *Myrianthusarbores* and *Alchorneacordifolia*. *Med Chem.* 2014;4:533-539
60. Collawn SS and Patel S. Adipose-Derived Stem Cells, their Secretome, and Wound Healing. *J Cell Sci Ther.* 2014;5:165
61. Wlian L, et al. Wound healing effects of a lipocalin-derived peptide. *J Clin Toxicol.* 2014;4:187
62. Marjeta U. Chemo proteomics, a valuable tool for biomarker and drug discovery. *Mol Biol.* 2014;3:e117
63. Kajhoj TQ, et al. Test of critical steps towards a combined cell and gene therapy approach for the treatment of duchenne muscular dystrophy. *J Mol Genet Med.* 2015;9:160.
64. Altaner C. Prodrug gene therapy for cancer mediated by mesenchymal stem/stromal cells engineered to express yeast cytosinedeaminase uracil phospho ribosyl transferase. *J Stem Cell Res Ther.* 2015;5:264
65. Venkatachalam KV. Science vision 2020 on sulfur metabolism:what is needed and what can be achieved. *J Genet Syndr Gene Ther.* 2015;6:e129
66. Peter N, et al. Left Ventricular Non-Compaction:Current Controversy and New Insights. *J Genet Syndr Gene Ther.* 2015;6:255.
67. Nwosu VC. Antibiotic resistance with particular reference to soil microorganisms. *Res Microbiol.* 2001;152:421-430.
68. Palumbo JD, et al. Mutagenesis of beta-1,3-glucanase genes in lysobacter enzymogenes strain C3 results in reduced biological control activity toward bipolaris leaf spot of tall fescue and pythium damping-off of sugar beet *Phytopathology* 2005;95:701-707.

69. Cytryn E. The soil resistome: The anthropogenic, the native, and the unknown. *Soil Biol Biochem* 2013; p 63:18-23.
70. D' Costa VM, et al. Expanding the soil antibiotic resistome: exploring environmental diversity. *Curr Opin Microbiol.* 2007;10:481-489.
71. Burgos JM, et al. Presence of multidrug-resistant enteric bacteria in dairy farm topsoil. *J Dairy Sci.* 2005;1391-1398.
72. Sengupta N, et al. Diversity and antibiotic susceptibility pattern of cultivable anaerobic bacteria from soil and sewage samples of India. *Infect Genet Evol.* 2011;11:64-77.
73. Dantas G, et al. Bacteria subsisting on antibiotics. *Science.* 2008;320:100-103.
74. Liu B and Pop M. ARDB-antibiotic resistance genes database. *Nucleic Acids Res* 2009;37:D443-D447.
75. Rodriguez-Verdugo A, et al. Evolution of *Escherichia coli* rifampicin resistance in an antibiotic-free environment during thermal stress. *BMC Evol Biol.* 2013;13.
76. Hsu JT, et al. Prevalence of sulfonamide-resistant bacteria, resistance genes and integron-associated horizontal gene transfer in natural water bodies and soils adjacent to a swine feedlot in northern Taiwan. *J Hazard Mater.* 2014;277:34-43.
77. Stokes HW and Hall RM. A novel family of potentially mobile DNA elements encoding site-specific gene-integration functions: integrons. *Mol Microbiol.* 1989;3:1669-1683.
78. Zhang Q and Dick WA. Growth of soil bacteria, on penicillin and neomycin, not previously exposed to these antibiotics. *Sci Total Environ.* 2014;493:445-453.
79. Zhang Z, et al. Chitinases from the plant disease biocontrol agent, *Stenotrophomonas maltophilia* c3, *Phytopathol.* 1996;91:204-211.
80. Aseel DG, et al. Two isolates of potato virus Y (PVY) and the response of different potato cultivars against the viral infection. *J Virol Antivir Res.* 2015;4:4.
81. Shi C, et al. The influence of porcine reproductive and respiratory syndrome virus infection on the expression of cellular prion protein in MARC-145 cells. *J Virol Antivir Res.* 2016;4:4.
82. Thorley-Lawson DA. Epstein-Barr virus: exploiting the immune system. *Nature Reviews* 2001;1:75-82.
83. Appolin CM, et al. Ribavirin has an in vitro antiviral effect in rabies virus infected neuronal cells but fails to provide benefit in experimental rabies in mice. *J Virol Antivir Res.* 2013;2:2.
84. Thorley-Lawson DA. EBV the prototypical human tumor virus -just how bad is it? *Molecular Mechanisms in Allergy and Clinical Immunology.* 2005;116:251-261.
85. Kho MML et al. Gastro-intestinal involvement of primary varicella zoster virus infection in a renal transplant recipient. *J Virol Antivir Res.* 2004;3:3.
86. Aliyari R, et al. RNA-based viral immunity initiated by the Dicer family of host immune receptors. *Immunol Rev.* 2009;227-176.
87. Gutierrez JA, et al. Vitamin D metabolites inhibit hepatitis C virus and modulate cellular gene expression. *J Virol Antivir Res.* 2014;3:3.
88. Crotty S and Andino R. Implications of high RNA virus mutation rates: Lethal mutagenesis and the antiviral drug ribavirin. *Microbes Infect.* 2002;4.
89. Cullen BR. Five questions about viruses and microRNAs. *PLoS Pathog* 2010;6.
90. Ding SW and Voinnet O. Antiviral immunity directed by small RNAs. *Cell.* 2007;130.
91. You DM, et al. Twice daily dosing of telaprevir for treatment-naive and treatment-experienced patients with hepatitis C infection. *J Virol Antivir Res* 2014; 3:4.
92. Grijalva-Chon JM and Longoria CR. Viral threats in aquaculture: the battle continues. *J Virol Antivir Res.* 2015; 4:1.
93. El-Wahab EWA, et al. Seroprevalence, immunostatus and factors associated with blood borne viral infections among Egyptian municipal solid waste workers. *J Virol Antivir Res* 2015;4:4.
94. Mishra KP, et al. Plant derived antivirals: a potential source of drug development. *J Virol Antivir Res* 2013;2:2.
95. McRae S. Treatment options for venous thromboembolism: Lessons learnt from clinical trials. *Thromb J.* 2014;12 :27.
96. Dasta JF, et al. Daily hospitalization costs in patients with deep vein thrombosis or pulmonary embolism treated with anticoagulant therapy. *Thromb Res.* 2015;135:303-310.

97. Grosse SD, et al. The economic burden of incident venous thromboembolism in the United States:A review of estimated attributable healthcare costs. *Thromb Res.* 2016;137:310.
98. Lefebvre P, et al. All-Cause and potentially disease-related health care costs associated with venous thromboembolism in commercial, medicare and medicaid beneficiaries. *J Manag Care Pharm.* 2012;18:363-374.
99. Deitel zweig SB, et al. Prevalence of clinical venous thromboembolism in the USA:current trends and future projections. *Am J Hematol.* 2011;86: 217-220.
100. Chamroonsawasdi K, et al. Monitoring and evaluation of a model development project and strategic campaign on HIV/AIDS prevention among Muslim communities in Thailand. *J AIDS Clin Res.* 2015; 5.