INVESTIGATION OF Kdr MUTATION IN DENGUE VECTOR Aedes aegypti CYPERMETHRIN-RESISTANT FROM DENGUE ENDEMIC AREA IN MEDAN CITY, NORTH SUMATERA

PROVINCE, INDONESIA

Background

In 2017, dengue affected > 1200 people in Medan City.
Cypermethrin insecticide, the most common used in dengue vector control and Its resistance status and molecular mechanism has not been reported

Objective

Investigating resistance status of *Ae. aegypti* againts cypermethrin and detection of kdr mutation in *Ae. aegypti* VGSC gene

Conclusion

Kdr Mutation were found in population of *Ae. aegypti* resistant and susceptible to cypermethrin from dengue endemic areas in Medan.





Discussion

- **S989P** : Generally associated with V1016G/I.
- □ V1016G : Commonly found in Asia and has been reported in Indonesia. Related to resistance pyretroid type II.
- □ **F1534C** : Distributed worldwide. Related to resistance pyrethroid type I.
- □ Combine Kdr Mutation : S989P+V1016 on DII S6 and F1534C on DIII S6 of VGSC gene are related to pyrethroid type I and II.

Eradication of mosquito breeding site by '1 house 1 Jumantik' is one of the family members living in one house who is assigned to do monitoring larva periodically.

Solution

References

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