**Supplementary Table 5. Polymorphism of *Aedes aegypti* ITS2haplotypes from Indonesia**

Cluster Cluster best hit a Location Haplotype % identity b

1a MH142327 Russia H1 99.65 %

KY382418 Sri Lanka H2 99.30 %

HE820724 Russia H3 99.30 %

H6 98.25%

H7 98.25%

H8 98.23%

H4 97.55%

H5 97.54%

H9 97.20%

H10 94.68%

1b MH142327 Russia H11 99.51%

MH142320 Russia

MH142318 Russia

KY328418 Sri Lanka

KF471584 Rockefeller Strain

KF471587 Rockefeller Strain

KF471579 New Caledonia

(France)

KP259840 India

HE820724 Russia

1c KF471584 Rockefeller Strain H12 91.29%

MH142327 Russia 91.23%c

KY382418 Sri Lanka 91.23%c

HE820724 Russia 91.23%c

1d KF471584 Rockefeller strain H13 88.24%

MH142327 Russia 88.11%c

KY382418 Sri Lanka 88.11%c

HE820724 Russia 88.11%c

1e KF471579 New Caledonia H14 94.93 %

(France)

KF471584 Rockefeller Strain H15 92.61%

1f KU497614 NA H16 99.65%

KU497614 NA H17 99.65%

KU497614 NA H18 99.65%

KU497614 NA H19 96.14%

a) All sequences displaying the same best hit score were reported with their respective accession number

b) The percentage of identity of a given haplotype is the same for each best hit sequence

c) Second best hit score