

1 **Appendix S1.** Description of comparator wild-type and transgenic Ag(PMB)1 strains

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3 **Mosquito strains**

4 G3 is a wild-type colony (stock number MRA-112) which has been maintained in the
5 laboratory for 43 years (Malaria Research and Reference Reagent Resource Center 2017). This
6 strain derives from a region in West Africa (the Gambia) where two incipient species, *An. gambiae*
7 and *An. coluzzii*, coexist. G3 strain is assumed to be an inter-specific hybrid between *An. gambiae*
8 and *An. coluzzii*, but no complete genome sequence data are still published to confirm this
9 assumption. However, as VectorBase reported “On the basis of both inversion polymorphisms and
10 rDNA characteristics, this colony has been suggested to represent a mix of *Anopheles gambiae* and
11 *Anopheles coluzzii*. Since this colony has been distributed broadly among different institutions
12 without any specific molecular or inversion efforts at quality control, it should probably simply be
13 recognized as a hybrid stock (*An. gambiae s.l.*)” ([https://www.vectorbase.org/organisms/anopheles-](https://www.vectorbase.org/organisms/anopheles-coluzzii/g3)
14 [coluzzii/g3](https://www.vectorbase.org/organisms/anopheles-coluzzii/g3)).

15 Ag(PMB)1 transgenic line was developed from the G3 strain and classified as a sex ratio
16 distorter - autosomal X-shredder based on I-PopI homing endonuclease (Galizi et al., 2014). This
17 strain is marked with the fluorescent marker *3XP3DsRed* and green fluorescence protein (GFP), the
18 latter of which was fused to I-PpoI. Both GFP and the I-PpoI enzyme were expressed in sperm. Due
19 to the sex ratio distortion conferred by the transgene, Ag(PMB)1 strain has been maintained in the
20 laboratory by continuously backcrossing Ag(PMB)1 females to G3 males. The transgenic
21 mosquitoes used for these experiments were screened for the expression of the *3xP3 DsRed*
22 fluorescent marker using a Complex Object Parametric Analyzer and Sorter (COPAS, Union
23 Biometrica, Boston, USA). This backcrossing procedure promoted a high genetic similarity
24 between transgenic individuals and wild-type counterparts, except for the transgene itself (Galizi et
25 al., 2014). Life-history parameters of wild-type and transgenic Ag(PMB)1 mosquito strains were

26 collected from literature or experiments (please see subchapter of the Material and Methods
27 *Additional survival measurements of wild-type and Ag(PMB)I for model parametrization*).

28 Mosquito strains were maintained and reared in a climate-controlled room at a temperature
29 of $27 \pm 1^\circ\text{C}$ and $75\% \pm 10\%$ relative humidity, with a 12:12h light and dark regime. The mosquitoes
30 were reared as described in Valerio et al. (2016) with the following modifications. Adult
31 mosquitoes were carried in plastic cages of $17.5 \times 17.5 \times 17.5$ cm and fed *ad libitum* with a 10%
32 sucrose solution with 0.1% of methylparaben (Benedict, Hood-Nowotny, Howell, & Wilkins 2008).
33 Females were blood fed weekly with defibrinated and heparinized sterile cow blood (Allevamento
34 Blood di Fiastra Maddalena, Teramo, Italy) using a Hemotek membrane feeder
35 (DiscoveryWorkshops, Accrington, UK). Eggs were collected and placed in a single tray with 350
36 mL of deionized water for hatching and provided 5 mL of 2% w/v larval diet. Subsequently, larvae
37 were allotted to three trays (each containing about 250 larvae) and maintained according to the
38 standard rearing protocol based on Damien's larval diet as the food source (Damiens, Benedict,
39 Wille, & Gilles 2012). Pupae were collected and placed in small crystallization dish glass inside a
40 fresh adult cage for emergence.

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42 **References**

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