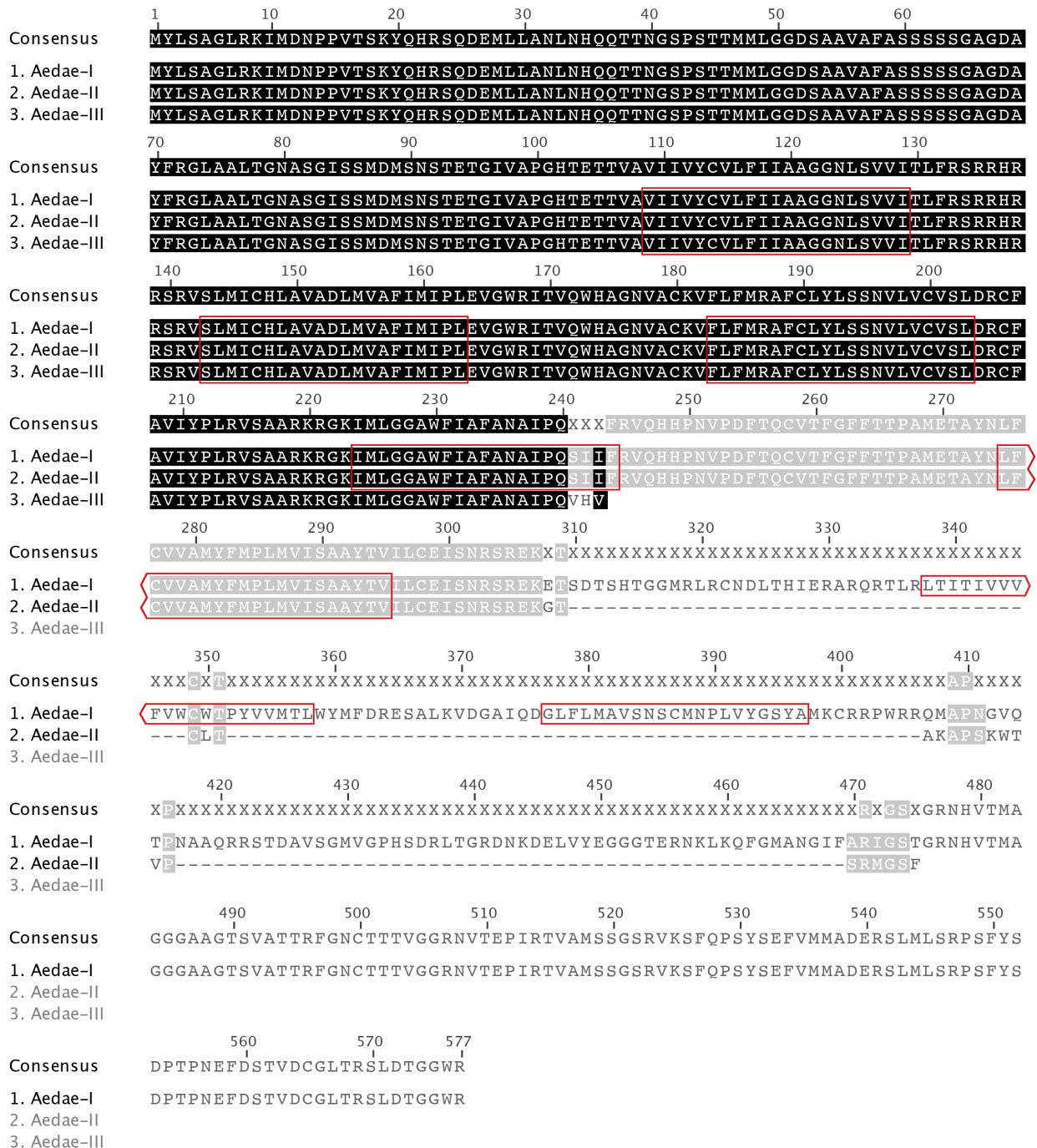


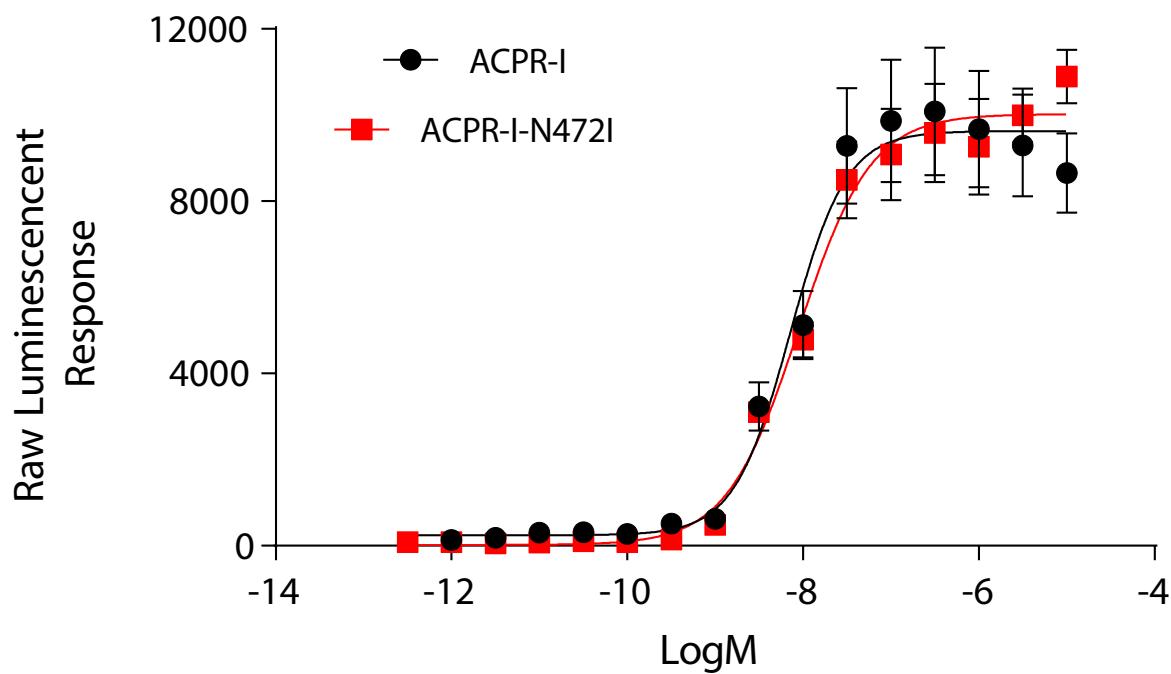
**Molecular identification, transcript expression, and functional deorphanization of  
the adipokinetic hormone/corazonin-related peptide receptor in the disease vector,  
*Aedes aegypti*.**

Azizia Wahedi and Jean-Paul Paluzzi

SUPPLEMENTARY INFORMATION



**Figure S1. Sequence alignment of *Aedes aegypti* ACP receptor variants I, II, and III.** Aligned amino acid sequences of ACPRs from *A. aegypti*, transcript variant I, (GenBank: MF461644), transcript variant II, (GenBank: MF461645), transcript variant III, (GenBank: MF461646). Residues outlined in red indicate predicted transmembrane domains based on the *A. aegypti* sequences. Highlighting of residues indicates % identity with black denoting 100% sequence identity, dark grey denotes 80-100% identity, and light grey represents amino acid positions with 60-80% sequence identity.



**Figure S2. Functional heterologous receptor assay of CHO-K1 aequorin cells transiently expressing the *AedaeACPR-I* and *AedaeACPR-I-N472I*.** Dose dependent effect on the bioluminescence response (mean 0-15s) after the addition of between  $10^{-12}$  –  $10^{-5}$  M doses of *AedaeACP* to *AedaeACPR-I* and *AedaeACPR-I-N472I*. Luminescence is normalized to the BSA control. Data represent the mean  $\pm$  standard error (n=4).

**Table S1.** GenBank accession numbers and references for receptor sequences in Fig. 3.

Receptor	Putative Ligand(s)	Accession Number	Reference(s)
<i>Acyrtosiphon pisum</i> AKHR	AKH	XP_003245941	<sup>1</sup>
<i>Aedes aegypti</i> AKHR1	AKH	XP_001655248	<sup>2</sup>
<i>Aedes albopictus</i> AKHR	AKH	XP_019540948	
<i>Anopheles gambiae</i> AKHR	AKH	ABD60146	<sup>3</sup>
<i>Anopheles darlingi</i>	AKH	ETN67429	<sup>4</sup>
<i>Anopheles sinensis</i>	AKH	KFB53098	<sup>5</sup>
<i>Culex pipiens</i> AKHR	AKH	AAWU01041683	<sup>1</sup>
<i>Apis mellifera</i> AKHR	AKH	NP_001035354	<sup>6</sup>
<i>Bombyx mori</i> AKHR	AKH, ACP	NP_001037049	<sup>7-11</sup>
<i>Musca domestica</i> AKHR		XP_019895730	
<i>Manduca sexta</i> AKHR	AKH	ACE00761	<sup>12</sup>
<i>Drosophila melanogaster</i> AKHR	AKH	AAC61523	<sup>7,13</sup>
<i>Locusta migratoria</i> AKHR		ANW09575	
<i>Polyrhachis vicina</i> AKHR		ADK55068	
<i>Nasonia vitripennis</i> AKHR	AKH	NP_001161243	<sup>14</sup>
<i>Tribolium castaneum</i> AKHR	AKH	NP_001076809	<sup>14</sup>
<i>Periplaneta Americana</i> AKHR	AKH	ABB20590	<sup>14</sup>
<i>Rhodnius prolixus</i> AKHR	AKH	AIJ49751	<sup>15</sup>
<i>Gryllus bimaculatus</i> AKHR	AKH	ADZ17179	<sup>16</sup>
<i>Glossina morsitans</i> AKHR	AKH	AEH25943	
<i>Aedes aegypti</i> CRZR	CRZ	AAGE02020874	<sup>2</sup>
<i>Aedes albopictus</i> CRZR		XP_019554625	
<i>Anopheles gambiae</i> CRZR	CRZ	AAQ67361	<sup>17</sup>
<i>Apis mellifera</i> CRZR	CRZ	NP_001137393	<sup>6</sup>
<i>Culex pipiens</i> CRZR	CRZ	AAWU01031675	<sup>1</sup>
<i>Bombyx mori</i> CRZR	CRZ	NP_001127719	<sup>8,14,18</sup>
<i>Drosophila melanogaster</i> CRZR	CRZ	AAM21341	<sup>19</sup>
<i>Nasonia vitripennis</i> CRZR	CRZ	AAZX01006363	<sup>20</sup>
<i>Bactrocera dorsalis</i> CRZR	CRZ	AQX83392	<sup>21</sup>
<i>Rhodnius prolixus</i> CRZR-A	CRZ	AND99324	<sup>22</sup>
<i>Manduca sexta</i> CRZR	CRZ	AAR14318	<sup>23</sup>
<i>Musca domestica</i> CRZR	CRZ	NP_001295994	
<i>Anopheles gambiae</i> ACPR	ACP	ABX52399	<sup>14</sup>
<i>Anopheles darlingi</i> ACPR	ACP	ETN61096	<sup>24</sup>
<i>Aedes albopictus</i> ACPR	ACP	XP_019559258	
<i>Culex pipiens</i> ACPR	ACP	XP_001842868	<sup>1</sup>
<i>Tribolium castaneum</i> ACPR	ACP	ABX52400	<sup>14</sup>
<i>Nasonia vitripennis</i> ACPR	ACP	NP_001164571	<sup>14</sup>
<i>Bombyx mori</i> ACPR1	ACP, AKH	NP_001127726	<sup>14,23</sup>
<i>Bombyx mori</i> ACPR2	ACP, AKH	NP_001127745	<sup>14,23</sup>
<i>Rhodnius prolixus</i> ACPR-C	ACP, AKH	AKO62858	<sup>25</sup>

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