

*An. gambiae*

G			C			A			A			G			
der	hets	wt													
2013	0.07	0.52	0.40	0.10	0.48	0.43	0.02	0.62	0.36	0.05	0.57	0.38	0.12	0.50	0.38
2015	0.12	0.40	0.48	0.08	0.42	0.50	0.18	0.31	0.51	0.12	0.37	0.51	0.13	0.38	0.50
2018	0.10	0.45	0.45	0.10	0.43	0.48	0.10	0.58	0.33	0.10	0.45	0.45	0.10	0.53	0.38

*An. coluzzii*

G			C			A			A			G			
der	hets	wt													
2011	0.04	0.30	0.65	0.04	0.26	0.70	0.04	0.13	0.83	0.13	0.13	0.74	0.17	0.30	0.52
2012	0.10	0.39	0.51	0.09	0.39	0.52	0.07	0.42	0.51	0.09	0.42	0.49	0.12	0.51	0.38
2014	0.07	0.37	0.56	0.09	0.37	0.54	0.13	0.33	0.54	0.14	0.30	0.56	0.15	0.48	0.37
2016	0.17	0.37	0.46	0.13	0.35	0.52	0.13	0.35	0.52	0.11	0.30	0.59	0.10	0.41	0.49
2018	0.24	0.44	0.32	0.18	0.39	0.42	0.12	0.59	0.29	0.18	0.36	0.45	0.31	0.47	0.22

der = derived

hets = heterozygote

wt = wild type

Supplementary Table 1: Sweep frequencies. Frequencies of derived (swept SNP), heterozygote and wild type SNPs and each locus, for each year for both *An. gambiae* and *An. coluzzii*.

**Supplementary Table 2: Graphical Data with corresponding statistics.** For each significant result, the p value and mean data are shown for each graphical figure in the paper. Each figure is separated by a bold underline, with the figure the statistics refer to specified above the said line.

#### Tiassale vs N'Gouso

Transcript	$\Delta\text{Act}_u$	p-value
SAP2	12.43	2.65E-04
CSP6	3.79	3.16E-02

#### Tiassale vs Kisumu

Transcript	$\Delta\text{Act}_u$	p-value
SAP2	6.48	1.19E-03
CSP1	2.56	3.64E-02
CSP5	4.24	1.83E-03
CSP6	5580.03	1.73E-03

Statistics and mean ddCT values used in Extended Data Figure 2

#### Tiassale

Transcript	Antennae		Head		Legs		Midgut		Malpighian Tubules		Reproductive Tissue		Abdomen Carcass	
	$\Delta\text{Act}_u$	p-value												
SAP1					83.41	0.024	0.0799	0.047						
SAP2	0.284	0.00813	4.1	<0.001	35.77	<0.001	0.015	<0.001	0.0365	<0.001	0.087	<0.001	0.184	<0.001
SAP3			5.03	0.00213	9.05	<0.001	0.0763	<0.001					0.066	<0.001
CSP1			5.04	0.00191	14.24	<0.001	0.301	0.00652	0.327	0.00891			3.49	0.00891
CSP3	513.86	<0.001	17.85	<0.001			0.149	<0.001	0.307	<0.001	0.0978	<0.001	0.26	<0.001
CSP4									0.307	<0.001	0.051	0.0284		
CSP5	20.28	<0.001	0.039	<0.001	42.03	<0.001	0.0787	<0.001	0.172	<0.001	0.317	0.0049		
CSP6					11.74	<0.001	0.0761	0.0094			3.66	0.0153		

#### N'Gouso

Transcript	Antennae		Head		Legs		Midgut		Malpighian Tubules		Reproductive Tissue		Abdomen Carcass	
	$\Delta\text{Act}_u$	p-value												
SAP1					45.65	<0.001	0.132	0.0112						
SAP2			6.84	<0.001	53.39	<0.001	0.0333	<0.001	0.151	0.0242	0.0689	0.0027	0.15	0.0196
SAP3			10.24	0.0157	19.16	0.0016	0.0268	<0.001	0.0454	<0.001			0.737	<0.001
CSP1					1.92	0.0447							4.46	0.0351
CSP3	2701.75	<0.001	40.22	<0.001			0.042	<0.001	0.0253	<0.001	0.0144	<0.001	0.0000746	<0.001
CSP4					26.92	0.0208	0.147	0.02438	0.1	0.00657				
CSP5					35.49	<0.001	0.0757	0.0029						
CSP6														

Statistics and mean ddCT values used in Figure 1a

#### Tiassale post-exposure

Transcript	30 minutes		1 hour		2 hours		4 hours		24 hours		48 hours			
	$\Delta\text{Act}_u$	p-value												
SAP1			0.298	0.0251	2.35	0.03913	11.32	<0.001	5.66	<0.001	0.263	0.0121		
SAP2							5.25	<0.001	4.42	0.0028	2.98	0.00879		
SAP3									5.83	0.00213	6.96	0.00102		
CSP1											0.213	0.0163		
CSP3												0.0055		
CSP4														
CSP5	0.208	0.0217	0.169	0.0016	0.525	0.0013			16.622	0.0077	17.16			
CSP6			12.75	0.0416	6.59									

Statistics and mean ddCT values used in Figure 1b; Extended Data 3a.

#### SAP2

Mortality	
Change	p-value
Deltamethrin	11.7% to 79
Permethrin	5.2% to 24.7
Alpha-cypermethrin	2.8% to 10.8
Banfora Del	13.19% to 48
	2.19E-02

#### CSP6

Mortality	
Change	p-value
Deltamethrin	11.7% to 31
	4.74E-02

#### Transgenics

Mortality	
Change	p-value
Permethrin	84.3% to 67
	2.29E-02

Statistics and mean mortality values used in Figure 2a; Extended Data Figure 5 and Figure 2b.

**Supplementary Table 3: Primer List.** Primers used for both synthesis of dsRNA constructs and qPCR.

**RNAi Primers**

ID	Forward	Reverse
dsSAP2	taatacgactcaatagggTTCTCGTCCGGTTGCTTC	taatacgactcaatagggTAGTAGACCCCATTCCCCACTT
dsSAP3	taatacgactcaatagggATGAAATTCTCGTCGTTGCG	taatacgactcaatagggTTCTCCGGGTGTAATTCTTCT
dsCSP4	taatacgactcaatagggTGTAGCCCTTGACGGTTT	taatacgactcaatagggTTCTCAGCACGCACATGAT
dsCSP6	taatacgactcaatagggGCACAGCACAGCAACCTTT	taatacgactcaatagggTTCCCCTTTCAGCCAGCAT
dsGFP	taatacgactcaatagggAGAACGTAAACGCCACAAGTTC	taatacgactcaatagggAGACTTGTACAGCTCGTCCATGCC

**qPCR Primers**

ID	Forward	Reverse
SAP1	ACGTCAACACAGAACGATCAAC	TTGCTGGTGTACTTATCCTGGG
SAP2	GCAGCTTGAGAGCGTCTTCT	GAAAGCGATGGCGACGAACA
SAP3	AGTGTAGCGAGAAAGCAGAAAGAG	GCGGTACTTGTGACGTAGATG
CSP1	AGCCTTTCGTGTCTGTTCA	CGGTACGCTGTCTAGATTCACT
CSP3	AACCTACGTACCCAAGTACGAT	ATCAGGGAAAGGTTGTCTTCAGC
CSP4	GCTATCAGCGGCAGTTATTGTG	AGCCGGTTAATATTCTGGCTGT
CSP5	GCACGGACACTCTACAGTACAA	TCGGGCAGAACATCGTTCAGAT
CSP6	GACAGTTCTGTTCTGTGCCG	CCTTGCAGTGGCTTTTGC
S7	AGAACCGAGCAGACCACCATC	GCTGCAAACCTCGGCTATT
EF	GGCAAGAGGCATAACGATCAATGCG	GTCCATCTGCGACGCTCCGG