

Possible role of an *Anopheles* transient receptor potential channel homologue in malaria parasite transmission

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Abstract

Midgut glycoproteins are potential receptors for pathogens such as arboviruses and malaria parasites that are transmitted by vector mosquitoes. An antiserum prepared against wheat germ agglutinin binding *Anopheles tessellatus* midgut proteins was used to screen an *An. gambiae* cDNA library. A clone homologous to the gene for a *Drosophila melanogaster* transient receptor potential-gamma cation channel protein was identified.

Introduction

Glycoproteins associated with the luminal surface of the mosquito midgut are potential receptors for mosquito transmitted pathogens such as malaria parasites [1-6] and arboviruses [7]. Antibodies against wheat germ agglutinin-binding midgut glycoproteins, ingested in an infective blood meal, block transmission of *Plasmodium vivax* and *P. falciparum* to the malaria vector *Anopheles tessellatus* [6]. To identify the relevant target midgut glycoproteins, a cDNA expression library of female *An. gambiae* abdomen was screened with the same antiserum against wheat germ agglutinin binding midgut glycoproteins.

Materials and Methods

Wheat germ agglutinin binding glycoproteins from sugar-fed female *An. tessellatus* midgut were purified and used to produce a rabbit antiserum as previously described [6]. The antiserum, pre-absorbed with *E. coli*, was used

to screen an *E. coli* lambda/bluescript (ZAP Express®/pBK-CMV; Stratagene, USA) cDNA library derived from female *An. gambiae* abdomen [8] according to standard procedures [9] modified by the manufacturer's instructions (Stratagene, USA). Cloned DNAs from antibody-reactive clones were excised and sequencing of these clones was performed from both the 5' and 3' ends. Nucleotide sequences were compared by BLAST search against the annotated *An. gambiae* Ensembl database (Wellcome Trust Sanger Institute, www.ensembl.org) and subsequently also with the *Drosophila melanogaster* genome.

Results

BLAST analysis of the 5'- and 3'-sequences of one unique clone obtained in the antibody screen, against the *An. gambiae* genome Ensembl database, showed that this was almost identical to a sequence in the contig CRA_x9P1GAV5CRW_68 present on the right arm of chromosome 3 of *An. gambiae* [Fig 1]. This *An. gambiae* sequence is homologous to the trp gamma gene product of *D. melanogaster*; (probability e-71 and 35% identity in amino acids; gbAAF53548.1(AE003652), which is a transient receptor potential cation channel protein, TRPgamma. The sequence of the predicted *Anopheles* protein showed several potential N-linked and O-linked glycosylation sites.

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Figure 1. Comparison of the DNA sequences of a part of the contig CRA_x9P1GAV5CRW_68 from the right arm of chromosome 3 of *An. gambiae* (Contig clone) and that of the antibody-reactive clone obtained using forward (clone E forw) and reverse (clone E rev) sequencing primers

Figure 1

Contig clone	18750	gttgatgaaggggctcaataatagccgcacggccggcgcggacgcggcgtc	Contig clone	19449	cgggacggaaaggaggcacgcgcaccactatcaactgcattcggtgcgt
clone E rev	14	-----gcggcgcacggacgcggggcc	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	18800	agcatttgcgcggcggtcgaggatattgcgcgcgcgtgcggcgcatttcg	Contig clone	19499	ggcagcaactgaagcgcgcggaaaaggttctccaacagccgcactcca
clone E rev	34	agcatttgcgcggcggtcgaggatattgcgcgcgcgtgcggcgcatttcg	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	18850	ccgcacgagctgttgcggcggtgttgcggcggtgtgcacaccgaagaa	Contig clone	19549	gccccgtgtggagccacccgtgcgcgcgtgcgcacgcgcgtcgc
clone E rev	84	ccgcacgagctgttgcggcggtgttgcggcggtgtgcacaccgaagaa	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	18900	ggcactgcaccagggtgaatccgcgcgcgcgcgcgcgcgcgcgcgcgc	Contig clone	19599	actacgcgcggtaatacagaacgcgtacgtggactttgcgcgttag
clone E rev	134	ggcactgcaccagggtgaatccgcgcgcgcgcgcgcgcgcgcgcgcgc	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	18950	gctcgtcggggagaacaacaacccgcgcgcgcgcgcgcgcgcgcgcgc	Contig clone	19649	ccggatgcgggcattgtttagggagcttttagatattatgttttttttt
clone E rev	184	gctcgtcggggagaacaacaacccgcgcgcgcgcgcgcgcgcgcgcgc	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	19000	ccgaaggtaactccaaaagtgcgcgcgcgcgcgcgcgcgcgcgcgcgc	Contig clone	19699	gctggcgcgtgcgcgcgcgcgcgcgcgcgcgcgcgcgcgcgcgcgc
clone E rev	234	ccgaaggtaactccaaaagtgcgcgcgcgcgcgcgcgcgcgcgcgcgc	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	19050	tccgttgcatttcataaaaaacccctaaaaaggcgcgcgcgcgcgcgc	Contig clone	19749	ttcattgc
clone E rev	282	-----	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	19100	tttccttcgcaggaaacgcgttattgtgcggccaaacatggcgcgcgc	Contig clone	19849	atcaagcttatgtgcgcgcgcgcgcgcgcgcgcgcgcgcgcgcgc
clone E rev	282	-----ggaacgcgttattgtgcggccaaacatggcgcgcgcgcgcgc	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	19150	cgaaggtaatcggcgctcccgatcgaaagactcggtgtgcgcgcgcgc	Contig clone	19899	acacgc
clone E rev	319	cgaaggtaatcggcgctcccgatcgaaagactcggtgtgcgcgcgcgc	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	19200	ccgggtcgccggcgatcgccgcgcgcgcgcgcgcgcgcgcgcgcgc	Contig clone	19949	cgc
clone E rev	369	ccgggtcgccggcgatcgccgcgcgcgcgcgcgcgcgcgcgcgcgc	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	19250	agccatcaggaggcacccgcgcgcgcgcgcgcgcgcgcgcgcgcgc	Contig clone	20049	cgaagggtcgccgcgcgcgcgcgcgcgcgcgcgcgcgcgcgcgc
clone E rev	419	agccatcaggaggcacccgcgcgcgcgcgcgcgcgcgcgcgcgcgc	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	19300	acgtcgagactgtccggatccacacaagcatcacaatcgcgcgcgcgc	Contig clone	20099	agcagccggccgcaggcggtgcgcgcgcgcgcgcgcgcgcgcgc
clone E rev	469	acgtcgagactgtccggatccacacaagcatcacaatcgcgcgcgcgc	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	19350	acggaccggggg-caccaagatgcgcggcgcgcgcgcgcgcgcgc	Contig clone	20149	cagaagatcatccggccgcggcggcgcgcgcgcgcgcgcgcgcgc
clone E rev	519	acggaccggggggcacaagatgcgcggcgcgcgcgcgcgcgcgc	clone E rev	606	-----
clone E forw	946	-----	clone E forw	946	-----
Contig clone	19399	cgc	Contig clone	20199	gcaggaggcatcgatcgtgcgtgcgtgcgtgcgtgcgtgcgtgc
clone E rev	569	cgc	clone E rev	615	-----
clone E forw	946	-----	clone E forw	946	-----

Contig clone	20249	tagggcagggtggagcggtgacacgtccggggaccggctcactgg	Contig clone	21099	gctacccaacaacggagactcatatgcgcacccctggagacaggggggg
clone E rev	615	-----	clone E rev	615	-----
clone E forw	946	-----	clone E forw	846	gttacccaacaacggagactcatatgcgcacccctggagacaggggggg
Contig clone	20299	cagtctggccacgtccacgtacgtcggtcgcaagattccggcg	Contig clone	21149	-----tcattcttttaatccgatagtctgttaaggacgggtcgatg
clone E rev	615	-----	clone E rev	615	-----
clone E forw	946	-----	clone E forw	796	gggggtcattcttttaatccgatagtctgttaaggacgggtcgatg
Contig clone	20349	aacgcactacggggcatttacactcccagggctggttatagggagac	Contig clone	21193	caagcgcgcacaccaaactacacacgatacatcttaggcacgataaga
clone E rev	615	-----	clone E rev	615	-----
clone E forw	946	-----	clone E forw	746	caaccgcgcacaccaaactaaacacgatacaccttaggcacgataaga
Contig clone	20399	ggtagccaaagggtgcgactatttgtatgtaaaatgtgtcg	Contig clone	21243	aagacatactacaacccgtccaatatccaaataaagttagttagcaga
clone E rev	615	-----	clone E rev	615	-----
clone E forw	946	-----	clone E forw	696	aagacatactacaacccgtccaatatccaaataaagttagttagcaga
Contig clone	20449	ctcaaggcgttgagcgcatacggaaatacgatgtgttaactagtc	Contig clone	21293	tagattacttatcagcataaagagtaaaaaccagtttagtaggtgc
clone E rev	615	-----	clone E rev	615	-----
clone E forw	946	-----	clone E forw	646	tagattacttatcagcataaagagtaaaaaccagtttagtaggtgc
Contig clone	20499	aaaagcatcaattctaaacgtgtggcgaagaagaatggita	Contig clone	21343	cgaggagtgtataaggagtgcacggggacacgcaatcgaggatggcaac
clone E rev	615	-----	clone E rev	615	-----
clone E forw	946	-----	clone E forw	596	cgaggagtgtataaggagtgcacggggacacgcaatcgaggatggcaac
Contig clone	20549	aagacgcaagacgagttgtaatgcatactcgcaactactctgcagatgg	Contig clone	21393	agtgttgaacgctaacacacatcagacatccgagtaattgaaatagcaa
clone E rev	615	-----	clone E rev	615	-----
clone E forw	946	-----	clone E forw	546	agtgttgaacgctaacacacatcagacatccgagtaattgaaatagcaa
Contig clone	20599	ctggacaacgcgttcgtgtatgtttaagaaaattctctgttgc	Contig clone	21443	aacttaacacaagaataaaaaggaggcctataatgttaacatgtaaa
clone E rev	615	-----	clone E rev	615	-----
clone E forw	946	-----	clone E forw	496	aacataacacaagaataaaaaggaggcctataatgttaacatgtaaa
Contig clone	20649	aggagagagataggatgttttaggcctaaaaggaaaccgtacgggtcg	Contig clone	21493	catgatgtatgttatcagtaaacaacacatcagacatccgagtaattgaaatagcaa
clone E rev	615	-----	clone E rev	615	-----
clone E forw	946	-----	clone E forw	446	catgatgtatgttatcagtaaacaacacatcagacatccgagtaattgaaatagcaa
Contig clone	20699	taacagagccgcattcgaccgaatgtcccatcatggacgggtcg	Contig clone	21543	cggaaccactataatgataacgtacgttctcgcttgcaggacgatta
clone E rev	615	-----	clone E rev	615	-----
clone E forw	946	-----catttg-----	clone E forw	396	cggaaccactataatgataatgtacgttctcgcttgcaggacgatta
Contig clone	20749	gctgtcgctatgccatacaggtaaccccttggcgacgatgtca	Contig clone	21593	gacagtaacaaaacaacaaaacaacgttgaattggatcgcatcat
clone E rev	615	-----	clone E rev	615	-----
clone E forw	938	-----	clone E forw	346	gacagtaacaaaacaacaaaacaacgttgaattggatcgcatcat
Contig clone	20799	gatcatggacacgatcggagcggaaatggagtgcacagtgcgtt	Contig clone	21643	tagagcagatgtatggaaacttcgtacaaaacgtaaaaaaaaa-cttagta
clone E rev	615	-----	clone E rev	615	-----
clone E forw	938	-----	clone E forw	296	tagagcagatgtatggaaacttcgtacaaaacgtaaaaaaaaactagta
Contig clone	20849	gaatgttgttagtgcataaggcatatatagatgtataacgatacaat	Contig clone	21692	agcaacattaaacaatttatgtcataattggagcggaaaaaccatgt
clone E rev	615	-----	clone E rev	615	-----
clone E forw	938	-----	clone E forw	246	agcaacattaaacaatttatgtcataattggagcggaaaaaccatgt
Contig clone	20899	acaataaaaaagggaccaggctttggagggtctcggtcgccga	Contig clone	21742	tgcgttagacgctaagacgataactatacgttagtggagacgatag
clone E rev	615	-----	clone E rev	615	-----
clone E forw	938	-----	clone E forw	196	tgcgttagacgactacgataactatacgttagtggagacgatag
Contig clone	20949	gcacaccaacatccgtggccgtggaggtagctgtttaactatc	Contig clone	21792	agagaaaaaggagagaaagagagaaagggtttgaaacgaataaaagat
clone E rev	615	-----	clone E rev	615	-----
clone E forw	938	-----	clone E forw	146	agagaaaaaggagagaaagagagaaagggtttgaaacgaataaaagat
Contig clone	20999	atttgttaagctatggaaactcgataggacaacaagagaggacagcg	Contig clone	21842	aaaattaaaaagggtatcgctgtctaaaagcaaaacttagtagctaa
clone E rev	615	-----	clone E rev	615	-----
clone E forw	938	-----taagctt-gaactcgatagg-acaaccaagagaggacagcg	clone E forw	96	aaaattaaaaagggtatcgctgtctaaaagcaaaacttagtgctaa
Contig clone	21049	gctagtgtatgttaagtaccgtctttccccataccgttgatgttt	Contig clone	21892	acacgacagatgtatgtgaaaatgaaacaaaaactataatatta
clone E rev	615	-----	clone E rev	615	-----
clone E forw	896	gctagtgtatgttaagtaccgtctttccccataccgttgatgttt	clone E forw	46	acacgatagatgtatgtgaaa-----

Discussion

Characterising midgut molecules involved in pathogen-mosquito interactions may help in developing new methods of controlling diseases transmitted by mosquitoes, e.g. by impairing vector competence [10] as achieved experimentally for *An. stephensi* through transgenesis [11]. The midgut proteins interacting with malaria parasites are likely to be located in the posterior midgut which is where the blood meal is digested and parasite development and invasion occurs. The midgut-specific genes from *An. gambiae* characterised so far include those encoding for structural and digestive proteins [12,13], immune response genes of *An. gambiae* that are activated during midgut invasion by ookinetes [8] as well as genes for defensive proteins such as cecropin [14] and defensin [15] that are expressed in the anterior midgut. A cell surface mucin specifically expressed in the midgut of *An. gambiae* has been characterised by screening a cDNA library of *An. gambiae* midgut with an antiserum against female *An. gambiae* peritrophic membrane proteins [13].

TRP cation channel protein in *Drosophila* has been detected in photoreceptors and nerve cell axons [16]. A human analogue has been reported in hepatocytes [17]. More recently a TRP cation channel protein has been identified as having a role in calcium signalling and fluid transport in the *Drosophila* Malpighian tubule epithelium [18]. The related *Anopheles* protein detected here could be a membrane glycoprotein cation transporter present on the luminal surface of the midgut cells, although this remains to be confirmed in expression studies. Implication of the gene in the interaction between mosquito and pathogen can be addressed by generating specific antibody against this gene and testing for invasion inhibition or interference RNA (RNAi) mediated gene silencing and subsequent *Plasmodium* infection or in vitro interaction assays between recombinant protein and *Plasmodium*. Since ingestion of the antiserum to midgut glycoproteins reduces infectivity of malaria parasites to *An. tessellatus* [6], proteins such as the TRP cation channel, and the cell surface mucin [8] warrant further investigation as potential targets for transmission blocking vaccines and other methods for interfering with malaria transmission.

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