Alignment of sequences of the block 10 of PvMSP-1 found in parasites infecting 41 patients. Five sequences (I, II, III, VI and VIII corresponding to 73% of all sequences) are identical to the sequences of recombinant antigens used in serology (BR07, BP13, BP29, BP01, BP39, respectively). Sequences IV and V are quite similar to BP29 antigen (74.0% and 67.1% of amino acid similarity, respectively), whereas VII sequence has 63.7% of identity with BP01 antigen.

	10 20 30 40 50 60 70 80 90         .
-	
I	
II	ST SVAVTVPGAVVPGVP TAAAAGS GASGAVPPA GSVPAT GPGAAA
III	ST SVAVTVPGAVVPGVP TA AAAGS GASGAVPPA AAAG SGAS GAVPPAG GP SPPAT GG VVP GVVE SAEAQ TKA Q
IV	ST SVA VTVPGAVVPG VP TA AAAGS GASGAVPPA GGPSPPAT GG VVP G VVESAEAQ T Q T Q
v	ST SVAVTVPGAVVPGVP TA AAAGS GASGAVPPA GGPSPPATGGVVP GGVVPGVVESAEAQ T QT Q
IV	GA SA PAAAA – VTVP GA GVPAAAAV T– VP GA GVPAA – – – – – – GV VP GA – PA – – – GAA–PA GAAP AGAA – – – – – – – P S AP GAQEQ T Q T Q
VII	GA SA PAAAA – VTVP GA GVPAA GAGV– VP GA AAAGVVP GA SAAGV VP GA – PA – – – GAA – – – – – – – – – – – – – –
VIII	GA S T TAA TLPVTVP SA –VPG G L PGAGVP GA AA–GL TPP– –– PPA G SV–– PA T G PGAAA–– – – – – – – – – – – – – – – – G S TEENVAA K