22,125 FcB1-schizont-ESTs sequenced
21,805 (98.5%) exploitable

est2genome, BLAST score > 700

19,459 ESTs matched on the *P. falciparum* genome
(PlasmoDB vs 4.4 and vs 5.3)
328 clusters

447 ESTs matched on the *P. falciparum* genome
(PlasmoDB vs 5.3)
24 clusters

9,254 ESTs: 243 protein-coding genes
10,611 ESTs: 4 rRNA loci
41 ESTs: 4 subtelomeric loci

2,346 non matched ESTs

est2genome, BLAST score > 500

839 ESTs <60nt and/or low complexity (excluded)

1,060 non matched ESTs

BLAST on UniProt

270 ESTs: MSP1 (K1-Type), Ebl-1

629 non matched ESTs

160 ESTs: mitochondrial genome
1 EST: apicoplast genome