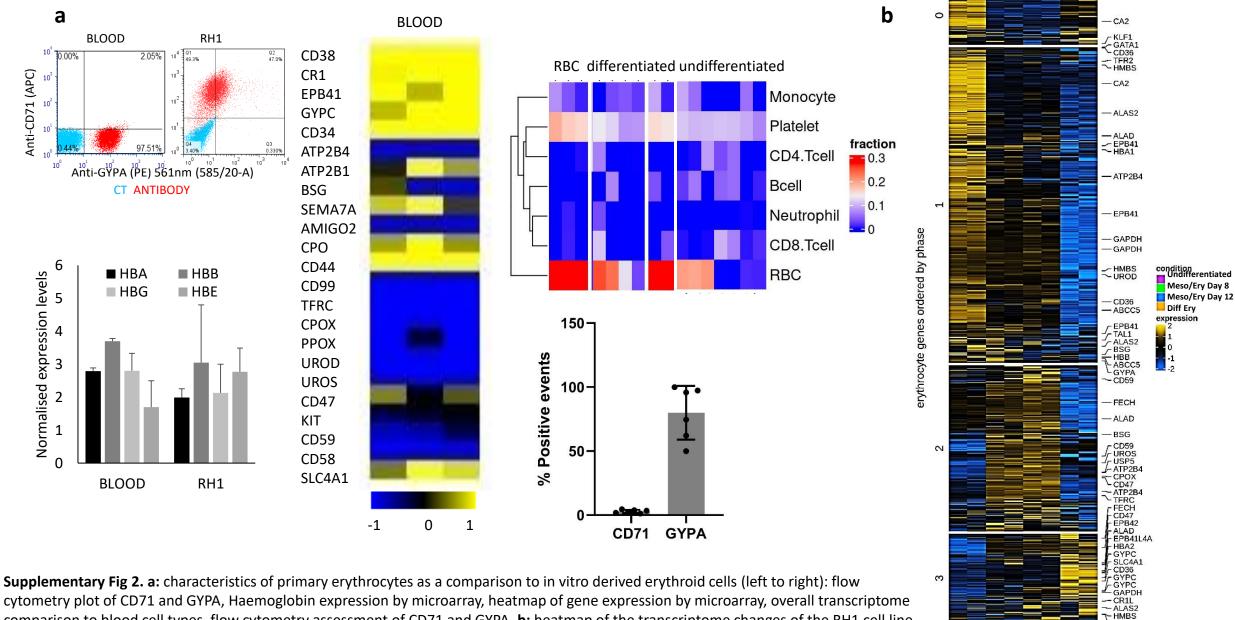
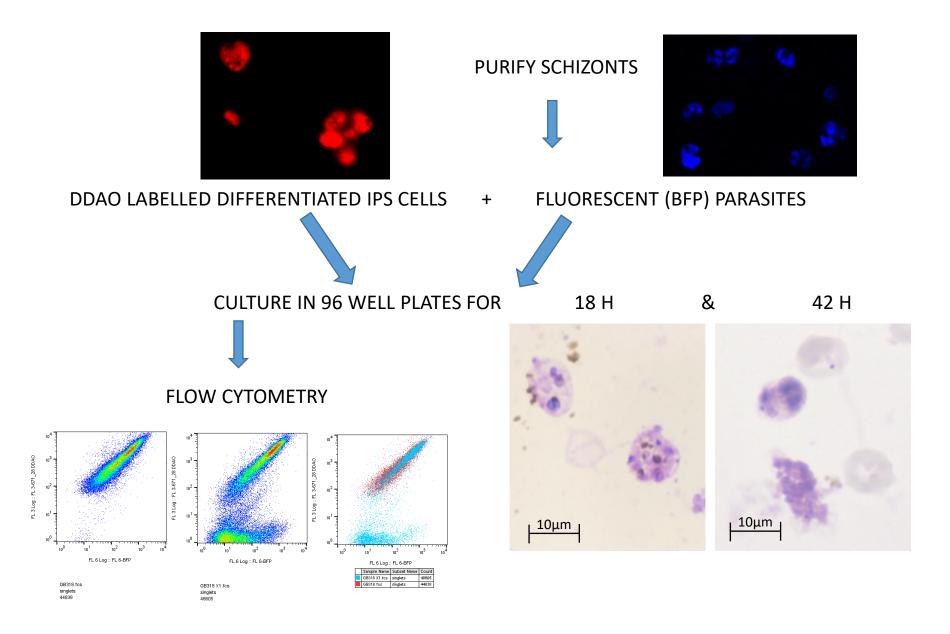


Supplementary Fig 1: schematic representation of the strategy to generate erythroid cells in vitro pluripotent stem cells of diverse origin to assess invasion by P. falciparum



HBG2 - ABCC5 - HBA2 - GYPB

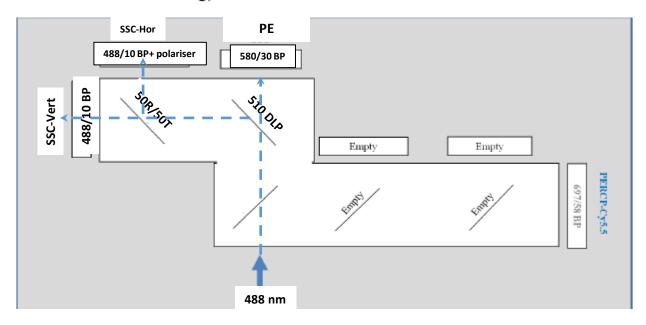
cytometry plot of CD71 and GYPA, Haemoglobin expression by microarray, heatmap of gene expression by microarray, overall transcriptome comparison to blood cell types, flow cytometry assessment of CD71 and GYPA. b: heatmap of the transcriptome changes of the RH1 cell line through the differentiation protocol assessed by microarray.

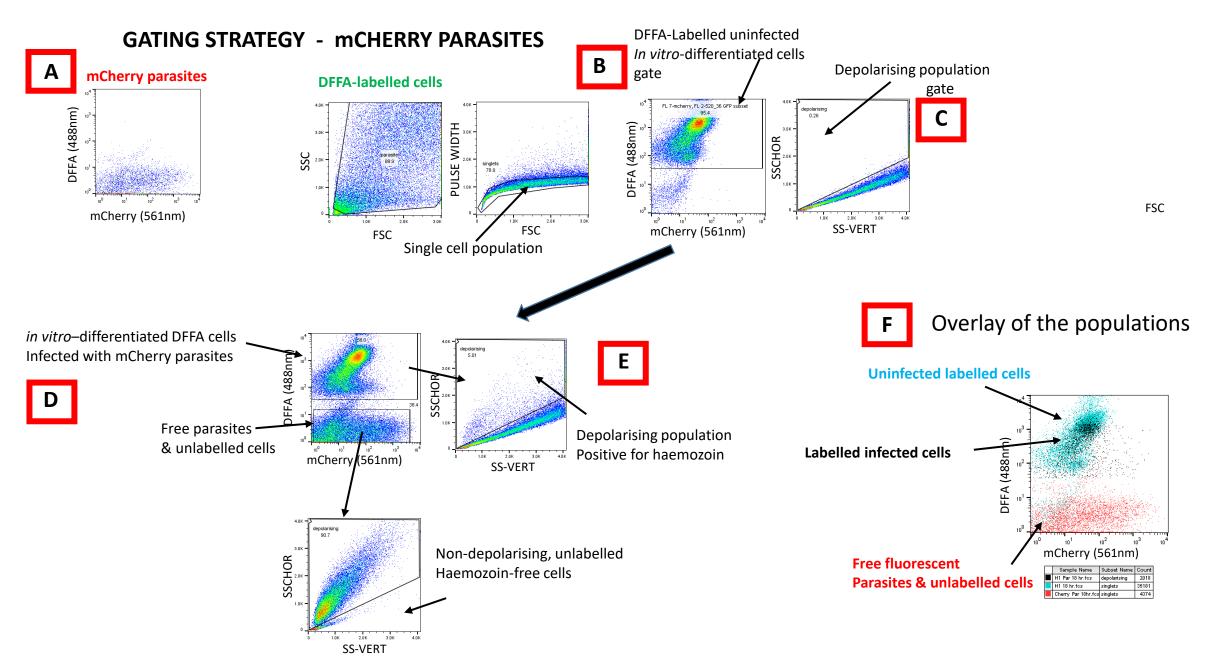


Supplementary Fig 3: design of the invasion assay developed to assess infectivity of *P. falciparum* into *in vitro* derived erythroid cells

MO-FLO FLOW CYTOMETRY OPTICS LAYOUT

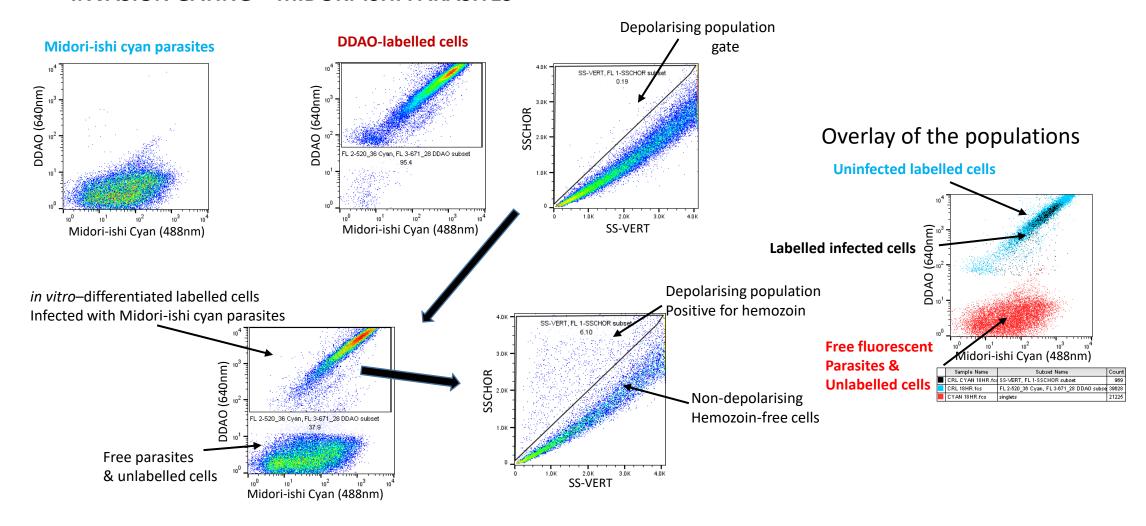
Laser Path 1 - Z Config, 488nm excitation (50mW)



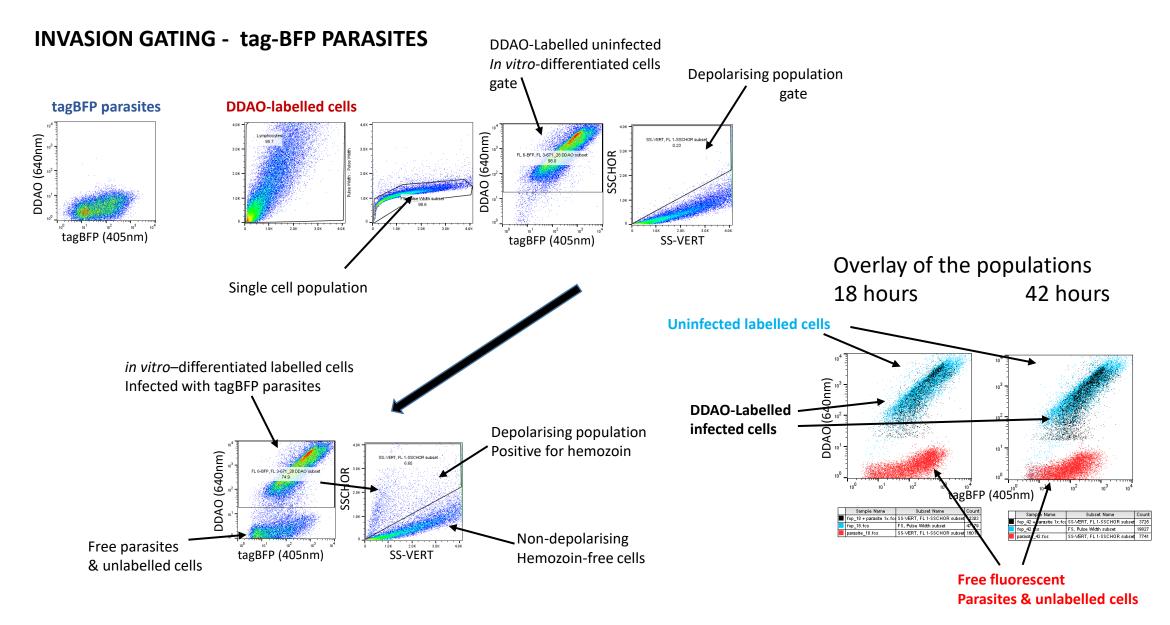


Supplementary Fig 5: gating of invasion assays. A: fluorescent parasites alone; B labelled target cells alone and C depolarisation gate; D co-culture of labelled cells and parasites applying the established gates to assess parasitaemia (E). F overlay of the population of labelled non-infected cells, labelled infected cells and parasites alone.

INVASION GATING - MIDORI-ISHI PARASITES

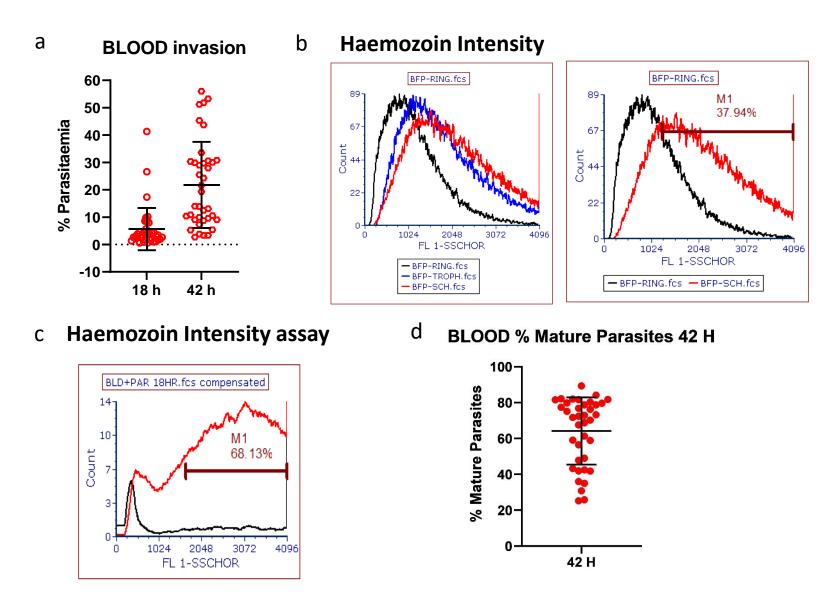


Supplementary Fig 6: gating of invasion assays. Same strategy as Figure 5 but using the far-red cell dye DDAO to label the target cells and parasites expressing Midori-ishi cyan fluorochrome



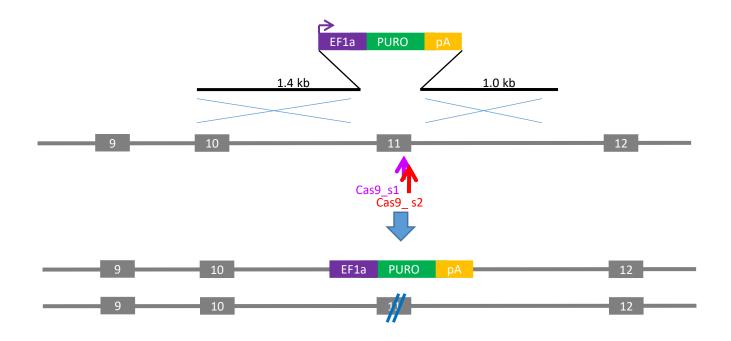
Supplementary Fig 7: gating of invasion assays. Same strategy as Figure 5 but using the far-red cell dye DDAO to label the target cells and parasites expressing tag-BFP fluorochrome

HAEMOZOIN GATE FOR MATURE PARASITES



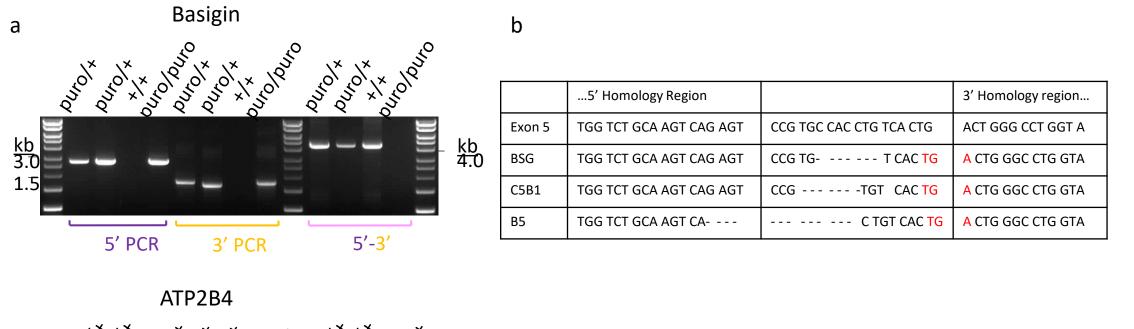
Supplementary Fig 8: invasion quantification in primary erythrocytes. a: Hz depolarisation of invasion assays at 18 and 42 hours. b: depolarisation intensity in synchronised cultures of parasites. c: determination of Hz intensity gate for assessment of growth. d: % of mature parasites at 42 hours of culture in primary erythrocytes are measured by Hz intensity. Results are presented as means and SD.

GENOME EDITING STRATEGY



Supplementary Fig 9: diagram of the genome editing strategy used to knock out Basigin and ATP2B4

GENOTYPING OF EDITED LINES



kb 3.0 1.5		5' Homology Region		3' Homology region
	Exon 11	ATG AGC ACC GTC ATC AGG	AAT CCC AAC GGT GGC TTC	CGT ATG ATC TGC ATA GCT
	ATP2B4 A1	ATG AGC ACC GTC ATC AGG	AAT CCC AAC GGT GC- T TC	GTA TGA TCT GCA TAG
	ATP2B4 E6	ATG AGC ACC GTC ATC AGG	AAT CCC AAC G- T G GCT TC	GTA TGA TCT GCA TAG
5' PCR 3' PCR 5'-3'				

Supplementary Fig 10: assessment of edited cell lines. a: genotyping across the homology arms to detect the insertion of the selection cassette. b: sequencing the second allele to detect damage leading to inactivation.