

Supporting Information:

Dissecting the essentiality of the bifunctional trypanothione synthetase-amidase in *Trypanosoma brucei* using chemical and genetic methods

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Table S1. Intracellular thiols of *T. brucei* transgenic cell lines Thiols were determined as described in Materials and Methods. Each value represents the mean of triplicate determinations.

Cell line	Thiols nmol (10^8 cells) $^{-1}$	
	GSH	T[SH] $_2$
WT	0.47 ± 0.05	0.38 ± 0.02
SKO	0.42 ± 0.02	0.25 ± 0.01
OE	0.56 ± 0.06	0.96 ± 0.05
cDKO + tetracycline	0.51 ± 0.06	0.42 ± 0.04

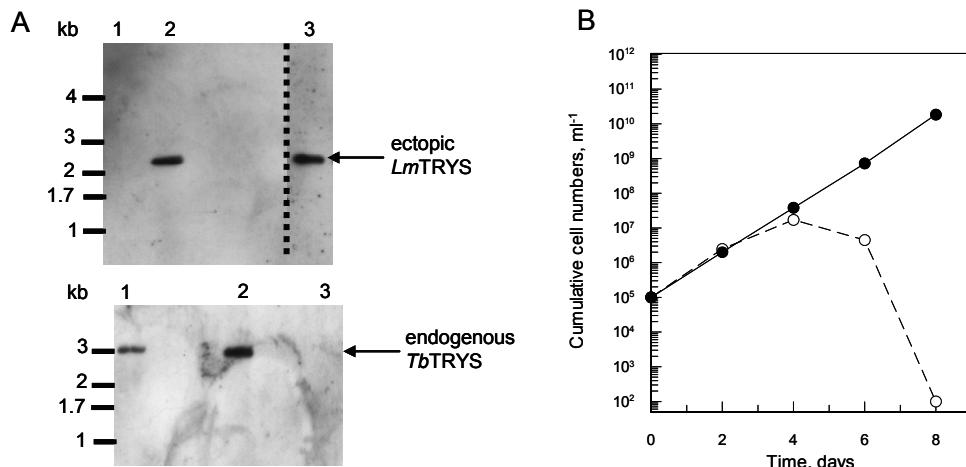


Fig. S1. Complementation of cDKO cells with ectopic *L. major* TRYS. (A) Confirmation of genotype of *T. brucei* TRYS conditional double knockout cell line. Southern-blot analysis of *PstI*-digested genomic DNA (~5 µg) from wild-type *T. brucei* cells (lane 1), *LmTRYSTiΔTRYS::PAC* (lane 2) and *LmTRYSTiΔTRYS::PAC/ΔTRYSTi::HYG* (lane 3). The *LmTRYS* ORF was used as a probe in the upper panel and shows the ectopic copy of *LmTRYSTi* at ~2.5 kb. The *TbTRYS* ORF was used as a probe in the lower panel and the allelic *TbTRYS* can be seen at ~3 kb. Dotted line indicates where the southern image was cropped for clarity. (B) The growth of the cDKO cell line expressing *L. major* TRYS in HMI9-T media was monitored in the presence (closed circles) and absence of tetracycline (open circles, dashed line).

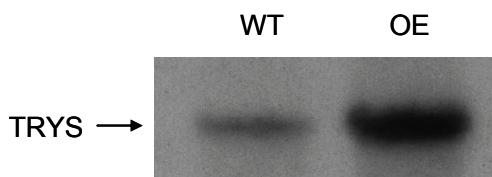


Fig. S2. Immunoblot analysis of a TRYS overexpressing cell line. Immunoblots of cell extracts of WT and OE (plus tetracycline) cells were probed with anti-serum to *T. brucei* TRYS (1×10^7 parasites in each lane).

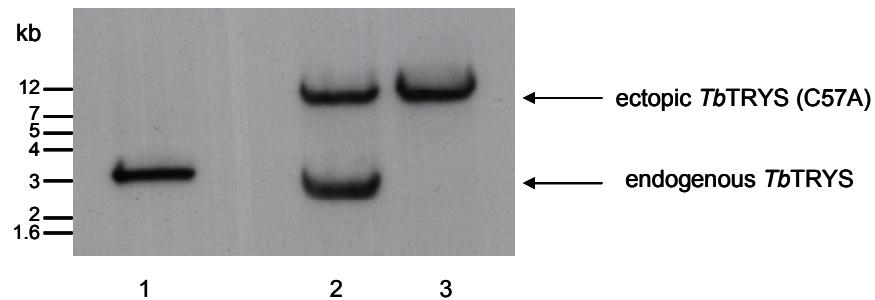


Fig. S3. Genotypic analysis of cDKO(C57A) cell line. Genotypic confirmation of the TRYS cDKO(C57A) cell line. Southern-blot analysis of PstI-digested genomic DNA (~5 µg) from wild-type *T. brucei* cells (lane 1), *TRYS(C57A)^{T1}ΔTRYS::PAC* (lane 2) and *TRYS(C57A)^{T1}ΔTRYS::PAC/ΔTRYS::HYG* (lane 3) cells; the TRYS ORF probe shows allelic *Tb*TRYS at 3 kb and the ectopic copy *Tb*TRYS(C57A)^{T1} at ~10 kb.