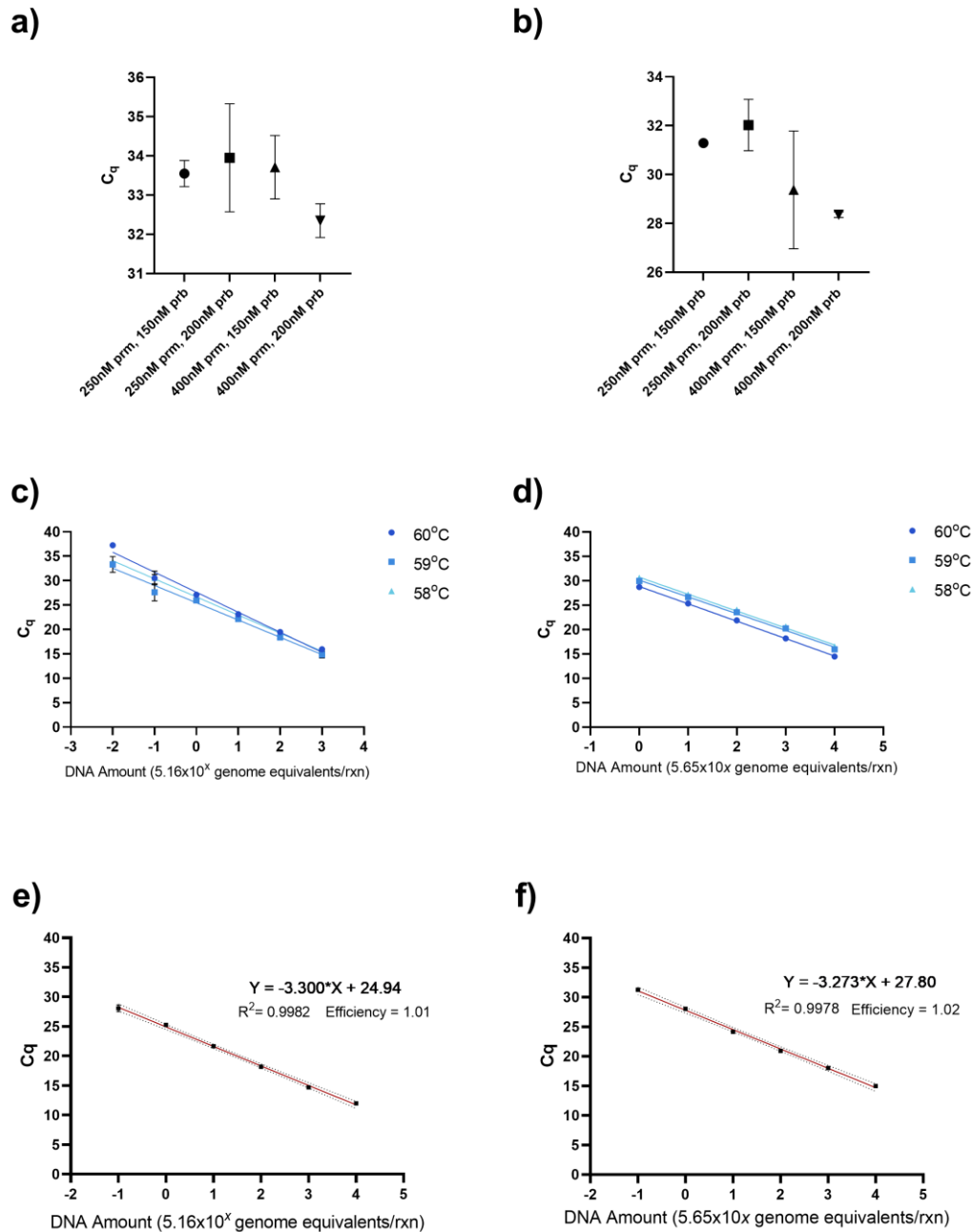


Consistent detection of *Trypanosoma brucei* but not *T. congolense* DNA in faeces of experimentally infected cattle

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S1: (a-b) Charts displaying Cq values obtained from TBR-qPCR (a) and TCS-qPCR (b) screening against four different combinations of two primer and probe concentrations in 10fg/μL respective target DNA in triplicate. Symbols represent the mean, error bars represent the standard deviation. (c-d) Charts displaying standard curve analyses of TBR-qPCR (c) and TCS-qPCR (d) Cq values at different annealing temperatures (58°C, 59°C and 60°C) over ten-fold dilution series (1ng/μL to 10fg/μL) of respective target DNA. Assays performed in triplicate. Symbols represent the mean and error bars represent the standard deviation. (e-f) XY scatter plot with standard curve displaying performance of optimised TBR-qPCR (e) and TCS-qPCR (f) performance over respective detectable ranges in spiked dilution series. In short, 10 ng/μL target DNA of *T. brucei* AnTat 1.1 (e) or *T. congolense* IL3000 (f) was spiked into a composite pre-inoculation faecal DNA sample and subsequently serially diluted 1:10 across five further composite faecal DNA samples. Linear regression best-fit line shown in red with 95% confidence bands displayed as dotted lines.

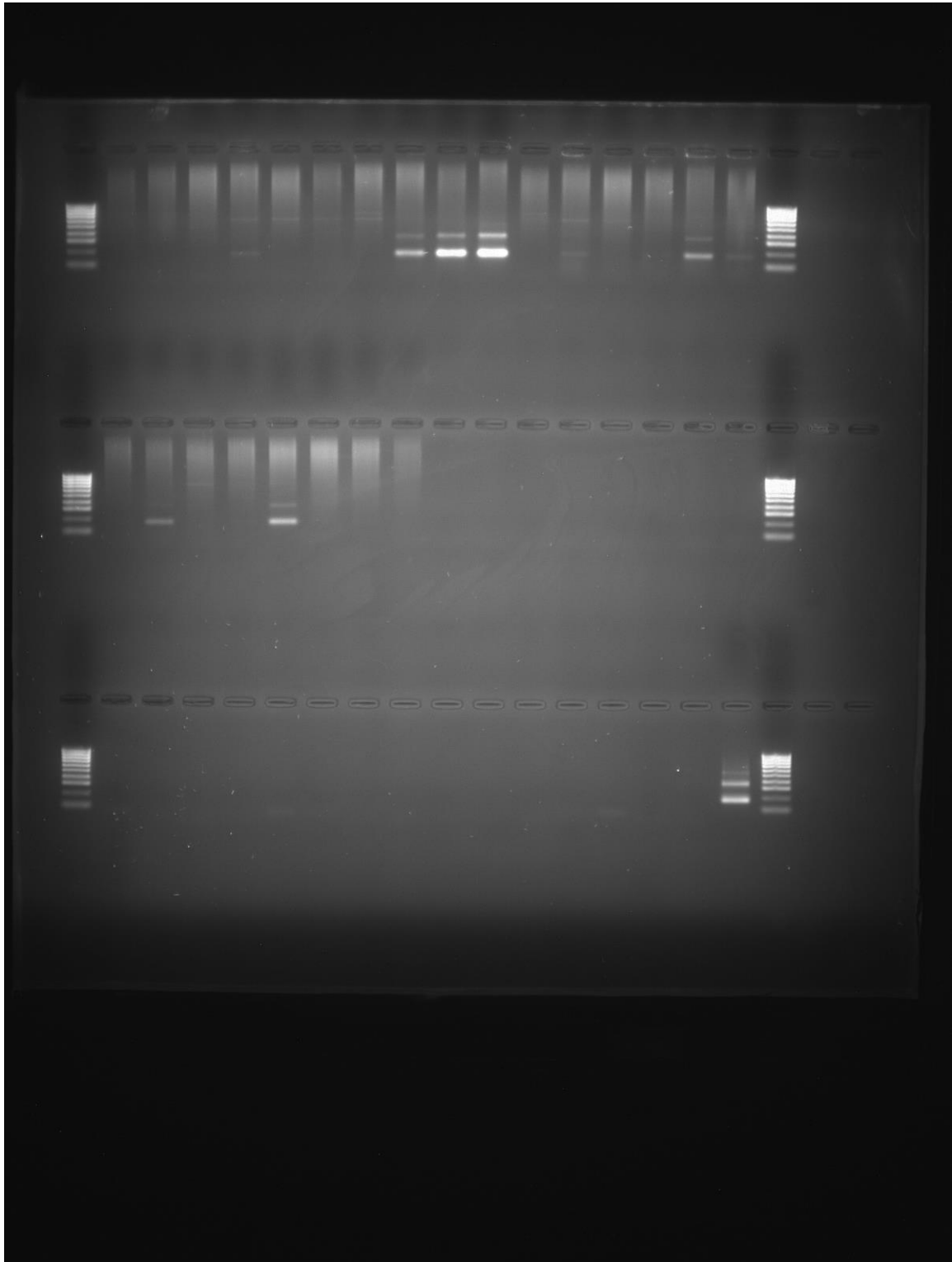
a)

Species	Sub-species	Strain	DNA conc.	Target (Y/N)	Amplification (total = 3)	Mean Cq.
<i>T. brucei</i>	<i>brucei</i>	AnTat 1.1	1ng/μL	Y	3	16.84
<i>T. brucei</i>	<i>gambiense</i>	ELIANE	1ng/μL	Y	3	10.69
<i>T. brucei</i>	<i>rhodesiense</i>	Z3	1ng/μL	Y	3	20.67
<i>T. congolense</i>	Savannah	IL3000	1ng/μL	N	0	-
<i>T. congolense</i>	Forest	ANR3	1ng/μL	N	0	-
<i>T. congolense</i>	Kilifi	WG84	1ng/μL	N	1	38.06
<i>T. godfreyi</i>		Ken7	1ng/μL	N	3	36.36
<i>T. simiae</i>		TV008	1ng/μL	N	2	36.70
<i>T. simiae</i>	Tsavo	114	1ng/μL	N	0	-
<i>T. vivax</i>		Miranda	1ng/uL	N	0	-

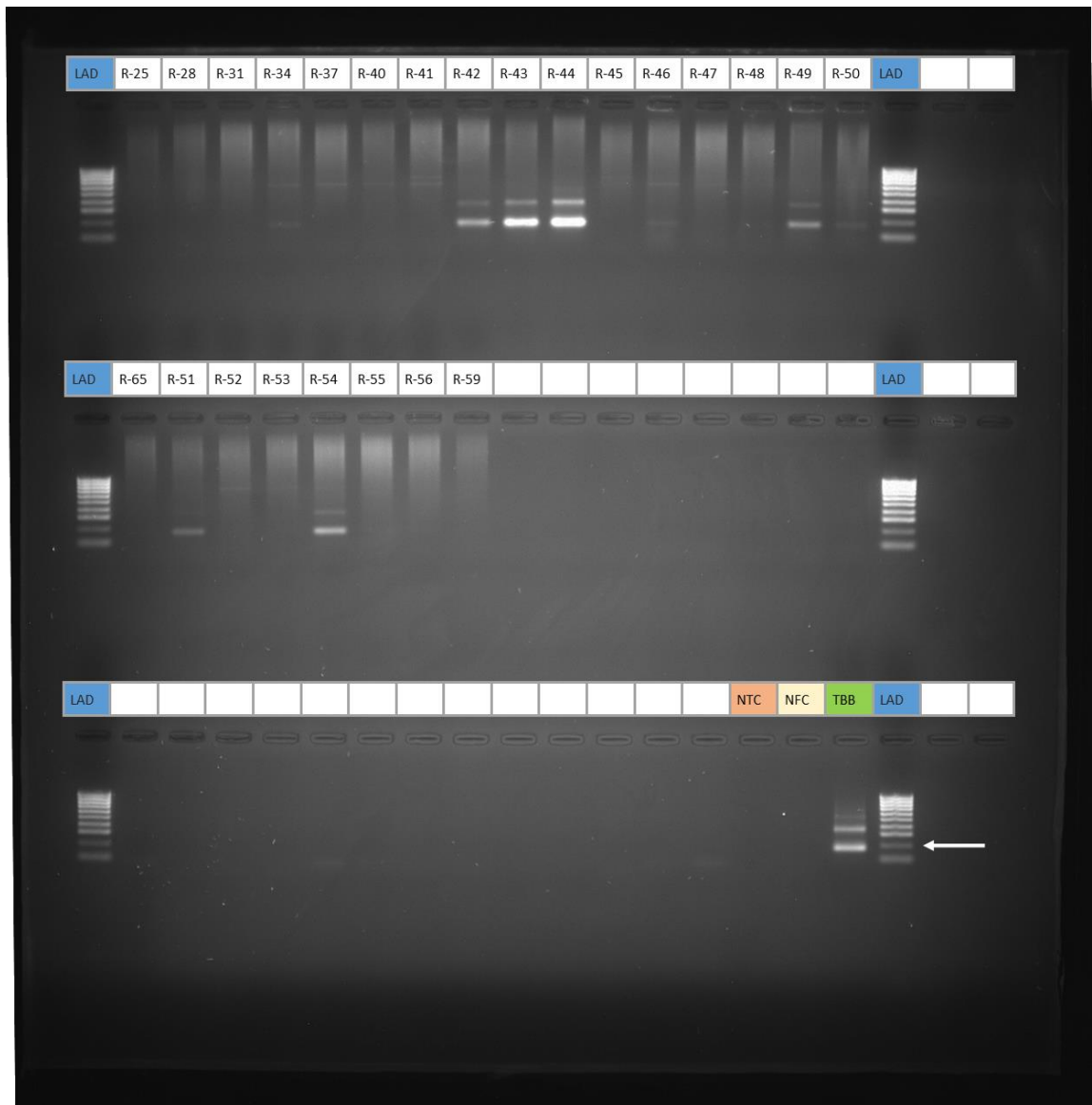
b)

Species	Sub-species	Strain	DNA Conc.	Target (Y/N)	Amplification (total = 3)	Mean Cq.
<i>T. brucei</i>	<i>brucei</i>	AnTat 1.1	1ng/μL	N	0	-
<i>T. brucei</i>	<i>gambiense</i>	ELIANE	1ng/μL	N	0	-
<i>T. brucei</i>	<i>rhodesiense</i>	Z3	1ng/μL	N	0	-
<i>T. congolense</i>	Savannah	IL3000	1ng/μL	Y	3	14.96
<i>T. congolense</i>	Forest	ANR3	1ng/μL	N	3	33.19
<i>T. congolense</i>	Kilifi	WG84	1ng/μL	N	0	-
<i>T. godfreyi</i>		Ken7	1ng/μL	N	0	-
<i>T. simiae</i>		TV008	1ng/μL	N	3	16.70
<i>T. simiae</i>	Tsavo	114	1ng/μL	N	2	36.89
<i>T. vivax</i>		Miranda	1ng/μL	N	0	-

S2: Tables displaying analytical specificity results of optimised TBR-qPCR (a) and TCS-qPCR (b) assays. (a) TBR-qPCR analytical specificity testing was carried out against a panel of 12 *Trypanosoma* sp. DNA samples screened in triplicate. Testing revealed successful amplification in 9/9 target DNA samples (*T. brucei* M249, *T. b. rhodesiense* Z212, *T. b. gambiense* ELIANE in triplicate). However, low-level amplification was also recorded in non-target *Trypanosoma* DNA samples (3/3 *T. godfreyi*, 2/3 *T. simiae* and 1/3 *T. congolense* Kilifi). (b) TCS-qPCR analytical specificity testing revealed amplification in 3/3 target DNA samples (*T. congolense* IL3000 in triplicate). However amplification was also recorded in non-target *Trypanosoma* DNA samples; 3/3 *T. simiae* DNA samples, along with low-level amplification in 3/3 *T. congolense* Forest and 2/3 *T. simiae* Tsavo DNA samples.



S3a: An unannotated and uncropped gel image displaying results of products resulting from TBR-PCR screening of DNA extracted from the first 23 post-inoculation cattle faecal samples. 100bp molecular ladder is present in wells 1 and 18 on each row.



S3b: An annotated, rotated and cropped version of S3a gel image displaying results of products resulting from TBR-PCR screening of DNA extracted from the first 23 post-inoculation cattle faecal samples (three to ten days post-innoculation). LAD = 100bp molecular ladder, NTC = no-template control (nuclease-free water), NFC = negative faeces control. TBB = *T. b. brucei* M249 DNA 1 ng/ μ L (positive control). A white arrow in the lower right corner indicates the position of diagnostic 173 bp TBR-PCR target product, visible in positive control in addition to several samples.

Species	Sub-species	Strain	DNA Concentration
<i>Trypanosoma brucei</i>	<i>brucei</i>	M249	1ng/μL
<i>Trypanosoma brucei</i>	<i>brucei</i>	AnTat 1.1	1ng/μL
<i>Trypanosoma brucei</i>	<i>gambiense</i>	ELIANE	1ng/μL
<i>Trypanosoma brucei</i>	<i>rhodesiense</i>	Z4	1ng/μL
<i>Trypanosoma congolense</i>	Savannah	IL3000	1ng/μL
<i>Trypanosoma congolense</i>	Savannah	GAM2	1ng/μL
<i>Trypanosoma congolense</i>	Forest	ANR3	1ng/μL
<i>Trypanosoma congolense</i>	Kilifi	WG84	1ng/μL
<i>Trypanosoma godfreyi</i>		Ken7	1ng/μL
<i>Trypanosoma simiae</i>		TV008	1ng/μL
<i>Trypanosoma simiae</i>	Tsavo	114	1ng/μL
<i>Trypanosoma vivax</i>		Y486	1ng/μL
<i>Trypanosoma vivax</i>		IL1392	1ng/μL
<i>Trypanosoma vivax</i>		Miranda	1ng/uL

S4: *Trypanosoma* sp. DNA controls used in study.