

CNCD147	MAKEDEVQTLVIDNGSGMCKAGFSGDEAPRSVFPSSVGRPKYKQQLVGGNAKDVFVGDEAC	60
TvACT1	MAEDEVQTLVIDNGSGMCKAGFSGDEAPRSVFPSSVGRPKYKQQLVGGNAKDVFVGDEAC	60

CNCD147	SKAGVLILKYPIEHGIVNNWDDMEKIWHHTFYNELRVDPTEHPVLLTEAPLNPKANREKM	120
TvACT1	SKAGVLILKYPIEHGIVNNWDDMEKIWHHTFYNELRVDPTEHPVLLTEAPLNPKANREKM	120

CNCD147	ISLMFDTFNVPSSFYVGIQAVLSLYSSGRTTGIVFDAGDGVSHTVPIYEGYSLPHAIMRLN	180
TvACT1	ISLMFDTFNVPSSFYVGIQAVLSLYSSGRTTGIVFDAGDGVSHTVPIYEGYSLPHAIMRLN	180

CNCD147	LAGRDLTAWMVKLTERGNATTAEKEIVRDIKEKLCYVALDFDAEMEKAATDSSINVN	240
TvACT1	LAGRDLTAWMVKLTERGNATTAEKEIVRDIKEKLCYVALDFDAEMEKAATDSSINVN	240

CNCD147	YTLPDGNVITIGNERFRCPEMPLFKPYFDGMEYDGIDKTLFDSIMKCIDDVRKDLYANIVL	300
TvACT1	YTLPDGNVITIGNERFRCPEMPLFKPYFDGMEYDGIDKTLFDSIMKCIDDVRKDLYANIVL	300

CNCD147	SGGTTMF N GIAERLDKEITALAPPTMKVKIVAPEERKYAVWVGGSLASLATFPQMVTIK	360
TvACT1	SGGTTMF Q GIAERLDKEITALAPPTMKVKIVAPEERKYAVWVGGSLASLATFPQMVTIK	360

CNCD147	EYDEAGPSIVHRKCF	376
TvACT1	EYDEAGPSIVHRKCF	376

Supplementary Figure S5. Amino acid alignment of the TvACT1 protein of the *T. vaginalis* CNCD147 isolate and the G3 *T. vaginalis* genome. The letters in red indicate the aa changes found after amplicon DNA sequencing.

* Represents identical amino acid.