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Tetrahymena      MREVISIHVGGGGIQLVGNACWELFCLEHGIQPDGQMPSDKTIGGGDDAFNTFFSETGAGK 60
Toxoplasma      MREVISIHVGGAGIQLVGNACWELFCLEHGIQPDGQMPSDKTIGGGDDAFNTFFSETGAGK 60
*****.***.*****.*****.*****.*****.*****.*****.*****.*****

Tetrahymena      HVPRVFLDLEPTVIVDEVRTGTYRQLFHPEQLISGKEDAANNFARGHYTIGKEIVDLCLD 120
Toxoplasma      HVPRCVFLDLEPTVVDEVRTGTYRHLFHPEQLISGKEDAANNFARGHYTIGKEIVDLSLD 120
****.*****.*****.*****.*****.*****.*****.*****.*****.***

Tetrahymena      RIRKLADNCTGLQGEIVFNLSVGGGTGSGLGSLLLERLSVDYGKSKLGFYIYPSPOVSTA 180
Toxoplasma      RIRKLADNCTGLQGEIVFNLSVGGGTGSGLGSLLLERLSVDYGKSKLNFCSWPSPOVSTA 180
*****.***.*****.*****.*****.*****.*****.*****.*****.*****

Tetrahymena      VVEPYNSILSTHSLLEHTDVAVMLDNEAIYDICRRNLDIERPTYTNLNRLIAQVISSITA 240
Toxoplasma      VVEPYNSVLSHSLLEHTDVAVMLDNEAIYDICRRNLDIERPTYTNLNRLIAQVISSITA 240
*****.*****.*****.*****.*****.*****.*****.*****.*****.***

Tetrahymena      SIRFDGALNVDVTEFQTNLVPYPRIHFMSSYAPIISAEKAYHEQLSVAEITNSAFEPAN 300
Toxoplasma      SIRFDGALNVDVTEFQTNLVPYPRIHFMSSYAPIISAEKAYHEQLSVAEITNSAFEPAS 300
*****.*****.*****.*****.*****.*****.*****.*****.*****.***

Tetrahymena      MMAKCDPRHGKYMACSMYRGDVVPKDVNASIATIKTKRTIQFVDWCPTGFKVGINYQPP 360
Toxoplasma      MMAKCDPRHGKYMACCLMYRGDVVPKDVNAAVATIKTKRTIQFVDWCPTGFKCGINYQPP 360
*****.***.*****.*****.***.*****.*****.*****.*****.*****

Tetrahymena      TVVPGDLAKVMRAVCMISNSTAIAEVFSRLDHKFDLMYAKRAVHWYVGEEMEEGEFSE 420
Toxoplasma      TVVPGDLAKVMRAVCMISNSTAIAEVFSRMDHKFDLMYAKRAVHWYVGEEMEEGEFSE 420
*****.*****.*****.*****.*****.*****.*****.*****.*****.*****

Tetrahymena      AREDLAALEKDYEEVGIETAEGEGEEEGY---- 449
Toxoplasma      AREDLAALEKDYEEVGIETAEGEGEEEGYGDEY 453
*****.*****.*****.*****.*****.*****.*****.*****.*****.*****

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Supplemental Figure S1: A clustal alignment of the amino acid sequence of *T. gondii* α 1-tubulin and *T. thermophila* α -tubulin. Residues that differ between the two proteins are highlighted in yellow. Locations of the point mutations H28Q, L136F, R243S, T239I and I252L that were introduced into *T. thermophila* α -tubulin are boxed. Individual amino acids are colored to reflect their chemical properties: blue: acidic; green: hydroxyl/amine/basic/Q; magenta: basic; and red: small, hydrophobic (including aliphatic Y).