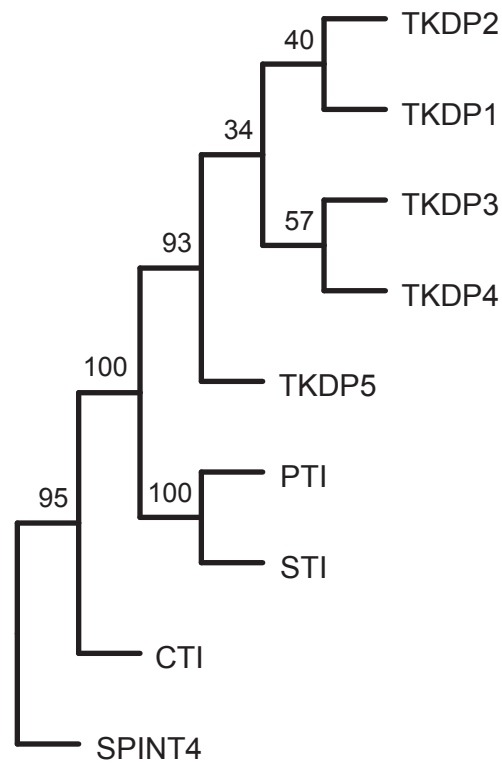




Additional file 8 - Figure S5. Relationship between bovine CTI, PTI, STI, the TKDPs and SPINT4

Phylogenetic relationship between the protein-coding regions of the *CTI*, *PTI*, *STI*, *TKDP1-5* and *SPINT4* transcripts. The tree was constructed using the maximum likelihood model of nucleotide substitution based upon the assumption of a molecular clock. The *SLPI* transcript (omitted) was used to root the tree.