

# BSgenome.Tgondii.ToxoDB.7.0

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Tgondii	<i>Toxoplasma gondii</i> ME49 (ToxoDB-7.0)
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## Description

*Toxoplasma gondii* ME49 genome Release 7.0 available at <http://www.toxodb.org>

## Note

This BSgenome data package was made from the following source data files:

-- information not available --

See [?BSgenomeForge](#) and the BSgenomeForge vignette (`vignette("BSgenomeForge")`) in the BSgenome software package for how to make a BSgenome data package.

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## See Also

[BSgenome-class](#), [DNAString-class](#), [available.genomes](#), [BSgenomeForge](#)

## Examples

```
if ("AGAPS" %in% masknames(Tgondii)) {  
  
  ## Check that the assembly gaps contain only Ns:  
  checkOnlyNsInGaps <- function(seq)  
  {  
    ## Replace all masks by the inverted AGAPS mask
```

```
    masks(seq) <- gaps(masks(seq)["AGAPS"])
    unique_letters <- uniqueLetters(seq)
    if (any(unique_letters != "N"))
      stop("assembly gaps contain more than just Ns")
  }

  ## A message will be printed each time a sequence is removed
  ## from the cache:
  options(verbose=TRUE)

  for (seqname in seqnames(Tgondii)) {
    cat("Checking sequence", seqname, "... ")
    seq <- Tgondii[[seqname]]
    checkOnlyNsInGaps(seq)
    cat("OK\n")
  }
}

## See the GenomeSearching vignette in the BSgenome software
## package for some examples of genome-wide motif searching using
## Biostrings and the BSgenome data packages:
if (interactive())
  vignette("GenomeSearching", package="BSgenome")
```

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