

Organism	Origin of data	Status (date)
<i>Acanthamoeba catellanii</i>	TBestDB <sup>1</sup>	ESTs and clustered ESTs
<i>Acropora millepora</i>	Ensembl <sup>2</sup>	traces (source date: 06/05/2007)
<i>Acropora palmata</i>	Ensembl	traces (source date: 06/05/2007)
<i>Acyrtosiphon pisum</i>	Ensembl	traces (dod: 17.07.07)
<i>Alternaria brassicicola</i>	WUSTL GSC <sup>3</sup>	scaffolds (v1.0)
<i>Aplysia californica</i>	NCBI traceDB <sup>4</sup>	traces (dod: 05.07.07)
<i>Aspergillus nidulans</i>	Broad <sup>5</sup>	scaffolds & putative genes (release 4)
<i>Aspergillus terreus</i>	Broad	scaffolds (source date: 05/05/2006) & CDS (source date: 05/09/2007) (assembly 1)
<i>Batrachochytrium dendrobatidis</i>	Broad	scaffolds (source date: 02/16/2007), CDS (source date: 02/28/2007) & excluded reads (source date: 02/16/2007) (assembly 1)
<i>Biomphalaria glabrata</i>	NCBI traceDB	traces (dod: 05.07.07)
<i>Blastocystis hominis</i>	TBestDB	ESTs and clustered ESTs
<i>Botrytis cinerea</i>	Broad	scaffolds (source date: 04/25/2006) & putative genes (dod: 16.05.2007) (assembly 1)
<i>Brugia malayi</i>	Ensembl	traces (dod: 21.02.2007)
<i>Caenorhabditis elegans</i>	Wormbase <sup>6</sup>	chromosomes - WS177
<i>Caenorhabditis japonica</i>	NCBI traceDB	traces (dod: 14.03.07)
<i>Caenorhabditis remanei</i>	NCBI traceDB	traces (dod: 14.03.07)
<i>Caenorhabditis</i> sp. 4 (CE-II)	NCBI traceDB	traces (dod: 14.03.07)
<i>Candida glabrata</i> CBS 138	Génolevures <sup>7</sup>	Génolevures 2 Release 2: chromosomes (source date: 2004/06/30) & CDS (source date: 2006/08/17)
<i>Capitella capitata</i>	JGI <sup>8</sup> (ftp)	contigs (dod: 28.06.07)
<i>Capitella</i> sp. I ECS-2004	Ensembl	traces (dod: 21.02.07)
<i>Cerebratulus lacteus</i>	Ensembl	traces (dod: 21.02.07)
<i>Chaetomium globosum</i> CBS 148.51	Broad	scaffolds (source date: 03/14/2006) & putative genes (source date: 04/26/2006)
<i>Chlamydomonas reinhardtii</i>	JGI	scaffolds (v3.1) & 1st milestone transcripts
<i>Ciona intestinalis</i>	JGI	scaffolds & transcripts (v2.0)
<i>Coprinopsis cinerea</i>	Broad	scaffolds (assembly 1) & putative genes (gene set v1.0)
<i>Cyanidioschyzon merolae</i> strain 10D	Univ. of Tokio <sup>9</sup>	scaffolds (source date: 16.11.2004) & coding regions (source date: 17.09.2005)
<i>Daphnia pulex</i>	Ensembl	traces (dod: 21.02.07)
<i>Dictyostelium discoideum</i>	NCBI	complete genome (dod: 09.05.07) & coding regions (source date: 10.06.07)
<i>Encephalitozoon cuniculi</i> GB-M1	NCBI	complete genome & coding regions (Build 1.1)
<i>Euglena gracilis</i>	TBestDB	ESTs and clustered ESTs
<i>Euprymna scolopes</i>	Ensembl	traces (dod: 21.02.07)
<i>Fusarium verticillioides</i>	Broad	scaffolds & transcripts (assembly 3)
<i>Helobdella robusta</i>	Ensembl	traces (dod: 21.02.07)
<i>Heterorhabditis bacteriophora</i>	Ensembl	traces (dod: 21.02.07)
<i>Hydra magnipapillata</i>	NCBI traceDB	traces (dod: 11.06.07)
<i>Ixodes scapularis</i>	Ensembl	traces (dod: 21.02.07)
<i>Karlodinium micrum</i>	TBestDB	ESTs and clustered EST
<i>Laccaria bicolor</i>	JGI	scaffolds & transcripts (v1.0)

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<sup>1</sup><http://tbestdb.bcm.umontreal.ca>

<sup>2</sup><http://www.ensembl.org>

<sup>3</sup><http://genome.wustl.edu>

<sup>4</sup><ftp://ftp.ncbi.nih.gov/pub/TraceDB>

<sup>5</sup><http://www.broad.mit.edu>

<sup>6</sup><ftp://ftp.wormbase.org/pub/wormbase>

<sup>7</sup><http://cbi.labri.fr/Genolevures>

<sup>8</sup><http://genome.jgi-psf.org>

<sup>9</sup><http://merolae.biol.s.u-tokyo.ac.jp>

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Organism	Origin of data	Status (date)
Leishmania major strain Friedlin	Sanger <sup>10</sup>	genome v5.2
Lottia gigantea	Ensembl	traces (dod: 21.02.07)
Magnaporthe grisea	Broad	scaffolds, excluded reads & predicted transcripts (assembly release 5)
Malus x domestica	TFT <sup>11</sup>	contigs (apple database v4.0)
Mastigamoeba balamuthi	TBestDB	ESTs and clustered EST
Monosiga brevicollis	JGI	scaffolds, cDNA reads & transcripts (v1.0)
Naegleria gruberi	JGI	scaffolds, transcripts & ESTs (v1.0)
Nectria haematococca	JGI	chromosomes & transcripts (v2.0)
Nematostella vectensis	JGI	scaffolds & transcripts (v1.0)
Neosartorya fischeri	TIGR <sup>12</sup>	scaffolds & coding regions (dod: 14.05.07)
Neurospora crassa	Broad	scaffolds (assembly 7) & CDS (release 3)
Oryza sativa	RAP-DB <sup>13</sup>	IRGSP build 4.0
Oscarella carmela	CIG UC Berkeley <sup>14</sup>	ESTs (dod: 15.07.07)
Ostreococcus lucimarinus	JGI	chromosomes & transcripts (v2.0)
Ostreococcus tauri	JGI	chromosomes & transcripts (v2.0)
Oxytricha trifallax	Ensembl	traces (dod: 21.02.07)
Paramecium tetraurelia	Genoscope <sup>15</sup>	scaffolds (v2.1) & mRNAs (v1.0)
Pediculus humanus	Ensembl	traces (dod: 21.02.07)
Petromyzon marinus	WUSTL GSC	scaffolds (assembly v3.0)
Phaeodactylum tricornutum	JGI	chromosomes & transcripts (v2.0)
Phanerochaete chrysosporium	JGI	scaffolds (v2.0) & transcripts (v2.1)
Physarum polycephalum	NCBI/TBestDB	traces/ESTs and clustered ESTs
Physcomitrella patens	JGI	scaffolds & transcripts (v1.1)
Phytophthora ramorum	JGI	scaffolds & transcripts (v1.1)
Plasmodium falciparum 3D7	NCBI	chromosomes & coding regions
Platynereis dumerilii	Ensembl	traces (dod: 21.02.07)
Polysphondylium pallidum	TBestDB	ESTs and clustered EST
Populus trichocarpa	JGI	scaffolds (v1.0) & transcripts (v1.1)
Porites lobata	Ensembl	traces (dod: 21.02.07)
Postia placenta	JGI	scaffolds & transcripts (v1.0)
Prunus dulcis	CUGI <sup>16</sup>	contigs (dod: 23.05.07)
Prunus persica	CUGI	contigs (dod: 23.05.07)
Puccinia graminis f. sp. tritici	Broad	scaffolds (assembly 1)
Pyrenophora tritici-repentis	Broad	scaffolds (assembly 1)
Reniera sp. JGI-2005	Ensembl	traces (dod: 21.02.07)
Saccharomyces cerevisiae S288c	Stanford <sup>17</sup>	chromosomes, ESTs & CDSs (dod: 05/03/2007)
Saccoglossus kowalevskii	Ensembl	traces (dod: 10.05.07)
Schistosoma mansoni	Sanger	scaffolds (assembly release v3.1) & predicted genes (v4.0)
Schizosaccharomyces pombe 972h-	Sanger	chromosomes & CDS (dod: 05/03/2007)
Schmidtea mediterranea	WUSTL GSC	scaffolds (v3.1)
Spisula solidissima	Ensembl	traces (dod: 21.02.07)
Stagonospora nodorum	Broad	scaffolds & putative genes (assembly 1)
Tetrahymena thermophila	TIGR	scaffolds (closed 30.09.06) & gene predictions (annotation release 31.08.06)
Thalassiosira pseudonana	JGI	chromosomes & transcripts (v3.0)
Trichoderma reesei	JGI	scaffolds & transcripts (v2.0)
Trichomonas vaginalis	TIGR	scaffolds and unplaced contigs (31.03.2005) & gene predictions (15.02.2007)

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<sup>10</sup><http://www.sanger.ac.uk>

<sup>11</sup><http://genomics.msu.edu/fruitdb>

<sup>12</sup><http://www.tigr.org>

<sup>13</sup><http://rapdb.lab.nig.ac.jp>

<sup>14</sup><http://cigbrowser.berkeley.edu/oscarella/>

<sup>15</sup><http://www.genoscope.cns.fr>

<sup>16</sup><ftp://ftp.genome.clemson.edu/pub>

<sup>17</sup><http://genome-www.stanford.edu>

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Organism	Origin of data	Status (date)
Trypanosoma brucei TREU927	NCBI	chromosomes & CDS (Build 1.1)
Trypanosoma cruzi	TeruziDB <sup>18</sup>	contigs & CDS (Release 5.1)
Uncinocarpus reesii	Broad	scaffolds & putative genes (assembly 2)
Ustilago maydis	Broad	scaffolds, excluded reads & putative genes (assembly 1 release 2)
Volvox carteri	JGI	scaffolds, ESTs & transcripts (v1.0)
Yarrowia lipolytica CLIB122	Génolevures	Génolevures 2 Release 2: chromosomes (source date: 03.07.2004) & CDS (source date: 17.08.2006)

Table 1: Overview of the origin and status of the screened sequences. dod: day of download.

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<sup>18</sup><http://tcruzidb.org/tcruzidb>