ID   AC010724; SV 6; linear; genomic DNA; STD; HUM; 157200 BP.
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AC   AC010724;
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DT   23-SEP-1999 (Rel. 61, Created)
DT   26-FEB-2004 (Rel. 78, Last updated, Version 13)
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DE   Homo sapiens BAC clone RP11-152F13 from 15, complete sequence.
XX
KW   HTG.
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OS   Homo sapiens (human)
OC   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia;
OC   Eutheria; Euarchontoglires; Primates; Haplorrhini; Catarrhini;
Hominidae;
OC   Homo.
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RN   [1]
RP   1-157200
RX   PUBMED; 9847074.
RA   Wilson R.;
RT   "Toward a complete human genome sequence";
RL   Genome Res. 8(11):1097-1108(1998).
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RN   [2]
RP   1-157200
RA   Cordes M., Maupin R., Myers M.;
RT   "The sequence of Homo sapiens BAC clone RP11-152F13";
RL   Unpublished.
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RN   [3]
RP   1-157200
RA   Waterston R.H.;
RT   ;
RL   Submitted (21-SEP-1999) to the INSDC.
RL   Genome Sequencing Center, Washington University School of Medicine,
4444
RL   Forest Park Parkway, St. Louis, MO 63108, USA
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RN   [4]
RP   1-157200
RA   Waterston R.H.;
RT   ;
RL   Submitted (26-JUL-2000) to the INSDC.
RL   Genome Sequencing Center, Washington University School of Medicine,
4444
RL   Forest Park Parkway, St. Louis, MO 63108, USA
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RN   [5]
RP   1-157200
RA   Waterston R.;
RT   ;
RL   Submitted (08-NOV-2000) to the INSDC.
RL   Department of Genetics, Washington University, 4444 Forest Park
Avenue, St.
RL   Louis, Missouri 63108, USA
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RP   1-157200
RA   Waterston R.;
RT   ;
RL   Submitted (09-MAY-2001) to the INSDC.

RL   Department of Genetics, Washington University, 4444 Forest Park Avenue, St.
RL   Louis, Missouri 63108, USA
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DR   EPD; EP41007; HS_RPS17.
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DR   Ensembl-Gn; ENSG00000184779; homo_sapiens.
DR   Ensembl-Gn; ENSG00000196648; homo_sapiens.
DR   Ensembl-Gn; ENSG00000214575; homo_sapiens.
DR   Ensembl-Gn; ENSG00000261086; homo_sapiens.
DR   Ensembl-Scaffolds; AC010724.6:1-157200; homo_sapiens.
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DR   GOA; H3BNX1.
DR   GOA; H3BPD6.
DR   GOA; H3BUA9.
DR   GOA; J3KN28.
DR   GOA; P0CW22.
DR   HGNC; HGNC:10397; RPS17.
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DR   HGNC; HGNC:40029; RPS17L.
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DR   UniProtKB/TrEMBL; H3BQG9; H3BQG9_HUMAN.
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CC   On Jul 26, 2000 this sequence version replaced gi:7630867.
CC   -------------- Genome Center
CC   Center: Washington University Genome Sequencing Center
CC   Center code: WUGSC
CC   Web site: http://genome.wustl.edu/gsc
CC   Contact: sapiens@watson.wustl.edu
CC   -------------- Summary Statistics
CC   Center project name: H_NH0152F13
CC   --------------.
CC   NOTICE:  This sequence may not represent the entire insert of this
CC   clone.  It may be shorter because we only sequence overlapping
CC   clone sections once, or longer because we provide a small overlap
CC   between neighboring data submissions.
CC   This sequence was finished as follows unless otherwise noted:
CC   all regions were double stranded, sequenced with an alternate
CC   chemistry, or covered by high quality data (i.e., phred quality >=
CC   30); an attempt was made to resolve all sequencing problems, such
CC   as compressions and repeats; all regions were covered by sequence
CC   from more than one subclone; and the assembly was confirmed by
CC   restriction digest.
CC   MAPPING INFORMATION:
CC   Mapping information for this clone was provided by Dr. John D.
CC   McPherson, Department of Genetics, Washington University, St. Louis
CC   MO.  For additional information about the map position of this
CC   sequence, see http://genome.wustl.edu/gsc
CC   SOURCE INFORMATION:
CC   The RPCI-11 human BAC library was made from the blood of one male
CC   donor, as described by Osoegawa,K., Woon,P.Y., Zhao,B., Frengen,E.,
CC   Tateno,M., Catanese,J.J. and de Jong,P.J. (1998) An improved
CC   approach for construction of bacterial artificial chromosome
CC   libraries.  Genomics 51:1-8.  The clone may be obtained either from
CC   Research Genetics, Inc. (http://www.resgen.com) or Pieter de Jong
CC   and coworkers at the Roswell Park Cancer Institute
CC   (http://bacpac.med.buffalo.edu)
CC   VECTOR:  pBACe3.6
CC   NEIGHBORING SEQUENCE INFORMATION:
CC   The clone sequenced to the right is RP11-96J23.  Actual start of
CC   this clone is at base position 1 of RP11-152F13; actual end is at
CC   base position 157200 of RP11-152F13.
CC   The sequence RP11-152F13 from base position 85582 to 85867 is
CC   represented by sequence derived from PCR from clone DNA.
CC   The sequence RP11-152F13 from base position 30413 to 40908 is the
CC   first copy of a duplication.  The region has single m13 subclone
CC   coverage, the sequence is believed to be correct.  The region
CC   corresponds to restriction digest information:  hindiii band size
CC   1137 real and 1133 in silico, 5742 real and 6043 in silico, 9770
CC   real and 9654 in silico; ecorv band size 5948 real and 5958 in
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