

Resource Summary Report

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ADAPT: A Database of Affymetrix Probesets and Transcripts

RRID:SCR_008688

Type: Tool

Proper Citation

ADAPT: A Database of Affymetrix Probesets and Transcripts (RRID:SCR_008688)

Resource Information

URL: <http://bioinformatics.picr.man.ac.uk/adapt/Welcome.adapt>

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Description: Library of many-to-many relationships between Affymetrix probesets transcripts and genes, by directly mapping every probe against publicly available mRNAs/cDNA sequences from RefSeq and Ensembl. You can search the database by AffymetrixProbeset ID, Transcript Accession, Gene Symbol or Gene Accession. Currently, the ADAPT database holds information on 23 Array Types containing 255771 unique Probesets, that themselves map onto 252788 entries in both RefSeq and Ensembl. ADAPT is clusterable, cross platform, runs on any Java Virtual machine after version 1.4, and utilises the following software: :- Apache Tomcat :- Apache Struts :- JBoss Application Server :- Postgresql Sponsors: ADAPT has been written with funding by Cancer Research UK. Keywords: Database, Gene, Transcript, Mapping, Probe, mRNA, cDNA, Sequence, Association, Array, Probeset,

Synonyms: ADAPT

Resource Type: database, data or information resource

Resource Name: ADAPT: A Database of Affymetrix Probesets and Transcripts

Resource ID: SCR_008688

Alternate IDs: nif-0000-36878

Record Creation Time: 20220129T080248+0000

Record Last Update: 20260725T191156+0000

Ratings and Alerts

No rating or validation information has been found for ADAPT: A Database of Affymetrix Probesets and Transcripts.

No alerts have been found for ADAPT: A Database of Affymetrix Probesets and Transcripts.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We have not found any literature mentions for this resource.