

Tfdp1 Cul4a And Cdc16 Identified As Targets For Amplification At 13q34 In Hepatocellular Carcinomas

**tfdp1, cul4a, and cdc16 identified as targets for ...** - tfdp1, cul4a, and cdc16 identified as targets for amplification at 13q34 in hepatocellular carcinomas kohichiroh yasui,1 shigeki arii,2 chen zhao,1 issei imoto,1 masakazu ueda,3 hisaki nagai,4

**ptk2 and eif3s3 genes may be amplification targets at ...** - recently we identified tfdp1, cul4a, and cdc16 as targets of amplification at 13q34 in hccs.7 cgh allows a global overview of changes in dna copy number throughout the genome of a tumor. in the work reported here, we examined 39 primary hcc tumors by cgh to explore genomic alterations that might affect initiation and/or progression of liver tumors. among losses and gains involving several ...

**de novo large rare copy-number variations contribute to ...** - article open de novo large rare copy-number variations contribute to conotruncal heart disease in chinese patients christopher cy mak 1, pak cheong chow , anthony py liu , kelvin yk chan2, yoyo wy ...

**list of lincnas - arraystar** - list of lincnas the lincnas collected for the mouse cancer pathways are listed in table 1. click an lincna accession number to see its detailed information and its potential target gene(s) in cancer.

**cd1d gene is a target for a novel amplicon at 1q22.1 in ...** - tfdp1, cul4a, cdc16 and eef1a2 have been found to be associated with increased copy numbers at 8q, 13q and 20q locus [14-17]. to investigate the regions with high-level amplification and their potential target genes within these amplicons in shi-guang zhang and wen-qin song contributed equally to this work. s.-g. zhang w.-q. song laboratory of chromosome research, college of life science ...

**table s1. gene sets used for gsea - cancer research** - table s1. gene sets used for gsea ets target set (in queried microarray dataset) myc upregulated target set abi2 abce1 adamtsl2 abl1 ahsa1 acat1 aldh3a2 aco1

**postnatal diagnosis of constitutive ring chromosome 13 ...** - genetics and molecular research 14 (1): 1692-1699 (2015) funpec-rp funpecrp postnatal diagnosis of constitutive ring chromosome 13 using both conventional and

**oligo gearray dna microarray - cosmo bio** - catalog number. format: ohs-020 hybtube (standard protocol) ehs-020 hybplate (higher throughput protocol) description . the oligo gearray human cell cycle microarray profiles the expression of 112 genes key to cell cycle regulation.

**case report smallest critical region for microcephaly in a ...** - genetics and molecular research 12 (2): 1311-1317 (2013) funpec-rp funpecrp case report smallest critical region for microcephaly in

**13q deletion in a girl contributing to antenatal stroke ...** - cdc16 cell division cycle 16 (source: hgnc symbol; acc: 1720) 13q34 upf3a upf3 regulator of nonsense transcripts homolog a (yeast) (source: hgnc symbol; acc: 20332) 13q34

**supplementary materials - liebertpub** - absorption 39 mt1x, mt1g, mt1f nucleotide excision repair 10 0.0072689 73 rfc5, pold4, rfc3, rfc4, cul4a, lig1, pold2, pcna, cdk7, ercc3 p53 signaling

**ohs-020 cell cycle - cosmo bio** - cell cycle progression is precisely controlled by cyclin-dependent

kinases (cdks) and proteins that regulate cdks. these cdks and cdk-modifying proteins, including cyclins, cdk inhibitors,

**supplementary material - the international journal of ...** - supplementary material corresponding to: genetic and epigenetic instability of human bone marrow mesenchymal stem cells expanded in autologous serum

**annotations for all recurrently lost regions - haematologica** - annotations for all recurrently lost regions cytoband 1p36.12 3p14.2 3q13.13 q value 0,01352 0,1513 0,085307 residual q value 0,01352 0,1513 0,085307

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