

Provider information for human full-length cDNA

cDNA sequencing project/ provider	# cDNA	Method	URL	Reference
FLJ/NEDO (Ver3.0)	54,020	Oligo-capping method and selection by one-pass sequences	http://flj.lifesciencedb.jp/	Ota <i>et al.</i> 2004
HUGE/KDRI	2,012	In vitro protein synthesis and selection by one-pass sequences	http://www.kazusa.or.jp/huge/	Kikuro <i>et al.</i> 2004
MGC/NIH (as of 2010/08/02)	27,628	Selecting gene candidates from 5'-EST libraries	http://mgc.nci.nih.gov/	Temple <i>et al.</i> 2009
MIPS/GSF	8,057	Selecting FLcDNA clones from 5'- and 3'-EST libraries	http://mips.gsf.de/	Mewes <i>et al.</i> 2008
/CHGC	738	Selecting FLcDNA clones from EST libraries	http://www.chgc.sh.cn/en/	Zhang <i>et al.</i> 2000
Total	92,455			

FLJ: Full-Length cDNA Japan
HUGE: Human Unidentified Gene-Encoded Large Proteins
MGC: Mammalian Gene Collection
MIPS: Munich Information Center for Protein Sequences

NEDO: New Energy and Industrial Technology Development Organization (Japan)
KDRI: Kazusa DNA Research Institute (Japan)
NIH: National Institutes of Health (USA)
GSF: German Research Center for Environment and Health
CHGC: Chinese National Human Genome Center