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DATASHEET

Dopamine Receptor D3 (DRD3) Antibody

Catalogue No.:abx320594

DRD3 Antibody is a Rabbit Polyclonal against DRD3.

Target:	DRD3
Reactivity:	Human
Host:	Rabbit
Clonality:	Polyclonal
Tested Applications:	ELISA, IHC
Recommended dilutions:	IHC: 1/20 - 1/200. Optimal dilutions/concentrations should be determined by the end user.
Immunogen:	Recombinant human D (3) dopamine receptor protein (213-324AA).
Purification:	Antigen Affinity Chromatography.
Form:	Liquid
Isotype:	IgG
Conjugation:	Unconjugated
Storage:	Aliquot and store at -20 °C. Avoid repeated freeze/thaw cycles.
Swiss Prot:	P35462
GeneID:	1814
Gene Symbol:	DRD3
OMIM:	126451
NCBI Accession:	NP_000787.2, NM_000796.5, NP_001269492.1, NM_001282563.2, NP_001277738.1, NM_001290809.1, NP_387512.3, NM_033663.3, UniGene,gene-oriented,nucleotide,sequence,clustersMore... UniGene#is 121478, Genome,annotation,databasesEnsembl,eukaryotic,genome,annotation,projectMore... EnsemblENST00000638373, ENSP00000371169, ENSG00000151577, ENST00000460779, ENSP00000419402, ENSG00000151577, ENST00000467632, ENSP00000420652, ENSG00000151577, Database,of,genes,from,NCBI,RefSeq,genomesMore... GeneID#1814, KEGG,Kyoto,Encyclopedia,of,Genes,and,GenomesMore... KEGG#hsa:1814, < p>UniProtKB,Keywords,constitute,a,< a,href="http://www.uniprot.org/keywords/"+controlled,vocabulary< /a> with,a,hierarchical,structure,. Keywords,summarise,the,content,of,a,UniProtKB,entry,and,facilitate,the,search,for,proteins,of,interest.< p>< a href="/help/keywords" target="_top">More... < /a>< p>Keywords - ,Coding,sequence,diversity<Alternative,splicing_Polymorphism< p>This,section,provides,links,to,the,UniProt,Reference,Clusters,< a href="http://www.uniprot.org/help/uniref">UniRef< /a>, UniRef,consists,of,clusters,for,UniProtKB,sequences,(including,isoforms),and,selected,UniParc,sequences,,in,order,to,obtain,complete,coverage,of,the,sequence,space,at,resolutions,of,100%,,90%,and,50%,identity.< p>< a href="/help/similar_proteins_section" target="_top">More... < /a>< p>Similar,proteins<links,to,similar,proteins,from,the,UniProt,Reference,Clusters,(UniRef),at,100%,,90%,and,50%,sequence,identity:100%UniRef100,combines,identical,sequences,and,sub-fragments,with,11,or,more,residues,from,any,organism,into,one,UniRef,entry.90%UniRef90,is,built,by,clustering,UniRef100,sequences,that,have,at,least,90%,sequence,identity,to,,and,80%,overlap,with,,the,longest,sequence,(a,k,a,seed,sequence).50%UniRef50,is,built,by,clustering,UniRef90,seed,sequences,that,have,at,least,50%,sequence,identity,to,,and,80%,overlap,with,,the,longest,sequence,in,the,cluster.< p>This,section,is,used,to,point,to,information,related,to,entries,and,found,in,data,collections,other,than,UniProtKB.< p>< a,href="/help/cross_references_section" target="_top">More... < /a>< p>Cross-references< p>This,subsection,of,the,< a,href="http://www.uniprot.org/manual/cross_references_section">Cross-references< /a> section,provides,links,to,various,web,resources,that,are,relevant,for,a,specific,protein.< p>< a,href="/help/web_resource" target="_top">More... < /a>< p>Web,resources<IMPD,The,Singapore,human,mutation,and,polymorphism,databaseSequence,databasesSelect,the,link,destinations:EMBL,nucleotide,sequence,databasesMore... EMBL/GenBank,nucleotide,sequence,databasesMore... GenBank/EMA,Data,Bank,of,Japan, a,nucleotide,sequence,databasesMore... DDBJ/Link,Updated0,30469,mRNA, Translation: AAA03543.1, , Frameshift,AHG00361,Genomic,DNA, Translation: AAA64369.1, ,Sequence,problems,U32499,mRNA, Translation: AAA73929.1 AC092896,Genomic,DNA, No,translation,available,CH471052,Genomic,DNA, Translation: EAW79610.1 BC058510,mRNA, Translation: AAH58510.1 BC128123,mRNA, Translation: AAI28124.1 The,Consensus,CDS,(CCDS),projectMore... CCDS|CCDS2878.1, [P35462-1]CCDS53829.1, [P35462-3]
Buffer:	PBS, pH 7.3, containing 0.02% sodium azide and 50% glycerol.
Note:	This product is for research use only.