

Advanced Search Query Builder

Full Text

Structure Attributes

Source Organism Taxonomy Name (Full Lineage)

x

is

Leishmania Viannia braziliensis

+ NOT

Count

x

OR

Source Organism Taxonomy Name (Full Lineage)

x

is

Leishmania braziliensis

+ NOT

Count

x

OR

Source Organism Taxonomy Name (Full Lineage)

x

is

Leishmania braziliensis braziliensis

+ NOT

Count

x

AND / OR

Add Attribute

Add Subquery

Remove Subquery

Add Subquery

Chemical Attributes

Sequence Similarity

Sequence Motif

Structure Similarity

Structure Motif

Chemical Similarity

Return

Structures

grouped by

No Grouping

Include Computed Structure Models (CSM)

Count

Clear

Search

Nº	ID no PDB	Enzima	Método - resolução	Ligantes	Observações
1	4KCE	Peroxiirredoxi na mitocondrial	DIFRAÇÃO DE RAIOS X - 2,79 Å	-	
a	4KB3	Peroxidoxin mitochondrial	X-RAY DIFFRACTION - 2.93 Å	-	
2	5TSQ	Inosine-uridine-preferring nucleoside hydrolase homolog (N-D-ribosylp urine ribohydrolase)	X-RAY DIFFRACTION - 1.53 Å	beta-D-ribofuranse (BDR)	COM REDOCKING
3	4WZH	Dihydroorotate dehydrogenase	X-RAY DIFFRACTION - 2.12 Å	<ul style="list-style-type: none"><li>FLAVIN MONONUCLEOTIDE</li><li>4-(2-HYDROXYETHYL)-1-PIPERAZINE ETHANESULFONIC ACID</li></ul>	

<b>4</b>	3CH7	6-phosphogluconolactonase	X-RAY DIFFRACTION - 2.29 Å	-	
a	3CSS	6-phosphogluconolactonase	X-RAY DIFFRACTION - 1.70 Å	PHOSPHATE ION	
<b>5</b>	6EDI	glucokinase	X-RAY DIFFRACTION - 1.85 Å	-	
<b>6</b>	1X6O	eukaryotic initiation factor 5a	X-RAY DIFFRACTION - 1.60 Å	DI(HYDROXYETHYL) ETHER / Dietilenoglicol	
<b>7</b>	4KPC	Nucleoside diphosphate kinase b	X-RAY DIFFRACTION - 2.70 Å	PHOSPHATE ION	
<b>8</b>	1S4H	60S acidic ribosomal protein P2 ()	SOLUTION NMR	-	