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Accession	_		· ·		Source spider monkey							
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0 1 1 5		7.0. 4000										
Created Da					Formal	F	Ateles sp.					
Seq-rev Da												
Txt-rev Dat	e 0	3-Mar-2000										
Reference												
Refinfo				Α	ccinfo							
Refid		0004		4	Label		MAR					
Authors					Acces							
Au	thor	Entry			Mol-t							
					Seq-s		1-104					
Citation	า				<b>Xrefs</b>							
Ту		other			Statu							
Cor	ntent		by Shelnutt, J.A., Rousseau, D.		Exp-s							
			argoliash, E., Biochemistry 20,	C	ontents	En	try					
		6485-6497										
Volume	9			Ν	lote							
Year	198	31										
Pages												
Title												
Xrefs				1								
Month				1								
Genetics				Kον	words							
Classification				КСУ	Keywor	·4 E	ntry					
Superfamily	( Ent	·m/			Reywor		acetylated amino end					
Superramily												
		ochrome c					hromoprotein					
	cyt	ochrome c homology					electron transfer					
							neme					
						_	ron					
							netalloprotein					
							nitochondrion					
						C	oxidative phosphorylation					
						r	espiratory chain					
Feature La	abel	Feature-type	Description					Seq-	Status			
								spec				
C	YC	domain	cytochrome c homology					4-98				
		modified-site	acetylated amino end (Gly)					1	predicted			
		binding-site	heme (Cys) (covalent)					14,17	predicted			
		binding-site	heme iron (His, Met) (axial liga	ınds	s)				predicted			
Summary	-				uence	0	DVEKGKRIEIMKCSOCHT\		CHKTGPNLHGLFGRKTGQASGFT			
Length 104	1			4	1401100		TEANKNKGIIWG	LICON	O. NEHOEF GIACTORASOFT			
	nplet	Δ					DTLMEYLENPKKYIPGTKM	IFVGIR	KKEFRADI IAYI KKATNE			
Status	iipiet						CTELLIN WITH OTHER	• • • • •				
Status			l									
ld C	CMS		l l	Prot	tein							
Header						cyto	ochrome c [validated]					
Uid	C	CMS		Ora	anism							
Accession	_	ntry	ļ.	- 3	Source	r	mouse					
71000001011		23057					nouse mouse					
		04604			Formal		Mus musculus					
		00009	Į.		TUTTIAL	Į.	vius musculus					
Created D.												
Created Da												
Seq-rev Da												
	O 129	3-Jul-2000										

Reference							
• Refinfo	А	ccinfo					
Refid A23057		Labe		LIM			
Authors		Acce			057		
Author Entry		Mol-1					
Limbach, K.J. Wu, R.		Seq-		1-10	J5		
Citation			Kref	Dh	Uid		
		,	riei				
Type Content					X01756		
Volume 13					g50618 CAA25899.1		
Year 1985					q50619		
Pages 617-630		Statu		FID	g30017		
Title Characterization of a mouse somatic cytochrome c ger	ane and			co stra	in BALB/c		
three cytochrome c pseudogenes.		ontents			III DALD/C		
Xrefs		Ontents	LIII	ıy			
Xref	N	ote					
Db MUID							
Uid 85215501							
Month							
• Refinfo	A	ccinfo	_				
Refid A04604		Labe		CAR			
Authors		Acce			604		
Author Entry		Mol-1					
Carlson, S.S.		Seq-		2-10	05		
Mross, G.A.		Xrefs					
Wilson, A.C.		Statu					
Mead, R.T.	0				in BALB/c		
Wolin, L.D.		ontents	EIII	ry			
Bowers, S.F. Foley, N.T.	N	ote					
Muijsers, A.O.	IN.	ote					
Margoliash, E.							
Citation							
Туре							
Content							
Volume 16							
Year 1977							
Pages 1437-1442							
Title Primary structure of mouse, rat, and guinea pig cytoch	chrome c.						
Xrefs							
Xref							
Db MUID							
Uid 77134768							
Month							
Genetics	Key	words					
Introns 57/1		Keywo					
Classification					ed amino end		
Superfamily Entry				hromop			
cytochrome c					transfer		
cytochrome c homology				eme			
				on			
				netallop			
				nitocho			
					phosphorylation		
			re	espirato	ory chain		
Feature Label Feature-type Description						Seq-	Status
MAT product						spec	our orine antal
MAT product cytochrome c	en /					2-105	experimental
CYC domain cytochrome c homolog		tura 6	·ma\			5-99	our orine antal
modified-site acetylated amino end		iture for	in)			2	experimental
binding-site heme (Cys) (covalent)		-)					experimental
binding-site heme iron (His, Met) (				ICDVE.	CKKIENOKOVOO		predicted
Summary Longth 105	Seq	uence			KGKKIFVQKCAQCF NKNKGITW	TIVEKG	GKHKTGPNLHGLFGRKTGQAAG
Length 105						MIFAG	IKKKGERADLIAYLKKATNE
Type complete Status			10	LUILIV	LILLINI KKIIFUII	WIII AO	MINOCIALLINATIVE
Status							
		tein					

leader				Name cytochrome c [validated]					
Uid	CCRT	Orga	anism						
Accession	Entry		Source	rat					
	A04605		Common	Norway rat					
	C28160		Formal	Rattus norvegicus					
	A00009								
Created Date	31-Dec-1990								
Seq-rev Date	30-Sep-1991								
Txt-rev Date	28-Jul-2000								

Deference	
Reference	A!
• Refinfo	Accinfo
Refid A04605	Label SCA
Authors	Accession A04605
Author Entry	Mol-type DNA
Scarpulla, R.C.	Seq-spec 1-105
Agne, K.M.	Xrefs
Wu, R.	
	Xref Db Uid
Citation	GB K00750
Туре	GB M28216
Content	NID g550511
Volume 256	PIDN AAA21711.1
Year 1981	PID g203699
Pages 6480-6486	Status
Title Isolation and structure of a rat cytochrome c gene.	Exp-source
Xrefs	Contents Entry
Xref	
Db MUID	Note
Uid 81215609	
Month	
Refinfo	Accinfo
Refid A28160	Label VIR
Authors	Accession C28160
Author Entry	Mol-type mRNA
Virbasius, J.V.	Seq-spec 1-105
Scarpulla, R.C.	Xrefs
Citation	Xref Db Uid
Type	GB M20622
Content	NID g203722
Volume 263	PIDN AAA41014.1
Year 1988	PID g203723
Pages 6791-6796	Status
Title Structure and expression of rodent genes encoding the testis-	Exp-source Exp-source
specific cytochrome c. Differences in gene structure and	Contents Entry
evolution between somatic and testicular variants.	
Xrefs	Note
Xref	
Db MUID	
Uid 88198250	
Month	
Refinfo	Accinfo
Refid A04604	Contents Entry
	COINCINS EITH Y
Authors	
Author Entry	Note peptide mapping, compositional analysis, and partial sequencing
Carlson, S.S.	indicate that rat cytochrome c is identical with that of mouse
Mross, G.A.	
Wilson, A.C.	
Mead, R.T.	
Wolin, L.D.	
	-
Bowers, S.F.	
Foley, N.T.	
Muijsers, A.O.	
Margoliash, E.	
Citation	
Туре	
Content	
Volume 16	
Year 1977	
Pages 1437-1442	
Title Primary structure of mouse, rat, and guinea pig cytochrome c.	
Xrefs	
Xref	
Db MUID	
Uid 77134768	
Month	

Gen	etics				Key	words						
	Introns !	57/1				Keyword	Entry					
Clas	sification						blocked amino end					
	Superfan	nily E	Intry				chromoprotein					
		C	cytochrome c				electron transfer					
		C	cytochrome c homology				heme					
			,				iron					
							metalloprotein					
							mitochondrion					
							oxidative phosphorylation					
							respiratory chain					
Fea	ture	Labe	el Feature-type	Description				Seq-	Status			
			3,1	, , , , , ,				spec				
		MAT	product	cytochrome c				2-105	experimental			
		CYC		cytochrome c homology				5-99	exportitiontal			
		0.0	modified-site	blocked amino end (Gly) (in m	atur	re form)	(probably acetylated)	2	experimental			
			binding-site	heme (Cys) (covalent)	iatai	ic ioiiii)	(probably acceptated)	15 18	experimental			
			binding-site	heme iron (His, Met) (axial lig	ando	-1			predicted			
Cur	nmary	1	biliding-site	Therne from (tris, wet) (axial lig		uence	MCDVEKCKKIENOKCYOCH		GKHKTGPNLHGLFGRKTGQAAG			
	Length 1	INE			seq	luerice	FSYTDANKNKGITW	IIVEKG	GKHK I GPINLHGLEGKK I GQAAG			
			loto				GEDTLMEYLENPKKYIPGTK	MIFAG	IKKKGEBADI IAYI KKATNE			
		compl	icic				OLD I LIVE I LEIVI KKI IF OT	AU	OEIVIDEIVITEIVATNE			
	Status											
Id		CCR	В		Prot	tein						
Hea	der						ytochrome c [validated]					
	Uid		CCRB		Ora	anism	, s [vandatou]					
	Accession	n	Entry		org	Source	rabbit					
	Accession		Entry									
	Crostod	Data	13-Jul-1981				Orvetolagus cupiculus					
			13-Jul-1981			Formai	Oryctolagus cuniculus					
		Jate	28-Jul-2000									
	erence											
•	Refinfo				A	ccinfo						
	Refic		400009			Label	NEE					
	Auth					Acces						
		Autho	or Entry			Mol-ty	rpe protein					
			Needleman, S.B.			Seq-s	pec 1-104					
			Margoliash, E.			Xrefs						
	Citat	ion				Status	3					
		Туре				Exp-s	ource					
		Conte			C	ontents						
		me 2										
	Year	_	1966		N	ote						
	Page	_	353-863									
	Title		Rabbit heart cytochrome c.									
	Xrefs		Cabbit Heart Cytochrome C.									
		Xref			-							
	-		DE MILID									
			Db MUID									
			Jid 66093127									
^	Mon	เท			14							
	etics				Key	words	de					
Clas	sification					Keyword						
	Superfan						acetylated amino end					
			cytochrome c				chromoprotein					
		C	cytochrome c homology				electron transfer					
							heme					
							iron					
							metalloprotein					
							mitochondrion					
							oxidative phosphorylation					
							respiratory chain					
Fea	ture	Labe	el Feature-type	Description				Seq-	Status			
								spec				
		CYC	domain	cytochrome c homology				4-98				
		510	modified-site	acetylated amino end (Gly)				1	experimental			
			binding-site	heme (Cys) (covalent)				14 17	experimental			
			binding-site	heme iron (His, Met) (axial lig	ando	(3)			predicted			
Sur	nmary	-	philang-arc	mente iron (ms, wet) (axial lig		uence	CDAEKCKKIEAOKCVOCHI		KHKTGPNLHGLFGRKTGQAVGF			
Juli	Length 1	104			seq	ucrice	SYTDANKNKGITWG	VERGG	KINTO NETIGEI GKKTGQAVGF			
			lata				EDTLMEYLENPKKYIPGTKN	MEAGIR	KKDERADI IAVI KKATNE			
		compl	lete				CO TEMETERINI KKI II OTKI		DEIVIDEIVITEINATIVE			
	Status											

14	CCGW		Protein						
Id Header	CCGVV		Name cytochrome c [validated]						
Uid	CCGW		Organism	Cytoci	irome c [validated]				
Accession		i i	Source	au	anaco				
Accession	A04608		Commo						
	A00009		Formal		ma guanicoe				
Created F	Date 31-Dec-1990		Torritar	Lai	na guarneoc				
	Date 31-Dec-1990								
	ate 28-Jul-2000								
Reference	atc  20-3di-2000								
Refinfo			Accinfo						
Refid	A04608		Label	1	NIE				
Autho			Acces		A04608				
	author Entry		Mol-t		protein				
Í	Niece, R.L.		Seq-s		1-104				
	Margoliash, E.		Xrefs						
	Fitch, W.M.		Statu						
Citati				source					
	уре		Contents						
	Content								
	ne 16		Note						
Year	1977								
Page	68-72								
Title	Complete amino acid sequer	nce of guanaco (Lama guanicoe)							
	cytochrome c.								
Xrefs									
>	(ref								
	Db MUID								
	Uid 77087753								
Mont	h								
Genetics			Keywords						
Classification			Keywor						
Superfam					etylated amino end				
	cytochrome c				omoprotein				
	cytochrome c homology			_	ctron transfer				
				her					
				iror					
					talloprotein				
					ochondrion				
					dative phosphorylation				
				res	piratory chain	_	Tax :		
Feature	Label Feature-type	Description				Seq- spec	Status		
	CYC domain	cytochrome c homology				4-98			
	modified-site	acetylated amino end (Gly)				1	experimental		
	binding-site	heme (Cys) (covalent)				14,17	experimental		
	binding-site	heme iron (His, Met) (axial liga	inds)				predicted		
Summary			Sequence	GD'	VEKGKKIFVQKCAQCHT\		KHKTGPNLHGLFGRKTGQAVGF		
Length 1	04		'		TDANKNKGITWG				
	omplete			EET	TLMEYLENPKKYIPGTKM	IFAGIK	KKGERADLIAYLKKATNE		
Status									

Id	CCCM			Protein						
Header	CCCIVI			Name cytochrome c						
Uid	C	CCM		Organism	yıoc	LIII OI II E C				
Accessio	_	ntry		Source	۸۲	rabian camel				
Accessio	_	04607			_	rabian camel				
		00009		Formal		amelus dromedarius				
Croated		1-Dec-1990		FUIIIdi	Co	arrielus urorrieuarius				
		1-Dec-1990								
		3-Mar-2000								
Reference	Jale U	3-IVIdI -2000								
Refinfo				Accinfo						
Refi	4 1	4607		Label		SOK				
Auth		4007		Acces	cion					
	Author	Entry		Mol-ty		protein				
	Author	Sokolovsky, M.		Seq-s						
		Moldovan, M.		Xrefs	pec	1-104				
Citat	ion	INOIGOVAII, IVI.		Statu	2					
	Туре			Exp-s		20				
	Conten	+		Contents						
	me 11			Contents	LIIII	y				
Year				Note						
Page	_	5-149								
Title			nrome c from the camel, Camelus							
Title		omedarius.	nome e nom the camer, cameras							
Xrefs										
	Xref									
-		MUID								
		72096652								
Mon										
Genetics				Keywords						
Classification				Keywor	d Er	ntry				
Superfan	nily En	try		,		cetylated amino end				
·		ochrome c				nromoprotein				
		ochrome c homology				ectron transfer				
					he	eme				
					irc	on				
					m	etalloprotein				
					mi	itochondrion				
					ОХ	xidative phosphorylation				
					re	spiratory chain				
Feature	Label	Feature-type	Description				Seq-	Status		
							spec			
	CYC	domain	cytochrome c homology				4-98			
		modified-site	acetylated amino end (Gly)				1	experimental		
		binding-site	heme (Cys) (covalent)					predicted		
		binding-site	heme iron (His, Met) (axial liga					predicted		
Summary				Sequence						
	Length 104				SYTDANKNKGITWG					
Type o					EE	ETLMEYLENPKKYIPGTKM	IFAGIK	KKGERADLIAYLKKATNE		
Status										

Id	CCWI	HC		Pro	tein				
Header					Name o	ytoch	nrome c		
Uid	(	CCWHC		Org	anism				
Accessio	n E	ntry			Source	Cal	lifornia gray whale		
	F	A04606			Commo	n Cal	lifornia gray whale		
	F	A00009			Formal	Esc	chrichtius robustus, Esc	hrichtiu	ıs gibbosus
Created	Date 3	31-Dec-1990							
Seq-rev	Date 3	31-Dec-1990							
Txt-rev I	Date 0	03-Mar-2000							
Reference									
<ul> <li>Refinfo</li> </ul>				Α	ccinfo				
Refi	d A	04606			Label		GOL		
Auth	nors				Acces	sion	A04606		
	Author	Entry			Mol-ty	/ре	protein		
		Goldstone, A.			Seq-s	рес	1-104		
		Smith, E.L.			Xrefs				
Cita	tion				Status	S			
	Туре				Exp-s				
	Conter			С	ontents	Entry			
Volu	ıme 24								
Year	r 19	966		Ν	ote				
Page		180-4486							
Title		mino acid sequence of whale	e heart cytochrome c.						
Xref									
	Xref								
	Db	MUID							
		67041932							
Mon	ith			_					
Genetics				Key	words				
Classification					Keyword				
Superfar							cked amino end		
		tochrome c		1			omoprotein		
	су	tochrome c homology					ctron transfer		
						her			
						iror			
							talloprotein		
							ochondrion		
							dative phosphorylation		
	1		I			res	piratory chain	-	In
Feature		Feature-type	Description					Seq- spec	Status
	CYC	domain	cytochrome c homology					4-98	
		modified-site	blocked amino end (Gly) (prol	bably	y acetyla	ted)		1	experimental
	binding-site heme (Cys) (covalent)								predicted
	binding-site heme iron (His, Met) (axial lig								predicted
Summary			Seq	uence		GDVEKGKKIFVQKCAQCHTVEKGGKHKTGPNLHGLFGRKTGQAVGF			
Length 104					SYTDANKNKGITWG EETLMEYLENPKKYIPGTKMIFAGIKKKGERADLIAYLKKATNE				
	comple	te				EE	ILIVIEYLENPKKYIPGIKM	IIFAGIK	KKGERADLIAYLKKATNE
Status									

Id CCPG	Protein					
Header	Name cytochrome c [validated]					
Uid CCPG	Organism					
Accession Entry	Source					
	Commo	n dor	mestic pig			
Created Date 17-Mar-1987	Formal	Sus	s scrofa domestica			
Seg-rev Date 17-Mar-1987		•				
Txt-rev Date 28-Jul-2000						
Reference						
Refinfo	Accinfo					
Refid A90743	Label		STE			
Authors	Acces	ssion	A00007			
Author Entry	Mol-t	Mol-type protein				
Stewart, J.W.	Seq-s	рес	1-104			
Margoliash, E.	Xrefs					
Citation	Statu					
Туре		ource				
Content	Contents	Entry				
Volume 43						
Year 1965	Note					
Pages 1187-1206						
Title The primary structure of the cytochrome c from various organs	5					
of the hog.						
Xrefs	_					
Xref						
Db MUID						
Uid 66072936						
Month Genetics	Keywords					
Classification	Keywor	d Ent	m.			
Superfamily Entry	Reywor		tylated amino end			
cytochrome c	chromoprotein					
cytochrome c homology			electron transfer			
cytochrome c nomology		hen				
		iron				
			talloprotein			
			ochondrion			
			dative phosphorylation			
			piratory chain			
Feature Label Feature-type Description		163	piratory criairi	Seq-	Status	
reature Eaber reature-type Description				spec	Status	
CYC domain cytochrome c homology				4-98		
modified-site acetylated amino end (Gly)				1	experimental	
binding-site heme (Cys) (covalent)				14.17	experimental	
	heme iron (His, Met) (axial ligands)				predicted	
	Sequence	GDV	VEKGKKIFVQKCAQCHT		KHKTGPNLHGLFGRKTGQAPGF	
Length 104	SY			SYTDANKNKGITWG		
Type complete		EET	EETLMEYLENPKKYIPGTKMIFAGIKKKGEREDLIAYLKKATNE			
Status						

Id CCBO	Protein				
Header	Name cytochrome c				
Uid CCBO	Organism				
Accession Entry	Source bovine				
A92022	Common cattle				
A00007	Formal Bos primigenius taurus				
Created Date 31-Mar-1992	3				
Seg-rev Date 31-Mar-1992					
Txt-rev Date 03-Mar-2000					
Reference					
Refinfo	Accinfo				
Refid A92022	Label NAK				
Authors	Accession A92022				
Author Entry	Mol-type protein				
Nakashima, T.	Seq-spec 1-104				
Higa, H.	Xrefs				
Matsubara, H.	Status				
Benson, A.	Exp-source				
Yasunobu, K.T.	Contents Entry				
Citation					
Туре	Note				
Content					
Volume 241					
<u>Year</u> 1966					
Pages 1166-1177					
Title The amino acid sequence of bovine heart cytochrome c.					
Xrefs					
Xref					
Db MUID					
Uid 66132521					
Month					
• Refinfo	Accinfo				
Refid A61297	Contents Entry				
Authors					
Author Entry	Note				
Tsunasawa, S.					
Narita, K.					
Citation					
Туре					
Content					
Volume 92					
Year 1982					
Pages 607-613					
Title					
Xrefs					
Month					
Genetics	Keywords				
Classification					
Feature Label Feature-type Description	Seq- Status				
	spec				
Summary	Sequence				