

OriGene Technologies, Inc.

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Product datasheet for PH306675

AKR7A3 (NM_012067) Human Mass Spec Standard

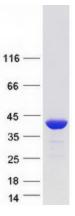
Product data:

Description:AKR7A 3M S Standard C13 and N15-labeled recombinant protein (NP_036199)Species:HumanSpecies:HumanExpression DDNA ClossRC20675Predicted MW:3.2 k DaProtein Sequence: Red-Cloning site Green=Tags(s)Sc206675 protein sequence Red-Cloning site Green=Tags(s)With TKATPLFGNSLKPDSLRPQLETSLKRLQPRVDLPYLHPPDHSTPLACHQLEGCKFMEL GLSNXAKPATVLGAMEMGRRNDAPTSAAVTRAFLEGCHTEIDTAFVYSEQSETILGGLGLRGGSDC KVKDTKATPLFGNSLKPDSLRPQLETSLKRLQPRVDLPYLHPPDHSTPLACHQLEGCKFMEL GLSNXAKPATVLGAMEMGRRNDAPTSAAVTRAFLEGCHTEIDTAFVYSEQSETILGGLGLRGGSDC KVKDTKATPLFGNSLKPDSLRPQLETSLKRLQPRVDLPYLHPPDHSTPLACHQLEGCKFMEL GLSNXAKPATVLGAMEMGRRNDAPTSAAVTRAFLEGCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATVLGAMEMGRRNDAPTSAAVTRAFLEGCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATVLGAMEMGRRNDAPTSAAVTRAFLEGCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATVLGAMEMGRRNDAPTSAAVTRAFLEGCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATVLGAMEMGRRNDAPTSAAVTRAFLEGCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATVLGAMEMGRRNDAPTSAAVTRAFLEGCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATVLGAMEMGRRNDAPTSAAVTRAFLEGCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATVLGAMEMGRRNDAPTSAAVTRAFLEGCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATVLGAMEMGRRNDAPTSAAVTRAFLEGCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATTPLAGGLTGNSCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATTPLAGGLTGNSCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATTPLAGGLTGNSCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATTPLAGGLTGNSCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATTPLAGGLTGNSCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATTPLAGGLTGNSCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATTPLAGGLTGNSCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATTPLAGGLTGNSCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATTPLAGGLTGNSCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSNXAKPATTPLAGGLTGNSCHTEPPCLARFLEGTAFVARGUQGG GLSNXAKPATTPLAGGLTGNSCHTEPPCLARFLEGTAFVARGUQGGKFMEL GLSN	Product Type:	Mass Spec Standards
Fxpression Host:HEK293Expression CDNA CloonRC206675Predicted MW:3.7. kDaPredicted MW:SRC206675 protein sequence Red=Cloning site Green=Tags(s)Wixtor KARPAPTVLGAMEMGRINDAPTSAAVTRAFLEGHTEIDTAFVSEGQSETLIGGLGRLGGSMC Red=Cloning site Green=Tags(s)MSRQLSRARPATVLGAMEMGRINDAPTSAAVTRAFLEGHTEIDTAFVSEGQSETLIGGLGRLGGSMC RCUDSCRPVGRFFGNTWAEMWRDAPTSAAVTRAFLEGHTEIDTAFVSEGQSETLIGGLGRLGGSMC RCUDSCRPVGRFFGNTWAEMWRDAPTSAAVTRAFLEGHTEIDTAFVSEGQSETLIGGLGRLGGSMC RCUDSCRPVGRFFGNTWAEMWRDAPTSAAVTRAFLEGHTEIDTAFVSEGQSETLIGGLGRLGGSMC RCUDSCRPVGRFFGNTWAEMWRDAPTSAAVTRAFLEGHTEIDTAFVSEGQSETLIGGLGRLGGSMC RCUDSCRPVGRFFGNTWAEMWRDAPTSAAVTRAFLEGHTEIDTAFVSEGQSETLIGGLGRLGGSMC RCUDSCRPVGRFFGNTWAEMWRDAPTSAAVTRAFLEGHTEIDTAFVSEGQSETLIGGLGRLGGSMC RCUDSVAAWEVAELICTLCSSNGULTITVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHLQGGSMC RCUDSVAAWEVAELICTLCSSNGULTITVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHLQGGSMC RCUDSVAAWEVAELICTLOSSNGULTIVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHLQGGSMC SCUDSVAAWEVAELICTLCSSNGULTIVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHLQGGSMC SCUDSVAAWEVAELICTLOSSNGULTIVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHLQGGSMC SCUDSVAAWEVAELICSSNGULTIVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHLQGGSMC SCUDSVAAWEVAELICSSNGULTIVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHLGGSA SCUDSVAAWEVAELICSSNGULTIVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHLGGSAGA SCUDSVAAWEVAELICSSNGULTIVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHT SCUDSVAAWEVAELICSSNGULTIVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHT SCUDSVAAWEVAELICSSNGULTIVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHT SCUDSVAAWEVAELICSSNGULTIVVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHTTag:SNBQ SAGERTMINED SDS-PAGE AND COOMASSIE BILD STAIL SCUDSCRVC SCUDSVAAWEVAELICSSNGULTIVGGNYMAITRQVETELFPCLRHFGLRPVAFNPLAGHTStage:SNBM TISHCLION SDGULTIVGGNYMAITRQVETELFPCLRHFGLRPVAFNFA SCUDSCRVC SCUDSVAAWEVAELICSSNGULTIVG	Description:	AKR7A3 MS Standard C13 and N15-labeled recombinant protein (NP_036199)
Spression cDNA GloomR2206675Predicted MW:37.2 kDaProtein Sequence:Rc206675 protein sequenceRed=Cloning site Green=Tags(s)MSRQLSRARPATVLGAMEMGRRMDAPTSAAVTRAFLERGHTEIDTAFVSEGQSETILGGLGLRLGGSDC KVKIDTKATPLFGNSLKPQLETSLKRLQCPRVDLFVLHMPDATPVEETLRAAQLHQEGKPNEL GLSNVAAWEVAEICTLCKSNGWILPTVQGMYNAITRQVEETLPPACHGHQEGKPNEL SUSDGKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGRRMDAPTSAAVTRAFLERGHTEIDTAFVSEGQSETILGGLGLRLGGSDC KVKIDTKATPLFGNSLKPQLETSLKRLQCPRVDLFVLHMPDATPVEETLRAAQLHQEGKPNEL SUSDGKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGRRMDAPTSAAVTRAFLERGHTEIDTAFVSEGQSETILGGLGLRGSDC KVKIDTKATPLFGNSLKPQLETSLKRLQCPRVDLFVLHMPDATPVEETLRAAQLHQEGKPNEL SUSDGKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGATGGLLTGKKK VEEDKOKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGATGGLLTGKKK VEEDKOKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGATGGLLTGKKK VEEDKOKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGATGGLLTGKKK VEEDKOKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGATGGLLTGKKK VEEDKOKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGATGGLLTGKKK VEEDKOKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGATGGLLTGKKK VEEDKOKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGATGGLLTGKKK VEEDKOKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGATGGLLTGKKK VEEDKOKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGATGGLLTGKKK VEEDKOKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGATGGLLTGKK VEEDKOKQPVQRFFGNTWAEHTFEGLAPVDAFVGAMEMGATGGLGGLGGSDC NGDAGTGLGASGLGASGSDTSATLRWMYHHSQLGA MAGZGLAGGLGASGGDC NGSGLGAGGDGLAGAGGLGAGGDGLGAGGDGLGAGGDGLGAGGDGLGAGGDGLGAGGDGLGAGGDGLGAGGDGLGAGGGDGLGAGGDGLGGGDGLGAGGDGLGAGGDGLGGGGDGGGGGGGG	Species:	Human
or AA Sequence:Predicted MW:37.2 kDaProtein Sequence:>Rc206675 protein sequenceRed=Cloning site Green=Tags(s)MSRQLSRARPATVLGAMEMGRRMDAPTSAAVTRAFLERGHTEIDTAFVYSEGQSETILGGLGLRLGGSDCRVKIDTKAIPLFGNSLKPOLETSLKRLQCPRVDEFVLHPPHRPHSTPVEETLRACHQLHQEGKFMEL GLSNYAAWEVAEICTLCKSNGWILPTVYQGWYNAITRQVETELPCLRHGLFYAFNPLAGGLLTGKYKRVKIDTKAIPLFGNSLKPOLETSLKRLQCPRVDEFVLHPPHRPHSTPVEETLACHQLHQEGKFMEL GLSNYAAWEVAEICTLCKSNGWILPTVYQGWYNAITRQVETELPCLRHGLFYAFNPLAGGLLTGKYKRVKIDTKAIPLFGNSLKPOLETSLKRLQCPRVDEFVLHPPHRPHSTPVEETLACHQLHQEGKFMEL GLSNYAAWEVAEICTLCKSNGWILPTVQGWYNAITRQVETELPCLRHGLFYAFNPLAGGLLTGKYKRVKIDTKAIPLFGNSLKPOLEFSLKRLQCPRVDFVLHPPHRPHSTPVEETLACHQLHQEGKFMEL GLSNYAAWEVAEICTLCKSNGWILPTVQGWYNAITRQVETELPCLRHGLFYAFNPLAGGLLTGKYKTATRPLEQKLISEEDLAANDILDYKDDDDKVTATRPLEQKLISEEDLAANDILDYKDDDDKVPurity:>0.05 µg/µL as determined by microplate BCA methodGoncentration:>0.05 µg/µL as determined by microplate BCA methodBuffer:>0.05 µg/µL as determined by microplate BCA methodBuffer:>1.80eled with [U-13C6, 15N4]-L-Arginine and [U-13C6, 15N2]-L-LysineBuffer:>2.5 mM Tris-HCl, 100 mM glycine, pH 7.3Storage:>1.60eled for 3 months from receipt of products under proper storage and handling conditionsRefSeqNP 036199RefSeq ORF:>93Spnonyms:>4.FAR2	Expression Host:	HEK293
Protein Sequence:Rc206675 protein sequence Red=Cloning site Green=Tags(s)WSRQLSRARPATVLGAMEMGRRMDAPTSAAVTRAFLERGHTEIDTAFVYSEGQSETILGGLGLRLGGSDC RVK.DTKAIPLFGNSLKPDSLRFQLETSLKRLQCPRVDLFYLHMPDHSTPVEETLRACHQLHQEGKFMEL GLSNYAAWEVAEICTLCKSNGWILPTVYQGMYNAITRQVETELPPCLRHFGLRFYAFNPLAGGLLTGKYK YEDKOGKQPVGRFFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUQPVGRFFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA MCDGXUQPVGRFFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUQPVGRFFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFGNTWAEMYRNYKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFGNTWAEMYRNRYKEHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFGNTWAEMYRNYKEHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HODGXUGPVGRFGNTWAEMYRNYKEHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HOTGXUGPVGRFGNTWAEMYRNRYKEHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HONGXUGPVGRFGNTWAEMYRNRYKEHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HONGXUGPVGRFGNTWAEMYRNRYKEHFEGIAPVEKALQAYGASGNTSATLRWMYHHSQLQGA HONGXUGPVGRFGNTWAEMYRNRYKEHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HONGXUGPVGRFGNTGNTWAEMYRNRYKEHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HONGXUGPVGRFGNTGNTYKEHPEGIAPVEKALQUAAYGANGANGANGTSATLRWMYHHSQLQGA HONGXUGPVGRFGNTGNTATUNGHTGNAGIAPVEKALQUAAYGANGANGTGNTYKEHFEGIAPVEKALQUAAYGANGAAYGNGANGTGNTSATLRWMYHSQLQGA H	•	RC206675
Red=Cloning site Green=Tags(s)MSRQL SRARPATVLGAMEMGRRMDAPTSAAVTRAFLERGHTEIDTAFVYSEGQSETILGGLGLRLGGSDC RVKIDTKAIPLFGNSLKPOSLRFQLETSLKRLQCPRVDLFYLHMPDHSTPVEETLRACHQLHQEGKFMEL GLSNYAAWEVAEICTLCKSNGWILPTVYQGMYNAITRQVETELFPCLRHFGLRFYAFNPLAGGLLTGKYK YEDKDGCQPVGRFGNTWAEMYRNRVWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HGDAVILGMSSLEQLEQNLAAAEEGPLEPAVVDAFNQAWHLVAHECPNYFRTag:C-Myc/DDKPurity:> 80% as determined by SDS-PAGE and Coomassie blue stainingConcentration:> 0.05 µg/µL as determined by microplate BCA methodLabeling Method:Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-LysineBuffer:25 mM Tris-HCl, 100 mM glycine, pH 7.3Storage:Store at -80°C. Avoid repeated freeze-thaw cycles.Stability:Stable for 3 months from receipt of products under proper storage and handling conditions.RefSeq:NP 036199RefSeq ORF:993Synonyms:AFAR2	Predicted MW:	37.2 kDa
RYKIDTKAIPLFGNSLKPDSLRFQLETSLKRLQCPRVDLFYLHMPDHSTPVETLRACHQLHQEGKFMEL GLSNYAAWEVAEICTLCKSNGWILPTVYQGMYNAITRQVETELFPCLRHFGLRFYARNPLAGGLLTGKYK SYEDKDGKQPVGRFFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA HGDAVILGMSSLEQLQNLAAAEEGPLEPAVVDAFNQAWHLVAHECPNYFRTarrepleckliseEDLAANDILDYKDDDDKVTag:C-Myc/DDKPurity:S0% as determined by SDS-PAGE and Coomassie blue stainingConcentration:>0.05 µg/µL as determined by microplate BCA methodLabeling Method:Labeled with [U-13C6, 15N4]-L-Arginine and [U-13C6, 15N2]-L-LysineBuffer:0.5 mM Tris-HCI, 100 mM glycine, pH 7.3Storage:Store at -80°C. Avoid repeated freeze-thaw cycles.Stability:Stable for 3 months from receipt of products under proper storage and handling conditions.RefSeq:NP 036199RefSeq ORF:933Synonyms:AFAR2	Protein Sequence:	
Tag:C-Myc/DDKPurity:> 80% as determined by SDS-PAGE and Coomassie blue stainingConcentration:> 0.05 µg/µ as determined by microplate BCA methodLabeling Method:Labeled with [U-13C6, 15N4]-L-Arginine and [U-13C6, 15N2]-L-LysineBuffer:25 mM Tris-HCI, 100 mM glycine, pH 7.3Storage:Store at -80°C. Avoid repeated freeze thaw cycles.Stability:Stable for 3 months from receipt of products under proper storage and handling conditions.RefSeq Size:NP 036199RefSeq ORF:93Synonyme:AFAR2		RVKIDTKAIPLFGNSLKPDSLRFQLETSLKRLQCPRVDLFYLHMPDHSTPVEETLRACHQLHQEGKFMEL GLSNYAAWEVAEICTLCKSNGWILPTVYQGMYNAITRQVETELFPCLRHFGLRFYAFNPLAGGLLTGKYK YEDKDGKQPVGRFFGNTWAEMYRNRYWKEHHFEGIAPVEKALQAAYGASAPSMTSATLRWMYHHSQLQGA
Purity:> 80% as determined by SDS-PAGE and Coomassie blue stainingPurity:> 80% as determined by microplate BCA methodConcentration:> 0.05 µg/µL as determined by microplate BCA methodLabeling Method:Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-LysineBuffer:25 mM Tris-HCl, 100 mM glycine, pH 7.3Storage:Store at -80°C. Avoid repeated freeze-thaw cycles.Stability:Stable for 3 months from receipt of products under proper storage and handling conditions.RefSeq:NP 036199RefSeq ORF:993Synonyms:AFAR2		TRTRPLEQKLISEEDLAANDILDYKDDDDKV
Concentration:>0.05 μg/μL as determined by microplate BCA methodLabeling Method:Labeled with [U-13C6, 15N4]-L-Arginine and [U-13C6, 15N2]-L-LysineBuffer:25 mM Tris-HCl, 100 mM glycine, pH 7.3Storage:Store at -80°C. Avoid repeated freeze-thaw cycles.Stability:Stable for 3 months from receipt of products under proper storage and handling conditions.RefSeq:NP 036199RefSeq ORF:993Synonyms:AFAR2	Tag:	C-Myc/DDK
Labeling Method:Labeled with [U-13C6, 15N4]-L-Arginine and [U-13C6, 15N2]-L-LysineBuffer:25 mM Tris-HCl, 100 mM glycine, pH 7.3Storage:Store at -80°C. Avoid repeated freeze-thaw cycles.Stability:Stable for 3 months from receipt of products under proper storage and handling conditions.RefSeq:NP 036199RefSeq ORF:993Synonyms:AFAR2	Purity:	> 80% as determined by SDS-PAGE and Coomassie blue staining
Buffer:25 mM Tris-HCl, 100 mM glycine, pH 7.3Storage:Store at -80°C. Avoid repeated freeze-thaw cycles.Stability:Stable for 3 months from receipt of products under proper storage and handling conditions.RefSeq:NP 036199RefSeq Size:1301RefSeq ORF:993Synonyms:AFAR2	Concentration:	>0.05 μg/μL as determined by microplate BCA method
Storage:Store at -80°C. Avoid repeated freeze-thaw cycles.Stability:Stable for 3 months from receipt of products under proper storage and handling conditions.RefSeq:NP 036199RefSeq Size:1301RefSeq ORF:993Synonyms:AFAR2	Labeling Method:	Labeled with [U- 13C6, 15N4]-L-Arginine and [U- 13C6, 15N2]-L-Lysine
Stability:Stable for 3 months from receipt of products under proper storage and handling conditions.RefSeq:NP 036199RefSeq Size:1301RefSeq ORF:993Synonyms:AFAR2	Buffer:	25 mM Tris-HCl, 100 mM glycine, pH 7.3
RefSeq: NP 036199 RefSeq Size: 1301 RefSeq ORF: 993 Synonyms: AFAR2	Storage:	Store at -80°C. Avoid repeated freeze-thaw cycles.
RefSeq Size: 1301 RefSeq ORF: 993 Synonyms: AFAR2	Stability:	Stable for 3 months from receipt of products under proper storage and handling conditions.
RefSeq ORF:993Synonyms:AFAR2	RefSeq:	<u>NP 036199</u>
Synonyms: AFAR2	RefSeq Size:	1301
	RefSeq ORF:	993
Locus ID: 22977	Synonyms:	AFAR2
	Locus ID:	22977



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	AKR7A3 (NM_012067) Human Mass Spec Standard – PH306675
UniProt ID:	<u>095154, A0A384MDN8</u>
Cytogenetics:	1p36.13
Summary:	Aldo-keto reductases, such as AKR7A3, are involved in the detoxification of aldehydes and ketones.[supplied by OMIM, Apr 2004]
Protein Families	: Druggable Genome
Product imag	ges:



Coomassie blue staining of purified AKR7A3 protein (Cat# [TP306675]). The protein was produced from HEK293T cells transfected with AKR7A3 cDNA clone (Cat# [RC206675]) using MegaTran 2.0 (Cat# [TT210002]).

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