

Cloning and characterization of GPCATs

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PpaGPCATH1 1 K-----
PpaGPCATH2 1 K-----
SmoGPCATH1 1 MAPDDAG-----
SmoGPCATH2 1 VQ-----
ZmaGPCATH1 1 MASEVEDD-----
ZmaGPCATH2 1 MASEVEDE-----
OsaGPCATH 1 MASEVEDD-----
MacGPCATH1 1 MGDGETGS-----
StuGPCATH1 1 MSSNEESM-----
StuGPCATH2 1 MEEYNGD-----
RcoGPCATH 1 MSNNEDI-----
AthGPCATH 1 MANNEDSN-----
BvuGPCATH 1 MNDSSEFHINILHHQLPF-----
NnuGPCATH 1 MSNKEDAE-----
EguGPCATH 1 MSEGAGL-----
NadGPCATH 1 GWFTLKKRT-----
AtrGPCATH 1 MESYESP-----
LtuGPCATH 1 GEKKKGRRK--LLKNRP-----
AfiGPCATH 1 NPARIQHSE-----
KfiGPCATH 1 MGLILEAA-----
EmuGPCATH 1 MVLAPLQNK-----
OviGPCATH 1 MAPSIANI-----
SpuGPCATH 1 MSLQNLNGCSLKQ-----
CgiGPCATH 1 MRDTLKENTKQFATA-----
LgiGPCATH1 1 MMVDKEDAKRKTQN-----
LgiGPCATH2 1 MTSQSMNYNQNILIV-----
LcoGPCATH 1 MAASMTKPPPELADSH----S-MSS-----
MciGPCATH1 1 MTPKEDTPGDSDLQDT----T-VAAAAA-E-----
MciGPCATH2 1 MLTEERLEHLSSHESD----A-----
MveGPCATH 1 MVTSSIDGPTHNPSST----M-SS-SDS---DSD--VPLTRISAPDPD-EHTYMDANATD-----PAWISE
UmaGPCATH 1 MRRASSTSTYSHVSD----V-SS-SGD---ALR--PTLSRASSTACSD-----DGSLSTPSFRTHHSSASASME-----
LbiGPCATH 1 MWHSTLVFADT-----
SpoGPCATH 1 MDHLEFDENTDSEYSI----F-E-----
SceGPCATH 1 MYKLDNNDIDDET-----
LthGPCATH 1 MDIPDTLDTDSDEGS-----
PkuGPCATH 1 MDIDNMEKDNGCSTP----T-SDES-----
NcrGPCATH 1 MADLAEPEISLPVSGL----S-HKSEGAPSTDGE--AILTPDTPSTPPLNSSPGEHDPNTPT-----LDVTT
CbaGPCATH 1 MSEPIRIPLSKTDITL---D-EM--AS-----H--DIAGSASTDTAGS-LTGALTPGYQSG-----YQSSS
AclGPCATH 1 MEGAGLDDHPGGGRTA----Q-VTGAGN-----S--PLLPPAAVDDMDA-VYDTDEYAPSST-----RSYSP
CglGPCATH 1 MADSEDRSKPISIPGN----N-NRWEA--STDSE--AILTPELSGSPD-DDSHEHAVVTPT-----LGPTV
BbaGPCATH 1 MDDEQPAAVSHPNPNA----T-FQDSIS-D---E--HHTEPGLTSPLSS-SPPDPTILDTPS-----LIATT
DdiGPCATH1 1 MEVLASEISSNEKSAF-----
DdiGPCATH2 1 MEKKDKNEKISSSPI----N-IFRRYS-I-----
AsuGPCATH 1 MTMTMMVIKVEKHSKV----G-----
EdiGPCATH 1 MSTSSEKYQREFQ-----
AcaGPCATH1 1 MDHKKVDKHEHNSRA----G-----
AcaGPCATH2 1 MADAVHNEPEADIRAY-----
PbrGPCATH 1 IQQTSADLAYVGTTRL--RRVYRVASADC---DPV--V-----CINKVT-----IY
NgrGPCATH1 1 MEKQEEIEITNNNFDS----E-----
NgrGPCATH2 1 MSQQTKKKKESKQEK----K-----
MbrGPCATH 1 MASESTVPADPAESPL----G-LSQ-----
BsaGPCATH 1 MWWEIFGVFGVC-----
OtaGPCATH 1 MRRTRSVRSDD-----
MpuGPCATH1 1 MASPIAAAAAT-----
MpuGPCATH2 1 MSPSFAEP-----
MspGPCATH1 1 MAEAVMRRRRR-----
MspGPCATH2 1 MNLEPPTL-----
CreGPCATH 1 MESFGVAAAEAA-----
CowGPCATH 1 MPANEAPLSPSINATT----ATSPATT--ETTTAAATNTNPSGAAAN-GVQDGGKAQ-----T
CmeGPCATH 1 MDDGGRALGSVDHF----D-ET-----

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Cloning and characterization of GPCATs

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PpaGPCATH1 2 -----
PpaGPCATH2 2 -----
SmoGPCATH1 9 -----
SmoGPCATH2 3 -----
ZmaGPCATH1 9 -----
ZmaGPCATH2 9 -----
OsaGPCATH 9 -----
MacGPCATH1 9 -----
StuGPCATH1 9 -----
StuGPCATH2 9 -----
RcoGPCATH 9 -----
AthGPCATH 9 -----
BvuGPCATH 20 ----- IE-----
NnuGPCATH 9 -----
EguGPCATH 9 -----
NadGPCATH 10 -----
AtrGPCATH 8 -----
LtuGPCATH 16 -----
AfiGPCATH 10 -----
KfiGPCATH 9 -----
EmuGPCATH 11 -----
OviGPCATH 10 -----
SpuGPCATH 15 -----
CgiGPCATH 16 -----
LgiGPCATH1 15 -----
LgiGPCATH2 16 -----
LcoGPCATH 21 ----- GS-----
MciGPCATH1 25 ----- TA-----
MciGPCATH2 18 ----- S-----
MveGPCATH 56 PTPP-----SLGPTS-----LVPN-----
UmaGPCATH 60 -----
LbiGPCATH 12 -----
SpoGPCATH 19 -----
SceGPCATH 15 -----
LthGPCATH 17 -----
PkuGPCATH 22 -----
NcrGPCATH 61 PKSP-----AVLS-----RNS-----
CbaGPCATH 53 PGSP-----YLLS-----RNP-----
AclGPCATH 55 TASP-----PSLS-----RNH-----
CglGPCATH 59 PTSP-----RLS-----RNP-----
BbaGPCATH 56 PKSP-----KLS-----RHP-----
DdiGPCATH1 17 ----- TEEEDDYEFSDSETDSL-----
DdiGPCATH2 25 ----- AD-----
AsuGPCATH 18 -----
EdiGPCATH 15 -----
AcaGPCATH1 18 -----
AcaGPCATH2 17 ----- EPQAADA-----
PbrGPCATH 40 PAS-----AAPKG-----IFVW-----MPLNRA-----RRRG
NgrGPCATH1 18 -----ERMDQRMDQGKEGMADHYGSS-----
NgrGPCATH2 18 -----QEQDP-----
MbrGPCATH 21 ----- ES-----
BsaGPCATH 13 -----LVVYAAVTRFTTLAAITQMGSIQ-----
OtaGPCATH 12 -----ATVDE-----L-----SRARERAK-----
MpuGPCATH1 12 -----AAVTG-----LV-----RRM-----RRR-----
MpuGPCATH2 9 -----
MspGPCATH1 12 -----GE-----
MspGPCATH2 9 -----
CreGPCATH 13 -----
CowGPCATH 52 ---SSTSENAVQTMERPRSAL-----LRPGQEEQEQAEREAHQTTTHQLHDGYAAA-----
CmeGPCATH 20 -----D-----

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Cloning and characterization of GPCATs

PpaGPCATH1	2	-----
PpaGPCATH2	2	-----
SmoGPCATH1	9	-----
SmoGPCATH2	3	-----
ZmaGPCATH1	9	-----
ZmaGPCATH2	9	-----
OsaGPCATH	9	-----
MacGPCATH1	9	-----
StuGPCATH1	9	-----
StuGPCATH2	9	-----
RcoGPCAT	9	-----
AthGPCAT	9	-----
BvuGPCATH	22	-----VR-E-----QI-LKSLLFQSI-
NnuGPCATH	9	-----
EguGPCATH	9	-----
NadGPCATH	10	-----
AtrGPCATH	8	-----
LtuGPCATH	16	-----
AfiGPCATH	10	-----
KfiGPCATH	9	-----
EmuGPCATH	11	-----
OviGPCATH	10	-----
SpuGPCATH	15	-----
CgiGPCATH	16	-----
LgiGPCATH1	15	-----
LgiGPCATH2	16	-----KP-----
LcoGPCATH	23	-----SS--NG-DD--LS-DF-----EHDLDVI-----EGW-
MciGPCATH1	27	-----SS--SG-----IK-NF-----G--NEDMFSL-----DEW-
MciGPCATH2	19	-----TL--LS-ED--ST-EY-----D--TDELTTV-----DSW-
MveGPCATH	70	-----ILTQLSFS-----
UmaGPCATH	60	-----GIDLLGSSASA-----AAL-
LbiGPCATH	12	-----
SpoGPCATH	19	-----E-DN-----D--YGLHGLD-----DSV-
SceGPCAT	15	-----NSV-----SLT-
LthGPCATH	17	-----EPMKV-----TLA-
PkuGPCATH	22	-----SS-----NAYADL-----S--SSDAPFF-----EGV-
NcrGPCATH	72	-----SF--SG--SISSTL-HD-----E--WDVFPPL-----DRL-
ChaGPCATH	64	-----SH--IG--S--QSL-QE-----D--W--EVPL-----DKL-
Ac1GPCATH	66	-----S-----FSY-QD-----D--WDTFPPL-----DKL-
Cg1GPCATH	69	-----SF--SG--S--SSY-QE-----D--WDPFPPL-----DRL-
BbaGPCATH	66	-----SF--SG--S--SSY-HD-----D--WDSLPPPL-----DRL-
DdiGPCATH1	35	-----DYN--LE--DSSN-D-ENSGDGDGGGVVESDDKNLKRKR--SKRPTTL-----GESN
DdiGPCATH2	27	-----VG--SGSSISRLQH-DS-----N--NN-----NSP-
AsuGPCATH	18	-----ELSPSSR-RL-----K--DK-----SKD-
EdiGPCATH	15	-----
AcaGPCATH1	18	-----EQ--VE-EL-----R--DENPSTV-----SED-
AcaGPCATH2	24	-----QH--LH--DESE-EQHHSQNDSE-----
PbrGPCATH	62	SADERMTE-DEVGASDVA--T--DSE-ED-----D--ER-----SASE-
NgrGPCATH1	39	-----
NgrGPCATH2	23	-----
MbrGPCATH	23	-----A-----S-Q-----VI-----EDI-
BsaGPCATH	36	-----ERAMPWA-----
OtaGPCATH	26	-----
MpuGPCATH1	25	---DALSDDDENGA-TTSSDEG-V--GS--P-----R--GH--VVAHHA-----
MpuGPCATH2	9	-----
MspGPCATH1	14	-----ENGY-MSPTSDG-E--DS--P-----R--EW--RR--RNPLAP-
MspGPCATH2	9	-----
CreGPCATH	13	-----
CowGPCATH	99	-----ESEQDHA-----
CmeGPCATH	21	-----SS--AF-PD--SE-AY-----YEPDL-----LSL-

Cloning and characterization of GPCATs

PpaGPCATH1	2	-----	
PpaGPCATH2	2	-----	
SmoGPCATH1	9	-----	
SmoGPCATH2	3	-----	
ZmaGPCATH1	9	-----	
ZmaGPCATH2	9	-----	
OsaGPCATH	9	-----	
MacGPCATH1	9	-----	
StuGPCATH1	9	-----	
StuGPCATH2	9	-----	
RcoGPCATH	9	-----	
AthGPCATH	9	-----	
BvuGPCATH	36	-----	
NnuGPCATH	9	-----	
EguGPCATH	9	-----	
NadGPCATH	10	-----	
AtrGPCATH	8	-----	
LtuGPCATH	16	-----	
AfiGPCATH	10	-----	
KfiGPCATH	9	-----	
EmuGPCATH	11	-----	
OviGPCATH	10	-----	
SpuGPCATH	15	-----	
CgiGPCATH	16	-----	
LgiGPCATH1	15	-----	
LgiGPCATH2	18	-----	
LcoGPCATH	43	-----TTADFISE-----	
MciGPCATH1	46	-----ETSDFISE-----	
MciGPCATH2	40	-----SSHDFVAK-----	
MveGPCATH	78	-----	
UmaGPCATH	74	-----	
LbiGPCATH	12	-----	
SpoGPCATH	33	-----	
SceGPCATH	21	-----	
LthGPCATH	25	-----	
PkuGPCATH	41	-----	
NcrGPCATH	95	-----	
CbaGPCATH	83	-----	
AcIGPCATH	83	-----	
CglGPCATH	90	-----	
BbaGPCATH	87	-----	
DdiGPCATH1	82	KDGETANSVSI SAGTSKSNISSPLVDGADILSKPNSSPTSSSLSGSNSAN--GFLSDDDDLLNGNEQLNENHNTNSTNG	
DdiGPCATH2	47	-----LVPTSC-----	
AsuGPCATH	33	-----SDKYLD-----	
EdiGPCATH	15	-----	
AcaGPCATH1	35	-----TNDAFVD-----	
AcaGPCATH2	42	-----NATQPERAEDLA-----	
PbrGPCATH	93	-----	
NgrGPCATH1	39	-----EDED-----	
NgrGPCATH2	23	-----VTTA-----	
MbrGPCATH	31	-----R--NFMSDL-----	
BsaGPCATH	43	-----SFS-----	
OtaGPCATH	26	-----	
MpuGPCATH1	56	-----	
MpuGPCATH2	9	-----	
MspGPCATH1	40	-----	
MspGPCATH2	9	-----	
CreGPCATH	13	-----	
CowGPCATH	106	-----SCD-----	
CmeGPCATH	39	-----	

Cloning and characterization of GPCATs

PpaGPCATH1	2	-----
PpaGPCATH2	2	-----
SmoGPCATH1	9	-----
SmoGPCATH2	3	-----
ZmaGPCATH1	9	-----
ZmaGPCATH2	9	-----
OsaGPCATH	9	-----
MacGPCATH1	9	-----
StuGPCATH1	9	-----
StuGPCATH2	9	-----
RcoGPCATH	9	-----
AthGPCATH	9	-----
BvuGPCATH	36	-----SLCKIPSI-----
NnuGPCATH	9	-----
EguGPCATH	9	-----
NadGPCATH	10	-----
AtrGPCATH	8	-----
LtuGPCATH	16	-----
AfiGPCATH	10	-----
KfiGPCATH	9	-----
EmuGPCATH	11	-----
OviGPCATH	10	-----
SpuGPCATH	15	-----
CgiGPCATH	16	-----
LgiGPCATH1	15	-----
LgiGPCATH2	18	-----
LcoGPCATH	51	-----NDFDMI--E-----
MciGPCATH1	54	-----NDFDMI--D-----
MciGPCATH2	48	-----DDYAMI--G-----
MveGPCATH	78	-----L--D-----
UmaGPCATH	74	-----PLSDF--E-----
LbiGPCATH	12	-----F--E-----
SpoGPCATH	33	-----GFTDLF--D-----
SceGPCATH	21	-----SLLEFL--D-----
LthGPCATH	25	-----NWVEFL--D-----
PkuGPCATH	41	-----TLDDL--Q-----
NcrGPCATH	95	-----TVLDDL--D-----
CbaGPCATH	83	-----TFFDIF--D-----
Ac1GPCATH	83	-----TVFDLL--D-----
Cg1GPCATH	90	-----TVLDIL--D-----
BbaGPCATH	87	-----TVLDDL--D-----
DdiGPCATH1	160	ATNNNNNNNNNSGNSPSHQSSSEL-----
DdiGPCATH2	53	-----D--S-----
AsuGPCATH	39	-----E-----
EdiGPCATH	15	-----
AcaGPCATH1	42	-----A-----
AcaGPCATH2	54	-----LQIASGK-----
PbrGPCATH	93	-----VFGQIV--D-----
NgrGPCATH1	43	-----TLSDQL-----
NgrGPCATH2	27	-----TITPT-----
MbrGPCATH	38	-----DFATQQ--D-----
BsaGPCATH	46	-----KPDA--GGGS-----
OtaGPCATH	26	-----
MpuGPCATH1	56	-----
MpuGPCATH2	9	-----
MspGPCATH1	40	-----FFK-----
MspGPCATH2	9	-----
CreGPCATH	13	-----
CowGPCATH	109	-----HLLHDDHAALLDAA--DDLLAPGIQLQGSVNDMNDRVESYVMQTLDRIASGLSDSDEDDPDGAD-----
CmeGPCATH	39	-----SLSDIF--T-----

Cloning and characterization of GPCATs

PpaGPCATH1	2	-----
PpaGPCATH2	2	-----
SmoGPCATH1	9	-----
SmoGPCATH2	3	-----
ZmaGPCATH1	9	-----
ZmaGPCATH2	9	-----
OsaGPCATH	9	-----
MacGPCATH1	9	-----
StuGPCATH1	9	-----
StuGPCATH2	9	-----
RcoGPCATH	9	-----
AthGPCATH	9	-----
BvuGPCATH	45	-----
NnuGPCATH	9	-----
EguGPCATH	9	-----
NadGPCATH	10	-----
AtrGPCATH	8	-----
LtuGPCATH	16	-----
AfiGPCATH	10	-----
KfiGPCATH	9	-----
EmuGPCATH	11	-----
OviGPCATH	10	-----
SpuGPCATH	15	-----
CgiGPCATH	16	-----
LgiGPCATH1	15	-----
LgiGPCATH2	18	-----
LcoGPCATH	58	-----
MciGPCATH1	61	-----
MciGPCATH2	55	-----
MveGPCATH	80	-----
UmaGPCATH	81	-----
LbiGPCATH	14	-----
SpoGPCATH	40	-----
SceGPCATH	28	-----
LthGPCATH	32	-----
PkuGPCATH	48	-----
NcrGPCATH	102	-----
CbaGPCATH	90	-----
AcIGPCATH	90	-----
CglGPCATH	97	-----
BbaGPCATH	94	-----
DdiGPCATH1	185	----- PLLNLFQKFLNREAI -----
DdiGPCATH2	55	-----
AsuGPCATH	40	-----
EdiGPCATH	15	-----
AcaGPCATH1	43	-----
AcaGPCATH2	62	----- SKKSFFERFTVLWAQ -----
PhrGPCATH	100	-----
NgrGPCATH1	49	----- VVEDEEE-YAYY-DW -----
NgrGPCATH2	33	----- TTT----- T-NN -----
MbrGPCATH	45	-----
BsaGPCATH	54	GHRSTT----- NTTATSSQQPDKATS ----- SEAT
OtaGPCATH	26	-----
MpuGPCATH1	56	-----
MpuGPCATH2	9	-----
MspGPCATH1	43	-----
MspGPCATH2	9	-----
CreGPCATH	13	AE-----
CowGPCATH	170	AVRDSSSHSDNPDNLQFYQQHDNEQTDRDASSIRRQQEKRASKRSVPSGADSDSEWVMSVPGSDARDLSDMDAGFTPSAS
CmeGPCATH	46	-----

Cloning and characterization of GPCATs

PpaGPCATH1	2	-----
PpaGPCATH2	2	-----
SmoGPCATH1	9	---SP-----
SmoGPCATH2	3	-----
ZmaGPCATH1	9	---AAVP-----GA--P-----L--VA-----
ZmaGPCATH2	9	---AATP-----L-----LA-----
OsaGPCATH	9	---AAAV-----LT--N-----GG-----
MacGPCATH1	9	---DEGE-----AF--QKVRH---RIA-DQS-----KR--VV-----
StuGPCATH1	9	---EDSN-----GHSF--GKVKQ---RFK-DRS-----
StuGPCATH2	9	---S-----F--GKVKQ---RFK-DRS-----
RcoGPCATH	9	---NEFV-----SNGDSF--EKVKQ---RLK-DRS-----KK--VA-----
AthGPCATH	9	---SNGL-----DSF--DAVKQ---RFK-DRS-----KK--VV-----
BvuGPCATH	68	KVEEVN-----GHSF--QKVKQ---RLK-DRT-----
NnuGPCATH	9	---DGGD-----PF--PTVKQ---RFS-DRT-----KK--V-----
EguGPCATH	9	---DD-----GEAF--RKVRQ---RIA-QDS-----KR--VV-----
NadGPCATH	28	EGEDCD-----AKPF--RKVRQ---KLV-----
AtrGPCATH	8	---E-----AQE---AFR-----
LtuGPCATH	39	IVGGGE-----KESFR-KAVRK---RLS-DRT-----KK--VA-----
AfiGPCATH	31	EVDGDR-----GQPF--RRVKQ---RFS-DRT-----KK--VA-----
KflGPCATH	9	---AEEATARSASTSDSET--EGVRQ---RRP-GDA-----EV--LA---QEPL-----WV---P-
EmuGPCATH	11	-----
OviGPCATH	10	-----
SpuGPCATH	15	-----TEVAE---QDDG-----TK--WAQ---PAKD-----
CgiGPCATH	16	-----EEE-VTY---E-----
LgiGPCATH1	15	-----YNTG-TSR---E-----
LgiGPCATH2	18	-----VDTVVTCT-NDDG-----NTKPVAS---VTINGFIP-RSS---S-----
LcoGPCATH	61	-----AA-L-EQVTH---QLN-DKQ-----RE--IR---AR-----STEWTTTKT-
MciGPCATH1	64	-----AA-L-DQVTN---QLE-DKQ-----RE--FI---SK-----SSEWTKTR-
MciGPCATH2	58	-----AA-L-EQVTL---QIR-DVK-----NE--IK---AL-----LSTEWTSMTQ-
MveGPCATH	84	-----QN-L-EEELT---DLP-VLANQIYKTLRQD---LK---KNKQ---K-LKHQM---K-
UmaGPCATH	85	-----PR-I-DLAGR---KWS-AFS---GDMRSR---AR---RRRE---E-LVRKA-----
LbiGPCATH	18	-----SR-V-DLQOR---KIQ-KHS-----DR--LK---MKAE---E-TFKIK---E-
SpoGPCATH	45	-----RV-Y-SWVHK---HYN-QKK-----GQ---LKHGVSQRKN---K-L-----
SceGPCATH	33	-----VV-S-KYYHG---SHLSKAE---QK---LR---NFEG---F-RRRKPHEHD-SH-
LthGPCATH	37	-----AQ-S-QVPR---KYLTRAR-----RR--LK---S-----ARPRQ-KL-
PkuGPCATH	53	-----LE-L-RKKKD---DTYKWVR---KV---YN---DK-----
NcrGPCATH	107	-----QQ-L-EKQOR---GIH-ART-----EK--MR---KSGE---A-LKEKSAIARERMV-
ChaGPCATH	95	-----AK-L-EKWQA---TLQ-AQK-----EK--LA---RQOE---R-IRSSGNAKDRAM-
Ac1GPCATH	95	-----QK-L-EKWQH---AIN-VQK-----EK--VR---KQRE---K-LKYTSMHAKDRVV-
Cg1GPCATH	102	-----QQ-L-EKQOK---GIS-AQT-----EK--VR---KSRQ---A-FGNKSRIARERMV-
BbaGPCATH	99	-----QQ-L-EKQOK---GIS-AQT-----DK--VR---RSRD---A-LKSKTQARHRMV-
DdiGPCATH1	200	-----KVQQ---RHS-AIN-----NR--LQ---R-----L-KKVVD
DdiGPCATH2	55	-----P-T-SNFK-----
AsuGPCATH	45	-----SP-I-GQFKK---SID-DRR-----
EdiGPCATH	15	-----
AcaGPCATH1	48	-----RE-L-DLAD---ALQ-SLP-----V-----GFK-
AcaGPCATH2	77	-----RKVID---KLL-T-S-----PH--VK---R-----
PhrGPCATH	105	-----VR-L-ENQOT---TLE-TG-----R-GA---GLVRD-----AR-
NgrGPCATH1	62	-----ETVGS---SLQ-NQE-----EE--LI---DLWM---RINDAFSDRKKSLVN
NgrGPCATH2	39	-----N-----NT--TT---TTSS---SP-----KQ
MbrGPCATH	48	-----AAPLSVNMNR---RLT-IL-----R-
BsaGPCATH	79	-----
OtaGPCATH	26	-----AVAR---AIR-RK-----SG--ELLKS-----AK-
MpuGPCATH1	56	-----RGVQ---FFK-RP-----VE--NV-----
MpuGPCATH2	9	---PNKV---A-----
MspGPCATH1	43	-----RPRRP---SLP-EG-----AR--AI-----
MspGPCATH2	9	---RSSE---E-----
CreGPCATH	15	-----
CowGPCATH	279	-----RS-R-RPSAA---HLD-RES-----EQ--AR---RLQK---SMSTSSI--GDRLID
CmeGPCATH	48	-----S-L-----

Cloning and characterization of GPCATs

PpaGPCATH1	30	-----YCHGVVSFGTFCFL-----	LGSKPADLPEKLYC-FFFTTVAPLRNIIYYRIRKWHYFL-----
PpaGPCATH2	29	-----YCHGVVSFGTFCFL-----	LGSRFDALPEKLYC-FFFTTVAPLRNIIYYRIRKWHYFL-----
SmoGPCATH1	48	-----YCHGVFCFGTFCV-----	MGSRPEHEIPYLYC-FFFTTVAPLRNIIYYRIRKWHYFL-----
SmoGPCATH2	30	-----YCHGVFCFGTFCFL-----	LGSRPQDIPYLYC-TFFTVAPLRNIIYYRIRKWHYFL-----
ZmaGPCATH1	58	-----HLIGVIGFGGFCFL-----	LGAREQDIPYVYC-FFYVIFVPLRNIIYYRIRKWHYFL-----
ZmaGPCATH2	55	-----HLIGVIGFGGFCFL-----	LGAREQDIPYVYC-FFYVIFVPLRNIIYYRIRKWHYFL-----
OsaGPCATH	57	-----HLIGVIGFGTFCFL-----	LGAREQDIPYVYC-FFYVIFVPLRNIIYYRIRKWHYFL-----
MacGPCATH1	69	-----HLIGVIGFGAFCFL-----	LGAREQDIPYVYC-FFYVIFVPLRNIIYYRIRKWHYFL-----
StuGPCATH1	61	-----HLMGVIGFGAFCFL-----	LGAREKDIRYVYC-FFYVIFVPLRNIIYYRIRKWHYFL-----
StuGPCATH2	55	-----HLIGVIGFGAFCFL-----	LGAREQDIRYAYS-FFYVIFVPLRNIIYYRIRKWHYFL-----
RcoGPCATH	73	-----HLIGVIGFGGFCFL-----	LGAREQDIPYVYC-FFYVIFVPLRNIIYYRIRKWHYFL-----
AthGPCATH	70	-----HLVGVIGFGGFCFL-----	LGAREQDIPLVYC-FFYVIFVPLRNIIYYRIRKWHYFL-----
BvuGPCATH	125	-----YICGVIGFGGFCFL-----	LGAREQDIPYVYC-FFYVIFVPLRNIIYYRIRKWHYFL-----
NnuGPCATH	67	-----HLIGVIGFGAFCFL-----	LGSRPQDIPYVYC-FFYVIFVPLRNIIYYRIRKWHYFL-----
EguGPCATH	73	-----HLMGVIGFGAFCFL-----	LGAREQDIPYVYC-FFYVIFVPLRNIIYYRIRKWHYFL-----
NadGPCATH	85	-----HLIGVIGFGAFCFL-----	LGSRPQDIPYVYC-VFYVTFVPLRNIIYYRIRKWHYFL-----
AtrGPCATH	54	-----HLIGVIGFGAFCFL-----	LGSRPQDIPYIYC-FFYVTFVPLRNIIYYRIRKWHYFL-----
LtuGPCATH	104	-----HLIGVIGFGAICFL-----	LGAREQDIPYVYC-FFYVTFVPLRNIIYYRIRKWHYFL-----
AfiGPCATH	106	-----HLIGVIGFGSFCFL-----	LGAREQDIPYVYC-FFYVTFVPLRNIIYYRIRKWHYFL-----
KflGPCATH	91	-----YSIGVICFGLFCFL-----	VGAKPQVLEPYEYC-FFLFVALPLRNIIYYRIRKWHYFL-----
EmuGPCATH	38	-----FVLSVTLITVVFVCF-----	AVVREPALILYIES-GFFVAVVFRVINYWRSNIILEM-----
OviGPCATH	20	-----FVLCATAFVLDSTFC-----	FVVREALILYIES-GFFVAVVFRVINYWRSNIILEM-----
SpuGPCATH	82	-----FVVSILITIVVCCS-----	MMSFQWLEPYLYT-AYLSVLIPIRAYKIIYRIRKWHYFL-----
CgiGPCATH	91	-----YVVAVITLIFLTHA-----	TVYAQWLEPYIYA-VSTPIILILRVIMYWKNKWQYFL-----
LgiGPCATH1	68	-----YVLSIGTILLLTHS-----	LMASQWLEPYIYT-ISTPCLLILRTIYWKIRKWHYFL-----
LgiGPCATH2	94	-----YVLSIGTILLLTHS-----	LMASQWLEPYIYT-ISTPCLLILRTIYWKIRKWHYFL-----
LcoGPCATH	136	-----FVVGGNACVSPAL-----	AARLPEWLEPYIYT-IQSLYLLSLRLIYRIRKWHYFL-----
MciGPCATH1	139	-----FVVGGNACVAPAL-----	ALRYPSLEPSIYS-SQLVLLILIRVVIYRIRKWHYFL-----
MciGPCATH2	134	-----FVVGVNTCITPVL-----	AARLPEWLEPYIYT-VQLCYLITLRFVYRIRKWHYFL-----
MveGPCATH	166	-----FTVGMFECCCTPFL-----	VAQYPEWLELVHT-VQVATLIMRVFLYRIRKWHYFL-----
UmaGPCATH	161	-----FVVGVCNIVVSSLV-----	FALRELEPTLYS-LLAYFPLRNVSYRIRKWHYFL-----
LbiGPCATH	92	-----FFFCVMSLLSALL-----	FGIAPHWLVHAYS-VQGLYLLPLRAYRYRIRKWHYFL-----
SpoGPCATH	115	-----FALGVSTCILTALL-----	VGMAPESVHLIYT-IQLFVYVPLRNIIYYRIRKWHYFL-----
SceGPCATH	116	-----YPTFLFNFFIGL-----	MGRPEWVHVIYT-ILFFVLMPIRYTYRIRKWHYFL-----
LthGPCATH	111	-----YPTFLFHFAIGL-----	IGKYPSEWVHVIYT-GMLVLLMPVRYTYRIRKWHYFL-----
PkuGPCATH	118	-----FSFSVYMLFFIGL-----	IGHSEPYHILYS-VMFALMPLRILTYRIRKWHYFL-----
NcrGPCATH	188	-----FIFGVNIFISGL-----	IGAPPEWFHVIYT-VQIYFMPIRYRYRIRKWHYFL-----
ChaGPCATH	176	-----FISAVNIFISGL-----	VGAVEQYFYWYT-IQLYFMPIRAYTYRIRKWHYFL-----
AclGPCATH	176	-----FISAVNIFISGL-----	IGAYPEYFYIES-VQLAYFMPIRYRYRIRKWHYFL-----
CglGPCATH	183	-----FICGVNIFVSGL-----	IGGYPEYFHLIYT-IQVYFMPIRYRYRIRKWHYFL-----
BbaGPCATH	180	-----FICGVNIFISGL-----	IGGYPEYFHLIYT-VQLYFMPIRMFTYRIRKWHYFL-----
DdiGPCATH1	268	-----FTVGLIIMVSEFV-----	LLKSPQLFYLYYT-VLIFPLMAREVMIYRIRKWHYFL-----
DdiGPCATH2	107	-----FVIGVNLICVISEF-----	CKKPYLYQYFES-GEYVLFATRIILYRIRKWHYFL-----
AsuGPCATH	104	-----FVNGVNLICVISEF-----	YGRAPKYLEDVFA-VEFCLGFLRLIYRIRKWHYFL-----
EdiGPCATH	55	-----FVCGVLTLLTQHL-----	LSALPQFMPIYV-ILIFPLGARFIYRIRKWHYFL-----
AcaGPCATH1	104	-----FVAGVTNALTFLL-----	LGRHPEWLEIYIYT-ILKAFLLITIRYVYRIRKWHYFL-----
AcaGPCATH2	138	-----FMMGLILMLVTEAL-----	L-TRKNAMYQBYT-VMFPLMVFREASYRIRKWHYFL-----
PhrGPCATH	170	-----FTVGLIGSLITVCL-----	LGRSSDLVKKIYT-YLIVLLISLRIYLYRIRKWHYFL-----
NgrGPCATH1	140	-----FTVGLIIMVSEFV-----	LAKYFYLLEDIYI-VMSVILMVRIFHYRIRKWHYFL-----
NgrGPCATH2	94	-----FTVGLIIMVSTTL-----	FCHSEKYLEIYI-LWMLPLMIYRIRKWHYFL-----
MbrGPCATH	116	-----FVACVSTLALALEFL-----	LLQETRYLPEYV-ALMFPILMARVFLYRIRKWHYFL-----
BsaGPCATH	102	-----YTSVINAVALTALL-----	FGVVEHYFHWHS-PKALMYLAHRNYTFIRKWHYFL-----
OtaGPCATH	84	-----FCVGVFGIMTEK-----	ATRAPEKFWMRYA-FAMPALVCRAYRYRIRKWHYFL-----
MpuGPCATH1	109	-----FTVGVVGLMVTQHL-----	ATTREGLFWLYL-VVAPFVAFANVATYRIRKWHYFL-----
MpuGPCATH2	50	-----FFVGVNIFLTAFL-----	LGNTPEYVWVQL-LKNVAVSTASVIRKWHYFL-----
MspGPCATH1	94	-----FTVGVVGLMVTQFL-----	ATEHEEYFWVYFL-CSAPLFIYRIRKWHYFL-----
MspGPCATH2	42	-----FALGLFNWGSALC-----	LGRFVQYVWVQV-VKSMNYLSIRKWHYFL-----
CreGPCATH	55	GAESDVVEFLDATTGGPGVVAEALDMHEVFHLEEQAAALQVGLGEAVVTE-----	-----QKRRR-----
CowGPCATH	394	-----FTGFVASFLVTEFL-----	LLRYEQWVWLYA-VCIPLVGTIRVLYRIRKWHYFL-----
CmeGPCATH	88	-----FVIGLNLVLFTEK-----	ILRQEARVWQVYL-YQLTPLLILRYSYRIRKWHYFL-----

Cloning and characterization of GPCATs

PpaGPCATH1	80	-----IDFCYYANAFVVFV-----ML-LF-----Y-----P-T--ND---KLFLLCFSESEGP
PpaGPCATH2	79	-----FDFCYYANTFVVV-----ML-LY-----F-----P-K--ND---KLFLLVCFSESEGP
SmoGPCATH1	98	-----IDFCYYANVFMA-----ML-LV-----F-----P-N--NE---KLFMVCFSSESEGP
SmoGPCATH2	80	-----FDFCYYANAFMA-----ML-LY-----F-----P-K--NE---KLFMVCFAFSESEGP
ZmaGPCATH1	108	-----IDFCYYANTFLLV-----ML-LF-----Y-----P-K--DE---KLFMVCFSSESEGP
ZmaGPCATH2	105	-----IDFCYYANTFLLV-----ML-LF-----Y-----P-E--DE---KLFMVCFSSESEGP
OsaGPCATH	118	ALQFQAIDCIDFCYYANTFLLV-----ML-LF-----Y-----P-K--DE---KLFMVCFSSESEGP
MacGPCATH1	119	-----IDFCYYANTFLLV-----ML-LF-----F-----P-K--NE---KLFMVCFSSESEGP
StuGPCATH1	111	-----IDFCYYANTFLLV-----ML-LF-----F-----P-T--KE---KLFMVCFSSESEGP
StuGPCATH2	105	-----IDFCYYANTFLLV-----ML-LG-----F-----P-T--NE---TFFMVCFSSESEGP
RcoGPCAT	123	-----IDFCYYANTFLLV-----DL-LL-----Y-----P-K--DE---KLFMVCFSSESEGP
AthGPCAT	120	-----IDFCYYANTFLLV-----DL-LL-----Y-----P-K--NE---KLFMVCFSSESEGP
BvuGPCATH	175	-----IDFCYYANTFLLV-----AL-LM-----Y-----P-K--NE---KLFMVCFSSESEGP
NnuGPCATH	117	-----IDFCYYANTFLLV-----ML-IL-----Y-----P-R--NE---KFFMVCFSSESEGP
EguGPCATH	123	-----IDFCYYANTFLLV-----VL-LF-----F-----P-K--NE---KLFMVCFSSESEGP
NadGPCATH	135	-----IDFCYYANTFLLV-----ML-LL-----Y-----P-R--NE---KLFMVCFSSESEGP
AtrGPCATH	104	-----IDFCYYANTFLLV-----ML-LL-----Y-----P-K--NE---KLFMVCFSSESEGP
LtuGPCATH	154	-----IDFCYYANTFLLV-----ML-LF-----Y-----P-R--NE---KLFMVCFSSESEGP
AfiGPCATH	156	-----IDFCYYANTFLLV-----ML-LF-----Y-----P-R--NE---KLFMVCFSSESEGP
KflGPCATH	141	-----IDFCYYANAFVVF-----L-LV-----Q-----P-D--NE---FLFMICFAFSESEGP
EmuGPCATH	88	-----IGPCYLVNVYTL-----F-LWA-----V-----P-D--SR---PVFLVLEGLANST
OviGPCATH	70	-----IDTCYLINLISL-----L-LWA-----H-----P-T--CH---KLFQVQGLANAH
SpuGPCATH	132	-----ADFCYYANAFSL-----F-LWA-----A-----P-Y--KS---NYFMVVEGVVNGP
CgiGPCATH	141	-----IDFCYYANIFWYS-----F-LWV-----V-----P-Y--NE---KFFAVVEALSNGP
LgiGPCATH1	118	-----IDFCYYGNVWYL-----F-LWI-----W-----P-Q--HG---NIFMASFAVANGP
LgiGPCATH2	144	-----IDFCYYGNIFWFA-----F-LWI-----A-----P-H--HG---DLFVAFALANGP
LcoGPCATH	186	-----FDLCYFVNAMTLL-----F-LWV-----F-----P-G--SR---AMFIATYCLTNGP
MciGPCATH1	189	-----FDICYFVNAMTLL-----F-LWA-----Y-----P-T--SK---LLFVASFMTNGP
MciGPCATH2	184	-----FDLCYFVNLTLL-----F-LWV-----F-----P-S--ST---LLYTAATFTLNGP
MveGPCATH	216	-----IDMCYFVNNAVLL-----Y-LYV-----F-----P-Q--SQ---ALLGAVLLSLGP
UmaGPCATH	211	-----FDLCYFVLNVALL-----F-LWV-----F-----P-S--SE---FLFTVVCYCAHGP
LbiGPCATH	142	-----FDLCYFVTINNFV-----Y-LWL-----F-----P-S--SP---ALFVACYCLSHCA
SpoGPCATH	165	-----ADFCYFVGNILLLV-----Y-LWI-----F-----P-E--SR---FLFLLSYSSISYGT
SceGPCAT	166	-----ADFCYFVNMLCCL-----F-LWI-----F-----P-Y--SY---SLFQSCFAFTFTCT
LthGPCATH	161	-----ADLCYFVNLMVLL-----F-VWV-----W-----P-D--SV---FLYQSCFAFTFTGT
PkuGPCATH	168	-----ADLCYFVNYLTL-----Y-LWI-----L-----P-D--SQ---SLYVACCSFSWGS
NcrGPCATH	238	-----ADLCYFVNFLLL-----S-LWV-----F-----P-G--SK---FLFTAVYCLAYGN
ChaGPCATH	226	-----ADLCYFVNFLLS-----S-LWA-----F-----P-R--SK---FLFVSAYCLAYGN
AclGPCATH	226	-----ADLCYFVNMLCML-----S-LWV-----F-----P-R--SK---FLFVSTFCLTFGN
CglGPCATH	233	-----ADLCYFVNFLML-----S-LWA-----F-----P-N--SK---FLFTAVYCLAFGN
BbaGPCATH	230	-----ADLCYFVNVLML-----S-LWA-----F-----P-S--SK---FLFTAFAFCLAFGN
DdiGPCATH1	318	-----IDFCYLCQIILLF-----Y-LFG-----HQYILGEK-V--LP---NLFKLVFALSNGP
DdiGPCATH2	157	-----FDFCYFANLMLLF-----F-LYG-----L-----P-D--RQ---WYFLVTCFCLANGP
AsuGPCATH	154	-----EDFCYFANFLLW-----Y-T-HGPE-----YV-----R-S--ST---VYFQLVFCFCNGP
EdiGPCATH	105	-----IDFCYFCQITLL-----C-LPS-----L-----N-TEYIS---FLFQVAFVESHGP
AcaGPCATH1	154	-----FDFCYYANIFVLL-----Y-LHF-----F-----P-K--SE---TMFICFAFCTGP
AcaGPCATH2	187	-----IDFCYFCQIILLF-----F-LYW-----H-----S-S--NP---IYFQVFECLSNGP
PhrGPCATH	220	-----IDFCYFVTGIVYM-----Y-LYL-----Y-----P-K--SS---TLFVHVFAFANGP
NgrGPCATH1	190	-----IDFCYFANFLLF-----Y-LGF-----Y-----T-N--SP---FLFLLNFVNSSGP
NgrGPCATH2	144	-----IDFCYFNLALMF-----F-LHF-----F-----P-T--SP---FLFVITFANNCGP
MbrGPCATH	166	-----ADFCYFNLWVTL-----C-LLL-----P-----SVF--GA---TARRSEVMCHGP
BsaGPCATH	152	-----YDFCYFANGTLL-----Y-CWV-----C-----P-W--SD---VYFQLVFCVCSNGP
OtaGPCATH	134	-----YDFCYFANVSVM-----F-LWT-----R-----R-R--SA---TAFRCIFAFANGP
MpuGPCATH1	159	-----IDFCYFANAFVFA-----Q-LV-----N-----P-G--SA---FLFRTTFAYATGP
MpuGPCATH2	100	-----IDLCHVLGFFYLV-----I-AFAGLLASASPHLHAIFAP-----T-S--SK---LFFVAFGIANGP
MspGPCATH1	144	-----IDFCYYANLACFV-----H-LG-----A-----P-T--SG---FLFRSVFAFANGP
MspGPCATH2	92	-----IDLCHVLSFAYSIVGSISLAAWAI-----GLKEGLAPV-----T-S--SP---YVFRACFCLANGP
CreGPCATH	114	-----YDFCYFANLMLV-----Q-LWL-----L-----P-R--WA---PLA-----
CowGPCATH	444	-----IDFCYFACCLLF-----Y-LFF-----Y-----P-T--SG---QLFEVFESSMNGP
CmeGPCATH	138	-----YDYCYFVQALVVA-----F-Y-FG-----W-----P-R--TQAWRQVVFVAFASANGP

Cloning and characterization of GPCATs

PpaGPCATh1	115	LAWAIVWRGSLVFS	SSIDK	ISVLIHLLP	----	TVFFIIRW	WDP-ITFAH	-----
PpaGPCATh2	114	LAWAIVWRGSLVFS	SSFDK	LSVLIHLLP	----	IVFYIIRW	WDP-ASFSH	-----
SmoGPCATh1	133	LAWAIVWRGSLVFS	SSIDK	ISVLIHLLP	----	KFHSIVFFI	IRWDP-ITYPH	-----
SmoGPCATh2	115	LAWAIIWRGSLVFS	SSIDK	ISVLIHLLP	----	IVFFIIRW	WDP-STFSL	-----
ZmaGPCATh1	143	LAWAIVWRGSLVFS	GSFDK	LSVLIHLLP	----	IVLETIRW	WDP-QTFAA	-----
ZmaGPCATh2	140	LAWAIVWRGSLVFS	SSFDK	LSVLIHLLP	----	IVLETIRW	WNP-QTFAA	-----
OsaGPCATh	162	LAWAIVWRGSLVFS	SSFDK	LSVLIHLLP	----	IVLETIRW	WNP-QTFAA	-----
MacGPCATh1	154	LAWAIVWRNSLVFS	SSLDK	LSVLIHLLP	----	IVFFTIRW	WNP-ETFAA	-----
StuGPCATh1	146	LAWAIVWRGSLVFS	SSVDK	LSVLIHLLP	----	IVFETIRW	WDP-EFFGA	-----
StuGPCATh2	140	LAWAIVWRGSLVFS	SSVDK	TVSVFIHLLP	----	IVFETIRW	WDP-VFFGA	-----
RcoGPCAT	158	LAWAIVWRGSLVFS	SSVDK	LSVLIHLLP	----	IVFETIRW	WNP-ATFEA	-----
AthGPCAT	155	LAWAIVWRGSLVFS	SSPDK	LSVLIHLLP	----	IVFETIRW	WNP-ATFAA	-----
BvuGPCATh	210	LAWAIVWRGSLVFS	NSIDK	TSVLIHLLP	----	IVFETIRW	WNP-STFAA	-----
NnuGPCATh	152	LAWAIVWRGSLVFS	SSPDK	LSVLIHLLP	----	IVFETIRW	WDP-ATFEA	-----
EguGPCATh	158	LAWAIVWRGSLVFS	SSLDK	LSVLIHLLP	----	IVFETIRW	WNP-ETFAA	-----
NadGPCATh	170	LAWAIVWRGSLVFS	SSVDK	LSVLIHLLP	----	IVFETIRW	WDP-STVAD	-----
AtrGPCATh	139	LAWAIVWRGSLVFS	SSLDK	LSVLIHLLP	----	IVFETIRW	WDP-TRLGD	-----
LtuGPCATh	189	LAWAIVWRGSLVFS	SSVDK	LSVLIHLLP	----	IVFETIRW	WDP-VTFAA	-----
NguGPCATh	191	LAWAIVWRGSLVFS	SSVDK	LSVLIHLLP	----	IVFETIRW	WDP-ATFAA	-----
KflGPCATh	176	LAWAIVWRGSLVFS	SSLDK	LSVLIHLLP	----	IVLFIIRW	WDP-SNFSH	-----
EmuGPCATh	123	VYSAIILFRNS	EVFNHYDK	TSVFIHLLP	----	LLSYCIRW	FPK-NSSNP	-----
OviGPCATh	105	AYSQAFILFRNAL	VLDFOK	VSSFIHLLP	----	LLSYLIRW	FPS-ETSTL	-----
SpuGPCATh	167	VTWASIVLFRN	CLVHSIER	TCLIIHLLP	----	LLTYLIRW	YPT-ETSQH	-----
CgiGPCATh	177	LIWAMVVYRNS	LVFHS	TDKVTSAVTHLLP	----	LLSFGIRW	YRE-EVSKY	-----
LgiGPCATh1	153	LLWAAVIFRNS	LVLSV	VDKTVSVYIHTLPA	----	FLSFAIRW	YSE-DTSMK	-----
LgiGPCATh2	179	LLWAMVYRNS	LVLSH	TDKTVSVYIHTLPA	----	FLSFEVIRW	YPE-ESSSK	-----
MciGPCATh1	224	VAAWAVITWRNS	LVFHS	LDKTVSVFIHFFP	----	IVMYCIRW	MPE-LQRDIYC	-----DNTPFVTSYRNE
MciGPCATh2	219	VLWAVITWRNS	LVFHS	LDKTVSVFIHFFP	----	IVTYTIRW	ETV-LHGDPDQ	-----A-----LVYRDQ
MveGPCATh	251	LAFSAITWRNS	LVLSH	LDKTVSVYIHMSP	----	FLTYTIRW	LYP-DPE-H	-----A
UmaGPCATh	246	LAFSAITWRNS	LVFHS	LEKTVSFIHLYP	----	FVFTTVH	FEMP-HDEAV	-----S
LbiGPCATh	177	LASAITWRNS	LVFHD	QDKTVSEFIHLYP	----	FTFTVIR	HEY-NAE	-----A
SpoGPCATh	200	LAWSVAVAWRNS	LVFHS	LDKTVSFIHFFP	----	IVLHTIV	HLLTN-KSYLK	-----D
SceGPCAT	201	LCEAVITWRNS	LVLSH	TDKTVSFIHLLP	----	CVMYVIR	YHGLP-LEYKI	-----E
LthGPCATh	196	VTWASITWRNS	LVLSH	VDKTVSFIHMP	----	IVTMYI	HHGLG-EDLKR	-----A
PhrGPCATh	203	LSEAVITWRNS	LVLSH	TEKTVSFIHLYP	----	IVTMYVIR	HQLS-HEYKL	-----E
NcrGPCATh	273	NAIAIIMWRNS	LVFHS	LDKTVSFIHMP	----	IVTLHCTIR	VHLLD-PAEQR	-----K
CbaGPCATh	261	NAVAIVMWRNS	LVFHS	LDKTVSFIHMPP	----	IVALHCTIR	VHLLTP-DDYLR	-----E
Ac1GPCATh	261	NAVAIVMWRNS	LVFHS	MDKTVSFIHMP	----	IVTLHCTIR	VHLLTP-SKVLR	-----E
Cg1GPCATh	268	NAVAIVMWRNS	LVFHS	FDKTVSFIHMPC	----	IVTLHCTIR	VHLID-PAAQQ	-----A
BbaGPCATh	265	NAVAIVMWRNS	LVFHS	FDKTVSFIHMPC	----	IVTLHCTIR	VHLYP-EDKLO	-----G
DdiGPCATh1	359	LAWETIVWRNS	LVFHD	VDKTVSVFIHLLCP	----	IVTYCIRW	FNP-VYPEQ	-----
DdiGPCATh2	192	LLGAVIPFRNS	LVFHS	LDKATSVIHFFP	----	IVTYSLIRW	NDE-FLFSK	-----
AsuGPCATh	192	LMWAVAVWRNS	LVFHS	LDKATSVIHLLP	----	IVTASIRW	HILPATHPI	-----
EdiGPCATh	142	LLTAVPMWRNS	LVFHD	DDRTSVYIHLFP	----	IVLYCIRW	VVL-TTVPQ	-----
AcaGPCATh1	189	LAWSVAVWRNS	LVFHS	LDKTVSFIHLLTPN	----	IVLYAIRW	LDE-DPDSL	-----A
AcaGPCATh2	222	LAGAVVMWRNS	LVFHD	DDKTVSFIHLYP	----	IVTYCIRW	HHS-DSFVV	-----
PhrGPCATh	255	LAWAVPLWRNS	LVFHD	DDKTVSCAIIHLLP	----	IVTFQIRW	FPS-DSSLSICGD	-----
NgrGPCATh1	225	LATAIVMWRNS	LVFHS	LEKTVSVFIHFFP	----	IVTFAIRW	LMF-NDNTQ	-----
NgrGPCATh2	179	ILLAVVWRNS	LVFHD	DFDKTVSFIHFFP	----	IVSFAIRW	YPE--QFPE	-----
MbrGPCATh	202	LATAIVAVWRNS	LVFHS	LDKTVSVYIHLPA	----	IVSFSWR	WDEL-TECHA	-----
BsaGPCATh	187	LAWAVLAFS	QLFHS	APHVTSVFIHTEM	----	IVTFAIRW	LLP-DEHKA	-----
OtaGPCATh	168	VLLAVPLWRNS	LVFHS	VDKTVSFIHLLCP	----	IVMTWCGR	WYGF-GDENRLSVA	-----
MpuGPCATh1	194	ILLAVPLWRNS	LVFHS	HDKTVSFIHTAP	----	IVTYCARW	YGO-DASPLVAIGAWSERV	-----LGT
MpuGPCATh2	151	LAWAVPTGNAL	VLSH	SPKHAVALIHLSP	----	IVTWAVIRW	HAD-AHERA	-----
MspGPCATh1	179	ILLAVPLWRNS	LVFHS	HDKTVSFIHTAP	----	IVTWCTR	WYGO-GDLTWSWWGL-GDATGWLFAARE	-----GDHGQA
MspGPCATh2	143	LGWAVPALANAL	VLSH	VEHTAAFIHISPP	----	IVVMWAVIRW	HAA-EHEAA	-----
CreGPCATh	139	---KVGAGAGS	AVRRAPHQ	TVSFIHLLP	----	IVSWSIRW	YGDPAARFGP	-----
CowGPCATh	479	LAWSVITWNS	EVFHD	LDKTVSSAIIHLLP	----	IVTYSLIRW	YTP-LEGQA	-----
CmeGPCATh	177	VLWAVAVWRNS	LVFHS	LDKTVSVFIHFFP	----	IVTYNARW	HSD-LIISRRI	-----G-----ICSHE-

End of DUF2838 |

Cloning and characterization of GPCATs

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PpaGPCATH1 161 H-----AVDDT-G---PWPAPW--LLEN-DRQ-LWTLWLVVPE-LIAYSWQILYLLVNVN---
PpaGPCATH2 160 H-----AIDDT-G---PWPAYP--LLEN-NQQ-LWTLWLVVPE-LIAYTWQILYLLVNVN---
SmoGPCATH1 183 H-----SLDAT-G---PWPAPW--LVHD-NRA-LWTLWLVVPE-LIAYSWQALYLLIVNVN---
SmoGPCATH2 161 H-----STELT-G---PWPAPW--LVQT-RTD-LWTLWLVVPE-LIAYTWQILYFLIVNVN---
ZmaGPCATH1 189 M-----HPEGR-D---ARVTWP--YVED-KSY-MWTLWLVVPE-LIAYTWQILYFLIVNVN---
ZmaGPCATH2 186 M-----HPEGR-D---ARVTWP--YVED-KSY-LWTLWLVVPE-LIAYTWQILYFLIVNVN---
OsaGPCATH 208 M-----HPEGR-A---ARVTWP--YVGD-KSY-LWTLWLVVPE-LIAYTWQILYFLIVNVN---
MacGPCATH1 200 M-----HPAGR-A---ARVSWP--YVED-KSY-LWTLWLVVPE-LIAYTWQILYFLIVDVN---
StuGPCATH1 192 M-----HPEGT-P---ERASWP--YVES-QSY-LWTLWLVVPE-LIAYFWQILYKLVVDV---
StuGPCATH2 186 M-----HPEGT-P---ERASWP--YVEN-QSY-LWTLWLVVPE-LIAYFFWQILYKLVIVEV---
RcoGPCATH 204 M-----HPEGT-S---GRSSWP--YVED-KSF-LWTLWLVVPE-LIAYFWQILYKLVIVNVN---
AthGPCATH 201 M-----HPVGT-D---RRVSWP--YVED-KAY-LWTLWLVVPE-LIAYTWQILYFLIVNVN---
BvuGPCATH 256 M-----HPEGT-A---RRVSWP--YIED-TSY-LWTLWLVVPE-LIAYTWQILYFLIVNVN---
NnuGPCATH 198 M-----HSDGNAG---RRASWP--YVEN-KSY-LWTLWLVVPE-LIAYTWQILYFLIVNVN---
EguGPCATH 204 M-----HPEGR-A---ARVFWP--YVED-KSY-LWTLWLVVPE-LIAYTWQILYFLIVNVN---
NadGPCATH 216 I-----HLEQKGG---HRTAWP--HVAD-KTT-LWTLWLVVPE-LIAYTWQILYFLIVNVN---
AtrGPCATH 185 V-----DAGQQAG---VKAAMP--SVED-NAE-LWTLWLVVPE-LIAYTWQILYFLIVNVN---
LtuGPCATH 235 M-----HPEGKAS---DRASWP--YVED-KSY-LWTLWLVVPE-LIAYTWQILYFLIVDVN---
AfiGPCATH 237 M-----HPTGEAG---GRSSWP--YVED-KAY-LWTLWLVVPE-LIAYTWQILYKLVIVNVN---
KflGPCATH 222 H-----SPDEF-GPWGHGHTFPW--VVED-TRK-VWTLWLVVPE-LIAYSWQALYKLVIVEV---
EmuGPCATH 169 WWKGFVN-----T-----GGSTNV--SVFN-PMD-DWTLWLVVPE-LIAYTWQILYKLVIVNVN---
OviGPCATH 151 WYTAFS-----T-----GAKPNI--YLWI-HDL-DWTLWLVVPE-LIAYTWQILYKLVIVNVN---
SpuGPCATH 213 WYQDFAN-----T-----EMA-TGW-ELWLVVPE-LIAYTWQILYKLVIVNVN---
CgiGPCATH 223 WFDEF-----V-----PEM-LEL-SPIWLVVPE-LIAYTWQILYKLVIVNVN---
LgiGPCATH1 199 WRKDF-----V-----PDI-NNW-SVQWLVVPE-LIAYTWQILYKLVIVNVN---
LgiGPCATH2 225 WPCHF-----I-----PLL-SEW-SIWLWLVVPE-LIAYTWQILYKLVIVNVN---
LcoGPCATH 273 RFPQVTN-----D-ITM-SKEAMLLS-TIAYLWQALYKLVIVNVN---
MciGPCATH1 284 RFPALDN-----L-ELP-SKDVMIYS-TIAYGLWQSLYKLVIVNVN---
MciGPCATH2 274 HFPALSK-----M-EVL-GWYTFVYS-TGFYLAQIFVCFVNVN---
MveGPCATH 297 RYPALKN-----M-EVL-PTAQSITAA-LIAYLWQILYKLVIVNVN---
UmaGPCATH 293 RFPALKN-----L-ITL-NGYTSWFEN-TIAYLWQILYKLVIVNVN---
LbiGPCATH 222 RFPALRE-----L-EYL-HPLRALLS-AGIYVWQFLYKLVIVNVN---
SpoGPCATH 247 RFPALVK-----V-KKI-DLLSSVEHA-SFFYALWQILYKLVIVNVN---
SceGPCATH 248 RFPALVK-----Q-SEL-DIKKNIWT-SIYLLWQSLYKLVIVNVN---
LthGPCATH 243 RFPALVK-----A-----G-----S-KKW-NLKNFWT-SIYLLWQSLYKLVIVNVN---
PkuGPCATH 250 RFPALVK-----L-QQW-DIMYGIIQT-SIYLLWQSLYKLVIVNVN---
NcrGPCATH 320 RFPALVK-----I-----KTSP--GS-PTAYA-NVISMLAWS-TIAYLWQILYKLVIVNVN---
CbaGPCATH 308 RFPALVK-----I-----KYSSK--GS-P-GHY-ELAAMLWA-TIAYLWQILYKLVIVNVN---
AclGPCATH 308 RFPALVK-----I-----KFSKP--GS-P-NHY-SLLSMLAWS-TIAYLWQILYKLVIVNVN---
CglGPCATH 315 RFPALVK-----I-----KTSPA--GS-STAYA-NVFSMLAWS-SIYLLWQILYKLVIVNVN---
BbaGPCATH 312 RFPALVK-----I-----KTAKP--GS-ATAYA-NIVSMLAWS-TIAYLWQILYKLVIVNVN---
DdiGPCATH1 405 LA-----CGGVE-CVL-SKEKFFVPE-LIAYTWQILYKLVIVNVN---
DdiGPCATH2 238 -Q---NQ-----I-EEP-NKNFYIYPE-LIAYTWQILYKLVIVNVN---
AsuGPCATH 239 -D---KE-----S-LTA-TTETFWPE-LIAYTWQILYKLVIVNVN---
EdiGPCATH 188 -----L-TIMNGLIIP-SFLYIHWQILYKLVIVNVN---
AcaGPCATH1 236 TYET-LR-----G-KDA-TTQMVFLPE-LIAYTWQILYKLVIVNVN---
AcaGPCATH2 268 CR---AD-CSI-SITASFVPE-LIAYTWQILYKLVIVNVN---
PbrGPCATH 305 -----EWP--S--C-GTV-ALKDAIYE-LIAYTWQILYKLVIVNVN---
NgrGPCATH1 271 TPKIITP-----I-----GESYSICVNED-CYV-SLVDIYVPH-CLFIWQILYKLVIVNVN---
NgrGPCATH2 224 YS-----VCIREN-CEV-TIYFVIVPH-LIAYTWQILYKLVIVNVN---
MbrGPCATH 248 -NPE-LE-----M-CQL-SLMDLVVPE-LIAYTWQILYKLVIVNVN---
BsaGPCATH 233 NFTIASA-----E-----AP----IYSDP-LNV-STFDLWKSVPYFYGWWVLYVWVNVN---
OtaGPCATH 219 -----DLD--SDVT-VLQ-GSLDLVYVPE-LIAYTWQILYKLVIVNVN---
MpuGPCATH1 253 RLPGLLG----GNA-NRLGVA-----ELSTP--M-TS-TRE-MLTEFYVPE-LIAYTWQILYKLVIVNVN---
MpuGPCATH2 197 F-PGVFG-----VPHANS-G-D---AKRGTS--Y--A-ESV-TTMDLVVPE-LIAYTWQILYKLVIVNVN---
MspGPCATH1 248 VAPGLMYADVAVHRAPNRLT-----VEQLT--MSYD-VVD-VVREFYIYPE-LIAYTWQILYKLVIVNVN---
MspGPCATH2 189 F-PGILP-----MPL-----DAA--N--G-SGL-CIVELTPA-LIAYTWQILYKLVIVNVN---
CreGPCATH 183 PPEATEQ-----E-----RE----RW--H-RA--GGLSNVLE-LIAYTWQILYKLVIVNVN---
CowGPCATH 525 SVCSQDD-----C-SIP-SVWSWSSP-LIAYTWQILYKLVIVNVN---
CmeGPCATH 231 -----AD-----C-IALGRRAITVPE-LIAYTWQILYKLVIVNVN---

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Cloning and characterization of GPCATs

PpaGPCATH1	207	--LHQRILKDPEVM	TSF	RELSRKA	----	SR--A--NN--	IWWRL	SGIL	-----	G	D	
PpaGPCATH2	206	--LRRQRLNDPEVM	TSF	SRELSRKA	----	QR--S--NN--	IWWRL	SGVL	-----	G	D	
SmoGPCATH1	229	--LRRQRLLRDPEVM	TSF	RELSRKA	----	AR--A--NN--	IWWWL	SGIL	-----	G	E	
SmoGPCATH2	207	--LRRQRLNDPEIM	TSY	RELSRKA	----	SR--A--NN--	IWWRL	SGIL	-----	G	A	
ZmaGPCATH1	235	--LRRQRLLRDPEVM	TSY	RELSKKA	----	QK--A--NN--	IWWRL	SGIL	-----	G	D	
ZmaGPCATH2	232	--LRRQRLLRDPEVM	TSY	RELSKKA	----	QK--A--NN--	IWWRL	SGIL	-----	G	D	
OsaGPCATH	254	--LRRQRLKDPEVM	TSY	RELSKKA	----	QK--A--NN--	IWWRL	SGIL	-----	G	D	
MacGPCATH1	246	--LRRQRLQDPEVM	TSY	RELSKKA	----	QK--A--NN--	IWWRL	SGIL	-----	G	D	
StuGPCATH1	238	--LRRQRFQDPEVM	TSY	RELSKKA	----	KK--A--NN--	VWWRV	SGIL	-----	G	D	
StuGPCATH2	232	--LRRQRLLRDPEVM	TSY	RELSKKA	----	KK--A--NN--	MWWRV	SGIL	-----	G	D	
RcoGPCATH	250	--LRRQRLLRDPEVM	TSY	RELSKKA	----	QK--A--NN--	VWWRV	SGIL	-----	G	D	
AthGPCATH	247	--LRRQRLLRDPEVM	TSY	RELSKKA	----	EK--A--NN--	KLWQL	SGIL	-----	G	D	
BvuGPCATH	302	--LRRQRLLRDPEVM	TSY	RELSKKA	----	QK--A--NN--	IWWRL	SGIL	-----	G	D	
NnuGPCATH	245	--LRRQRLLRDPEVM	TSY	RELSKKA	----	QK--A--NN--	IWWRL	SGIL	-----	G	D	
EguGPCATH	250	--LRRQRLLRDPEVM	TSY	RELSKKA	----	QK--A--NN--	IWWRL	SGIL	-----	G	D	
NadGPCATH	263	--LRRQRLLRDPEVM	TSY	RELSKKA	----	KK--A--NN--	VWWRV	SGVL	-----	G	D	
AtrGPCATH	232	--LRRQRLLRDPEVM	TSY	RELSKKA	----	QK--A--NN--	IWWRL	SGVL	-----	G	D	
LtuGPCATH	282	--LRRQRLLRDPEVM	TSY	RELSKKA	----	KK--A--NN--	IWWRL	SGIL	-----	G	D	
AfiGPCATH	284	--LRRQRLLRDPEVM	TSF	RELSKKA	----	QK--A--NN--	VWWRV	SGIL	-----	G	D	
KflGPCATH	272	--LRRQRLLRDPOVIL	TSF	RELSRKA	----	ER--A--NN--	LFWWASGLL		-----	G	D	
EmuGPCATH	216	--LTKP---N-EKYN	DNY	RYLRTKY	----	FS--K--VD--	FFKR---	I---	-----	P	K	
OviGPCATH	198	--CVKP---S-EKH	DSY	RYMHKKK	----	LL--A--HL--	IWNR---	F---	-----	H	A	
SpuGPCATH	256	--ALGSTFSS-SEYL	NNY	TFIAYQS	----		NN---	VFSRAIRCK	-----	G	G	
CgiGPCATH	261	--MAKP---S-EEYV	TSF	NYLGKKQ	----		DS---	CLYKTFNVC	-----	G	E	
LgiGPCATH1	237	--FP--P---P-EEY	TSY	KCLTAQE	----		SS---	FMFAVDLIL	-----	G	P	
LgiGPCATH2	263	--CP--P---P-EHYQ	TSY	RYLTSST	----		ST---	FIYRAINLC	-----	G	P	
LcoGPCATH	310	--RRREKVES-GLRW	TSY	SWLLDDS	----	HG--R--KG--		LIQKAAFVF	-----	G	A	
MciGPCATH1	321	--GRRDKVES-GLRW	TSY	SWLLDDT	----	HG--K--KG--		FIQKMAFMF	-----	G	P	
MciGPCATH2	311	--AKRDKVES-GSRT	TSY	TTLNLS	----	PENKN--KS--		FILATASLF	-----	G	E	
MveGPCATH	334	--RWREKVES-GKRW	TSY	TWLLNDP	----	K-----SG--		MIAKTAHTF	-----	G	E	
UmaGPCATH	330	--RKKSKIET-GERL	NSY	STMSXGK	----	G-----		PVANLKGKA	-----	P	P	
LbiGPCATH	259	--DRRAKIQS-GQRT	TSF	SWLLNDK	----	Q-----G--		VIGRSLAST	-----	P	P	
SpoGPCATH	284	--GKQKIQE-G-RP	TSF	TWLSKAY	----	S-----KT--		KLGRAVAKL	-----	P	Q	
SceGPCATH	285	--KSSKIKS-GERL	TSF	EYLTHQ	----	F-----KN--		FW---AVKL	-----	R	S	
LthGPCATH	282	--RSSKIKS-QORA	TSF	EYLTHQ	----	F-----KN--		FW---AAKL	-----	P	E	
PkuGPCATH	287	--RADKIKK-G-KV	TSF	EYLRSMS	----	N-----TH--		KFVRFINSL	-----	P	D	
NcrGPCATH	366	--RRRDKIAA-G-RP	TSF	TWLRKSY	----	S-----KN--		WLGKAILNL	-----	P	D	
CbaGPCATH	353	--RAEKIAA-G-RP	TSF	TWLRKSY	----	A-----KT--		WIGKIVNSL	-----	P	E	
AclGPCATH	353	--RAEKIAA-G-RP	TSF	TWLRKSY	----	A-----KT--		WIGKIVINL	-----	P	E	
CglGPCATH	361	--RRREKIAA-G-RP	TSF	TWLRKSY	----	S-----KV--		WIGKIVLGL	-----	P	S	
BbaGPCATH	358	--LRRKIAA-G-RP	TSF	TWLRKSY	----	A-----RT--		WIGKIVLSM	-----	P	S	
DdiGPCATH1	441	--VDSDKTANDKELM	TSS	RWMSVQ	----	P-----H--		PIYRWQCKN	-----	G	Y	
DdiGPCATH2	271	--FKKDTLKE-NNYT	TSL	SFLSTIK	----	EG-EK--GS--		LISRYLNFA	-----	K	P	
AsuGPCATH	271	--LQNTLRE-RNYT	TTL	NELNLQ	----	S-----NS--		IYSRLNLI	-----	P	T	
EdiGPCATH	215	--FKKETLKK-NHYT	TSL	IWLSQEH	----	P-----H--		PVYLYLLKK	-----	G	F	
AcaGPCATH1	272	--TSAKKVEE-QDYQ	TTF	RWFSSMD	----	K-----GG--		IGKLIKKA	-----	S	P	
AcaGPCATH2	302	--MDKKLDSANIM	TSL	RWFIRED	----	K-----PH--		PIYKALLVK	-----	G	I	
PhrGPCATH	342	RLSFPVSVADDDLQ	TSF	KWLLSAN	----	E-----GP--		VFRVLRQKF	-----	PT	L	
NgrGPCATH1	319	--LDREYLQKQNNIQ	TSY	NLLISK	----	A-----F--		IQTLSDKLY	-----	G	K	
NgrGPCATH2	261	--LDNLIKERSHVI	TSF	RWTEMD	----	K-----KS--		KSYKLGQKF	-----	S	A	
MbrGPCATH	283	--NADHRTDQALE	TSL	RWLASP	----	GG--M--AAVAKTFC		RAVGLM	-----	K	PDENF	
BsaGPCATH	278	--MGPRKRR-GYK	TLE	DRVSEGG	----	P-----T--		KFLA-KISK	-----	N	D	
OtaGPCATH	255	--VDAKLDTPALM	TSL	RWLASDR	----	K-----SA--		ANRFAKLTMRGIGLFGTDEFFDPK	-----	T	---	
MpuGPCATH1	306	--HDVKRLDEDPELV	TSL	RWLAGDY	----	K-----LA--		VTRSTLRTRRRLGFKVDELFDAS	-----	S	---	
MpuGPCATH2	248	--FGRHKGAKKTAVYGT	TPR	AELVSDTVYHYTMRT	----	--VKPFGD--		VCGYEV	-----	---	ARPSK	
MspGPCATH1	307	--FYVKRLDNDPELV	TSL	RWLANDY	----	K-----AP--		VTRWALRTSRRVGLFAQDELFDHN	-----	S	---	
MspGPCATH2	232	--HGRYHG-HSNTSG	SSH	DTVFM	----	LR--G--NK--		VLSRWVGYD	-----	---	NDRPAD	
CreGPCATH	224	--FSRAKIQQR-GYR	TLE	NYVTSQK	----	K-----G--		TFHARR	-----	P	P	
CowGPCATH	562	--VDHAKIAANVR	K	TSY	RWLSSSR	----	G-----H--		ISYRMNKF	-----	D	R
CmeGPCATH	264	--LSRKLDRNPELIT	TSL	RWLTRHS	----	G-----S--		AVSRVLRNLF	-----	G	E	

Cloning and characterization of GPCATs

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PpaGPCATH1 312 -----
PpaGPCATH2 312 -----
SmoGPCATH1 339 K-TESSPENPSKNVISIQ-----EESSGG--TAPRGLLSTNC-----NGSVSSVSS--E-
SmoGPCATH2 317 -----
ZmaGPCATH1 345 I-EQGSWTQGAPAD-GTY-----IG-----
ZmaGPCATH2 342 M-EQGSSTQGAPGG-DGG-----TLGNHH-----QHT-SEE---
OsaGPCATH 364 M-EEANSSQNAEES-QGD-----LSANG-----QHS-SEH---
MacGPCATH1 356 I-VEEPDRLSADLS-TSS-----HATSG-----EPS-G-----
StuGPCATH1 349 V-PQNDDEISLQES-ATE-----TNNTESE-----TVH-S-----
StuGPCATH2 342 V-TEQQHENPLQEN-SMK-----TP-----
RcoGPCATH 360 A-HTQQYHSEPKQD-QSP-----NSMEIR-----MKT-IHS---
AthGPCATH 358 I-EEQILHHEAVSH-PTE-----NEPKST-----E-----
BvuGPCATH 412 L-EAQPQLDSSSPL-DSP-----VSAQYI-----QQS-----
NnuGPCATH 355 L-PIQPNTVATVE-PSE-----DAKEPV-----K-IDH---
EguGPCATH 360 I-LEQPRQPVVDLS-TNH-----QHTSDT-----PDE-----
NadGPCATH 373 I-LSESTQAQKEV-AVP-----DGLEK-----SGDTNNA--QK
AtrGPCATH 342 I-TIPIDQPVESVP-MNM-----EKNECN-----SLDPPDA--EP
LtuGPCATH 392 I-PMQQQLDLSLAT-DTL-----THSD-----
AfiGPCATH 394 I-PTADQPVFDSHA-END-----YTSVGD-----YPSRVDG---
KflGPCATH 382 K-QGDAAKDTISQP-NSP-----GEVCQK-----CGS-LHSSASK
EmuGPCATH 318 P-VNSKSPTEMEDENED-----EE-----VD-----E-----
OviGPCATH 300 LEVLNPEHELLKNIDEG-----KN-----PG-----G--MCE---
SpuGPCATH 362 PE-----
CgiGPCATH -----
LgiGPCATH1 -----
LgiGPCATH2 -----
LcoGPCATH 419 S-GKQMDKSKT-----
MciGPCATH1 430 K-LKA-----
MciGPCATH2 422 E-KNQNSKILT-----KQRSLR-----R-----KQ-----QKHLDD---
MveGPCATH 441 S-AVAANQLAHDADKRD-----H-----EE-----EELHKQ---
UmaGPCATH 436 L-IRTAEKVVADAGKHK-----SG-----ME-----QEGHVE---
LbiGPCATH 366 E-STARSSGSSTPTTICS-----GP-----SE-----TDLTAM---
SpoGPCATH 390 E-TPTNSGSSSALS-----SV-----ND-----KDYLNV---
SceGPCATH 389 N-LQQKLQPDSDAVISDA-----SG-----VL-----
LthGPCATH 386 D-LQQLSQKANSSSNNY-----AT-----GI-----E-----
PkuGPCATH 393 A-MQDDIEERDKSGLSKV-----HA-----GA-----QEEENK---
NcrGPCATH 472 K-WQHSPDGMVHSPTLTP-----DS-----DV-----N-LNTA---
CbaGPCATH 459 K-WQNSPGAMFSPNGPV-----EN-----AS-----AAGARI---
AclGPCATH 459 R-WQASPAGTTSPTLLDS-----AV-----VS-----FVASPA---
CglGPCATH 467 K-WQHSPDVVPHEDGSGT-----QT-----ES-----VVIDPQ---
BbaGPCATH 464 K-WQTSPELMLTSPLMQP-----KF-----LA-----L-FLTS---
DdiGPCATH1 549 N-KDQPLKITSIK--AFL-----SSTNQST-----DHYQQLTPPNLKHQKGHET-----
DdiGPCATH2 381 K-IIMSSIGNTTSFNSNCDSPIMEYSKQQQSNTLNSTHHKKEKPK-----SSTNQST-----
AsuGPCATH 377 K-TIVSSFGAGDMSPMQ-----ST-----SP-----N-ALSE---
EdiGPCATH 322 R-SESNTSIQTPKILDER-----DAEKSQ-----
AcaGPCATH1 378 R-LRRGSLALQAINIGD-----PT-----KN-----S-FLSF---
AcaGPCATH2 411 S-EAKEKAKEKAKKGL-----
PhrGPCATH 452 R-ELSAVSQPDVYHGS-----
NgrGPCATH1 426 M-ENKEQTSAITQ-----
NgrGPCATH2 369 H-PPQTNQK--Q-----EEQAS-KDAQ-E-----QR-----
MbrGPCATH 404 V-AQEAKIEAIAELA-----GA-----KK-----D-----
BsaGPCATH 383 S-QGADGTTPLLS--SPR-----MD-----FT-----DSEIE---
OtaGPCATH 377 C-QMVNELTESELETMQT-----RK-----NT-----PPEGD---
MpuGPCATH1 428 L-RHEDQQDGIIVTEVTG-----K-----
MpuGPCATH2 368 --EDGGKKK-----AG-----SG-----DLARR---
MspGPCATH1 429 V-RDDDNADAVAEAVLA-----VE-----
MspGPCATH2 345 --RDNAPSEAPKKH-EH-----A-----GK-----E-----
CreGPCATH 330 A-PAHVSASATKK--D-----
CowGPCATH 668 V-PSVVSPPA--D--V-----A-----GK-----E-----
CmeGPCATH 372 K-IETNASSNPE-----KSSM-----

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Cloning and characterization of GPCATs

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PpaGPCATH1 -----
PpaGPCATH2 -----
SmoGPCATH1 383 E-----KKA-----ICS--LCQQL-SSRKFE---LQK--
SmoGPCATH2 -----
ZmaGPCATH1 -----
ZmaGPCATH2 370 -----
OsaGPCATH 391 -----
MacGPCATH1 381 -----
StuGPCATH1 375 -----
StuGPCATH2 360 -----
RcoGPCAT 388 -----
AthGPCAT 381 -----
BvuGPCATH -----
NnuGPCATH 381 -----
EguGPCATH 385 -----
NadGPCATH 403 A-----
AtrGPCATH 373 L-----
LtuGPCATH 412 -----
AfiGPCATH 423 Q-----N-----GHP--VVDAN-ESKLVA---DAN--
KflGPCATH 414 DSLNGSLSGSGELET--NGDG-----LKG--QMDAE-----
EmuGPCATH 340 -----
OviGPCATH 326 -----S--DADVPCRLV-----SCESAHKIT--ELT---CH-----SAH
SpuGPCATH -----
CgiGPCATH -----
LgiGPCATH1 -----
LgiGPCATH2 -----
LcoGPCATH 430 -----
MciGPCATH1 434 -----
MciGPCATH2 448 -----K-----
MveGPCATH 467 -----G--REQ-----E-----EE--
UmaGPCATH 463 -----E--DRD-----EDKE-----QQAAA---AAA---AAAS--
LbiGPCATH 393 -----V--SAS-----KQPG-----PAS--DISE-----DSPT--
SpoGPCATH -----
SceGPCAT 416 -----N--RDE-----DFDD-----SS--
LthGPCATH 407 -----
PkuGPCATH 415 -----
NcrGPCATH 499 -----R--LGE-----P--TKNNNNSDSNDSSGSSTSSL----DNIP--LLTEEKSELPASATATATG-
CbaGPCATH 485 -----K--D-----DSSKGHDKRQSI---DKIP--LLEDTGQPQQN-----
AclGPCATH 486 -----L--DDS-----TSAHKECDKASL---DQIP--PLDAHAATTGS-----
CglGPCATH 494 -----A--GAS-----TTGSLADGISGKRLDDDMRGVSGV---DNIP--LLNDERPAAVL---ATGAD-
BbaGPCATH 491 -----P--KKA-----SGTPSFEPQAQTARDAMARTTSL---DRIP--LLDDYTTANST-----
DdiGPCATH1 573 -----
DdiGPCATH2 445 -----QFSDSTS-----YI---G-----HE--
AsuGPCATH 401 -----TSDNA-----I---G-----
EdiGPCATH 348 -----T-----
AcaGPCATH1 -----
AcaGPCATH2 437 -----C--S-F-----FVPALTI-----
PbrGPCATH -----
NgrGPCATH1 -----
NgrGPCATH2 -----
MbrGPCATH 430 -----APEQD-----V---Q-----
BsaGPCATH 403 -----
OtaGPCATH 403 -----
MpuGPCATH1 454 -----
MpuGPCATH2 -----
MspGPCATH1 455 -----S--G-----
MspGPCATH2 -----
CreGPCATH 343 -----
CowGPCATH 683 -----
CmeGPCATH -----

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Cloning and characterization of GPCATs

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PpaGPCATH1 -----
PpaGPCATH2 -----
SmoGPCATH1 404 -----A-----V-EEA-----TSSVRDLG-----LEPALFSRIK-----
SmoGPCATH2 -----
ZmaGPCATH1 -----
ZmaGPCATH2 370 -----
OsaGPCATH 391 -----
MacGPCATH1 381 -----
StuGPCATH1 375 -----
StuGPCATH2 360 -----
RcoGPCATH 388 -----
AthGPCATH 381 -----
BvuGPCATH -----
NnuGPCATH 381 -----
EguGPCATH 385 -----
NadGPCATH 404 -----E-----
AtrGPCATH 374 -----C-----
LtuGPCATH 412 -----
AfiGPCATH 442 -----Q-----V-D-----LLS-----
KflGPCATH 441 -----M-KLRLGNGKSEEEAKGEKVLGKNGV
EmuGPCATH 340 SLRHN----LE-----TAL-EED----RDS--D-----I-----
OviGPCATH 353 SVDSSTNVMLE----YDSMAS-KER----NGD--N-----I-----
SpuGPCATH -----
CgiGPCATH -----
LgiGPCATH1 -----
LgiGPCATH2 -----
LcoGPCATH 430 -----
MciGPCATH1 434 -----
MciGPCATH2 449 -----
MveGPCATH 474 -----GV-----VV-EDK--K--NI-----
UmaGPCATH 483 -----GL--P--Q-----AG-ENK--K--DQ-----
LbiGPCATH 412 -----SL--DPTL-----NL-DFK--K--DS-----
SpoGPCATH -----
SceGPCATH 426 -----SV-SS-----KS--D-----
LthGPCATH 407 -----E-----
PkuGPCATH 415 -----S-----
NcrGPCATH 547 -----AEV--D--GGAKD-VA-RAR--R-PGA--E-LFG--AG-----AAPS-----
CbaGPCATH 514 -----GN--A--NGGPN-SEANAR--A-TGA--E-ILYEAKGNVLDKKNVSAPL-----
AclGPCATH 517 -----DK--A--NITTD-TM-RER--R-----
CglGPCATH 542 -----MDM--D--GGARD-VA-RER--RMGSG--E-S-----R-----
BbaGPCATH 533 -----GF--D--GGARD-VA-RER--K--ST-----E-----
DdiGPCATH1 576 -YLK-----LIL-----
DdiGPCATH2 457 -----DL--A--PDTDD-VH-HHYNYS--S-----KPN-----
AsuGPCATH 408 -----EVG--S-----SPQ-----
EdiGPCATH 349 -----SDA--E-----
AcaGPCATH1 -----
AcaGPCATH2 447 -LYF-----LIERAL-AFA-----T-----
PbrGPCATH -----
NgrGPCATH1 -----
NgrGPCATH2 -----
MbrGPCATH 437 -----ELNE--ST--E-----DPF-----
BsaGPCATH 403 -----Q-----
OtaGPCATH 403 -----QILMTPV-----
MpuGPCATH1 454 -----GRR-----
MpuGPCATH2 -----
MspGPCATH1 457 -----GGD-LA-RRS-----GGG--D-L-----ARRSGGPQ-----
MspGPCATH2 -----
CreGPCATH 343 -----A-----
CowGPCATH 683 -----Q-----
CmeGPCATH -----

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PpaGPCATH1 -----
PpaGPCATH2 -----
SmoGPCATH1 427 -----PT-----D-----AI-SCAS
SmoGPCATH2 -----
ZmaGPCATH1 -----
ZmaGPCATH2 370 -----HI-----Q-----E-----
OsaGPCATH 391 -----S-----
MacGPCATH1 381 -----M-----
StuGPCATH1 375 -----E-----
StuGPCATH2 360 -----Q-----
RcoGPCATH 388 -----AE-----E-----Q-----
AthGPCATH 381 -----T-----
BvuGPCATH -----
NnuGPCATH 381 -----F-----
EguGPCATH 385 -----QN-----N-----S-----
NadGPCATH 405 -----VN-----N-----S-----
AtrGPCATH 375 -----RT-----N-----SL-----
LtuGPCATH 412 -----Y-----
AfiGPCATH 448 -----SG-----D-----E-----
KflGPCATH 466 ENGNGLHQRSIARANSCGTESGDKVSVEMNGKSSSLGLAR
EmuGPCATH 358 -----EV-----
OviGPCATH 378 -----AV-----
SpuGPCATH -----
CgiGPCATH -----
LgiGPCATH1 -----
LgiGPCATH2 -----
LcoGPCATH 430 -----A-----
MciGPCATH1 434 -----K-----
MciGPCATH2 449 -----LN-----
MveGPCATH -----
UmaGPCATH -----
LbiGPCATH -----
SpoGPCATH -----
SceGPCATH -----
LthGPCATH -----
PkuGPCATH -----
NcrGPCATH -----
CbaGPCATH 552 -----TGA-----TM
AclGPCATH -----
CglGPCATH -----
BbaGPCATH -----
DdiGPCATH1 -----
DdiGPCATH2 477 -----Q-----
AsuGPCATH 415 -----LLSV-----
EdiGPCATH -----
AcaGPCATH1 -----
AcaGPCATH2 -----
PbrGPCATH -----
NgrGPCATH1 -----
NgrGPCATH2 -----
MbrGPCATH 447 -----KL-N-----
BsaGPCATH -----
OtaGPCATH 410 -----D-----
MpuGPCATH1 457 -----E-----T-----
MpuGPCATH2 -----
MspGPCATH1 478 -----KSL-----R-----
MspGPCATH2 -----
CreGPCATH -----
CowGPCATH -----
CmeGPCATH -----

```

Appendix S1. Sequence alignment analysis of GPCATs.

The protein sequences of GPCATs were aligned using the T-Coffee PSI/TM-Coffee web-service (<http://tcoffee.org.cat/apps/tcoffee/do:tmcoffee>) and BoxShade (<http://www.ch.embnet.org>) was used for shading background according to conservation. Sequence numbering is according to the full-length protein sequences. See Fig. 3 for species abbreviations. The proteins for which catalytic GPCAT activity was determined (Fig. 7) have names highlighted in green. Sequence IDs: AcaGPCATH1, XP_004338657.1; AcaGPCATH2, XP_004368210.1; AclGPCATH, XP_001270843.1; AfiGPCATH, b3_c2279; AsuGPCATH, XP_012748160.1; AthGPCATH, NP_198396.1; AtrGPCATH, XP_011622896.1; BbaGPCATH, KGQ05851.1; BsaGPCATH, CUE73576.1; BvuGPCATH, XP_010666443.1;

CbaGPCATH, KIW96763.1; CgiGPCATH, EKC38105.1; CglGPCATH, XP_001229114.1; CmeGPCATH, XP_005537575.1; CowGPCATH, XP_004348404.1; CreGPCATH, XP_001693172.1; DdiGPCATH1, XP_636790.1; DdiGPCATH2, XP_643621.1; EdiGPCATH, XP_001738020.1; EguGPCATH, XP_010936661.1; EmuGPCATH, CDS42477.1; KflGPCATH, kfl00299_0090; LbiGPCATH, XP_001876784.1; LcoGPCATH, CDH53915.1; LgiGPCATH1, ESO91084.1; LgiGPCATH2, XP_009058350.1; LthGPCATH, XP_002553792.1; LtuGPCATH, b4_c3038; MacGPCATH, XP_009407293.1; MbrGPCATH, XP_001747618.1; MciGPCATH1, EPB88260.1; MciGPCATH2, EPB92530.1; MpuGPCATH1, XP_003062419.1; MpuGPCATH2, XP_003058481.1; MspGPCATH1, XP_002501336.1; MspGPCATH2, XP_002508227.1; MveGPCATH, KFH63290.1; NadGPCATH, b3_c6890; NcrGPCATH, XP_011394510.1; NgrGPCATH1, XP_002675713.1; NgrGPCATH2, XP_002673223.1; NnuGPCATH, XP_010273972.1; OsaGPCATH, EAY74703.1; OtaGPCATH, XP_003078006.1; OviGPCATH, XP_009166064.1; PbrGPCATH, CEO94537.1; PkuGPCATH, KGK40379.1; PpaGPCATH1, XP_001771632.1; PpaGPCATH2, XP_001756364.1; RcoGPCATH, XP_002514086.1; SceGPCATH, NP_011665.1; SmoGPCATH1, XP_002985957.1; SmoGPCATH2, XP_002979497.1; SpoGPCATH, NP_596320.2; SpuGPCATH, XP_003729899.1; StuGPCATH1, XP_006342345.1; StuGPCATH2, XP_006338125.1; UmaGPCATH, XP_011391556.1; ZmaGPCATH1, XP_008656604.1; ZmaGPCATH2, NP_001130177.1