



COPA (NM_004371.4) - cDNA + Protein - 2026-05-25

GTCGCTGACG TGGAGGCGTC CGAAGGGCAG CAGGGTGTGT CGGGGCTCGG -31

ATTAAGACAT CGGAGTCGGA GACCTGAGAG ATGTTAACCA AATTCGAGAC 20

MetLeuThrL ysPheGluTh 7

CAAGAGCGCG CGGGTCAAAG GGCTCAGCTT TCACCCCAA AGACCTTGGA 70

rLysSerAla ArgValLysG lyLeuSerPh eHisProLys ArgProTrpI 24

TCCTGACTAG TTTACATAAT GGGGTCATCC AGTTATGGGA CTATCGGATG 120

leLeuThrSe rLeuHisAsn GlyValIleG lnLeuTrpAs pTyrArgMet 40

TGCACTCTCA TTGACAAGTT TGATGAACAT GATGGTCCAG TGCGAGCAT 170 p.G56S

CysThrLeuI leAspLysPh eAspGluHis AspGlyProV alArgGlyIl 57

TGACTTCCAT AAGCAGCAGC CACTGTTCGT CTCTGGAGGA GATGACTATA 220

eAspPheHis LysGlnGlnP roLeuPheVa lSerGlyGly AspAspTyrL 74

AGATTAAGGT TTGGAATTAC AAGCTTCGGC GCTGTCTTTT CACATTGCTT 270

ysIleLysVa lTrpAsnTyr LysLeuArgA rgCysLeuPh eThrLeuLeu 90

GGGCACTTAG ATTATATTCG CACCACGTTT TTTCATCATG AATATCCCTG 320

GlyHisLeuA spTyrIleAr gThrThrPhe PheHisHisG luTyrProTr 107

GATTCTGAGT GCCTCCGATG ATCAGACCAT CCGAGTGTGG AACTGGCAAT 370

pIleLeuSer AlaSerAspA spGlnThrIl eArgValTrp AsnTrpGlnS 124

CTAGAACCTG TGTTTGTGTG TTAACAGGGC ACAACCATTA TGTGATGTGT 420

erArgThrCy sValCysVal LeuThrGlyH isAsnHisTy rValMetCys 140

GCTCAGTTCC ACCCCACAGA AGACTTGGTA GTATCAGCCA GCCTGGACCA 470

AlaGlnPheH isProThrGl uAspLeuVal ValSerAlaS erLeuAspGl 157

GACTGTGCGC GTTTGGGATA TTTCTGGTCT GAGGAAAAA AACCTGTCCC 520

nThrValArg ValTrpAspI leSerGlyLe uArgLysLys AsnLeuSerP 174

CTGGTGCGGT GGAATCGGAT GTGAGAGGAA TAACTGGGGT TGATCTATTT 570

roGlyAlaVa lGluSerAsp ValArgGlyI leThrGlyVa lAspLeuPhe 190

GGAACTACAG ATGCAGTGGT GAAGCATGTA CTAGAGGGTC ACGATCGTGG 620 [p.H199R](#) [L201P](#)

GlyThrThrA spAlaValVa lLysHisVal LeuGluGlyH isAspArgGl 207

AGTAAACTGG GCTGCCTTCC ACCCCACTAT GCCCCTTATT GTATCTGGGG 670 [p.A211V](#)

yValAsnTrp AlaAlaPheH isProThrMe tProLeuIle ValSerGlyA 224

CAGATGATCG TCAAGTGAAG ATCTGGCGCA TGAATGAATC AAAGGCATGG 720 [p.R227C](#) [p.K230N](#) [R233C](#) [p.R233H](#) [p.R233L](#) [p.K238E](#) [p.A239P](#) [p.W240R](#) [p.W240L](#) [p.W240S](#)

laAspAspAr gGlnValLys IleTrpArgM etAsnGluSe rLysAlaTrp 240

GAGGTTGATA CCTGCCGGGG CCATTACAAC AATGTATCTT GTGCCGTCTT 770 [p.E241K](#) [p.E241A](#) [E241D](#) [p.V242G](#) [p.D243N](#) [p.D243G](#) [G247D](#)

GluValAspT hrCysArgGl yHisTyrAsn AsnValSerC ysAlaValPh 257

CCACCCTCGC CAAGAGTTGTA TCCTCAGCAA TTCTGAGGAC AAGAGTATTC 820 [H258R](#) [L263S](#)

eHisProArg GlnGluLeuI leLeuSerAs nSerGluAsp LysSerIleA 274

GAGTCTGGGA TATGTCTAAG CGGACTGGGG TTCAGACTTTT CCGCAGAGAC 870 [p.R281W](#) [p.Q285H](#)

rgValTrpAs pMetSerLys ArgThrGlyV alGlnThrPh eArgArgAsp 290

CATGATCGTT TCTGGGTCCT AGCTGCTCAC CCTAACCTTA ACCTCTTTGC 920

HisAspArgP heTrpValLe uAlaAlaHis ProAsnLeuA snLeuPheAl 307

AGCAGGCCAT GATGGTGGTA TGATTGTGTT TAAGCTGGAA CGGGAACGGC 970

aAlaGlyHis AspGlyGlyM etIleValPh eLysLeuGlu ArgGluArgP 324

CAGCCTATGC TGTTTCATGGC AATATGCTAC ACTATGTCAA GGACCGATTC 1020 [p.R339Q](#)

roAlaTyrAl aValHisGly AsnMetLeuH isTyrValLy sAspArgPhe 340

TTACGACAGC TGGATTTCAA CAGCTCCAAA GATGTAGCTG TGATGCAGTT 1070

LeuArgGlnL euAspPheAs nSerSerLys AspValAlaV alMetGlnLe 357

GCGGAGTGGT TCCAAGTTTC CAGTATTCAA TATGTCATAC AATCCAGCAG 1120 [R358W](#) [p.Y370S](#)

uArgSerGly SerLysPheP roValPheAs nMetSerTyr AsnProAlaG 374

AAAATGCAGT CCTGCTTTTGT ACAAGAGCTA GCAATCTAGA GAATAGTACC 1170

luAsnAlaVa lLeuLeuCys ThrArgAlaS erAsnLeuGl uAsnSerThr 390

TATGACCTGT ACACCATCCC TAAAGATGCT GACTCCCAGA ATCCTGATGC 1220

TyrAspLeuT yrThrIlePr oLysAspAla AspSerGlnA snProAspAl 407

GCCTGAAGGG AAACGATCCT CAGGCCTGAC AGCCGTTTGG GTCGCTCGAA 1270 [p.P408S](#)

aProGluGly LysArgSerS erGlyLeuTh rAlaValTrp ValAlaArgA 424

ATCGGTTTGC TGTCC TAGAT CGGATGCATT CGCTTCTGAT CAAGAATCTG 1320
snArgPheAl aValLeuAsp ArgMetHisS erLeuLeuIl eLysAsnLeu 440

AAGAATGAGA TCACCAAAAA GGTACAGGTG CCCAACTGTG ATGAGATCTT 1370
LysAsnGluI leThrLysLy sValGlnVal ProAsnCysA spGluIlePh 457

CTATGCTGGC ACAGGCAATC TCCTGCTTCG AGATGCGGAC TCTATCACAC 1420
eTyrAlaGly ThrGlyAsnL euLeuLeuAr gAspAlaAsp SerIleThrL 474

TCTTTGACGT ACAGCAGAAG CGGACTCTGG CATCTGTGAA GATTTCTAAA 1470
euPheAspVa lGlnGlnLys ArgThrLeuA laSerValLy sIleSerLys 490

GTGAAATACG TTATCTGGTC AGCAGACATG TCACATGTAG CACTACTAGC 1520
ValLysTyrV alIleTrpSe rAlaAspMet SerHisValA laLeuLeuAl 507

CAAACACGCC ATTGTGATCT GTAACCGCAA ACTGGATGCT TTATGTAACA 1570 [p.D519Y](#)
aLysHisAla ileValIleC ysAsnArgLy sLeuAspAla LeuCysAsnI 524

TTCATGAGAA CATTCTGTGC AAGA GTGGGG CCTGGGATGA GAGTGGGGTA 1620 [S532I](#)
leHisGluAs nIleArgVal LysSerGlyA laTrpAspGl uSerGlyVal 540

TTTATCTATA CCACAAGCAA CCACATCAA TATGCTGTCA CCACTGGGGA 1670
PheIleTyrT hrThrSerAs nHisIleLys TyrAlaValT hrThrGlyAs 557

CCACGGGATC ATTCGAAGCT TGGATTTACC CATCTATGTC ACACGGGTGA 1720
pHisGlyIle ileArgThrL euAspLeuPr oIleTyrVal ThrArgValL 574

AGGGCAACAA TGTATACTGC CTAGACAGGG AGTGTCGTCC CCGGGTACTC 1770

ysGlyAsnAs nValTyrCys LeuAspArgG luCysArgPr oArgValLeu 590

ACCATTGATC CCACTGAGTT CAAATTCAAG CTGGCCCTGA TCAACAGAAA 1820

ThrIleAspP roThrGluPh eLysPheLys LeuAlaLeuI leAsnArgLy 607

ATATGATGAG GTACTGCACA TGGTGAGGAA TGCCAAACTA GTTGGCCAGT 1870

sTyrAspGlu ValLeuHisM etValArgAs nAlaLysLeu ValGlyGlnS 624

CTATTATTGC TTATCTCCAG AAGAAGGGCT ATCCTGAAGT GGCACCTGCAT 1920

erIleIleAl aTyrLeuGln LysLysGlyT yrProGluVa lAlaLeuHis 640

TTTGTCAAGG ATGAGAAAAC TCGCTTTAGT CTGGCACTGG AGTGTGGAAA 1970

PheValLysA spGluLysTh rArgPheSer LeuAlaLeuG luCysGlyAs 657

CATTGAGATT GCTCTGGAAG CAGCCAAAGC ACTGGATGAC AAGAACTGCT 2020

nIleGluIle AlaLeuGluA laAlaLysAl aLeuAspAsp LysAsnCysT 674

GGGAAAAGCT GGGAGAAGTG GCCCTGCTGC AGGGGAACCA CCAGATTGTG 2070 G678E

rpGluLysLe uGlyGluVal AlaLeuLeuG lnGlyAsnHi sGlnIleVal 690

GAAATGTGCT ATCAGCGTAC CAAAACTTT GACAACTTT CCTTCCTGTA 2120 Y694C T697I

GluMetCysT yrGlnArgTh rLysAsnPhe AspLysLeuS erPheLeuTy 707

TCTTATCACT GGCAACTTAG AAAAACTTCG CAAGATGATG AAGATTGCTG 2170

rLeuIleThr GlyAsnLeuG luLysLeuAr gLysMetMet LysIleAlaG 724

AGATCAGAAA GGACATGAGT GGCCACTATC AGAATGCCCT ATACCTGGGT 2220

IleArgLy sAspMetSer GlyHisTyrG InAsnAlaLe uTyrLeuGly 740

GATGTGTCAG AGCGTGTGCG GATCCTGAAG AACTGTGGAC AGAAGTCCCT 2270 [R745H](#)

AspValSerG luArgValAr gIleLeuLys AsnCysGlyG InLysSerLe 757

GGCCTATCTC ACAGCTGCTA CCCATGGCTT AGATGAAGAA GCTGAGAGCC 2320

uAlaTyrLeu ThrAlaAlaT hrHisGlyLe uAspGluGlu AlaGluSerL 774

TAAAGGAGAC ATTTGACCCA GAGAAGGAGA CAATCCCAGA CATTGACCCT 2370 [T777I](#)

euLysGluTh rPheAspPro GluLysGluT hrIleProAs pIleAspPro 790

AATGCCAAGC TGCTCCAGCC ACCTGCACCT ATCATGCCAT TGGATACCA 2420 [p.Asn807Thr](#)

AsnAlaLysL euLeuGlnPr oProAlaPro IleMetProL euAspThrAs 807

TTGGCCTTTA TTGACTGTAT CCAAAGGATT TTTTGAAGGC ACCATTGCCA 2470

nTrpProLeu LeuThrValS erLysGlyPh ePheGluGly ThrIleAlaS 824

GCAAAGGGAA GGGAGGAGCA CTGGCTGCTG ACATTGACAT TGACACTGTT 2520 [p.G828E](#)

erLysGlyLy sGlyGlyAla LeuAlaAlaA spIleAspIl eAspThrVal 840

GGTACAGAGG GCTGGGGAGA GGATGCAGAG CTGCAGTTGG ATGAAGATGG 2570 [G844D](#)

GlyThrGluG lyTrpGlyGl uAspAlaGlu LeuGlnLeuA spGluAspGl 857

GTTTGTGGAG GCTACAGAAG GTTTGGGGGA TGATGCTCTT GGCAAGGGAC 2620

yPheValGlu AlaThrGluG lyLeuGlyAs pAspAlaLeu GlyLysGlyG 874

AGGAAGAAGG AGGTGGCTGG GATGTAGAAG AAGATCTGGA GCTCCCTCCT 2670

InGluGluGl yGlyGlyTrp AspValGluG luAspLeuGl uLeuProPro 890

GAGCTGGATA TATCCCCTGG GGCAGCTGGT GGGGCTGAAG ATGGTTTCTT 2720
GluLeuAspI leSerProGl yAlaAlaGly GlyAlaGluA spGlyPhePh 907

TGTGCCCCCA ACCAAGGGAA CAAGTCCAAC TCAGATCTGG TGTAATAACT 2770
eValProPro ThrLysGlyT hrSerProTh rGlnIleTrp CysAsnAsnS 924

CTCAGCTTCC AGTTGATCAC ATCCTGGCAG GCTCTTTCGA AACAGCCATG 2820
erGlnLeuPr oValAspHis ileLeuAlaG lySerPheGl uThrAlaMet 940

CGGCTCCTTC ATGACCAAGT AGGGGTAATC CAGTTTGGCC CCTACAAGCA 2870
ArgLeuLeuH isAspGlnVa lGlyValIle GlnPheGlyP roTyrLysGl 957

ACTGTTCCCTA CAGACATACG CCCGAGGCCG CACAACCTAT CAGGCTCTGC 2920
nLeuPheLeu GlnThrTyrA laArgGlyAr gThrThrTyr GlnAlaLeuP 974

CCTGCCTACC CTCCATGTAT GGCTATCCTA ATCGCAACTG GAAGGATGCA 2970
roCysLeuPr oSerMetTyr GlyTyrProA snArgAsnTr pLysAspAla 990

GGGCTGAAGA ATGGTGTACC AGCTGTGGGC CTGAAGCTTA ATGACCTCAT 3020
GlyLeuLysA snGlyValPr oAlaValGly LeuLysLeuA snAspLeuIl 1007

CCAACGGTTG CAGCTGTGCT ACCAGCTCAC CACAGTTGGC AAATTTGAGG 3070 [p.C1013S](#)
eGlnArgLeu GlnLeuCysT yrGlnLeuTh rThrValGly LysPheGluG 1024

AGGCTGTGGA AAAATTCCGT TCCATCCTTC TCAGTGTGCC ACTTCTTGTT 3120
luAlaValGl uLysPheArg SerIleLeuL euSerValPr oLeuLeuVal 1040

GTGGACAATA AACAAGAGAT TGCAGAGGCC CAGCAGCTCA TCACCATTTG 3170

ValAspAsnL ysGlnGluIl eAlaGluAla GlnGlnLeuI leThrIleCy 1057

CCGTGAGTAC ATTGTGGGTT TGTCCGTGGA GACAGAAAGG AAGAAGCTGC 3220 [p.R1058C](#)

sArgGluTyr IleValGlyL euSerValGl uThrGluArg LysLysLeuP 1074

CCAAAGAGAC TCTAGAACAG CAGAAGCGCA TCTGTGAGAT GGCAGCCTAT 3270

roLysGluTh rLeuGluGln GlnLysArgI leCysGluMe tAlaAlaTyr 1090

TTCACCCACT CAAACCTGCA GCCTGTGCAC ATGATCCTGG TGCTGCGTAC 3320

PheThrHisS erAsnLeuGl nProValHis MetIleLeuV alLeuArgTh 1107

AGCCCTCAAT CTGTTCTTCA AGCTCAAGAA CTTCAAGACA GCTGCCACCT 3370

rAlaLeuAsn LeuPhePheL ysLeuLysAs nPheLysThr AlaAlaThrP 1124

TTGCTCGGCG CCTACTAGAA CTCGGGCCCA AGCCTGAGGT GGCCCAACAG 3420

heAlaArgAr gLeuLeuGlu LeuGlyProL ysProGluVa lAlaGlnGln 1140

ACCCGAAAAA TCCTGTCTGC CTGTGAGAAG AATCCCACAG ATGCCTACCA 3470 [p.R1142X](#)

ThrArgLysI leLeuSerAl aCysGluLys AsnProThrA spAlaTyrGl 1157

GCTCAATTAT GACATGCACA ACCCCTTTGA CATTGTGCT GCATCATATC 3520

nLeuAsnTyr AspMethHisA snProPheAs pIleCysAla AlaSerTyrA 1174

GGCCCATCTA CCGTGGAAG CCAGTAGAAA AGTGTCCACT CAGTGGGGCC 3570

rgProIleTy rArgGlyLys ProValGluL ysCysProLe uSerGlyAla 1190

TGCTATTCCC CTGAGTTCAA AGGTCAAATC TGCAGGGTCA CCACAGTGAC 3620

CysTyrSerP roGluPheLy sGlyGlnIle CysArgValT hrThrValTh 1207

AGAGATTGGC AAAGATGTGA TTGGTTTAAG GATCAGTCCT CTGCAGTTTC 3670

rGluIleGly LysAspValI leGlyLeuAr gIleSerPro LeuGlnPheA 1224

GC**TAA**GGCCC CCTTTGTGTG CATGGGTCAG TCACCATATG TTCCCCCAG *45

rgStop

AGAATGTGTC TATATCCTCC TTCTAACAGC ACCTTCCCCC TGCAGCTACT *95

CTTCAGATCT GGCTCTCTGT ACCCTAAAAC CTAGTATCTT TTTCTCTTCT *145

ATGGAAAATC CGAAGGTCTA AACTTGACTT TTTTGAGGTC TTCTCAACTT *195

GACTACAGTT GTGCTCATAA TTGTCCTTGC CTTTCCAGCT TAATTATTTT *245

AAGGAACAAA TGAAAACCTCT GGGCTGGGTG GAGTGGCTCA TACCTGTAAT *295

CCCAGCACCT TGGGAGGCTA CGGTGGGCAG ATCATCTGAG GCCAGGAGTT *345

CGAGACCTGC CTGGCCAACA TGGCAACACC CCGTCTCTAA TAAAAATATA *395

AAAATTAGCC TGGCATGGTA GCATGCGCCT ATAGTCCCAG CTGCTCAGGA *445

GGCTGAGGCA TGAGAATCGC TTGAACCTAG GAGGTGGAGG TTGCATTCAA *495

CTGAGATCAT ACCACTTCAT TCCAGCCTGG GTGACAGAGC AAGACTCTGT *545

CTCAAAAAAA AAAAAAAGGA AAACCTCTGT ATGGACATTT GTTTAGTAAA *595

TCCCTTCAGT ATTTATCCCT CCTTTCCCCA CAGCAGCTTT CTTTCCGTGC *645

AACTAGAAAG GAGCAGGATG TAATAAATAC ATTTTGGTGT GACTAGGCCA *695

CACCAACTCT TAATCATCTC CCATTTTCCT TAGACATTTA AATTTCAAGG *745

CAGGTACCCT CTGTGTACTC AGAAATTTGA AGAAGTTATT TGGTTTTCCA *795

AAATGCACAC TGCGGGTTAT TGATTTGTTT TTTACAATA TTGTTCTCAT *845

ATTTCTCACA CTAAATAAAT CTCTATGAGA GCTTCTTGAC TTGGCCATTT *895

ATTTCTTGGA CACTCTCATG TTCTTGTTCA CCCATGCAGG CACCCACCA *945

AAGTACATAT CTTCCCTCCA GTAATAATTT TTAATTACAA AATAAACATC *995

CACTATTGGA AAAAAAAAAA AAAAGCTAGC CGGGCATGGT GGTGGGTGCC *1045

TGTAATCCCA GCTACTCTGG AGGCTGAGGC AGAGGATTGC TTGAACCCGG *1095
GAGGCGGAGG TTGCAGTAAG CTGAGATCGC GCCACCGCAC TCCAGCCTGG *1145
GCGACAGAGT GAGACTCCAT CTCAAAAAAA AAGAAAGAAA AAAAGAAGCA *1195
CATGTTTTTC ATAGGGTATA TATGAGGACC TAAACTGCTG TGAAAATGAT *1245
AGAAAGCAAG TAGCTCCCTT ATTCTGTTTT TGATTGCAGC CTTTTATCTT *1295
TTGCTAATTA TAGCAATATT TATTGAGCAC CTGCCATGTG ACTGTCACTG *1345
TTCTAGATAT TTTACATGTA ATATACAGAT AAAAGAATAG TACTTTATAT *1395
ATATTACAAT GATACAATGA TTACATTAAC AATACAATAT TTTGCTTGTC *1445
ATATGCTAAG AATAATTGGG TAGAGTGACA TTAGTGTGCC TTCGATTAAA *1495
ATAAGTACTT TTTTGC GTGT TAAATTCATG TTTTCAATAA ATAATAAATG *1545
CATATAGTTG AAAAATCA

COPA (NM_004371.4) - cDNA + Protein - 2026-05-25

