



PSMA6 (NM\_002791.3) - cDNA + Protein - 2026-06-27

GGAAGCAGTC GCTGCAACTT CCGGGAGGTG CTTGTGTGCC TGGTGC GGGA -46

GCTACGGGGC CCAGGGATTG TGTTTAAAGT AGTGCTTCTA CCAACATGTC 5

MetSe 2

CCGTGGTTCC AGCGCCGGTT TTGACCGCCA CATTACCATT TTTTCACCCG 55

rArgGlySer SerAlaGlyP heAspArgHi sIleThrIle PheSerProG 19

AGGGTCGGCT CTACCAAGTA GAATATGCTT TTAAGGCTAT TAACCAGGGT 105

luGlyArgLe uTyrGlnVal GluTyrAlaP heLysAlaIl eAsnGlnGly 35

GGCCTTACAT CAGTAGCTGT CAGAGGGAAA GACTGTGCAG TAATTGTCAC 155

GlyLeuThrS erValAlaVa lArgGlyLys AspCysAlaV alIleValTh 52

ACAGAAGAAA GTACCTGACA AATTATTGGA TTCCAGCACA GTGACTCACT 205 EX3Del

rGlnLysLys ValProAspL ysLeuLeuAs pSerSerThr ValThrHisL 69

TATTCAAGAT AACTGAAAAC ATTGGTTGTG TGATGACCGG AATGACAGCT 255

euPheLysIl eThrGluAsn IleGlyCysV alMetThrGl yMetThrAla 85

GACAGCAGAT CCCAGGTACA GAGGGCACGC TATGAGGCAG CTAAGTGGAA 305

AspSerArgS erGlnValGl nArgAlaArg TyrGluAlaA laAsnTrpLy 102

ATACAAGTAT GGCTATGAGA TTCCTGTGGA CATGCTGTGT AAAAGAATTG 355

sTyrLysTyr GlyTyrGluI leProValAs pMetLeuCys LysArgIleA 119

CCGATATTTTC TCAGGTCTAC ACACAGAATG CTGAAATGAG GCCTCTTGGT 405

laAspIleSe rGlnValTyr ThrGlnAsnA laGluMetAr gProLeuGly 135

TGTTGTATGA TTTTAATTGG TATAGATGAA GAGCAAGGCC CTCAGGTATA 455

CysCysMetI leLeuIleGl yIleAspGlu GluGlnGlyP roGlnValTy 152

TAAGTGTGAT CCTGCAGGTT ACTACTGTGG GTTTAAAGCC ACTGCAGCGG 505

rLysCysAsp ProAlaGlyT yrTyrCysGl yPheLysAla ThrAlaAlaG 169

GAGTTAAACA AACTGAGTCA ACCAGCTTCC TTGAAAAAAAA AGTGAAGAAG 555

lyValLysGl nThrGluSer ThrSerPheL euGluLysLy sValLysLys 185

AAATTTGATT GGACATTTGA ACAGACAGTG GAAACTGCAA TTACATGCCT 605

LysPheAspT rpThrPheGl uGlnThrVal GluThrAlaI leThrCysLe 202

GTCTACTGTT CTATCAATTG ATTTCAAACC TTCAGAAATA GAAGTTGGAG 655

uSerThrVal LeuSerIleA spPheLysPr oSerGluIle GluValGlyV 219

TAGTGACAGT TGAAAATCCT AAATTCAGGA TTCTTACAGA AGCAGAGATT 705

alValThrVa lGluAsnPro LysPheArgI leLeuThrGl uAlaGluIle 235

GATGCTCACC TTGTTGCTCT AGCAGAGAGA GACTAAACAT TGTCGTTAGT \*14

AspAlaHisL euValAlaLe uAlaGluArg AspStop

TTACCAGATC CGTGATGCCA CTTACCTGTG TGTTTGGTAA CAACAAACCA \*64

ACATCATGGA GGTCCCTGGA TTGAAAAAGG AGCCTCTCCC ACTCCTCCTA \*114

CCACCGAAGT GGTTAGGACT CTATATAAAT AAAACAAGG CTTTGGAAA \*164

ATAATTGCAT CCTGTTGTTT TCA

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