



PSMA5 (NM_002790.4) - cDNA - 2026-06-27

AGTTGGTCCT TAGTACTGCG GCCGTGTGGG TGAGTTGGCT GCCGGTGAGT -69
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CGTGAATACT TTTTCTCCCG AAGGAAGATT ATTTCAAGTG GAATATGCCA 82
TTGAGGCTAT CAAGCTTGGT TCTACAGCCA TTGGGATCCA GACATCAGAG 132
GGTGTGTGCC TAGCTGTGGA GAAGAGAATT ACTTCCCCAC TGATGGAGCC 182
CAGCAGCATT GAGAAAATTG TAGAGATTGA TGCTCACATA GTTTGTGCCA 232
TGAGTGGGCT AATTGCTGAT GCTAAGACTT TAATTGATAA AGCCAGAGTG 282
GAGACACAGA ACCACTGGTT CACCTACAAT GAGACAATGA CAGTGGAGAG 332
TGTGACCCAA GCTGTGTCCA ATCTGGCTTT GCAGTTTGGG GAAGAAGATG 382
CAGATCCAGG TGCCATGTCT CGTCCCTTTG GAGTAGCATT ATTATTTGGA 432
GGAGTTGATG AGAAAGGACC CCAGCTGTTT CATATGGACC CATCTGGGAC 482
CTTTGTACAG TGTGATGCTC GAGCAATTGG CTCTGCTTCA GAGGGTGCCC 532 p.Arg168*
AGAGCTCCTT GCAAGAAGTT TACCACAAGT CTATGACTTT GAAAGAAGCC 582
ATCAAGTCTT CACTCATCAT CCTCAAACAA GTAATGGAGG AGAAGCTGAA 632
TGCAACAAAC ATTGAGCTAG CCACAGTGCA GCCTGGCCAG AATTTCCACA 682
TGTTACACAA GGAAGAACTT GAAGAGGTTA TCAAGGACAT TTAAGGAATC *6
CTGATCCTCA GAACTTCTCT GGGACAATTT CAGTCTAAT AATGTCCTTA *56
AATTTTATTT CCAGCTCCTG TTCCTTGAA AATCTCCATT GTATGTGCAT *106
TTTTTAAATG ATGTCTGTAC ATAAAGGCAG TTCTGAAATA AAGAAAATTT *156
TAAATATTT GTTAATAGAC TGTTCTCTTC TAATAGTCTT TTTTTTTTTT *206
TTTTTTTTAA GAGATGAGGT CTCACTATAT TGCCCAGGCT GTTTCAAAC *256

TCCTGGGTTT AAGTGATCTT CCCGCCTCTG CCTCCTGAAG TGCTAGGATT *306
ATAGGCGAGA GCCACTGTGC CCAGCCATTG TAATACAGTC TTTTGTGTTGA *356
AATGCAAATG TTAGTGGGCC AAAAGGACAG TGTATTAAAT ATCCCTATTC *406
TATGAAGATG TCTGTTCTAA AACTGTTTTA GTGATGCTCT ATAACAGAAT *456
CAACTATATC CATTAAATCCC ATGGTGTGTA TAAATGAACT AGTCCAGCTC *506
TTCCCAGTGG TTAATAGATG TTAACAGCAC TGTAATCCA TTTCGGCCCT *556
ATAATTTCTG GAGCAAACGT GTATTTTTGC TGCTTTTCTT ATAAGGAGAT *606
GCTACTAATT GACTATCAGA AGAACATGTT TTGAGGTCCT CCTCAAAAACA *656
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TGATAGACAA TTTGGAAAGA GATTTCTTG TTTACGAAGT GAAGGACTAA *756
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AATGAATGAT GAACTTTCCC CATTAGAGCT GCCATCATAT GATTGGTGGA *906
GATTGTTTAA AGAATCATTC TATCTCAGCC TCAAAAATTA CCACGGGAAG *956
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CATGGACACA CAAAATTAAG CTGTTGGGAC TTGAACAGCT CAAATTTTCT *1056
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GAAACTTCTA GTGGGGAGAG CCAAACAATA AACAAATATC CTTTTTAACT *1706
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TTCTGCTATA GAATGTTTCC TGGAGAAAGG TACGGTTTTT AGTAATAAGA *2106
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CACAAAGGAG AACTGACAGT GCCACTTTTA TATTTAATGT CAATCTCATT *2206
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GGAAATATTT TAATGTAAAT ATCGGGTTGG GTGCATTGTA TCATTTTGCT *2906
CTACAATGAC TATATCCTCA TTTGGGTTTA TGGCCAGTTT ATAATTA AAA *2956
GTAATCTTTA CTTGC

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