

**S2 Table. Oligonucleotides used in the present study for mung bean nuclease protection assay.**

No.	Sequence (5'-->3')
18S-01	GGCTTAATCTTTGAGACAAGCATATGACTACTGGCAGGATCAACCAGATA
18S-02	GCCATTCGCAGTTTCACTGTATAAATTGCTTATACTTAGACATGCATGGC
18S-03	TAGTAAAGGAAGTATCAAATAAACGATAACTGATTTAATGAGCCATTCGC
18S-04	AAGCATGTATTAGCTCTAGAATTACCACAGTTATACCATGTAGTAAAGGA
18S-05	TTATCTAATAAATACATCTCTTCCAAAGGGTCGAGATTTTAAGCATGTAT
18S-06	GTTATTATGAATCATCAAAGAGTCCGAAGACATTGATTTTTTATCTAATA
18S-07	GAATGAACCATCGCCAGCACAAGGCCATGCGATTCGAAAAGTTATTATGA
18S-08	CACTATCCTACCATCGAAAGTTGATAGGGCAGAAATTTGAATGAACCATC
18S-09	CGAACCTTATTCCCCGTTACCCGTTGAAACCATGGTAGGCCACTATCCT
18S-10	CTTGGATGTGGTAGCCGTTTCTCAGGCTCCCTCTCCGGAATCGAACCTT
18S-11	CTGAATTAGGATTGGGTAATTTGCGCGCCTGCTGCCTTCCTTGGATGTGG
18S-12	CCGAATGGGCCCTGTATCGTTATTTATTGCTACTACCTCCCTGAATTAGG
18S-13	GTTAAGGTATTTACATTGTACTCATTCCAATTACAAGACCCGAATGGGCC
18S-14	CGCGGCTGCTGGCACCAGACTTGCCTCCAATTGTTCTCGTTAAGGTAT
18S-15	CTGCAACAACCTTAATATACGCTATTGGAGCTGGAATTACCGCGGCTGCT
18S-16	CCGGCAACCGGGCCAAAGTTCAACTACGAGCTTTTAACTGCAACAAC
18S-17	GAAAGGCCCGTTGGAAATCCAGTACACGAAAAAATCGGACCGGCCAAC
18S-18	GGTTCGCCAAGAGCCACAAGGACTCAAGGTTAGCCAGAAGGAAAGGCCCC
18S-19	GCCTGCTTTGAACACTCTAATTTTTTCAAAGTAAAAGTCTGTTTCGCCA
18S-20	TCCTATTCTATTATTCATGCTAATATATTCGAGCAATACGCCTGCTTTG
18S-21	CATTACGATGGTCTAGAAACCAACAAAATAGAACCAACGTCCTATTCT
18S-22	CAATTGAATACTGATGCCCCGACCGTCCCTATTAATCATTACGATGGTC
18S-23	GTAGTTAGTCTTCAATAAATCCAAGAATTTACCTCTGACAATTGAATAC
18S-24	TTCTTGATTAATGAAAACGTCCTTGGCAAATGCTTTCGCAGTAGTTAGTC
18S-25	CTACGACGGTATCTGATCATCTTCGATCCCCTAACTTTGTTCTTGATTA
18S-26	CACCACCGATCCCTAGTCGGCATAGTTTATGGTTAAGACTACGACGGTA
18S-27	CTTTGATTTCTCGTAAGGTGCCGAGTGGGTCATTAACCAACACCCACCG
18S-28	GTTTCAGCCTTGCAGCATACTCCCCCAGAACCCAAAGACTTTGATTC
18S-29	AGGCTCCACTCCTGGTGGTGCCTTCCGTCAATTCCTTTAAGTTTCAGCC
18S-30	GACCTGGTGAAGTTTCCCGTGTGAGTCAAATTAAGCCGAGGCTCCACT
18S-31	CAAGAAAGAGCTCTCAATCTGTCAATCCTTATTGTGTCTGGACCTGGTGA
18S-32	CACCAACTAAGAACGGCCATGCACCACCACCCACAAAATCAAGAAAGAGC
18S-33	GGTCTCGTTGTTATCGCAATTAAGCAGACAAATCACTCCACCAACTAAG
18S-34	GATAACCAGCAAATGCTAGCACCCTATTTAGTAGGTTAAGGTCTCGTTC
18S-35	CTTCCATCGGCTTGAAACCGATAGTCCCTCTAAGAAGTGGATAACCAGCA
18S-36	GAACGTCTAAGGGCATCACAGACCTGTTATTGCCTCAAACCTCCATCGGC
18S-37	GACTCGCTGGTCCGTCAAGTGTAGCGCGCTGCGGCCAGAACGTCTAAG
18S-38	GGAGTTTACAAGATTACCAAGACCTCTCGGCCAAGGTTAGACTCGCTGG
18S-39	TTGAAGAGCAATAATTACAATGCTCTATCCCAGCACGACGGAGTTTCAC
18S-40	CAACGCAAGCTGATGACTTGCCTTACTAGGAATTCCTGTTGAAGAGCA
18S-41	ACTAGCGACGGGCGGTGTGTACAAAGGGCAGGGACGTAATCAACGCAAGC
18S-42	CTAAGCAGATCCTGAGGCCTCACTAAGCCATTCAATCGGTAAGTAGCGACG
18S-43	GTCCAAATTCTCCGCTCTGAGATGGAGTTGCCCTTCTCTAAGCAGATC
18S-44	CTTGTACGACTTTTAGTTCTCTAAATGACCAAGTTTGTCCAAATTCTC
18S-45	TAATGATCCTTCCGCAGGTTACCTACGGAACCTTGTTACGACTTTTAG
25S-01	CCTCCGCTTATTGATATGCTTAAGTTCAGCGGGTACTCCTACCTGATTTGAGGTCAAAC

25S-02	GCTTTTGCCGCTTCACTCGCCGTTACTAAGGCAATCCCGTTGGTTCTTTTCTCCGC
25S-03	CCCTCTCAAATTACAACCTCGGGCACCGAAGGTACCAGATTTCAAATTTGAGCTTTTGCC
25S-04	CTCTATGACGTCTGTCCAAGGAACATAGACAAGGAACGGCCCAAAGTTGCCCTCTCC
25S-05	CGAAGGCACTTTACAAAGAACCGCACTCTCGCCACACGGGATTCTCACCTCTATGACG
25S-06	GATGGAATTTACCACCCACTTAGAGCTGCATTCCCAAACAACCTCGACTCTTGAAGGCAC
25S-07	CATCACTGTA CTGTTCGCTATCGGTCTCTCGCCAATATTTAGCTTTAGATGGAATTTAC
25S-08	CAATTTACGTACTTTTTCACTCTTTTTCAAAGTTCTTTTCATCTTTCCATCACTGTAC
25S-09	GAGCAGAGGGGCACAAAACCCATGTCTGATCAAATGCCCTTCCCTTTCAACAATTTACG
25S-10	CACCAAACTGATGCTGGCCAGTAAAATGCGAGATTCCCCTACCCACAAGGAGCAGAGG
25S-11	GGCTATAATACTTACCGAGGCAAGCTACATTCTATGGATTTATCTGCCACCAAACTG
25S-12	CCTTGACTTACGTGCGAGTCTCAGTCCCAGCTGGCAGTATTCCACAGGCTATAATAC
25S-13	CCTGGTCCGTGTTTCAAGACGGGCGGCATATAACCATTATGCCAGCATCCTTGACTTAC
25S-14	CTTTCATTACGCGTATGGGTTTTACACCCAAACTCGCATAGACGTTAGACTCCTTGGT
25S-15	CATCAGGATCGGTGATTGTGCACCTTTGCGAGGCCCAAACCTACGTTCACTTTTATTAC
25S-16	CCATCTTTCGGGTCCCAACAGCTATGCTCTTACTCAAATCCATCCGAAGACATCAGGATC
25S-17	GAGCCTCCACCAGAGTTTCTCTGGCTTACCCTATTAGGCATAGTTACCATCTTTGCG
25S-18	CGCCCTATACCCAAATTCGACGATCGATTTGCAGTCCAGAACCGCTACGAGCCTCCACC
25S-19	GAGGGAAACTTCGGCAGGAACAGCTACTAGATGGTTCGATTAGTCTTTGCCCCATAC
25S-20	CTAATCATTGCTTTACCTCATAAACTGATACGAGCTTCTGCTATCCTGAGGGAACTTC
25S-21	CATATTTAAAGTTTGAGAATAGGTCAAGGTCAATTCGACCCGGAACCTTAATCATTG
25S-22	GCTCTCATTCAAATGTCCACGTTCAATTAAGTAACAAGGACTTCTTACATATTTAAAG
25S-23	CGGTTTATCCCGCATCGCCAGTTCTGCTTACCAAAAATGGCCCACTAAAAGCTCTTCATTC
25S-24	CCTTTTGTTGTTGCTGATGAGCGTGTATTCCGGCACCTTAACTCTACGTTTCGGTTCATCC
25S-25	GCGGATTCGACTTCCATGGCCACCGTCCGGCTGTCTAGATGAACTAACCTTTTGTGG
25S-26	CCATTTTCAGGGCTAGTTCAATTCGGCCGGTGAGTTGTACACACTCCTTAGCGGATTCCG
25S-27	CATCATATCAACCCTGACGGTAGAGTATAGGTAACACGCTTGAGCGCCATCCATTTTACG
25S-28	CTTACGGTCTAGGCTTCTGCTACTGACCTCCACGCTGCCTACTCGTCAGGGCATCATATC
25S-29	GAATATTTGCTACTACCACCAAGATCTGCACTAGAGGCCGTTTCGACCCGACCTTACGGTC
25S-30	CTGCTGTTGACGTGGAACCTTTCCCACTTCAAGTTCTCATTGAAATATTTGC
25S-31	CTTTGAAACGGAGCTTCCCATCTCTTAGGATCGACTAACCACGTCCAAGTCTGTTGAC
25S-32	CGGAATCTTAAACGGATTCCCTTTCGATGGTGGCTGCATAAAATCAGGCCTTTGAAACG
25S-33	CGTCTCCACATTACGTTACGTTACCGTGAAGAATCCATATCCAGGTTCCGGAATCTTAAAC
25S-34	GGTGATAAGCTGTTAAGAAGAAAAGATAACTCCTCCAGGGCTCGCGCCGACGCTCCAC
25S-35	GTGCTGGCCTCTCCAGCCATAAGACCCATCTCCGGATAAACCAATTCGGGGTGATAAG
25S-36	CTTCTGTGGATTTTACGGGCCGTACAAGCGCACCGGAGCCAGCAAAGGTGCTGGCCTC
25S-37	CCTTGGAGACCTGCTGCGGTTATCAGTACGACCTGGCATGAAAATCTTCTTCTGTGG
25S-38	GCCGACTTCCCTTATCTACATTATTCTATCAACTAGAGGCTGTTACCTTGAGACC
25S-39	CTACCCGACCTTAGAGCCAATCCTTATCCCGAAGTTACGGATCTATTTGCCGACTTCC
25S-40	CCCACCAAGCAGTCCACAAGCACGCCGCTGCGTCTGACCAAGGCCCTCACTACCCGACC
25S-41	CAAGGCCGTCTACAACAAGGCACGCAAGTAGTCCGCCTAGCAGAGCAAGCCCCACCAAGC
25S-42	GTTCTAAGTTGATCGTTAATTGTAGCAAGCGACGGTCTACAAGAGACCTACCAAGGCCGT
25S-43	CGCAATGCTATGTTTTAATTAGACAGTCAAGTCCCTTGTCCGTACCAGTTCTAAGTTG
25S-44	CAGAGCACTGGGCAGAAATCACATTGCGTCAACATCACTTTCTGACCATCGCAATGCTATG
25S-45	GTTACTCCCGCCGTTTACCCGCGCTTGGTTGAATTTCTCACTTTGACATTCAGAGCACTG
25S-46	GCGCGTCACTAATTAGATGACGAGGCATTTGGCTACCTAAGAGAGTCATAGTTACTCCC
25S-47	GTTTCGCTAGATAGTAGATAGGGACAGTGGGAATCTCGTTAATCCATTCATGCGGTCAC
25S-48	CAGGGTCTTCTTCCCGCTGATTCTGCCAAGCCGTTCCCTTGGCTGTGGTTTCGCTAG
25S-49	GAAGACCCTGTTGAGCTTACTCTAGTTTACATTGTGAAGAGACATAGAGGGTGTAG

25S-50	GAAACTATAAAGGTAGTGGTATTTCACTGGCGCCGAAGCTCCCACTTATTCTACACCCTC
25S-51	CTAGAACGTGGAAAATGAATTCAGCTCCGCTTCATTGAATAAGTAAAGAACTATAAAG
25S-52	CCACCTGACAATGTCTCAACCCGGATCAGCCCCGAATGGGACCTTGAATGCTAGAACGTG
25S-53	CTTAGGACATCTGCGTTATCGTTTAAACAGATGTGCCGCCAGCCAAACTCCCCACCTGAC
25S-54	GCTTTTACCCTTTTGTCTACTGGAGATTTCTGTTCTCCATGAGCCCCCTTAGGACATC
25S-55	GGCCACACTTTCATGGTTTGTATTCACACTGAAAATCAAATCAAGGGGGCTTTTACCC
25S-56	GTAACTTTTCTGGCACCTCTAGCCTCAAATTCGAGGGACTAAAGGATCGATAGGCCACAC
25S-57	GCAATGTCGCTATGAACGCTTGACTGCCACAAGCCAGTTATCCCTGTGGTAACTTTTCTG
25S-58	CGAATTCTGCTTCGGTATGATAGGAAGAGCCGACATCGAAGAATCAAAAAGCAATGTCGC
25S-59	CCCAGCTCAGTTCCTATTAGTGGGTGAACAATCCAACGCTTACCGAATTCTGCTTCGG
25S-60	GGTAACATTCATCAGTAGGGTAAACTAACCTGTCTCACGACGGTCTAAACCCAGCTCAC
25S-61	CAATTATCCGAATGAACTGTTCTCTCGTACTAAGTTCAATTACTATTGCGGTAACATTC
25S-62	CCAGCGGATGGTAGCTTCGCGGCAATGCCTGATCAGACAGCCGCAAAAACCAATTATCCG
25S-63	GAAATCACCGCTTCTAGCATGGATTCTGACTTAGAGGCGTTCAGCCATAATCCAGCGGATG
25S-64	GACGCCACAAGGACGCCTTATTCGTATCCATCTATATTGTGTGGAGCAAAGAAATCACCG
25S-65	CCCAAGGCCTTCCGCCAAGTGACCGTTGCTAGCCTGCTATGGTTCAGCGACGCCACAAG
25S-66	CAAATGATTTATCCCCACGAAAATGACATTGCAATTCGCCAGCAAGCACCCAAGGCC
25S-67	CAAGGCTACTCTACTGCTTACAATACCCCGTTGTACATCTAAGTCGTATACAAATG
25S-68	ACAAATCAGACAACAAAGGCTTAATCTCAGCAGATCGTAACAACAAGGCTACTCTACTGC
18S-561	CAGACTTGCCCTCCAATTGTTCTCGTTAAGGTATTTACATTGTAICTCAT
18S-562	ACAACCTTAATATACGCTATTGGAGCTGGAATTACCGCGGCTGCTGGCAC
25S-2791	CTAGCCTCAAATTCGAGGGACTAAAGGATCGATAGGCCACACTTTCATG
25S-2793	GAACGCTTGACTGCCACAAGCCAGTTATCCCTGTGGTAACTTTTCTGGCACC
25S-2265	CATTCATGCGCGTCACTAATTAGATGACGAGGCATTTGGCTACCTTAAGAG
18S-11a	CCTGCTGCCTTCTTGATGTGGTAGCCGTTTCTCAGGCTCCCTCTCCGG
18S-11b	CGTTATTTATTGCTACTACCTCCCTGAATTAGGATTGGGTAATTTGCGCGC
25S-60a	CGGTAACATTCATCAGTAGGGTAAAATAACCTGTCTCACGACGGTCTAAAC
Oligo- 491	TCCTCGTTAAGGTATTTACATTGTAICTATTCCAATTACAAGACCCGAATG
Oligo-542	GAGCTGGAATTACCGCGGCTGCTGGCACCAGACTTGCCCTCCAATTGT