```
                    *        20         *        40         *        60         *        80         *       100         *
Prcmoter_R : TCTTCAATTGCAAGTAGCCAACATGATCAAGATCCCACTTGATATAGTAGCTGATTTTAACACATGCCTTAGCTGATTTAACAACCGCCCGTTTCTTATTTTCGGGAATAC : 110
Prcmoter_S : ---CAATTGCAAGTAGCCAACATGATCAAGATCCCACTTGATATAGTAGCTGATTTTAACACATGCCTTAGCTGATTTAACAACCGCCCGTTTCTTATTTTCGGGAATAC : 106
              CAATTGCAAGTAGCCAACATGATCAAGATCCCACTTGATATAGTAGCTGATTTTAACACATGCCTTAGCTGATTTAACAACCGCCCGTTTCTTATTTTCGGGAATAC

                   120         *       140         *       160         *       180         *       200         *       220
Prcmoter_R : GCAAAAGGCTTGCATCTCATTACACAACTGCCTATTGTTCCTATGGAGTAATTATATTTTGCAGTTGCAGCCATCAGTGTCACATCAAAAGTGCTAAGCAAAGAGGAGCA : 220
Prcmoter_S : GCAAAAGGCTTGCATCTCATTACACAACTGCCTATTGTTCCTATGGAGTAATTATATTTTGCAGTTGCAGCCATCAGTGTCACATCAAAAGTGCTAAGCAAAGAGGAGCA : 216
              GCAAAAGGCTTGCATCTCATTACACAACTGCCTATTGTTCCTATGGAGTAATTATATTTTGCAGTTGCAGCCATCAGTGTCACATCAAAAGTGCTAAGCAAAGAGGAGCA

                    *       240         *       260         *       280         *       300         *       320         *
Prcmoter_R : CCACCTAAGCACAACCGACAAAAGGCAGCTACTGCAACTTTTGGTCCTATGTAAAGATGGCCTAGAAGAACAGAAGCTGAGTGTCACACTCACACCAGAGCAAGACGAGA : 330
Prcmoter_S : CCACCTAAGCACAACCGACAAAAGGCAGCTACTGCAACTTTTGGTCCTATGTAAAGATGGCCTAGAAGAACAGAAGCTGAGTGTCACACTCACACCAGAGCAAGACGAGA : 326
              CCACCTAAGCACAACCGACAAAAGGCAGCTACTGCAACTTTTGGTCCTATGTAAAGATGGCCTAGAAGAACAGAAGCTGAGTGTCACACTCACACCAGAGCAAGACGAGA

                   340         *       360         *       380         *       400         *       420         *       440
Prcmoter_R : AAGAGATAAGCATTCATTTGCATGCACCAATTATGAAATGCAATTCCAAGGTAGGCATGGGCAAATGTGTATCATCACTTATACTTGGCACTAACTTGTTGAGGTGTGCA : 440
Prcmoter_S : AAGAGATAAGCATTCATTTGCATGCACCAATTATGAAATGCAATTCCAAGGTAGGCATGGGCAAATGTGTATCATCACTTATACTTGGCACTAACTTGTTGAGGTGTGCA : 436
              AAGAGATAAGCATTCATTTGCATGCACCAATTATGAAATGCAATTCCAAGGTAGGCATGGGCAAATGTGTATCATCACTTATACTTGGCACTAACTTGTTGAGGTGTGCA

                    *       460         *       480         *       500         *       520         *       540         *
Prcmoter_R : GACTGCAGGCTACAGGAATGTCCTACTCTTTCCCTGAGAAGGTGATCTGAATTCTGAATTCATCATACAAAAGGTTGTGCCTTTCTTACAGGGGATTTGCTTCTTTTGTG : 550
Prcmoter_S : GACTGCAGGCTACAGGAATGTCCTACTCTTTCCCTGAGAAGGTGATCTGAATTCTGAATTCATCATACAAAAGGTTGTGCCTTTCTTACAGGGGATTTGCTTCTTTTGTG : 546
              GACTGCAGGCTACAGGAATGTCCTACTCTTTCCCTGAGAAGGTGATCTGAATTCTGAATTCATCATACAAAAGGTTGTGCCTTTCTTACAGGGGATTTGCTTCTTTTGTG

                   560         *       580         *       600         *       620         *       640         *       660
Prcmoter_R : TGAGTACTTGAAGTGTAGTGGCCATATTGGGCTGGAAGAGGCAGCATACCAACAGTAAGCAGAGCTGTTGGTGCAGCAGTAACAGTAAACTGGTGAGTATAGCTTGATAC : 660
Prcmoter_S : TGAGTACTTGAAGTGTAGTGGCCATATTGGGCTGGAAGAGGCAGCATACCAACAGTAAGCAGAGCTGTTGGTGCAGCAGTAACAGTAAACTGGTGAGTATAGCTTGATAC : 656
              TGAGTACTTGAAGTGTAGTGGCCATATTGGGCTGGAAGAGGCAGCATACCAACAGTAAGCAGAGCTGTTGGTGCAGCAGTAACAGTAAACTGGTGAGTATAGCTTGATAC

                    *       680         *       700         *       720         *       740         *       760         *
Prcmoter_R : ATTGAGCACTGTTGGTTTGTGCCAGTGAAGCACATGTGGCTAGTTGCTGTGTGTGTGACAGTGCAAGGAGGCAGGCCACCCACTTGGCATCCCTCTTTGATTCCTCCTGT : 770
Prcmoter_S : ATTGAGCACTGTTGGTTTGTGCCAGTGAAGCACATGTGGCTAGTTGCTGTGTGTGTGACAGTGCAAGGAGGCAGGCCACCCACTTGGCATCCCTCTTTGATTCCTCCTGT : 766
              ATTGAGCACTGTTGGTTTGTGCCAGTGAAGCACATGTGGCTAGTTGCTGTGTGTGTGACAGTGCAAGGAGGCAGGCCACCCACTTGGCATCCCTCTTTGATTCCTCCTGT

                   780         *       800         *       820         *       840         *       860         *       880
Prcmoter_R : TCATTTGAGGCTCAGGTGGGGCACAGTGAGAAAGAGAGAGAGAGAGAGAGAGAGAGAGAGGCCATCTTGTGTGTGTTGGATTATTGTGAGTCTTGTCTTTCTTATC : 880
Prcmoter_S : TCATTTGAGGCTCAGGTGGGGCACAGTGAGAAAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGGCCATCTTGTGTGTGTTGGATTATTGTGAGTCTTGTCTTTCTTATC : 876
              TCATTTGAGGCTCAGGTGGGGCACAGTGAGAAAGAGAGAGAGAGAGAGAGAGAGAGAGAGGCCATCTTGTGTGTGTTGGATTATTGTGAGTCTTGTCTTTCTTATC

                    *       900         *       920         *       940         *       960         *       980         *
Prcmoter_R : TATAGTTGCTTTTCTTCATTGTCTTTTGGTTTAAGGTCCTCATTTTCCACCAGCGCTTGCCAGGGAAAGTAATTCTGGAGGAAGACACAGATTCTTGAGGGAAAAGAGGT : 990
Prcmoter_S : TATAGTTGCTTTTCTTCATTGTCTTTTGGTTTAAGGTCCTCATTTTCCACCAGCGCTTGCCAGGGAAAGTAATTCTGGAGGAAGACACAGATTCTTGAGGGAAAAGAGGT : 986
              TATAGTTGCTTTTCTTCATTGTCTTTTGGTTTAAGGTCCTCATTTTCCACCAGCGCTTGCCAGGGAAAGTAATTCTGGAGGAAGACACAGATTCTTGAGGGAAAAGAGGT

                  1000         *      1020         *      1040         *      1060         *      1080         *      1100
Prcmoter_R : GAGATTCTTGATTTCAGATACAAGAGGAGTAGAGTGGTTCTTTGTCCCCTAGCCCTAGGTAGGTGAAAAAAAGCTTCCTTTGCTAATTGCTATAGGTTTCTTTGTATAGG : 1100
Prcmoter_S : GAGATTCTTGATTTCAGATACAAGAGGAGTAGAGTGGTTCTTTGTCCCCTAGCCCTAGGTAGGTGAAAAAAAGCTTCCTTTGCTAATTGCTATAGGTTTCTTTGTATAGG : 1096
              GAGATTCTTGATTTCAGATACAAGAGGAGTAGAGTGGTTCTTTGTCCCCTAGCCCTAGGTAGGTGAAAAAAAGCTTCCTTTGCTAATTGCTATAGGTTTCTTTGTATAGG

                    *      1120         *      1140         *      1160         *      1180         *      1200         *
Prcmoter_R : TGCCCCACAGAAAAGAAAGAAAGAAGCTACTTGACTTGACTTGGTATAGGTTCTTTTCCTGCCCCTCTGATTTGATTTGCATATGCTGCTGCCACCATTGCAGTGCTGCT : 1210
Prcmoter_S : TGCCCCACAGAAAAGAAAGAAAGAAGCTACTTGACTTGACTTGGTATAGGTTCTTTTCCTGCCCCTCTGATTTGATTTGCATATGCTGCTGCCACCATTGCAGTGCTGCT : 1206
              TGCCCCACAGAAAAGAAAGAAAGAAGCTACTTGACTTGACTTGGTATAGGTTCTTTTCCTGCCCCTCTGATTTGATTTGCATATGCTGCTGCCACCATTGCAGTGCTGCT

                  1220         *      1240         *      1260         *      1280         *      1300         *      1320
Prcmoter_R : AACATGGAAGAAATAATGCAGCAGCCTTGAGGTCAGGTAGTAGTAGTTCTTGGTGGTGTGGAACTGTGAAAGATCTGAGCTTGTAAAATCTTTCATGGTGGTGAGAAGTG : 1320
Prcmoter_S : AACATGGAAGAAATAATGCAGCAGCCTTGAGGTCAGGTAGTAGTAGTTCTTGGTGGTGTGGAACTGTGAAAGATCTGAGCTTGTAAAATCTTTCATGGTGGTGAGAAGTG : 1316
              AACATGGAAGAAATAATGCAGCAGCCTTGAGGTCAGGTAGTAGTAGTTCTTGGTGGTGTGGAACTGTGAAAGATCTGAGCTTGTAAAATCTTTCATGGTGGTGAGAAGTG

                    *      1340         *      1360         *      1380         *      1400         *      1420         *
Prcmoter_R : AAAGTGTGAAACTGTGATAAGATTTGAGATTGTGAAATTTTGGGGGTGGTTCTCTTGTCATGATGTATAGTAGGGTCCAGCACTGGACTGGTTGATGTTGTTGGCTTCAA : 1430
Prcmoter_S : AAAGTGTGAAACTGTGATAAGATTTGAGATTGTGAAATTTTGGGGGTGGTTCTCTTGTCATGATGTATAGTAGGGTCCAGCACTGGACTGGTTGATGTTGTTGGCTTCAA : 1426
              AAAGTGTGAAACTGTGATAAGATTTGAGATTGTGAAATTTTGGGGGTGGTTCTCTTGTCATGATGTATAGTAGGGTCCAGCACTGGACTGGTTGATGTTGTTGGCTTCAA

                  1440         *      1460         *      1480         *      1500         *      1520         *      1540
Prcmoter_R : AGGTTAGGGATAGAGAAAGCCTGAAGAGGGAGAGGTGTAGTAGGATATAAGATCTGAGAAAGCCAGAGATGAGAGTGAGGGAGAAGGCAGTGGTGTGTGGCCAAGCTTGAAGTGC : 1540
Prcmoter_S : AGGTTAGGGATAGAGAAAGCCTGAAGAGGGAGAGGTGTAGTAGGATATAAGATCTGAGAAAGCCAGAGATGAGAGTGAGGGAGAAGGCAGTGGTGTGTGGCCAAGCTTGAAGTGC : 1536
              AGGTTAGGGATAGAGAAAGCCTGAAGAGGGAGAGGTGTAGTAGGATATAAGATCTGAGAAAGCCAGAGATGAGAGTGAGGGAGAAGGCAGTGGTGTGTGGCCAAGCTTGAAGTGC

                    *      1560         *      1580         *      1600         *      1620         *      1640         *
Prcmoter_R : TCCAGTGAGGCTGCTTCTCTGGCTGCCAAGCTGATGGACGCGCCGCCATCGCCAGATCGGCTCGCCGGATCAGTGGAGCTCCATGCTTCTCCAGGCTGCTGCTGATTGAG : 1650
Prcmoter_S : TCCAGTGAGGCTGCTTCTCTGGCTGCCAAGCTGATGGACGCGCCGCCATCGCCAGATCGGCTCGCCGGATCAGTGGAGCTCCATGCTTCTCCAGGCTGCTGCTGATTGAG : 1646
              TCCAGTGAGGCTGCTTCTCTGGCTGCCAAGCTGATGGACGCGCCGCCATCGCCAGATCGGCTCGCCGGATCAGTGGAGCTCCATGCTTCTCCAGGCTGCTGCTGATTGAG

                  1660         *      1680         *      1700         *      1720         *      1740         *      1760
Prcmoter_R : AGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAAACAGTTTTTTTTTTTCAGGAGGAATTCATCAGGAATCTGCTGTGCTCCTTTTCTCTGGGAGGAAGCAAAGAGC : 1760
Prcmoter_S : AGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAAACAGTTTTTTTTTTTCAGGAGGAATTCATCAGGAATCTGCTGTGCTCCTTTTCTCTGGGAGGAAGCAAAGAGC : 1756
              AGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAGAAACAGTTTTTTTTTTTCAGGAGGAATTCATCAGGAATCTGCTGTGCTCCTTTTCTCTGGGAGGAAGCAAAGAGC

                    *      1780         *      1800         *      1820         *      1840         *      1860         *
Prcmoter_R : AGGTACTTCACTACCCCCTCCCCCCCACTGTTGAGGTAGTTGTGTGCTTTGCCTTTCTTGCATCTGAAATGATGATGTTGCTGTTCTTGCCATGATTGATGATTCCCCTGATA : 1870
Prcmoter_S : AGGTACTTCACTACCCCCTCCCCCCCACTGTTGAGGTAGTTGTGTGCTTTGCCTTTCTTGCATCTGAAATGATGATGTTGCTGTTCTTGCCATGATTGATGATTCCCCTGATA : 1866
              AGGTACTTCACTACCCCCTCCCCCCCACTGTTGAGGTAGTTGTGTGCTTTGCCTTTCTTGCATCTGAAATGATGATGTTGCTGTTCTTGCCATGATTGATGATTCCCCTGATA

                  1880         *      1900         *      1920         *      1940         *      1960         *      1980
Prcmoter_R : AAGCTAAGCTAGAGTCTGCATATATGGGCATTCATTGATACTCTTGGTAGCACATCTTGTGGTGGTTGTACTTCTCTCTCAAACTTCAATGATGCACAGTTGAGTCAAGT : 1980
Prcmoter_S : AAGCTAAGCTAGAGTCTGCATATATGGGCATTCATTGATACTCTTGGTAGCACATCTTGTGGTGGTTGTACTTCTCTCTCAAACTTCAATGATGCACAGTTGAGTCAAGT : 1976
              AAGCTAAGCTAGAGTCTGCATATATGGGCATTCATTGATACTCTTGGTAGCACATCTTGTGGTGGTTGTACTTCTCTCTCAAACTTCAATGATGCACAGTTGAGTCAAGT

                    *      2000         *      2020         *      2040         *      2060         *      2080         *
Prcmoter_R : GTCAGCTATGCCATAAATCTGTCCTGATTTCTGCAACTTCTAATTGAGTTATATTTTCTTTTTGAGTATCTGTCCTTCTCATATTTCATTTTGTTGATTTAAAGGTTGGA : 2090
Prcmoter_S : GTCAGCTATGCCATAAATCTGTCCTGATTTCTGCAACTTCTAATTGAGTTATATTTTCTTTTTGAGTATCTGTCCTTCTCATATTTCATTTTGTTGATTTAAAGGTTGGA : 2086
              GTCAGCTATGCCATAAATCTGTCCTGATTTCTGCAACTTCTAATTGAGTTATATTTTCTTTTTGAGTATCTGTCCTTCTCATATTTCATTTTGTTGATTTAAAGGTTGGA

                  2100         *
Prcmoter_R : GCAGCAGCAGCTGGGATCTGTAACTGGC : 2118
Prcmoter_S : GCAGCAGCA------------------- : 2095
              GCAGCAGCA
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Promoter_R：水稻基因组数据库公布的 *OsSPL3* 启动子序列；Promoter_S：克隆到的 *OsSPL3* 启动子测序序列。
Promoter R: Promoter sequence of *OsSPL3* published in the Rice Genome Database; Promoter S: Promoter sequence of *OsSPL3* obtained by sequencing.

附图 1　*OsSPL3* 启动子测序序列与在水稻基因组数据库（Rice Genome Annotation Project）中公布序列的比对分析

Attached Fig. S1　Alignment analysis of promoter sequences of *OsSPL3* between obtained by sequencing and published in the Rice Genome Database - Rice Genome Annotation Project