



| | | |
|---|---|---|
| | | <div><div></div><div>145014601470148014901500151015201530</div></div> |
| | | TGTACTTAGCCTTTTATTTCATTGGGGTGAATTGCAAGTTCACCATAGTTTTTCCACACAACCAAAAAGGAAACTGGAAAAATGTTCCCTCTCAATT |
| G0299590-1.seq (1>5802) | → | TGTACTTAGCCTTTTATTTCATTGGGGTGAATTGCAAGTTCACCATAGTTTTTCCACACAACCAAAAAGGAAACTGGAAAAATGTTCCCTCTCAATT |
| B01-E75159-G0299590-1_CMV-F.ab1 (1>785) | → | TGTACTTAGCCTTTTATTTCATTGGGGTGAATTGCAAGTTCACCATAGTTTTTCCACACAACCAAAAAGGAAACTGGAAAAATGTTCCCTCTC |
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| | | ACCATTATTGCCCGTCAAGCTCAGATTTAAATTGGCATAATGACTTAATAGGCACAGCCATACAAGTCAAAATGCCCAAGAGTCACAAGGCTATTTC |
| G0299590-1.seq (1>5802) | → | ACCATTATTGCCCGTCAAGCTCAGATTTAAATTGGCATAATGACTTAATAGGCACAGCCATACAAGTCAAAATGCCCAAGAGTCACAAGGCTATTTC |
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| | | AAGCAGACGGTTGGATGTGTTCATGCTTCCAATGGGTCACTACTTGTGATTTCGGCTGGTATGACCGGAAGTATATAACACAGTCCATCCGATCCT |
| G0299590-1.seq (1>5802) | → | AAGCAGACGGTTGGATGTGTTCATGCTTCCAATGGGTCACTACTTGTGATTTCGGCTGGTATGACCGGAAGTATATAACACAGTCCATCCGATCCT |
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| | | TCATCCATCTGTAGAACAATGCAAGGAAAGCATTGAACAAACGAAACAAGGAACCTTGGCTGAATCCAGGCTTCCCTCCTCAAAGTTGTGGATATG |
| G0299590-1.seq (1>5802) | → | TCATCCATCTGTAGAACAATGCAAGGAAAGCATTGAACAAACGAAACAAGGAACCTTGGCTGAATCCAGGCTTCCCTCCTCAAAGTTGTGGATATG |
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| | | CAACTGTGACGGATGCCGAAGCAGTGATTGTCCAGGTGACTCCTCACCATGTGCTGGTTGATGAATACACAGGAGAATGGGTTGATTACAGTTCA |
| G0299590-1.seq (1>5802) | → | CAACTGTGACGGATGCCGAAGCAGTGATTGTCCAGGTGACTCCTCACCATGTGCTGGTTGATGAATACACAGGAGAATGGGTTGATTACAGTTCA |
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| | | TCAACGAAAAATGCAGCAATTACATATGCCCCACTGTCCATAACTCTACAACCTGGCATTCTGACTATAAGGTCAAAGGCTATGTGATTCTAACC |
| G0299590-1.seq (1>5802) | → | TCAACGAAAAATGCAGCAATTACATATGCCCCACTGTCCATAACTCTACAACCTGGCATTCTGACTATAAGGTCAAAGGCTATGTGATTCTAACC |
| | | <div><div></div><div>2020203020402050206020702080209021002110</div></div> |
| | | TCATTTCCATGGACATCACCTTCTTCTCAGAGGACGGAGAGCTATCATCCCTGGGAAAGGAGGGCACAGGGTTCAGAAGTAAC TACTTTGCTTATG |
| G0299590-1.seq (1>5802) | → | TCATTTCCATGGACATCACCTTCTTCTCAGAGGACGGAGAGCTATCATCCCTGGGAAAGGAGGGCACAGGGTTCAGAAGTAAC TACTTTGCTTATG |
| | | <div><div></div><div>212021302140215021602170218021902200</div></div> |
| | | AAACTGGAGGCAAGGCCTGCAAAATGCAATACTGCAAGCATTGGGGAGTCAGACTCCCATCAGGTGTCTGGTTCGAGATGGCTGATAAGGATCTCT |
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| | | <div><div></div><div>2210222022302240225022602270228022902300</div></div> |
| | | TTGCTGCAGCCGATTCCCTGAATGCCCAGAAGGGTCAAGTATCTCTGCTCCATCTCAGACCTCAGTGGATGTAAGTCTAATT CAGGACGTTGAGA |
| G0299590-1.seq (1>5802) | → | TTGCTGCAGCCGATTCCCTGAATGCCCAGAAGGGTCAAGTATCTCTGCTCCATCTCAGACCTCAGTGGATGTAAGTCTAATT CAGGACGTTGAGA |
| | | <div><div></div><div>2310232023302340235023602370238023902400</div></div> |
| | | GGATCTTGATTATTCCCTCTGCCAAGAAACCTGGAGCAAAATCAGAGCGGGTCTTCCAATCTCTCCAGTGGATCTCAGCTATCTTGCTCCTAAAA |
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| | | ACCCAGGAACCGTCTGTCTTTCACCATAATCAATGGTACCCTAAAATACTTTGAGACCAGATACATCAGAGTCGATATTGCTGCTCCAATCCTCT |
| G0299590-1.seq (1>5802) | → | ACCCAGGAACCGTCTGTCTTTCACCATAATCAATGGTACCCTAAAATACTTTGAGACCAGATACATCAGAGTCGATATTGCTGCTCCAATCCTCT |
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| | | CAAGAAATGGTCGGAATGATCAGTGGAACTACCACAGAAAGGGAACGTGGGATGACTGGGCACCATATGAAGACGTGGAAATTGGACCCAATGGAG |
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| | | TTCTGAGGACCAGTTCAGGATATAAGTTTCCTTTATACATGATTGGACATGGTATGTTGGACTCCGATCTTCATCTTAGCTCAAAGGCTCAGGTGT |
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| | | TCGAACATCCTCACATTCAAGACGCTGCTTCGCAACTTCCTGATGATGAGAGTTTATTTTTTGGTGATACTGGGCTATCCAAAAATCCAATCGAGC |
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| | | <div><div></div><div>289029002910292029302940295029602970</div></div> |
| | | GTATCCATCTTTGCATTAAATTAAAGCACACCAAGAAAAGACAGATTTATACAGACATAGAGATGAACCGACTTGGAAAGTAAC TCAAATCCTGCA |
| G0299590-1.seq (1>5802) | → | GTATCCATCTTTGCATTAAATTAAAGCACACCAAGAAAAGACAGATTTATACAGACATAGAGATGAACCGACTTGGAAAGTAAC TCAAATCCTGCA |



