

Formatted Alignments

H5 245467 MiSeq	1	ATGGAGAACATAGTACTACTTCTTGCAATAGTTAGCCTTGTTAAAAGTGATCAGATTTC	60
H5 245467 Method A	1	ATGGAGAACATAGTACTACTTCTTGCAATAGTTAGCCTTGTTAAAAGTGATCAGATTTC	60
H5 245467 Method S	1	ATGGAGAACATAGTACTTCTTCTTGCAATAGTTAGCCTTGTTAAAAGTGATCAGATTTC	60
H5 245467 Method E	1	ATGGAGAACATAGTACTACTTCTTGCAATAGTTAGCCTTGTTAAAAGTGATCAGATTTC	60
H5 245467 Method K	1	ATGGAGAACATAGTACTACTTCTTGCAATAGTTAGCCTTGTTAAAAGTGATCAGATTTC	60
H5 245467 Method N	1	ATGGAGAACATAGTACTACTTCTTGCAATAGTTAGCCTTGTTAAAAGTGATCAGATTTC	60

H5 245467 MiSeq	61	ATTGGTTACCATGCAAACAATTCGACAGAGCAAGTTGACACGATAATGGAAAAGAATGTT	120
H5 245467 Method A	61	ATTGGTTACCATGCAAACAATTCGACAGAGCAAGTTGACACGATAATGGAAAAGAATGTT	120
H5 245467 Method S	61	ATTGGTTACCATGCAAACAATTCGACAGAGCAAGTTGACACGATAATGGAAAAGAACTC	120
H5 245467 Method E	61	ATTGGTTACCATGCAAACAATTCGACAGAGCAAGTTGACACGATAATGGAAAAGAATGTT	120
H5 245467 Method K	61	ATTGGTTACCATGCAAACAATTCGACAGAGCAAGTTGACACGATAATGGAAAAGAATGTT	120
H5 245467 Method N	61	ATTGGTTACCATGCAAACAATTCGACAGAGCAAGTTGACACGATAATGGAAAAGAATGTT	120

H5 245467 MiSeq	121	ACTGTTACACATGCCCAAGACATACTGGAAAAAACACACAACGGGAAGCTCTGTGATCTA	180
H5 245467 Method A	121	ACTGTTACACATGCCCAAGACATACTGGAAAAAACACACAACGGGAAGCTCTGTGATCTA	180
H5 245467 Method S	121	ACTGTTACACATGCCCAAGACATACTGGAAAAAACACACAACGGGAAGCTCTGTGATCTA	180
H5 245467 Method E	121	ACTGTTACACATGCCCAAGACATACTGGAAAAAACACACAACGGGAAGCTCTGTGATCTA	180
H5 245467 Method K	121	ACTGTTACACATGCCCAAGACATACTGGAAAAAACACACAACGGGAAGCTCTGTGATCTA	180
H5 245467 Method N	121	ACTGTTACACATGCCCAAGACATACTGGAAAAAACACACAACGGGAAGCTCTGTGATCTA	180

H5 245467 MiSeq	181	AATGGGGTGAAGCCTCTGATTTTAAAGGATTGTAGTGTAGCTGGATGGCTCCTCGGAAAC	240
H5 245467 Method A	181	AATGGGGTGAAGCCTCTGATTTTAAAGGATTGTAGTGTAGCTGGATGGCTCCTCGGAAAC	240
H5 245467 Method S	181	AATGGGGTGAAGCCTCTGATTTTAAAGGATTGTAGTGTAGCTGGATGGCTCCTCGGAAAC	240
H5 245467 Method E	181	AATGGGGTGAAGCCTCTGATTTTAAAGGATTGTAGTGTAGCTGGATGGCTCCTCGGAAAC	240
H5 245467 Method K	181	AATGGGGTGAAGCCTCTGATTTTAAAGGATTGTAGTGTAGCTGGATGGCTCCTCGGAAAC	240
H5 245467 Method N	181	AATGGGGTGAAGCCTCTGATTTTAAAGGATTGTAGTGTAGCTGGATGGCTCCTCGGAAAC	240

H5 245467 MiSeq	241	CCAATGTGCGACGAATTCATCAGAGTGCCTGAATGGTCATACATAGTGGAGCGGGCTAAC	300
H5 245467 Method A	241	CCAATGTGCGACGAATTCATCAGAGTGCCTGAATGGTCATACATAGTGGAGCGGGCTAAC	300
H5 245467 Method S	241	CCAATGTGCGACGAATTCATCAGAGTGCCTGAATGGTCATACATAGTGGAGCGGGCTAACT	300
H5 245467 Method E	241	CCAATGTGCGACGAATTCATCAGAGTGCCTGAATGGTCATACATAGTGGAGCGGGCTAAC	300
H5 245467 Method K	241	CCAATGTGCGACGAATTCATCAGAGTGCCTGAATGGTCATACATAGTGGAGCGGGCTAAC	300
H5 245467 Method N	241	CCAATGTGCGACGAATTCATCAGAGTGCCTGAATGGTCATACATAGTGGAGCGGGCTAAC	300

H5 245467 MiSeq	301	CCAGCTAATGACCTCTGTTACCCAGGGAGCCTCAATGACTATGAAGAACTGAAACACATG	360
H5 245467 Method A	301	CCAGCTAATGACCTCTGTTACCCAGGGAGCCTCAATGACTATGAAGAACTGAAACACATG	360
H5 245467 Method S	301	CCAGCTAATGACCTCTGTTACCCAGGGAGTCTCAATGACTATGAAGAACTGAAACACCTG	360
H5 245467 Method E	301	CCAGCTAATGACCTCTGTTACCCAGGGAGCCTCAATGACTATGAAGAACTGAAACACATG	360
H5 245467 Method K	301	CCAGCTAATGACCTCTGTTACCCAGGGAGCCTCAATGACTATGAAGAACTGAAACACATG	360
H5 245467 Method N	301	CCAGCTAATGACCTCTGTTACCCAGGGAGCCTCAATGACTATGAAGAACTGAAACACATG	360

H5 245467 MiSeq	361	TTGAGCAGAATAAATCATTTTGTGAGAAGATTCTGATCATCCCCAAGAGTTCCTGGCCAAAT	420
H5 245467 Method A	361	TTGAGCAGAATAAATCATTTTGTGAGAAGATTCTGATCATCCCCAAGAGTTCCTGGCCAAAT	420
H5 245467 Method S	361	TTGAGCAGAATAAATCATTTTGTGAGAAGATTCTGATCATCCCCAAGAGTTCCTGGCCAAAT	420
H5 245467 Method E	361	TTGAGCAGAATAAATCATTTTGTGAGAAGATTCTGATCATCCCCAAGAGTTCCTGGCCAAAT	420
H5 245467 Method K	361	TTGAGCAGAATAAATCATTTTGTGAGAAGATTCTGATCATCCCCAAGAGTTCCTGGCCAAAT	420
H5 245467 Method N	361	TTGAGCAGAATAAATCATTTTGTGAGAAGATTCTGATCATCCCCAAGAGTTCCTGGCCAAAT	420

H5 245467 MiSeq	421	CATGAAACATCACTAGGGGTGAGCGCAGCTTGTCCATACCAGGGAGCGCCCTCCTTTTTTC	480
H5 245467 Method A	421	CATGAAACATCACTAGGGGTGAGCGCAGCTTGTCCATACCAGGGAGCGCCCTCCTTTTTTC	480
H5 245467 Method S	421	CATGAAACATCACTAGGGGTGAGCGCAGCTTGTCCATACCAGGGAGCGCCCTCCTTTTTTC	480
H5 245467 Method E	421	CATGAAACATCACTAGGGGTGAGCGCAGCTTGTCCATACCAGGGAGCGCCCTCCTTTTTTC	480
H5 245467 Method K	421	CATGAAACATCACTAGGGGTGAGCGCAGCTTGTCCATACCAGGGAGCGCCCTCCTTTTTTC	480
H5 245467 Method N	421	CATGAAACATCACTAGGGGTGAGCGCAGCTTGTCCATACCAGGGAGCGCCCTCCTTTTTTC	480

H5 245467 MiSeq	481	AGAAATGTGGTGTGGCTTATCAAAAAGAACGATGCATACCCAACAATAAAGATAAGCTAC	540
H5 245467 Method A	481	AGAAATGTGGTGTGGCTTATCAAAAAGAACGATGCATACCCAACAATAAAGATAAGCTAC	540
H5 245467 Method S	481	AGAAATGTGGTGTGGCTTATCAAAAAGAACGATGCATACCCAACAATAAAGATAAGCTAC	540
H5 245467 Method E	481	AGAAATGTGGTGTGGCTTATCAAAAAGAACGATGCATACCCAACAATAAAGATAAGCTAC	540
H5 245467 Method K	481	AGAAATGTGGTGTGGCTTATCAAAAAGAACGATGCATACCCAACAATAAAGATAAGCTAC	540
H5 245467 Method N	481	AGAAATGTGGTGTGGCTTATCAAAAAGAACGATGCATACCCAACAATAAAGATAAGCTAC	540

H5 245467 MiSeq	541	AATAATACCAATCGGGAAGATCTCTTGATACTGTGGGGGATTTCATCATTCCAACAATGCA	600
H5 245467 Method A	541	AATAATACCAATCGGGAAGATCTCTTGATACTGTGGGGGATTTCATCATTCCAACAATGCA	600
H5 245467 Method S	541	AATAATACCAATC A GGGAAGATCTCTTGATACTGTGGGGGATTTCATCATTCCAACAATGCA	600
H5 245467 Method E	541	AATAATACCAATCGGGAAGATCTCTTGATACTGTGGGGGATTTCATCATTCCAACAATGCA	600
H5 245467 Method K	541	AATAATACCAATCGGGAAGATCTCTTGATACTGTGGGGGATTTCATCATTCCAACAATGCA	600
H5 245467 Method N	541	AATAATACCAATCGGGAAGATCTCTTGATACTGTGGGGGATTTCATCATTCCAACAATGCA	600

H5 245467 MiSeq	601	GAAGAGCAGACAAATCTCTACAAAAACCCAACCACCTACATTTTCAGTTGGAACATCAACT	660
H5 245467 Method A	601	GAAGAGCAGACAAATCTCTACAAAAACCCAACCACCTACATTTTCAGTTGGAACATCAACT	660
H5 245467 Method S	601	GAAGAGCAGACAAATCTCTA T AAAAACCCAACCACCTACATTTTCAGTTGGAACATCAACT	660
H5 245467 Method E	601	GAAGAGCAGACAAATCTCTACAAAAACCCAACCACCTACATTTTCAGTTGGAACATCAACT	660
H5 245467 Method K	601	GAAGAGCAGACAAATCTCTACAAAAACCCAACCACCTACATTTTCAGTTGGAACATCAACT	660
H5 245467 Method N	601	GAAGAGCAGACAAATCTCTACAAAAACCCAACCACCTACATTTTCAGTTGGAACATCAACT	660

H5 245467 MiSeq	661	TTAAACCAGAGGTTGGCACCAAAAATAGCTACTAGATCCCAAGTAAACGGGGCAACGTGGA	720
H5 245467 Method A	661	TTAAACCAGAGGTTGGCACCAAAAATAGCTACTAGATCCCAAGTAAACGGGGCAACGTGGA	720
H5 245467 Method S	661	TTAAACCAGAGGTTGG T ACCAAAAATAGCTACTAGATCCCAAGTAAACGGGGCAACGTGGA	720
H5 245467 Method E	661	TTAAACCAGAGGTTGGCACCAAAAATAGCTACTAGATCCCAAGTAAACGGGGCAACGTGGA	720
H5 245467 Method K	661	TTAAACCAGAGGTTGGCACCAAAAATAGCTACTAGATCCCAAGTAAACGGGGCAACGTGGA	720
H5 245467 Method N	661	TTAAACCAGAGGTTGGCACCAAAAATAGCTACTAGATCCCAAGTAAACGGGGCAACGTGGA	720

H5 245467 MiSeq	721	AGAATGGACTTCTTCTGGACAATCTTAAAACCAGATGATGCAATCCATTTTCGAGAGTAAT	780
H5 245467 Method A	721	AGAATGGACTTCTTCTGGACAATCTTAAAACCAGATGATGCAATCCATTTTCGAGAGTAAT	780
H5 245467 Method S	721	AGAATGGACTTCTTCTGGACAAT T TTAAAACC G GATGATGCAATCCATTTTCGAGAGTAAT	780
H5 245467 Method E	721	AGAATGGACTTCTTCTGGACAATCTTAAAACCAGATGATGCAATCCATTTTCGAGAGTAAT	780
H5 245467 Method K	721	AGAATGGACTTCTTCTGGACAATCTTAAAACCAGATGATGCAATCCATTTTCGAGAGTAAT	780
H5 245467 Method N	721	AGAATGGACTTCTTCTGGACAATCTTAAAACCAGATGATGCAATCCATTTTCGAGAGTAAT	780

H5 245467 MiSeq	781	GGAAATTTTCATTGCTCCAGAATATGCATACAAAATTGTCAAGAAAGGGGACTCAACAATT	840
H5 245467 Method A	781	GGAAATTTTCATTGCTCCAGAATATGCATACAAAATTGTCAAGAAAGGGGACTCAACAATT	840
H5 245467 Method S	781	GGAAATTTTCATTGCTCCAGAATATGCATACAAAATTGTCAAGAAAGGGGACTCAACAATT	840
H5 245467 Method E	781	GGAAATTTTCATTGCTCCAGAATATGCATACAAAATTGTCAAGAAAGGGGACTCAACAATT	840
H5 245467 Method K	781	GGAAATTTTCATTGCTCCAGAATATGCATACAAAATTGTCAAGAAAGGGGACTCAACAATT	840
H5 245467 Method N	781	GGAAATTTTCATTGCTCCAGAATATGCATACAAAATTGTCAAGAAAGGGGACTCAACAATT	840

H5 245467 MiSeq	841	ATGAAAAGTGGAGTGGAATATGGCCACTGCAACACCAAATGTCAAACCCCAGTAGGTGCG	900
H5 245467 Method A	841	ATGAAAAGTGGAGTGGAATATGGCCACTGCAACACCAAATGTCAAACCCCAGTAGGTGCG	900
H5 245467 Method S	841	ATGAAAAGTGGAGTGGAATATGGCCACTGCAACACCAAATGTCAAACCCCAGTAGG A GCG	900
H5 245467 Method E	841	ATGAAAAGTGGAGTGGAATATGGCCACTGCAACACCAAATGTCAAACCCCAGTAGGTGCG	900
H5 245467 Method K	841	ATGAAAAGTGGAGTGGAATATGGCCACTGCAACACCAAATGTCAAACCCCAGTAGGTGCG	900
H5 245467 Method N	841	ATGAAAAGTGGAGTGGAATATGGCCACTGCAACACCAAATGTCAAACCCCAGTAGGTGCG	900

H5 245467 MiSeq	901	ATAAATTCTAGTATGCCATTCCACAACATACATCCTCTCACCATTGGGGAATGCCCCAAA	960
H5 245467 Method A	901	ATAAATTCTAGTATGCCATTCCACAACATACATCCTCTCACCATTGGGGAATGCCCCAAA	960
H5 245467 Method S	901	ATAAATTCTAGTATGCCATTCCACAACATACATCCTCTCACCATTGGGGAATGCCCCAAA	960
H5 245467 Method E	901	ATAAATTCTAGTATGCCATTCCACAACATACATCCTCTCACCATTGGGGAATGCCCCAAA	960
H5 245467 Method K	901	ATAAATTCTAGTATGCCATTCCACAACATACATCCTCTCACCATTGGGGAATGCCCCAAA	960
H5 245467 Method N	901	ATAAATTCTAGTATGCCATTCCACAACATACATCCTCTCACCATTGGGGAATGCCCCAAA	960

H5 245467 MiSeq	961	TACGTGAAGTCAAACAAGTTGGTCCTTGCGACTGGGCTCAGAAATAGTCCTCTAAGAGAA	1020
H5 245467 Method A	961	TACGTGAAGTCAAACAAGTTGGTCCTTGCGACTGGGCTCAGAAATAGTCCTCTAAGAGAA	1020
H5 245467 Method S	961	TACGTGAAGTCAAACAAGTTGGTCCTTGCGACTGGGCTCAGAAATAGTCCTCTAAGAGAA	1020
H5 245467 Method E	961	TACGTGAAGTCAAACAAGTTGGTCCTTGCGACTGGGCTCAGAAATAGTCCTCTAAGAGAA	1020
H5 245467 Method K	961	TACGTGAAGTCAAACAAGTTGGTCCTTGCGACTGGGCTCAGAAATAGTCCTCTAAGAGAA	1020
H5 245467 Method N	961	TACGTGAAGTCAAACAAGTTGGTCCTTGCGACTGGGCTCAGAAATAGTCCTCTAAGAGAA	1020

H5 245467 MiSeq	1021	AAGAGAAGAAAAAGAGGCCTGTTTGGGGCGATAGCAGGGTTTATAGAGGGAGGATGGCAG	1080
H5 245467 Method A	1021	AAGAGAAGAAAAAGAGGCCTGTTTGGGGCGATAGCAGGGTTTATAGAGGGAGGATGGCAG	1080
H5 245467 Method S	1021	AAGAGAAGAAAAAGAGGCCTGTTTGGGGCGATAGCAGGGTTTATAGAGGGAGGATGGCAG	1080
H5 245467 Method E	1021	AAGAGAAGAAAAAGAGGCCTGTTTGGGGCGATAGCAGGGTTTATAGAGGGAGGATGGCAG	1080
H5 245467 Method K	1021	AAGAGAAGAAAAAGAGGCCTGTTTGGGGCGATAGCAGGGTTTATAGAGGGAGGATGGCAG	1080
H5 245467 Method N	1021	AAGAGAAGAAAAAGAGGCCTGTTTGGGGCGATAGCAGGGTTTATAGAGGGAGGATGGCAG	1080

H5 245467 MiSeq	1081	GGAATGGTTGATGGTTGGTATGGGTACCATCATAGCAATGAGCAGGGGAGTGGGTACGCT	1140
H5 245467 Method A	1081	GGAATGGTTGATGGTTGGTATGGGTACCATCATAGCAATGAGCAGGGGAGTGGGTACGCT	1140
H5 245467 Method S	1081	GGAATGGTTGATGGTTGGTATGGGTACCATCATAGCAATGAGCAGGGGAGTGGGTACGCT	1140
H5 245467 Method E	1081	GGAATGGTTGATGGTTGGTATGGGTACCATCATAGCAATGAGCAGGGGAGTGGGTACGCT	1140
H5 245467 Method K	1081	GGAATGGTTGATGGTTGGTATGGGTACCATCATAGCAATGAGCAGGGGAGTGGGTACGCT	1140
H5 245467 Method N	1081	GGAATGGTTGATGGTTGGTATGGGTACCATCATAGCAATGAGCAGGGGAGTGGGTACGCT	1140

H5 245467 MiSeq	1141	GCGGACAAAGAATCCACCCAAAAGGCAATAGATGGAGTTACCAATAAGGTCAACTCAATT	1200
H5 245467 Method A	1141	GCGGACAAAGAATCCACCCAAAAGGCAATAGATGGAGTTACCAATAAGGTCAACTCAATT	1200
H5 245467 Method S	1141	GC[A]GACAAAGAATCCACCCAAAAGGCAATAGATGGAGTTACCAATAAGGTCAACTCAAT[C]	1200
H5 245467 Method E	1141	GCGGACAAAGAATCCACCCAAAAGGCAATAGATGGAGTTACCAATAAGGTCAACTCAATT	1200
H5 245467 Method K	1141	GCGGACAAAGAATCCACCCAAAAGGCAATAGATGGAGTTACCAATAAGGTCAACTCAATT	1200
H5 245467 Method N	1141	GCGGACAAAGAATCCACCCAAAAGGCAATAGATGGAGTTACCAATAAGGTCAACTCAATT	1200

H5 245467 MiSeq	1201	ATTGACAAAATGAACACTCAATTTGAGGCAGTTGGAAGGGAGTTTAAATAACTTAGAAAGG	1260
H5 245467 Method A	1201	ATTGACAAAATGAACACTCAATTTGAGGCAGTTGGAAGGGAGTTTAAATAACTTAGAAAGG	1260
H5 245467 Method S	1201	ATTGACAAAATGAACACTCAATTTGAGGCAGTTGGAAGGGAGTTTAAATAACTTAGAAAGG	1260
H5 245467 Method E	1201	ATTGACAAAATGAACACTCAATTTGAGGCAGTTGGAAGGGAGTTTAAATAACTTAGAAAGG	1260
H5 245467 Method K	1201	ATTGACAAAATGAACACTCAATTTGAGGCAGTTGGAAGGGAGTTTAAATAACTTAGAAAGG	1260
H5 245467 Method N	1201	ATTGACAAAATGAACACTCAATTTGAGGCAGTTGGAAGGGAGTTTAAATAACTTAGAAAGG	1260

H5 245467 MiSeq	1261	AGGATAGAGAATTTGAACAAGAAAATGGAAGACGGATTCCCTAGATGTCTGGACCTATAAT	1320
H5 245467 Method A	1261	AGGATAGAGAATTTGAACAAGAAAATGGAAGACGGATTCCCTAGATGTCTGGACCTATAAT	1320
H5 245467 Method S	1261	AGGATAGAGAATTTGAACAAGAAAATGGAAGACGGATTCCCTAGATGTCTGGACCTATAAT	1320
H5 245467 Method E	1261	AGGATAGAGAATTTGAACAAGAAAATGGAAGACGGATTCCCTAGATGTCTGGACCTATAAT	1320
H5 245467 Method K	1261	AGGATAGAGAATTTGAACAAGAAAATGGAAGACGGATTCCCTAGATGTCTGGACCTATAAT	1320
H5 245467 Method N	1261	AGGATAGAGAATTTGAACAAGAAAATGGAAGACGGATTCCCTAGATGTCTGGACCTATAAT	1320

H5 245467 MiSeq	1321	GCTGAACTTCTAGTTCTCATGGAAAACGAGAGGACTCTAGATTTCCATGATTCAAATGTC	1380
H5 245467 Method A	1321	GCTGAACTTCTAGTTCTCATGGAAAACGAGAGGACTCTAGATTTCCATGATTCAAATGTC	1380
H5 245467 Method S	1321	GCTGAACTTCTAGTTCTCATGGAAAACGAGAGGACTCTAGATTTCCATGATTCAAATGTC	1380
H5 245467 Method E	1321	GCTGAACTTCTAGTTCTCATGGAAAACGAGAGGACTCTAGATTTCCATGATTCAAATGTC	1380
H5 245467 Method K	1321	GCTGAACTTCTAGTTCTCATGGAAAACGAGAGGACTCTAGATTTCCATGATTCAAATGTC	1380
H5 245467 Method N	1321	GCTGAACTTCTAGTTCTCATGGAAAACGAGAGGACTCTAGATTTCCATGATTCAAATGTC	1380

H5 245467 MiSeq	1381	AAGAACCTTTACGACAAAGTCAGATTACAGCTTAGGGATAATGCAAAGGAGCTGGGTAAAC	1440
H5 245467 Method A	1381	AAGAACCTTTACGACAAAGTCAGATTACAGCTTAGGGATAATGCAAAGGAGCTGGGTAAAC	1440
H5 245467 Method S	1381	AAGAACCTTTACGACAAAGTCAGATCAGCTTAGGGATAATGCAAAGGAGCTGGGTAAAC	1440
H5 245467 Method E	1381	AAGAACCTTTACGACAAAGTCAGATTACAGCTTAGGGATAATGCAAAGGAGCTGGGTAAAC	1440
H5 245467 Method K	1381	AAGAACCTTTACGACAAAGTCAGATTACAGCTTAGGGATAATGCAAAGGAGCTGGGTAAAC	1440
H5 245467 Method N	1381	AAGAACCTTTACGACAAAGTCAGATTACAGCTTAGGGATAATGCAAAGGAGCTGGGTAAAC	1440

H5 245467 MiSeq	1441	GGCTGTTTCGAATTCTATCACAAATGTGATAATGAATGTATGGAAAGTGTGAGAAATGGG	1500
H5 245467 Method A	1441	GGCTGTTTCGAATTCTATCACAAATGTGATAATGAATGTATGGAAAGTGTGAGAAATGGG	1500
H5 245467 Method S	1441	GGCTGTTTCGAATTCTATCACAAATG C GATAATGAATGTATGGAAAGTGTGAGAAATGGG	1500
H5 245467 Method E	1441	GGCTGTTTCGAATTCTATCACAAATGTGATAATGAATGTATGGAAAGTGTGAGAAATGGG	1500
H5 245467 Method K	1441	GGCTGTTTCGAATTCTATCACAAATGTGATAATGAATGTATGGAAAGTGTGAGAAATGGG	1500
H5 245467 Method N	1441	GGCTGTTTCGAATTCTATCACAAATGTGATAATGAATGTATGGAAAGTGTGAGAAATGGG	1500

H5 245467 MiSeq	1501	ACATATGACTACCCTCAGTATTCAGAAGAAGCAAGATTAAAAAGAGAAGAAATAAGCGGA	1560
H5 245467 Method A	1501	ACATATGACTACCCTCAGTATTCAGAAGAAGCAAGATTAAAAAGAGAAGAAATAAGCGGA	1560
H5 245467 Method S	1501	AC C TATGACTACCCTCAGTATTCAGAAGAAGCAAGATTAAAAAGAGAAGAAATAAGCGGA	1560
H5 245467 Method E	1501	ACATATGACTACCCTCAGTATTCAGAAGAAGCAAGATTAAAAAGAGAAGAAATAAGCGGA	1560
H5 245467 Method K	1501	ACATATGACTACCCTCAGTATTCAGAAGAAGCAAGATTAAAAAGAGAAGAAATAAGCGGA	1560
H5 245467 Method N	1501	ACATATGACTACCCTCAGTATTCAGAAGAAGCAAGATTAAAAAGAGAAGAAATAAGCGGA	1560

H5 245467 MiSeq	1561	GTGAAATTAGAATCAGTAGGAACTTACCAGATACTGTCAATTTATTCAACAGCGGCAAGT	1620
H5 245467 Method A	1561	GTGAAATTAGAATCAGTAGGAACTTACCAGATACTGTCAATTTATTCAACAGCGGCAAGT	1620
H5 245467 Method S	1561	GTGAAATTAGAATCA A TAGGAACTTACCAGATACTGTCAATTTATTCAACAGCGGC C AGT	1620
H5 245467 Method E	1561	GTGAAATTAGAATCAGTAGGAACTTACCAGATACTGTCAATTTATTCAACAGCGGCAAGT	1620
H5 245467 Method K	1561	GTGAAATTAGAATCAGTAGGAACTTACCAGATACTGTCAATTTATTCAACAGCGGCAAGT	1620
H5 245467 Method N	1561	GTGAAATTAGAATCAGTAGGAACTTACCAGATACTGTCAATTTATTCAACAGCGGCAAGT	1620

H5 245467 MiSeq	1621	TCCCTAGCACTGGCAATCATGATGGCTGGTCTATCTTTATGGATGTGCTCCAATGGGTCG	1680
H5 245467 Method A	1621	TCCCTAGCACTGGCAATCATGATGGCTGGTCTATCTTTATGGATGTGCTCCAATGGGTCG	1680
H5 245467 Method S	1621	TCCCTAGCACTGGCAATCATGATGGCTGGTCTATCTTTATGGATGTGCTCCAATGGGTCG	1680
H5 245467 Method E	1621	TCCCTAGCACTGGCAATCATGATGGCTGGTCTATCTTTATGGATGTGCTCCAATGGGTCG	1680
H5 245467 Method K	1621	TCCCTAGCACTGGCAATCATGATGGCTGGTCTATCTTTATGGATGTGCTCCAATGGGTCG	1680
H5 245467 Method N	1621	TCCCTAGCACTGGCAATCATGATGGCTGGTCTATCTTTATGGATGTGCTCCAATGGGTCG	1680

H5 245467 MiSeq	1681	TTACAGTGCAGAAATTTGCATTTAG	1704
H5 245467 Method A	1681	TTACAGTGCAGAAATTTGCATTTAG	1704
H5 245467 Method S	1681	TTACAGTGCAGAAATTTGCATTTA	1704
H5 245467 Method E	1681	TTACAGTGCAGAAATTTGCATTTAG	1704
H5 245467 Method K	1681	TTACAGTGCAGAAATTTGCATTTAG	1704
H5 245467 Method N	1681	TTACAGTGCAGAAATTTGCATTTAG	1704