

61 SRR1974789\_2\_prinseq\_good\_AYNE\_fastqc.html

Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTACCAGCTCGATAATTG	2116
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGCACTACACATGCTAGC	2036

62. SRR1974753\_2\_prinseq\_good\_FCMA\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	6367	0.24219374967429183
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	4956	0.1885208455137098
CGTCGCATTGGATGACGATGCCTAGTACTGTGCGCCAATTAGGTCGTCAT	2726	0.10369407281484522

63. SRR1974753\_1\_prinseq\_good\_Ov9m\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	7344	0.2793577662333908
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	4988	0.18973809068248274
CGTCGCATTGGATGACGATGCCTAGTACTGTGCGCCAATTAGGTCGTCAT	2744	0.10437877322228

64. SRR1974744\_2\_prinseq\_good\_g9p0\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	7176	0.13169144781864017

65. SRR1974744\_1\_prinseq\_good\_7bxh\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	8834	0.16211848523270167
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	5577	0.10234715781557362

66. SRR1974734\_2\_prinseq\_good\_Zn0j\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	4129	0.10301770262788704

67. SRR1974734\_1\_prinseq\_good\_XlrH\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	4976	0.12415017880270426

68. SRR1974727\_2\_prinseq\_good\_pER8\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	8456	0.1676356865767746
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	6693	0.1326851525849518
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	6439	0.12764973815844982

69. SRR1974727\_1\_prinseq\_good\_wP82\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	9630	0.19090960994966172
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	7354	0.1457891247736046
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	6522	0.1292951688568737
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	5114	0.10138232038240604

70. SRR1974721\_2\_prinseq\_good\_bdl\_\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	6493	0.13266241389610692
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	5044	0.10305701766394014

71. SRR1974721\_1\_prinseq\_good\_w0Do\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	7414	0.15147992247431644
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	5435	0.11104577537738196

72. SRR1974717\_1\_prinseq\_good\_FA9k\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	6185	0.10565336914672221

73. SRR1974716\_1\_prinseq\_good\_o5Fi\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	6299	0.10771577741156199

74. SRR1974711\_2\_prinseq\_good\_2g\_a\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	11412	0.20654250346816574
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	8645	0.15646336684913184

75. SRR1974711\_1\_prinseq\_good\_d08p\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	12874	0.2330028206842942
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	8733	0.15805605352151167

76. SRR1974708\_2\_prinseq\_good\_T5vR\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	7304	0.11240194838251381

77. SRR1974708\_1\_prinseq\_good\_965w\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	8120	0.12495944973521525

78. SRR1974702\_2\_prinseq\_good\_p8zj\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	9681	0.18487779465101428
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	7989	0.15256571650314565
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	5417	0.10344830220272123

79. SRR1974702\_1\_prinseq\_good\_\_9cT\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	12474	0.23821563996247827
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	8188	0.15636601411037132
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	5237	0.10001084708060755

80. SRR1974696\_1\_prinseq\_good\_DD17\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	5218	0.10036836999882089

81. SRR1974676\_1\_prinseq\_good\_K7Ho\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	2735	0.1314749615910441

82. SRR1974673\_2\_prinseq\_good\_zlhH\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2766	0.1607297572608767
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	2201	0.12789811848560725
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2038	0.11842633597167997
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1863	0.10825724431562304
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	1855	0.10779237155420329

83. SRR1974673\_1\_prinseq\_good\_0VO5\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	3325	0.19321274146508138
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	2612	0.15178095660354662
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2376	0.13806721014166418
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2235	0.12987382772164116
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	2129	0.12371426363282954
ATAGCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTC	2025	0.11767091773437287
CTGCATCCTATTTAGGGCTATTGATATTTAACAAATATCCAGCAAAGGTT	1874	0.10889644436257517
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	1866	0.10843157160115545

84. SRR1974672\_2\_prinseq\_good\_ItYa\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2406	0.15544714312572805
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1946	0.12572740670102528
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1635	0.10563428055301968
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	1624	0.10492359120373332
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1556	0.1005302388626903

85. SRR1974672\_1\_prinseq\_good\_1yPH\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2845	0.1838101089745205
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	2179	0.14078109928136387
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1884	0.12172170309595663
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	1880	0.12146327060530704
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1769	0.11429176898978093
CTGCATCCTATTTAGGGCTATTGATATTTAACAAATATCCAGCAAAGGTT	1690	0.10918772729945155
ATAGCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTC	1594	0.10298534752386139

86. SRR1974671\_1\_prinseq\_good\_ktAN\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2435	0.12004363968371583
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2307	0.11373333747446916
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2294	0.11309244740634254
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	2251	0.11097258025792374

87. SRR1974670\_2\_prinseq\_good\_s\_Cm\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2032	0.11909325024938079
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1997	0.1170419393444948
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	1869	0.10954000232091177

88. SRR1974668\_2\_prinseq\_good\_Qbjr\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	1645	0.11004087901590676

89. SRR1974670\_1\_prinseq\_good\_BSUP\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2520	0.14769438515179115
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2201	0.12899815147583027
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2168	0.12706405833693776
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1711	0.10027979880742645

90. SRR1974669\_2\_prinseq\_good\_FEm7\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1077	0.11194088024820317
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1039	0.10799124844743091

91. SRR1974668\_1\_prinseq\_good\_rZ\_P\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2042	0.13659785711275477
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1715	0.11472346961232832

92. SRR1974667\_1\_prinseq\_good\_hjgT\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	2075	0.11402897050249133

93. SRR1974666\_2\_prinseq\_good\_Mx3i\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2055	0.1038143042471417

94. SRR1974666\_1\_prinseq\_good\_U8k7\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2472	0.12488027255422593

95. SRR1974665\_2\_prinseq\_good\_mA9u\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	2741	0.12669467415525715

96. SRR1974665\_1\_prinseq\_good\_h5D1\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	3783	0.174858063600634
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2200	0.10168853817641944

97. SRR1974664\_2\_prinseq\_good\_DBh9\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	3530	0.20698325136092954
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2986	0.17508554916819705
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	2926	0.17156742024988098
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2601	0.1525108886090022
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	2393	0.14031470835883977
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	2049	0.12014410256049424
TAGTTTATAAGGAATATATCCCATTTTTAGTTATAATGATGCCTTATGTG	1994	0.11691915105203782
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	1975	0.11580507689457106
CTGCATCCTATTTAGGGCTATTGATATTTAACAAATATCCAGCAAAGGTT	1953	0.1145150962911885
ATAGCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTC	1887	0.1106451544810408
TATTAATAATCCCACTATACCAAGAATATCCCAATTATCCATAAACTG	1874	0.10988289321540565

98. SRR1974664\_1\_prinseq\_good\_5DgG\_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	4201	0.24632765976409773
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	3606	0.21143954799079656
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	3516	0.20616235461332244
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2971	0.17420601693861804
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	2891	0.16951517838086325
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	2824	0.16558660108874362
ATAGCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTC	2454	0.14389147275912778
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	2442	0.14318784697546458
CTGCATCCTATTTAGGGCTATTGATATTTAACAAATATCCAGCAAAGGTT	2412	0.14142878251630653
TAGTTTATAAGGAATATATCCCATTTTTAGTTATAATGATGCCTTATGTG	1981	0.11615688978640266
GTCGGCATCCAGACCGTCGGCTGATCGTGGTTTTACTAGGCTAGACTAGC	1896	0.11117287381878821

Sequence	Count	Percentage
GAGCATAAACCAACAACCTCTCAAAGAAGATGGGAAGCTATACTATATAGG	1873	0.10982425773343371
TATTAAATCCCAACTATACCAAGAATATCCCAATTATCCATAAAACTG	1738	0.10190846766722254

99. SRR1974625\_2\_prinseq\_good\_ljkk\_fastqc.html

#### Overrepresented sequences

Sequence	Count
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAACCTACCAATTGGAG	3605
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGCTACCAGCTCGATAATTG	3121
AGGCAGTTTTTGATCACGTTTATTGTAAGCCGTCAGCATCGGGATATCATCTGCTTCAAGCTCCT	3015
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGCACTACACATGCTAGC	2396
TGATCCCTTCCGGCGGTCCGTATAAAATCAGGCAGTTTTTGATCACGTTTATTGTAAGCCGTCAG	2128
CTATGGCGAATGACAGTGGAAGCTGTGTGTTGATTTCATGAATGCGATTCTGATATGGCGGCG	2045
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGAGCCTCACTTAGT	1933
GTCGGCATCCAGACCGTCGGCTGATCGTGGTTTTACTAGGCTAGACTAGCGTACGAGCACTATGG	1933
CTGCATCCTATTTAGGGCTATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTT	1879
AAGCCGTCAGCATCGGGATATCATCTGCTTCAAGCTCCTCAAGCAGCCGAAGCACTGTTTTTTCA	1874
TACCAGTACTGATTGCCGAGCAAATCTGTGAATGGCGACTATGCGCCTAGTCAACCAGATGTGTG	1771
TATTCAGGATCTTCCGACGACATTGATTGCTGCATTCCGCTCAACGCTTGAGGAAGTAAAGAAG	1759
ATCTTGGATATGATATGGCGCACTACACATGCTAGCCGCTGGGGAGATTAGCTCGAGTTGCCCT	1742
CTGGCAATTCGCTCCACACTAGAAGTCGACGAACAACGAAGAGCGACGATGCCCGTTTCAGGTG	1738
GATTGCCGAGCAAATCTGTGAATGGCGACTATGCGCCTAGTCAACCAGATGTGTGCTAGCCCGTC	1688
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGAGCCTCACTTAGTTAC	1630
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATAACCCAGCACTTAGA	1618

100. SRR1974625\_1\_prinseq\_good\_G9D0\_fastqc.html

#### Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGCTACCAGCTCGATAATTG	3612
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAACCTACCAATTGGAG	3083
AGGCAGTTTTTGATCACGTTTATTGTAAGCCGTCAGCATCGGGATATCATCTGCTTCAAGCTCCT	2855
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGCACTACACATGCTAGC	2514
TGATCCCTTCCGGCGGTCCGTATAAAATCAGGCAGTTTTTGATCACGTTTATTGTAAGCCGTCAG	2195
GTCGGCATCCAGACCGTCGGCTGATCGTGGTTTTACTAGGCTAGACTAGCGTACGAGCACTATGG	2067
TAGATATTTTTGCAAAGCGGGCGAGAACGAGAGAAGGCAAACCTCAAATTGAGCTGGCTCAGCTG	2032
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGAGCCTCACTTAGT	1958
CTGCATCCTATTTAGGGCTATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTT	1869
GATTGCCGAGCAAATCTGTGAATGGCGACTATGCGCCTAGTCAACCAGATGTGTGCTAGCCCGTC	1868
CTATGGCGAATGACAGTGGAAGCTGTGTGTTGATTTCATGAATGCGATTCTGATATGGCGGCG	1824
AAGCCGTCAGCATCGGGATATCATCTGCTTCAAGCTCCTCAAGCAGCCGAAGCACTGTTTTTTCA	1755
TACCAGTACTGATTGCCGAGCAAATCTGTGAATGGCGACTATGCGCCTAGTCAACCAGATGTGTG	1634



101. SRR1974590\_2\_prinseq\_good\_5Mpi\_fastqc.html

**Overrepresented sequences**

Sequence	Count
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAAGCTCTACCAATTGGAG	3297

102. SRR1974590\_1\_prinseq\_good\_tSLY\_fastqc.html

**Overrepresented sequences**

Sequence	Count
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAAGCTCTACCAATTGGAG	3273

103. SRR1974577\_2\_prinseq\_good\_nXJA\_fastqc.html

**Overrepresented sequences**

Sequence	Count
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAAGCTCTACCAATTGGAG	1946

104. SRR1974577\_1\_prinseq\_good\_GyQN\_fastqc.html

**Overrepresented sequences**

Sequence	Count
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAAGCTCTACCAATTGGAG	1803

**In TOTAL, 932 html files were each checked manually for overrepresented sequences.**

**A fraction of 143 out of the 932 files had overrepresented sequences**

**The other 789 files were well represented.**