

1. SRR1974928_2_prinseq_good_la_i_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2150	0.13075472845587788
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	2125	0.1292343246366235
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2080	0.1264975977619656

2. SRR1974928_1_prinseq_good_C6w8_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	2483	0.15100650732834642
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2365	0.14383020130146568
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2349	0.14285714285714285
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1770	0.1076445904032111

3. SRR1974917_2_prinseq_good_PLYf_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	1947	0.12631521582579788
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1578	0.10237566028408272
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1542	0.10004009388976905

4. SRR1974917_1_prinseq_good_whSH_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	2155	0.13980959943738802
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1608	0.10432196561267747
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1597	0.10360832032552607
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1569	0.1017917686855043

5. SRR1974913_1_prinseq_good_G8FD_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	2279	0.10486868636670264

6. SRR1974911_2_prinseq_good_DtEw_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	1889	0.10614577878073396

7. SRR1974911_1_prinseq_good__2aq_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	2602	0.14621033159739003

8. SRR1974910_1_prinseq_good_LIoM_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	1538	0.1001364016654676

9. SRR1974908_2_prinseq_good_jjpV_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2245	0.11922356516165282
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	2082	0.11056724394947046

10. SRR1974908_1_prinseq_good_e5Kb_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2486	0.1320221750520574
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	2353	0.12495904179303743
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	2028	0.10769950563377813

11. SRR1974906_2_prinseq_good_MrFp_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2243	0.10833953600925064

12. SRR1974906_1_prinseq_good_Ez3f_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2635	0.12727359669388116

13. SRR1974904_2_prinseq_good_xl5B_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	1176	0.1089266097704205
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1161	0.10753723974783859
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1125	0.10420275169364206

14. SRR1974904_1_prinseq_good_gqAN_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	1372	0.12708104473215723
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1264	0.11707758056956762
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1148	0.10633311906160095

15. SRR1974903_1_prinseq_good_VNXc_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2238	0.10166761232385906

16. SRR1974902_2_prinseq_good_wUV8_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	1883	0.12330149408932058

17. SRR1974902_1_prinseq_good_vshD_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	2403	0.1573518270295472

18. SRR1974899_2_prinseq_good_sqQS_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	4030	0.16166994218795194
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	3334	0.13374878095648432
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	2738	0.10983928082149191
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	2659	0.10667006855527648

19. SRR1974899_1_prinseq_good_kFiw_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	4850	0.19456556317904886
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	3993	0.1601856275822561
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	2978	0.11946726745303247
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	2963	0.11886551828856119
ATAGCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTC	2957	0.11862481862277269
CTGCATCCTATTTAGGGCTATTGATATTTAACAAATATCCAGCAAAGGTT	2768	0.11104277915043449
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2686	0.10775321705132478

20. SRR1974895_2_prinseq_good_zCfe_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	4518	0.1652985086913429
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	4477	0.16379845582362598
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	3208	0.11736999023502169
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	3164	0.11576017740137427

21. SRR1974889_2_prinseq_good_nLRV_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	3709	0.15548481161981723

Sequence	Count	Percentage
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	3053	0.12798466699253222
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	2601	0.10903639660909803
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2505	0.10501198520022703
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	2413	0.10115525760005903

22. SRR1974895_1_prinseq_good_Sw8k_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	5173	0.1892627679195035
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	5143	0.18816516826019844
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	3778	0.13822438376181795
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	3468	0.1268825206156656
ATAGCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTC	3429	0.12545564105856902
CTGCATCCTATTTAGGGCTATTGATATTTAACAAATATCCAGCAAAGGTT	3150	0.1152479642270319
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	3022	0.11056487234733028
TGCCTCTTTAAATATCTAAGTGCTGGGGTTATGAGTAGGGATGAGCATA	2909	0.10643058029728121
ATATCCCAATTATCCATAAACTGTAAGTAAGTGAGGCTCTCTCATTGGT	2889	0.10569884719107783
CATTTTTAGTTATAATGATGCCTTATGTGATAGATGCCTCTTTAAATAT	2863	0.10474759415301345

23. SRR1974889_1_prinseq_good_ozq9_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	4575	0.1917883562040075
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	3889	0.1630305830114503
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	3080	0.12911653270127715
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	3039	0.12739777366207186
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	2778	0.11645640514420388
ATAGCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTC	2720	0.114024989918011
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2603	0.10912023851344949
CTGCATCCTATTTAGGGCTATTGATATTTAACAAATATCCAGCAAAGGTT	2436	0.10211943950010102

24. SRR1974884_2_prinseq_good_cFz8_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	3574	0.13045333446242022
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	3398	0.124029219502883

25. SRR1974877_2_prinseq_good_LISD_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2571	0.10692198188689227
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	2490	0.10355337802347793

26. SRR1974884_1_prinseq_good_OVa5_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	4043	0.14757214080345968
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	3935	0.1436300702601073
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTATGCTCATCCCTACTCATA	3138	0.11453904967629397
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	2957	0.10793243145086082

27. SRR1974877_1_prinseq_good_gQEw_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	3200	0.13308064645587525
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	2836	0.11794272292151944

28. SRR1974872_2_prinseq_good_OtXG_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1599	0.1379668568923893
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	1521	0.13123676631227277
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1498	0.12925225242326402

29. SRR1974872_1_prinseq_good_0cFj_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1924	0.16600890097620827
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	1904	0.16428323672489634
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1777	0.15332526872906552

Sequence	Count	Percentage
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1332	0.11492923913737495

30. SRR1974869_2_prinseq_good_wQtE_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	1335	0.1322093721094674
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1069	0.10586653092510907

31. SRR1974870_2_prinseq_good_d682_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2149	0.19379897392055595
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2094	0.1888390187946227
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	1747	0.1575462110000983
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1208	0.10893865076595236

32. SRR1974870_1_prinseq_good_y9nV_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2399	0.21634422449297985
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2295	0.2069654002548515
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2033	0.18333797765495127
GTGCGCATCCAGACCGTCGGCTGATCGTGGTTTTACTAGGCTAGACTAGC	1377	0.1241792401529109
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1332	0.1201210950498746

33. SRR1974869_1_prinseq_good_GvNR_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	1520	0.15053052105347597
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1242	0.12299928101869549
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1156	0.11448242259066987
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1083	0.10725299625060163

34. SRR1974865_2_prinseq_good_tOMx_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2018	0.11414575927010065

35. SRR1974863_2_prinseq_good_cU57_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1077	0.10118319513418753

36. SRR1974865_1_prinseq_good_Auai_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	2537	0.1435023742657311
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1925	0.10888532536914954
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1798	0.10170172208505499

37. SRR1974863_1_prinseq_good_7ZzO_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1234	0.11593320593833556
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1217	0.11433607101049786

38. SRR1974862_2_prinseq_good_TZ2d_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	1699	0.10498587108921452

39. SRR1974862_1_prinseq_good_RveT_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	1930	0.11925999482176809
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAA	1906	0.11777696897942488

40. SRR1974861_2_prinseq_good_EgLL_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGCTCTA	3610	0.20292823760479475
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	3441	0.1934282730188639
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	3430	0.19280993212865538
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	2924	0.16436625117906364
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	2341	0.13159418399801232
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	2186	0.12288119872689234
GTGCGCATCCAGACCGTCGGCTGATCGTGGTTTTACTAGGCTAGACTAGC	2032	0.11422442626397311
ATAGCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTC	1979	0.11124514742933207
AGGCAGTTTTTGATCACGTTTATTGTAAGCCGTCAGCATCGGGATATCAT	1791	0.10067713948758654
TAGTTTATAAGGAATATATCCATTTTATGTTATAATGATGCCTTATGTG	1781	0.10011501140557877

41. SRR1974861_1_prinseq_good_K0Wm_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGCTCTA	4245	0.2386233708122863
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	4018	0.2258630633507106
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	3769	0.21186607410871783
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	3456	0.1942714651418755
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	2748	0.1544727969357274
GTGCGCATCCAGACCGTCGGCTGATCGTGGTTTTACTAGGCTAGACTAGC	2738	0.15391066885371965
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	2552	0.14345508652837566
ATAGCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTC	2403	0.1350793781064603
CTGCATCCTATTTAGGGCTATTGATATTTAACAAATATCCAGCAAAGGTT	2025	0.11383093660656768
GAGCATAAACCAACAACCTCTCAAAGAAGATGGGAAGCTATACTATATAGG	1956	0.10995225284071426
GATTGCCGAGCAAATCTGTGAATGGCGACTATGCGCCTAGTCAACCAGAT	1903	0.10697297400607322
TAGTTTATAAGGAATATATCCATTTTATGTTATAATGATGCCTTATGTG	1889	0.10618599469126239
AGGCAGTTTTTGATCACGTTTATTGTAAGCCGTCAGCATCGGGATATCAT	1857	0.10438718482883762

42. SRR1974860_2_prinseq_good_T5au_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	2446	0.1879228456097812
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	2148	0.165027911843749
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGCTCTA	2019	0.1551170176967082
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1951	0.1498926703943921
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1672	0.12845748072753646
CGTTGATACCACTGCTTACTCTGCGTTGATACCACTGCTTACTCTGCGTT	1554	0.11939170158528209

43. SRR1974857_2_prinseq_good_5Xzt_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1245	0.13718690807154876
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1185	0.13057549081508857
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	993	0.109418955594416

44. SRR1974857_1_prinseq_good__3FR_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	1503	0.16561600227432755
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1391	0.15327469006226854
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	1123	0.12374369298341306

45. SRR1974860_1_prinseq_good_yHja_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	3197	0.24562115184565436
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	2546	0.1956057092896578
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTA	2509	0.19276304972810346
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2206	0.16948397277807742
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAG	1873	0.14390003672408838
CGTTGATACCACTGCTTACTCTGCGTTGATACCACTGCTTACTCTGCGTT	1760	0.13521840076582786
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGA	1483	0.11393686837256971
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCATA	1460	0.1121698097261981
CTGCATCCTATTTAGGGCTATTGATATTTAACAAATATCCAGCAAAGGTT	1444	0.11094055153741786
ATAGCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTC	1424	0.10940397880144254
GTGCGCATCCAGACCGTCGGCTGATCGTGGTTTTACTAGGCTAGACTAGC	1316	0.10110648602717583

46. SRR1974859_1_prinseq_good_Zrzd_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
TGGTATCAACGCAGAGTAAGCAGTGGTATCAACGCAGAGTAAGCAGTGGT	1000	0.13815322297653881

47. SRR1974858_2_prinseq_good_ziAU_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGCTCTA	2125	0.13440588172788406
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	1965	0.12428590945660808
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	1785	0.11290094065142262

48. SRR1974858_1_prinseq_good_ziPD_fastqc.html

Overrepresented sequences

Sequence	Count	Percentage
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGCTCTA	2409	0.15236883250939892
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGC	2238	0.14155311214447272
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGAA	2066	0.13067414195285104

49. SRR1974808_1_prinseq_good_Kvin_fastqc(1).html

Overrepresented sequences

Sequence	Count
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGAGCCTCACTTAGT	4992
GGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGAGCCTCACTTAGTTAC	3022

50. SRR1974808_2_prinseq_good_ZBXn_fastqc.html

Sequence	Count	Percentage	Possible Source
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGAGCCTCACTTAGT	4710	0.15601693727821	No Hit
GCTTCCCATCTTCTTTGAGAGTTGTTGGTTTATGCTCATCCCTACTCAT AACCCAGCACTTAGA	3229	0.10695938226567729	No Hit
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGA AACTCTACCAATTGGAG	3140	0.10401129151880667	No Hit

51. SRR1974798_2_prinseq_good_Qepy_fastqc.html

Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGCTACCAAGCTCGATAATTG	2205
ATTGATATTTAACAAATATCCAGCAAAGGTTTTTCCAGGAGATGTTGGA AACTCTACCAATTGGAG	1931

52. SRR1974798_1_prinseq_good_tqzr_fastqc.html

Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTACCAGCTCGATAATTG	2512
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGCACTACACATGCTAGC	1817

53. SRR1974797_1_prinseq_good_SM7b_fastqc.html

Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTACCAGCTCGATAATTG	1743

54. SRR1974794_2_prinseq_good_a1gu_fastqc(1).html

Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTACCAGCTCGATAATTG	1575

55. SRR1974794_1_prinseq_good_gfwv_fastqc.html

Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTACCAGCTCGATAATTG	1727
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGCACTACACATGCTAGC	1241

56. SRR1974791_1_prinseq_good_tr4R_fastqc.html

Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTACCAGCTCGATAATTG	1946

57. SRR1974790_1_prinseq_good_DBG9_fastqc.html

Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTACCAGCTCGATAATTG	1666

58. SRR1974789_1_prinseq_good_oRX5_fastqc.html

Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTACCAGCTCGATAATTG	2434
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGCACTACACATGCTAGC	2020

59. SRR1974788_2_prinseq_good_tgXD_fastqc.html

Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTACCAGCTCGATAATTG	3552
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGCACTACACATGCTAGC	2881
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAAGCTCTACCAATTGGAG	2028
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGAGCCTCACTTAGT	1651

60. SRR1974788_1_prinseq_good__bMy_fastqc.html

Overrepresented sequences

Sequence	Count
GTGCGCCAATTAGGTCGTCATTGCGCCAGCTCGTCAGCGCTGGTCGTCTACCAGCTCGATAATTG	4012
TCGTCATCCAATGCGACGAGTCCTACACTATCTTGGATATGATATGGCGCACTACACATGCTAGC	2890
ATTGATATTTAACAAATATCCAGCAAAGGTTTTCCAGGAGATGTTGGAAGCTCTACCAATTGGAG	1843
GGTGGCTATCTATCCCTACCAAGGCTTATATTGAAGTATAAACCAATGAGAGAGCCTCACTTAGT	1726