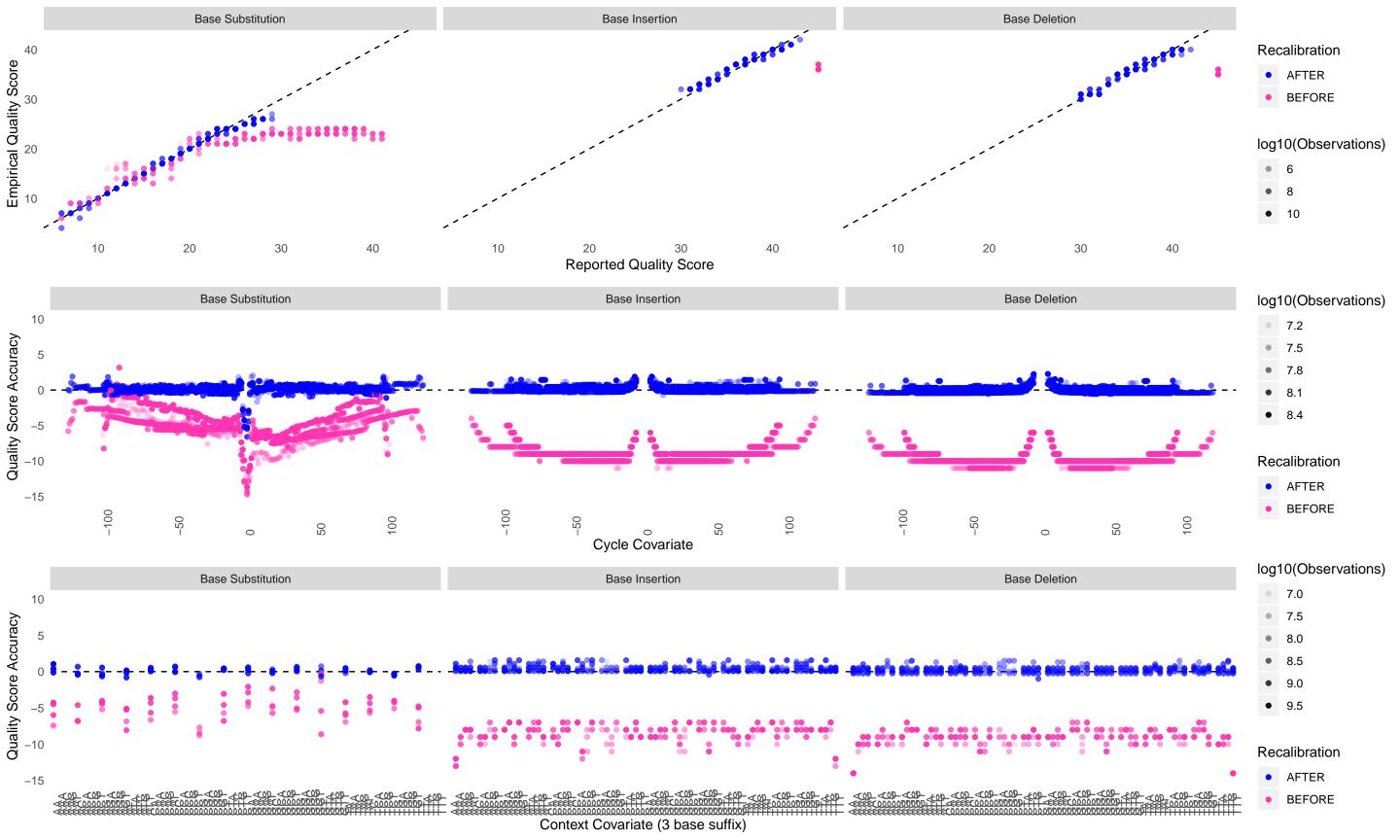
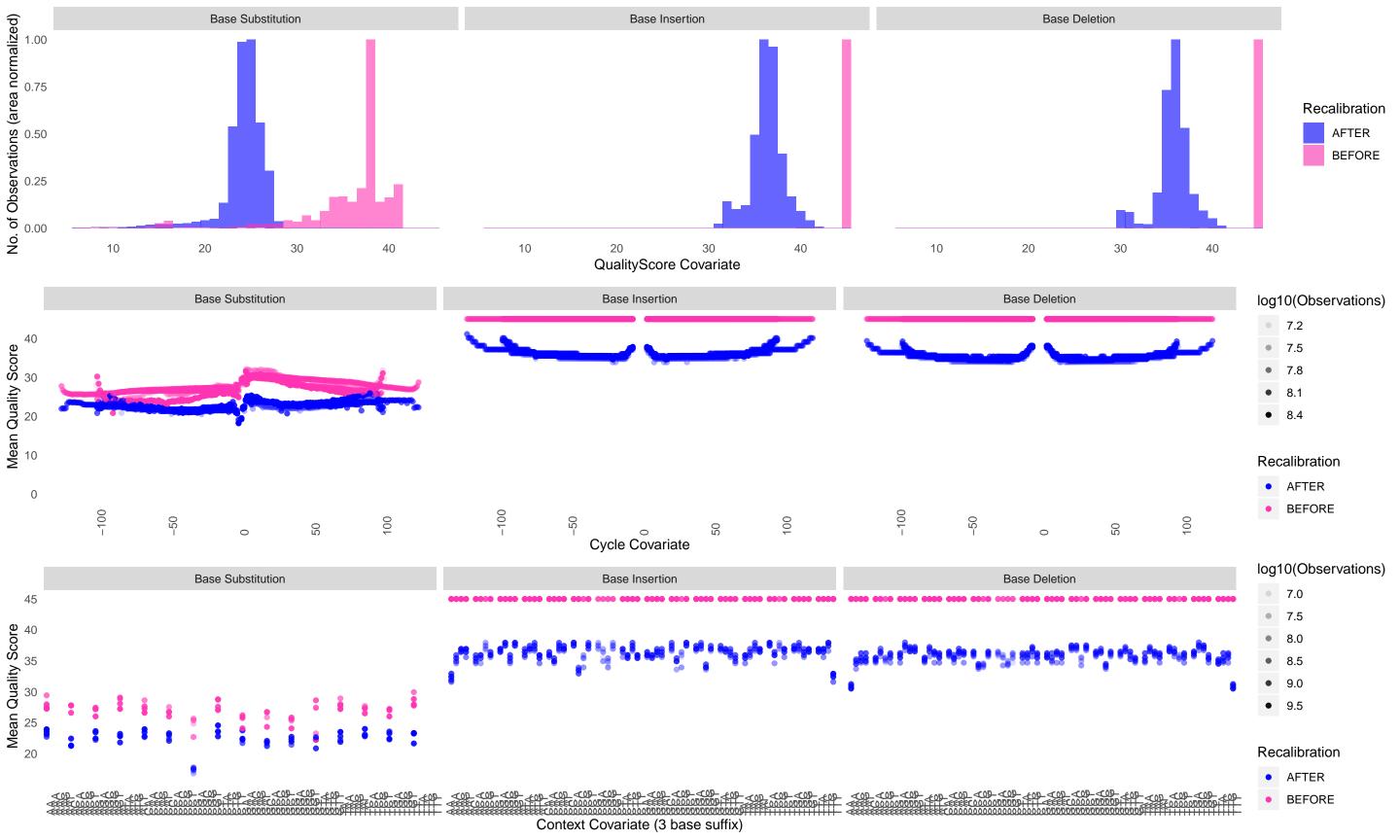
GATK BaseRecalibration report

Argument	Value
binary_tag_name	None
covariate	ReadGroup, QualityScore, Context, Cycle
default_platform	None
deletions_default_quality	45
force_platform	None
indels_context_size	3
insertions_default_quality	45
low_quality_tail	2
maximum_cycle_value	500
mismatches_context_size	2
mismatches_default_quality	—1
no_standard_covs	false
quantizing_levels	16
run_without_dbsnp	false
solid_nocall_strategy	THROW_EXCEPTION
solid_recal_mode	SET_Q_ZERO





Overall error rates by event type

ReadGroup	SAMN04316320	SAMN04316320	SAMN04316320	SAMN04316319	SAMN04316319	SAMN04316319	C2V1GACXX.3.Malaya	C2V1GACXX.3.Malaya	C2V1GACXX.3.Malaya	C6GFVANXX.2.Malaya	C6GFVANXX.2.Malaya	C6GFVANXX.2
EventType	M	I	D	M	1	D	M	1	D	M	i I	
EmpiricalQuality	22.0	36.0	35.0	22.0	37.0	36.0	22.0	36.0	35.0	22.0	36.0	
stimatedQReported	27.5	45.0	45.0	25.9	45.0	45.0	27.4	45.0	45.0	27.3	45.0	
Observations	2.66e+10	2.66e+10	2.66e+10	4.51e+10	4.51e+10	4.51e+10	5.26e+09	5.26e+09	5.26e+09	2.31e+10	2.31e+10	2
Frrors	1 696±08	6.410+06	7 640+06	2 800+08	1 016+07	1 200+07	3 696+07	1 390+06	1 686+06	1 480+08	5 530+06	6

Error rates by event type and initial quality score

ReadGroup	QualityScore	EventType	EmpiricalQuality	Observations	Errors
SAMN04316320 SAMN04316320	11 12	M	16 17	18395 22797	406.15 319.79
SAMN04316320 SAMN04316320	13 14	M M M	14 14	56378655 159084682	2221435.96 6932872.37 17997180.06
SAMN04316320 SAMN04316320	15 16	M	14 14 13	484034176 596951949	27888279.38
SAMN04316320 SAMN04316320	18 19	M M	14	33349230 10906	1453129.84 59.25
SAMN04316320	20	M	21 21	23034	138.31
SAMN04316320 SAMN04316320	21 22	M M	23 22 23 21	16883199 1876358	91604.32 11126.65
SAMN04316320 SAMN04316320	23 24	M M	23 21	77160286 40690166	430044.31 314124.77
SAMN04316320 SAMN04316320	25 26	M M	22 22	216493092 94423543	1512321.68 546263.70
SAMN04316320 SAMN04316320	27 28	M M	22 23	185810027 209779591	1097373.52 1000017.41
SAMN04316320	29 30	M	23	387582496	1770299.98
SAMN04316320 SAMN04316320 SAMN04316320	30 31 32	M M M	23 23 24 22	41580923 392360232	187826.20 1696716.51 690151.22
SAMN04316320	33	M	22 23	122212037 769748688	3750185.64
SAMN04316320 SAMN04316320	34 35	M M	23 23 24 24 24 24 24 36	1200396764 611849920	5645390.57 2573230.77
SAMN04316320 SAMN04316320	36 37	M	24	1079539688 1579946462	4489942.44 6135145.05
SAMN04316320 SAMN04316320 SAMN04316320	38 45	M	24	18194483835 26552691131	80347608.88 6414534.49
SAMN04316320	45	D	35	26552691131	7644750.55
SAMN04316319 SAMN04316319	6 7	M M	6 9 9	16924192 111891183	4008283.77 13503118.49
SAMN04316319 SAMN04316319	8	M M	9	145503553 105652147	20186352.69 12572750.37
SAMN04316319	9 10	M M	9 9	121243776	16086116.07
SAMN04316319 SAMN04316319	11 12	M	12 16	57497959 21708306	3704372.30 589434.00
SAMN04316319 SAMN04316319	13 14	M M	17 15	73933281 27935196	1639819.66 839352.80
SAMN04316319 SAMN04316319	15 16	M M	16 16	68361008 79167884	1643257.93 2224168.85
SAMN04316319 SAMN04316319	17 18	M M	17 17	87461380 179865487	1727925.65 3229903.55
SAMN04316319 SAMN04316319	19 20	M M	18 22	130418102 126745578	2135085.56 865637.36
SAMN04316319	21	M	20	75852629	759001.42
SAMN04316319 SAMN04316319	22 23	M M	21 21 21	156538210 201204057	1230622.03 1675276.05
SAMN04316319 SAMN04316319	24 25	M M	21	281421631 363719863	2299500.12 2716721.02
SAMN04316319	26	M	21 23 22 23	535536602	2930908.50 3123806.75 2124417.91
SAMN04316319 SAMN04316319	27 28	M	23	480209843 411803454 662047157	2124417.91
SAMN04316319 SAMN04316319	29 30	M M	24 23	1047256026	2869762.74 5129482.90
SAMN04316319 SAMN04316319	31 32	M M	23 24	1546636841 1159802807	7665082.69 4819570.05
SAMN04316319 SAMN04316319	33	M M	24 24 24 24 24 24	1575801695 3265286849	6897599.36
SAMN04316319 SAMN04316319	34 35 36	M M	24	4508865439 2698917665	13775359.75 17171642.87 10535530.33
SAMN04316319	37	M	24	4137031768	15285841.81
SAMN04316319 SAMN04316319	38 39	M M	24 24 23	3327474900 4587977893	14817886.54 17905273.81
SAMN04316319 SAMN04316319	40 41	M M	23 23	5248428483 7473643827	23857999.15 37302796.40
SAMN04316319 SAMN04316319	45 45	I D	23 37 36	45099766671 45099766671	37302796.40 10084437.35 12045411.49
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	6 7	M	6 9	1367818 10292203	328381.52 1262048.74
C2V1GACXX.3.Malaya	8	M	9	11672427	1512165.24
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	9 10	M M	10 10	7206629 7577130	744279.50 780870.71
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	11 12	M M M	12 14 16	3236364 1844009	197670.46 66675.31 144523.37
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	13 14	M	15	5804538 2045528	144523.37 64553.02
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	15 16	M M	16 16	5317054 6752409	125423.03 175212.98
C2V1GACXX 3 Malaya	17 18	M M	17	7159909	153374.83
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	19	M	17 18	16501136 10322431	304963.19 181872.25
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	20 21	M M	21 19	11393820 6294261	95382.99 71768.57
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	22 23	M M	21 21 21 21 21	13873313 15574889	117763.50 133120.89
C2V1GACXX.3.Malaya	24 25	M M	21 21	22224413 32066396	187541.81 229191.49
C2V1GACXX.3.Malaya	26 27	M M	22	45737369 40345086	285710.24 275330.00
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	28 29	M M	22 22 22 22 23	36535936 66240508	208714.10 346679.87
C2V1GACXX.3.Malava	30	M	23 23 22	92131392	486868.49
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	31 32	M M	22 23	149825495 114855645	868304.59 560481.13
C2V1GACXX 3 Malaya	33	M M	23 23 23 23 23 23 23 23	151327486 359787861	807581.44 1935466.76
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	34 35	M	23	500300704 294373965	2470766.44 1551765.80
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	36 37 38	M M M	23 23 22	483802496 354065983	2442182.55 2195524.72
C2V1GACXX.3.Malaya	39	M	23	556788627	3083211.90
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	40 41	M M	22 22	712986558 1106317612	4603678.80 7875802.28
C2V1GACXX.3.Malaya C2V1GACXX.3.Malaya	45 45	I D	36 35	5263949400 5263949400	1394155.43
C6GFVANXX.2.Malaya C6GFVANXX.2.Malaya	13 14	M M	14 13	9921931 120913859	1677297.50 406592.43 5692416.72
C6GFVANXX.2.Malaya C6GFVANXX.2.Malaya	15	M	14	354869019	13330898.07
C6GFVANXX.2.Malaya	16 18	M M	15 16	754694468 28790878	26320001.02 714008.08
C6GFVANXX.2.Malaya C6GFVANXX.2.Malaya	21 22	M M	22 23 22 21	1807014 1019956	11675.17 5689.72
C6GFVANXX.2.Malaya C6GFVANXX.2.Malaya	23 24	M M	22 21	20702814 31437807	137850.33 249394.59
C6GFVANXX.2.Malaya C6GFVANXX.2.Malaya	25 26	M M	22	251960032 73754421	1555881.32 359678.40
C6GFVANXX.2.Malaya	27 28	M	22	170067384	1016294.32
C6GFVANXX.2.Malaya C6GFVANXX.2.Malaya	28 29 30	M M	22 23 22 23 23 23 23 23 22 22	202031779 427781912	1037197.50 2108914.94
C6GFVANXX.2.Malaya C6GFVANXX.2.Malaya	30 31 32	M M M	23 23	44726394 369880708	213136.19 1719811.30 685118.48
C6GFVANXX.2.Malaya C6GFVANXX.2.Malaya	33	M	22 23	117820545 776144441	4122344.01
C6GFVANXX.2.Malaya C6GFVANXX.2.Malaya	34 35	M M	23 23 23 23 23	1215943940 537079360	6113011.85 2484910.76
C6GFVANXX.2.Malaya C6GFVANXX.2.Malaya	36	M M	23	1006987381 1616902201	4579028.56 6721430.00
C6GFVANXX.2.Malava	37 38	M	24 23	14986951335	68844026.01
C6GFVANXX.2.Malaya C6GFVANXX.2.Malaya	45 45	D D	36 35	23122189579 23122189579	5531713.32 6793255.86