

Want to compare gene expression levels between sample groups?

Then... Don't miss our **SPECIAL SALE** on
RNA-seq Data Analysis Service!

Until March-31-2014

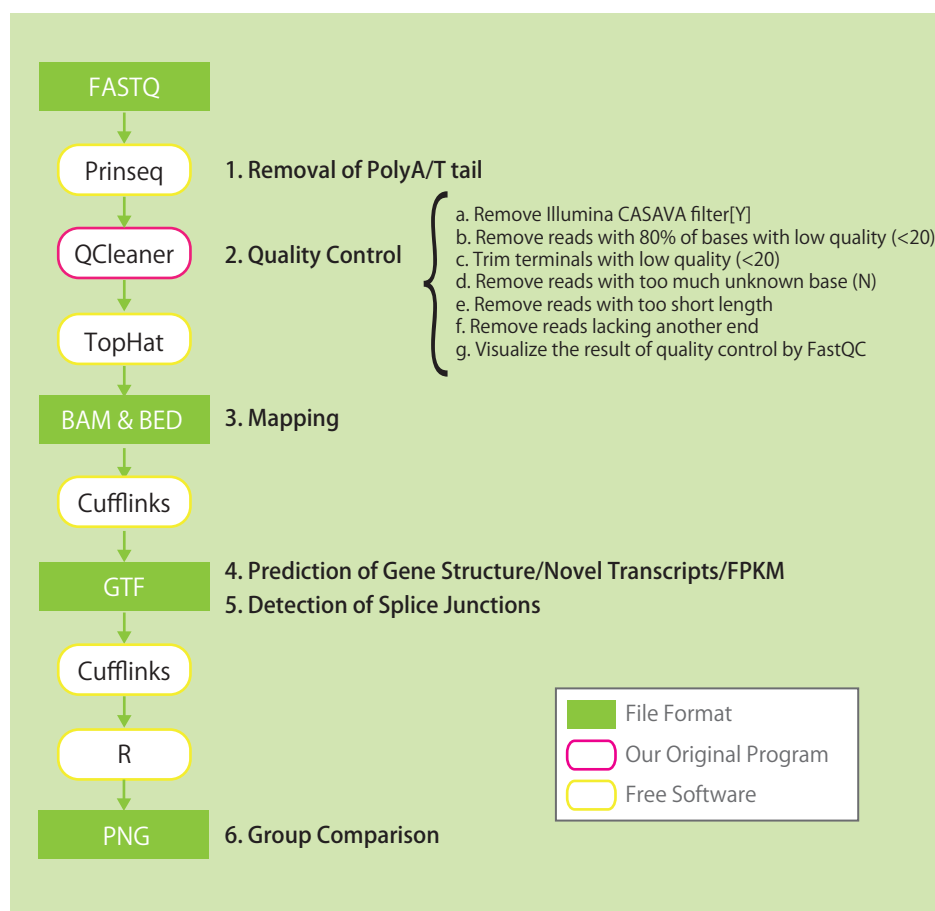
■ Features

• Sample Group Comparison

• Quick Service (2 weeks)

¥ 116,000 /sample

■ Analysis Pipeline



■ Price

Analysis Step 1~4	¥ 80,000
Step 5	¥ 16,000
Step 6 (/ combination)	¥ 20,000
Total	¥ 116,000

■ Required Data

- Sequence data: FASTQ
- Information of target groups

■ Data We Provide

- Quality report: HTML
- Result of transcript detection: EXCEL
- Result of expression comparison: EXCEL
- Summary report: PDF

※ This offer is only applied to 3 samples or more.

※ Tax is not included in the prices shown above.

※ The total number of base must be less than 3Gb/sample.

※ Please contact us if you live outside Japan.

※ Only paired-end data that are sequenced by Illumina platform are accepted.

※ The delivery term may be extended if the # of sample is more than 8.

※ The analysis result will be sent back to you by your HDD.

※ Please backup your original data before you send us your data.

※ Please send us extra HDD if the HDD with data has small available space (<15GBxSamples).

※ We also can detect fusion genes, but with extra charge.

※ We store your data for no more than 2 months.

Powering Bio-Research By IT



Amelieff Corporation <http://amelieff.jp/enquiry>

US contact:
InfiniteBio Inc.

www.infinitebio.com

natsuhikof@infinitebio.com



Need a help to search disease-associated genes?

Then... Don't miss our **SPECIAL SALE** on

Human Exome Data Analysis Service!

Until March-31-2014

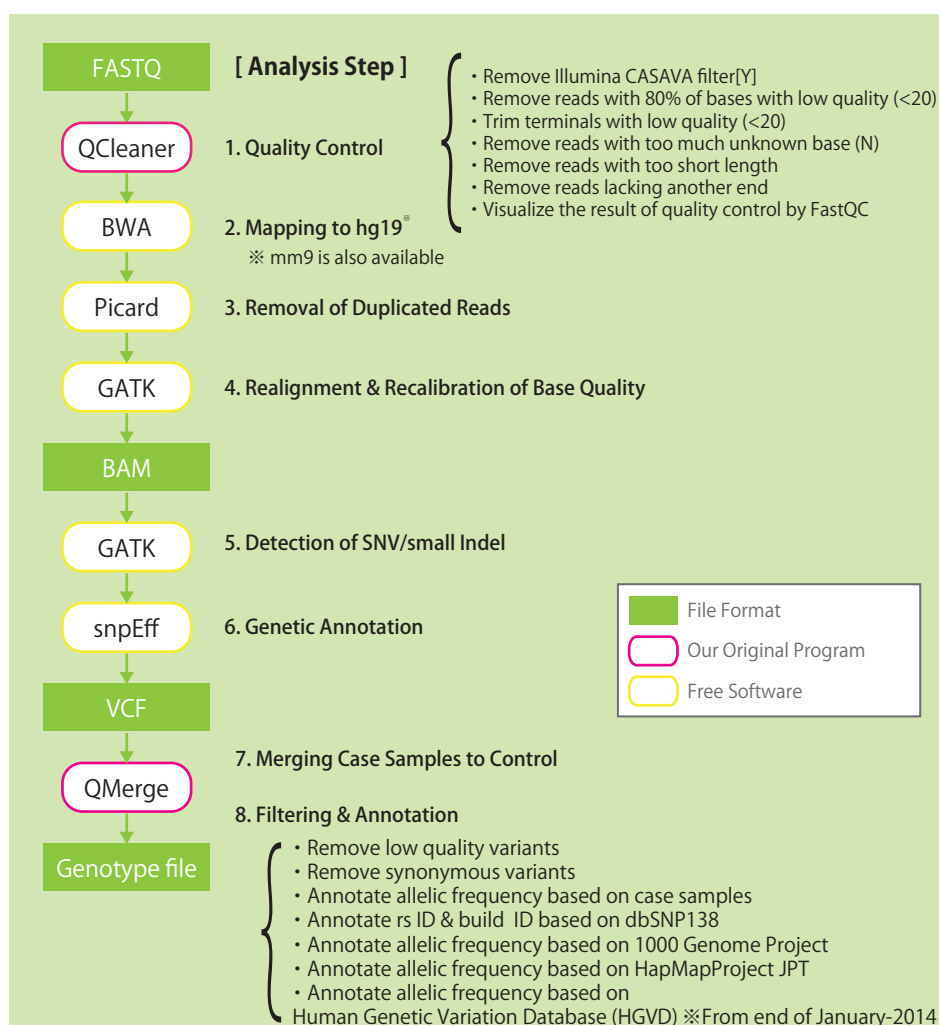
■ Features

• Case-specific Variants Detection

• Fast Service (2 weeks)

¥ 169,000/sample

■ Analysis pipeline



■ Price

Analysis Step1~2	¥ 49,000
Step3~6	¥ 60,000
Step7~8	¥ 60,000
Total	¥ 169,000

■ Required Data

- Sequence data: FASTQ
- Exome capture target region: BED
- Case/Control information

■ Data We Provide

- Quality report: HTML
- Mapping result: BAM
- Variant information: VCF
- Filtering result: EXCEL
- Summary report: PDF

※ This offer is limited to only if 3 samples or more.
 ※ Tax is not included in the prices shown above.
 ※ The total number of base must be less than 8Gb/sample.
 ※ Please contact us if you live outside Japan.

※ Only paired-end data that are sequenced by Illumina platform are accepted.

※ The delivery term may be extended if the # of sample is more than 8.

※ The analysis result will be sent back to you by your HDD.

※ Please backup your original data before you send us your data.

※ Please send us extra HDD if the HDD with data has small available space (<15GBxSamples).

※ We also can detect variants based on genetic model and can detect compound heterozygosity, but with extra charge.

※ We store your data for no more than 2 months.

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