



## Application

# Quantitative analysis of gene expression in human bone marrow-derived mesenchymal stem cells (primer check)

| Product Name | FastGene™ RNA Premium Kit(FG-81050, FG-81250)          |
|--------------|--|
| Manufacturer | NIPPON Genetics EUROPE                                 |
| Product Name | Transcriptor Universal cDNA Master(05893151001)        |
| Manufacturer | Roche Diagnostics                                      |
| Product Name | KAPA SYBR <sup>®</sup> FAST qPCR Kits (KK4610, KK4611) |
| Manufacturer | Kapa Biosystems, Inc.                                  |

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#### Purposes of these experiments

- In preparation for planned experiments in which hMSC (human mesenchymal stem cells) would be induced to differentiate into cartilage and bone, we checked primers targeting the following 10 genes using hMSC-BM (human bone marrow-derived mesenchymal stem cells).
  - Genes: ACTB, GAPDH, RUNX2 (2 types), ALPL, Collagen Type I, SP7/Osterix, BGLAP (2 types), SOX9, Aggrecan, and Collagen Type II (11 types)
- 2) We verified whether 2.5 µg of RNA, which is a recommended maximum amount of input RNA for reverse transcription reaction, can be used for qPCR analysis (Sample A).

### Experimental conditions

■ RNA extraction : FastGene<sup>™</sup> RNA Premium Kit

- Starting sample: hMSC-BM (human bone marrow-derived mesenchymal stem cells)  $1.8 \times 10^6$  cells
- $\bullet$  The amount of lysis buffer RL: 600  $\mu L$
- Pretreatment of the sample: Lysis buffer RL (600 μL) was added onto the cells, and the cells were not homogenized but lysed by vortexing alone.
- Elution buffer RE: 50 µL
- Measurement of yield and purity: NanoDrop One (Thermo Fisher Scientific, Inc.)

#### Reverse transcription : Transcriptor Universal cDNA Master

- The amount of input RNA: Total RNA 1.0 μg (Sample B) or 2.5 μg (Sample A)
- The type of primers: Random hexamer primers

| The composition of the reaction mixture:     |             | <ul> <li>Reaction program:</li> </ul> |              |
|--|-------------|---------------------------------------|--------------|
| RNA  | 1.0 µg      | Annealing                             | 25℃ · 5 min  |
|  | or 2.5 µg   | Ļ                                     |              |
| Transcriptor Universal Reaction Buffer       | 4 µL        | Enzyme reaction                       | 55℃ · 10 min |
| Transcriptor Universal Reverse Transcriptase | 1 µL        | $\downarrow$                          |              |
| Distilled Water                              | up to 20 µL | Denaturation and inactivation         | 85℃ · 5 min  |
| total  | 20 µL       |                                       |              |

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#### Quantitative analysis of gene expression : KAPA SYBR<sup>®</sup> FAST qPCR Kit

- The amount of input cDNA: 2 µL (per 10-µL reaction) of the reverse transcription reaction solution diluted by 10-fold with DEPC-treated water
- Real-time machine: LightCycler® 96

| 8,   |       |     |    |
|--|-------|-----|----|
| The composition of the reaction mixture:       |       |     |    |
| cDNA   |       | 2   | μL |
| KAPA SYBR <sup>®</sup> FAST qPCR Master Mix (2 | 2X)   | 5   | μL |
| 10 µM forward primer                           |       | 0.2 | μL |
| 10 μM reverse primer                           |       | 0.2 | μL |
| PCR-grade water                                | Up to | 10  | μL |
| total  |       | 10  | μL |
|  |       |     |    |

| Reaction program:    |               |
|----------------------|---------------|
| Initial denaturation | 95℃ · 3 min   |
| $\downarrow$         |               |
| Denaturation         | 95℃·10 sec    |
| Annealing            | 60°C · 20 sec |
| Extension reaction   | 72℃ · 1 sec   |
| Ļ                    |               |
| Melting              |               |
| Ļ                    |               |
| Cooling              | 40°C · 10 sec |

## Results

## 1. RNA extraction

| The volume of elution buffer ( $\mu$ L) | Concentration (ng/µL) | A260/A280 | A260/A230 | A260  | A280 |  |
|---|-----------------------|-----------|-----------|-------|------|--|
| 50                                      | 474.82                | 2.07      | 2.23      | 11.87 | 5.74 |  |

# 2. Quantitative analysis of gene expression

Sample A (RNA 2.5 µg)

As shown below, all the primers detected respective targets.

## Amplification curve



#### • Cq values and positive/negative judgements

| Primers   | No. | Cq value | Result   | Primers    | No. | Cq value | Result   | Primers       | No. | Cq value | Result   | Primers | No. | Cq value | Result   |
|-----------|-----|----------|----------|------------|-----|----------|----------|---------------|-----|----------|----------|---------|-----|----------|----------|
| GAPDH     | 1   | 17.85    | Positive | SP7        | 1   | 33.48    | Positive | Col2 9/4 2    | 1   | 27.30    | Positive | Col2 A  | 1   | 28.41    | Positive |
|           | 2   | 17.89    | Positive |            | 2   | 32.48    | Positive | 0012 074_2    | 2   | 27.30    | Positive |         | 2   | 28.39    | Positive |
| ACTB      | 1   | 15.28    | Positive | Palan1     | 1   | 28.53    | Positive | Col2 9/4_3    | 1   | 31.76    | Positive | Col2 B  | 1   | 32.38    | Positive |
|           | 2   | 15.27    | Positive | Dgiapi     | 2   | 28.66    | Positive |               | 2   | 31.53    | Positive |         | 2   | 34.27    | Positive |
| RUNX2_1 - | 1   | 22.98    | Positive | Bglap3     | 1   | 30.24    | Positive | Col2 9/4_4    | 1   | 31.32    | Positive | Col2 C  | 1   | 33.62    | Positive |
|           | 2   | 22.98    | Positive |            | 2   | 30.73    | Positive |               | 2   | 30.95    | Positive |         | 2   | 33.17    | Positive |
| BLINY2 2  | 1   | 23.41    | Positive | SOX9       | 1   | 22,53    | Positive | Col2 8/30_1 - | 1   | 39.03    | Positive | Col2 D  | 1   | 33.48    | Positive |
| HONX2_2   | 2   | 23.39    | Positive |            | 2   | 22.54    | Positive |               | 2   | 31.74    | Positive |         | 2   | 33.41    | Positive |
|           | 1   | 25.59    | Positive | Aggrecan   | 1   | 37.41    | Positive | Col2 8/30_2   | 1   | 31.43    | Positive |         |     |          |          |
|           | 2   | 25.63    | Positive | Aggreean   | 2   | 35.06    | Positive |               | 2   | 32.34    | Positive |         |     |          |          |
| Col1      | 1   | 14.26    | Positive | Col2 9/4 1 | 1   | 31.00    | Positive | Col2 8/30 3   | 1   | 35.11    | Positive |         |     |          |          |
|           | 2   | 14.30    | Positive | 0012 9/4_1 | 2   | 30.48    | Positive | 0012 0/30_3   | 2   | 37.93    | Positive |         |     |          |          |



#### Sample B (RNA 1.0 µg)

As shown below, most of the primers detected respective targets.

#### Amplification curve



Cq values and positive/negative judgements

| Primers  | No. | Cq value | e Result | Primers    | No. | Cq valu | e Result | Primers     | No. | Cq value | e Result | Primers | No. | Cq value | Result   |
|----------|-----|----------|----------|------------|-----|---------|----------|-------------|-----|----------|----------|---------|-----|----------|----------|
| GAPDH    | 1   | 18.66    | Positive | SP7        | 1   | 34.60   | Positive | Col2 9/4 2  | 1   | 28.12    | Positive |         | 1   | 29.66    | Positive |
|          | 2   | 18.67    | Positive |            | 2   | -       | Negative | 0012 0/4_2  | 2   | 28.17    | Positive |         | 2   | 30.15    | Positive |
| ACTB -   | 1   | 16.24    | Positive | Bglap1     | 1   | 30.51   | Positive | Col2 9/4_3  | 1   | 32.22    | Positive | Col2 B  | 1   | 34.47    | Positive |
|          | 2   | 16.22    | Positive |            | 2   | 29.61   | Positive |             | 2   | 32.47    | Positive | OOIZ D  | 2   | 34.67    | Positive |
| RUNX2_1  | 1   | 24.15    | Positive | Bglap3     | 1   | 30.95   | Positive | Col2 9/4_4  | 1   | 33.04    | Positive | Col2 C  | 1   | 36.38    | Positive |
|          | 2   | 24.21    | Positive |            | 2   | 31.28   | Positive |             | 2   | 31.69    | Positive | 0012 0  | 2   | 34.41    | Positive |
| BLINX2 2 | 1   | 24.60    | Positive | SOX9       | 1   | 23.90   | Positive | Col2 8/30_1 | 1   | 32.56    | Positive | Col2 D  | 1   | 34.11    | Positive |
| nonx2_2  | 2   | 24.61    | Positive |            | 2   | 23.81   | Positive |             | 2   | 34.34    | Positive |         | 2   | 33.22    | Positive |
|          | 1   | 26.45    | Positive | Aggrecan   | 1   | 34.99   | Positive | Col2 8/30_2 | 1   | 31.92    | Positive |         |     |          |          |
| ALPL     | 2   | 26.61    | Positive |            | 2   | -       | Negative |             | 2   | 31.29    | Positive |         |     |          |          |
| Col1     | 1   | 15.39    | Positive | Col2 9/4_1 | 1   | 30.70   | Positive | Col2 8/30_3 | 1   | 34.37    | Positive |         |     |          |          |
|          | 2   | 15.37    | Positive |            | 2   | 30.85   | Positive |             | 2   | 33.96    | Positive |         |     |          |          |

## **General evaluation**

- LightCycler<sup>®</sup> 96 system (high performance, simple operation)
  - → Because its melt curve analysis does not take time, the experimental time can be reduced, which is good.
    - · Because of its simple operation using a touchscreen, the machine is easy for beginners to use.
    - · Its software is easy to use, for example it is compatible with Excel.
- FastGene<sup>™</sup> RNA Premium Kit (low price, high quality)
  - To be honest, I felt it burdensome to eliminate gDNA after RNA was eluted, instead of doing it on column. However, because you do it carefully, even when you are not accustomed to it, there is a sense of security that gDNA can be sufficiently eliminated.
    - · Although there are multiple columns, they were color-coded in the protocol and easy to identify.
    - $\cdot$  Reaction time is much reduced than the previous products, and it became easy to process many samples.
    - · I am satisfied because its quality is equal to or better than Qiagen's RNeasy Mini Kit and its price is cheaper.
- Transcriptor Universal cDNA Master (convenience, high performance)
  - cDNA can be easily synthesized by mixing 2 premixed reagents and RNA, and is high-quality. I am very satisfied.
     Because all of our laboratory members are beginners of PCR, I am thankful that it can be done by simple operation.
- KAPA SYBR<sup>®</sup> FAST qPCR Kit (low price, high sensitivity)
  - → Clearer calibration curve can be drawn than Cells to Ct kit



When we purchased LightCycler<sup>®</sup>, we consulted Nippon Genetics, which is selling reagents from Roche, and they recommended these products because of their low price and high quality. Although the RNA purification step is burdensome, I am satisfied because RNA has been sufficiently purified in our laboratory which has many beginners.

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