

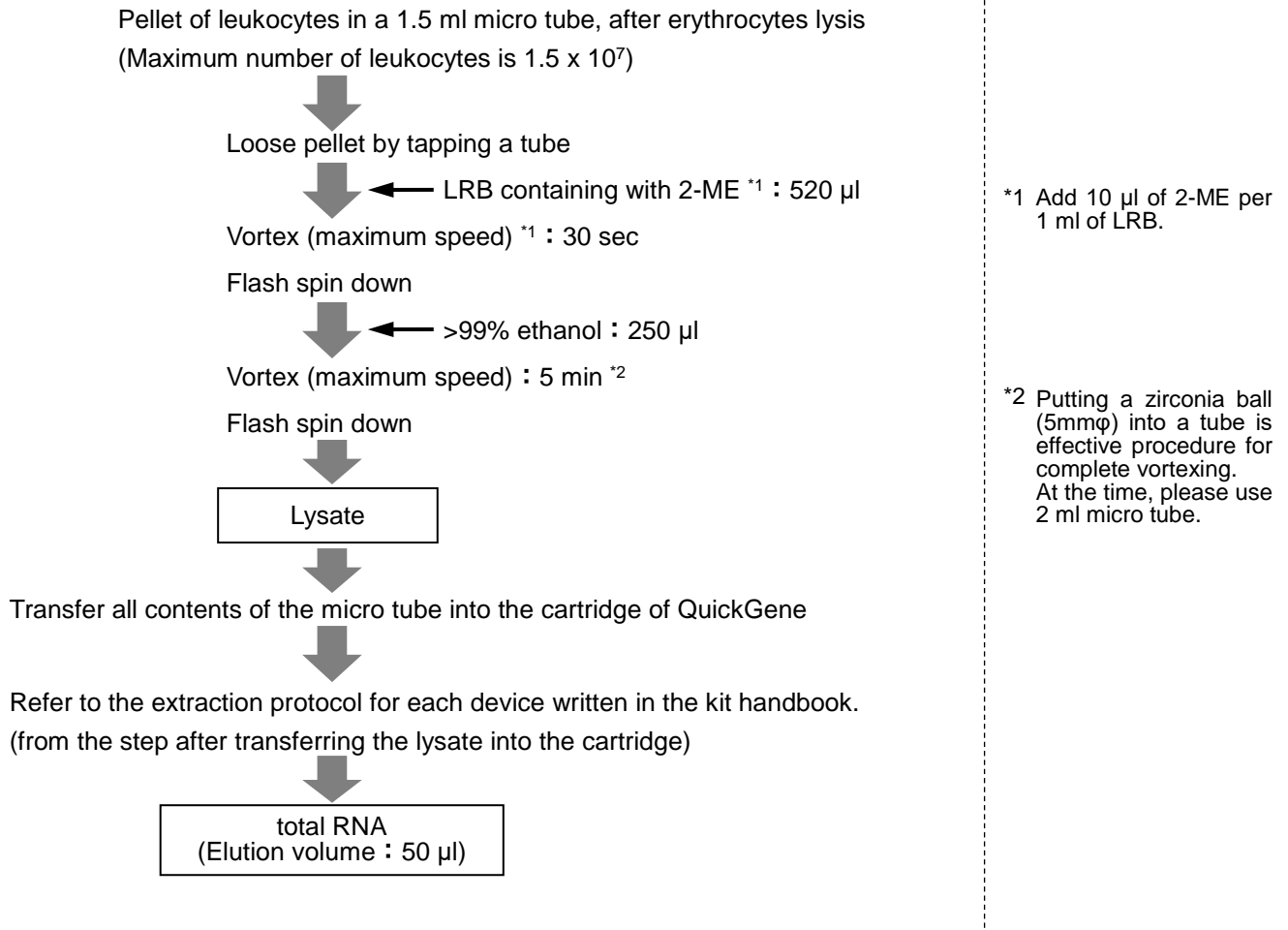


## **11. Total RNA Extraction from Blood of Animal**

RA-a-1

## Total RNA Extraction from Leukocyte

### Protocol



### Results

#### The yield of total RNA / Protein contamination : A260/280

	Number of leukocytes (cells)	QuickGene		Spin column method (company A) *1		Automatic magnetic bead method *2	
		( $\mu$ g)	A260/280	( $\mu$ g)	A260/280	( $\mu$ g)	A260/280
With DNase treatment	$2 \times 10^6$	0.6	2.20	0.4	2.04	0.7	2.46
	$1 \times 10^7$	4.5	2.21	3.8	2.09	-	-
	$1.5 \times 10^7$	6.5	2.10	-	-	-	-
Without DNase treatment	$1.0 \times 10^7$	5.0	2.17	4.2	2.10	-	-

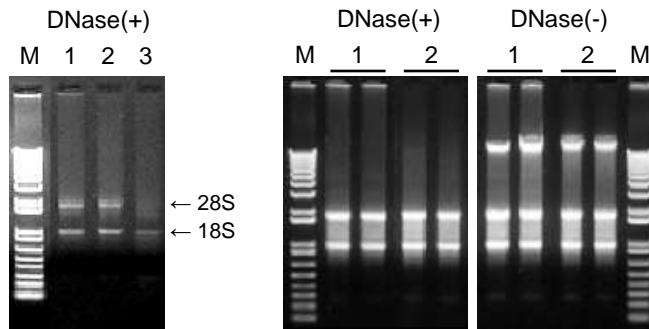
\*1 : For spin column method, maximum number of leukocytes is  $1 \times 10^7$ .

\*2 : For automatic magnetic bead method, maximum number of leukocytes is  $2 \times 10^6$ .

## Electrophoresis of total RNA

Number of leukocytes :  $2 \times 10^6$

Number of leukocytes :  $1 \times 10^7$

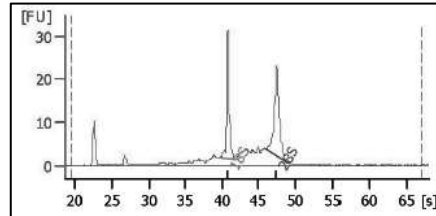
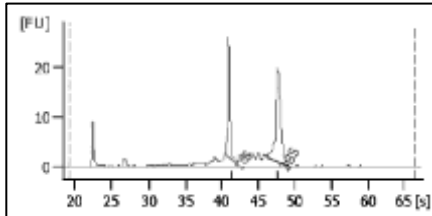


Electrophoresis condition : 1% Agarose / 1 x TAE  
 M : Marker (1Kb Plus DNA Ladder : Invitrogen)  
 1 : QuickGene  
 2 : Spin column method (A company)  
 3 : Automatic magnetic bead method

## The quality of total RNA (with DNase treatment)

QuickGene (Number of leukocytes :  $1 \times 10^7$ )

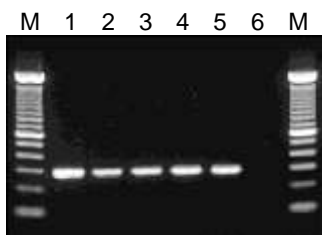
Spin column method (company A) (Number of leukocytes :  $1 \times 10^7$ )



	Number of leukocytes	QuickGene	Spin column method (company A) <sup>*1</sup>	Automatic magnetic bead method <sup>*2</sup>
RIN	$2 \times 10^6$	7.7	6.5	5.0
	$1 \times 10^7$	9.2	8.8	-
28S / 18S	$2 \times 10^6$	1.5	0.8	0.0
	$1.0 \times 10^7$	1.6	1.2	-

## Other

### RT-PCR



M : Marker (100bp DNA Ladder : Invitrogen)  
 1 : Positive control  
 2,3 : QuickGene  
 4,5 : Spin column method (A company)  
 6 : Negative control

### Real Time PCR

Number of copied *GAPDH* per  $1\mu\text{g}$  of total RNA (For isolation from  $1 \times 10^7$  leukocytes)

QuickGene	$3.15 \times 10^7$
Spin column method (company A)	$1.11 \times 10^7$

Used model : Real Time PCR system Roche LightCycler  
 Used reagents : LightCycler FastStart DNA Master SYBR Green I  
 LightCycler Human GAPDH Primer Set

## Common protocol is usable for the following

No Data

Depending on sample and storage conditions, nucleic acid may not be extractable.  
 Therefore, we cannot guarantee accurate data.  
 The extracted nucleic acid contains unintended acid (ex: when extracting DNA, RNA is also extracted).

Depending on sample and storage conditions, nucleic acid may not be extractable.  
Therefore, we cannot guarantee accurate data.  
The extracted nucleic acid contains unintended acid (ex: when extracting DNA, RNA is also extracted).