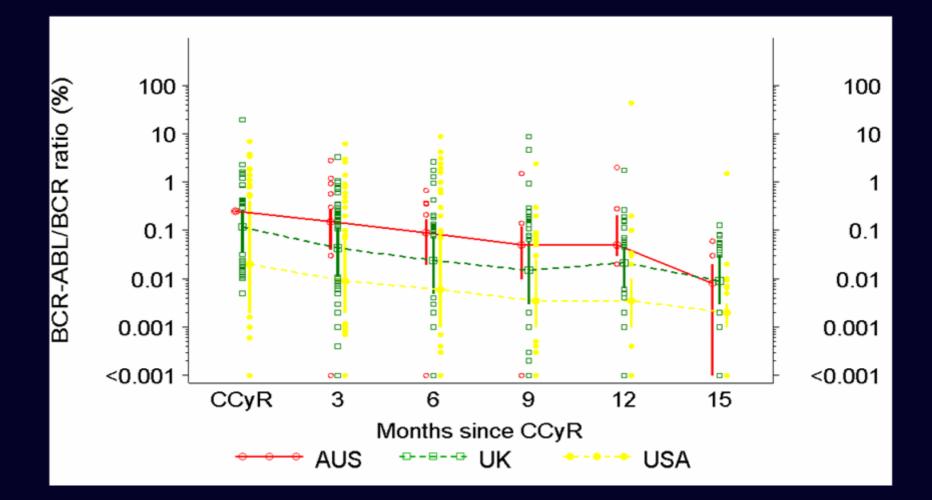
### Standardization of RQ-PCR for BCR-ABL

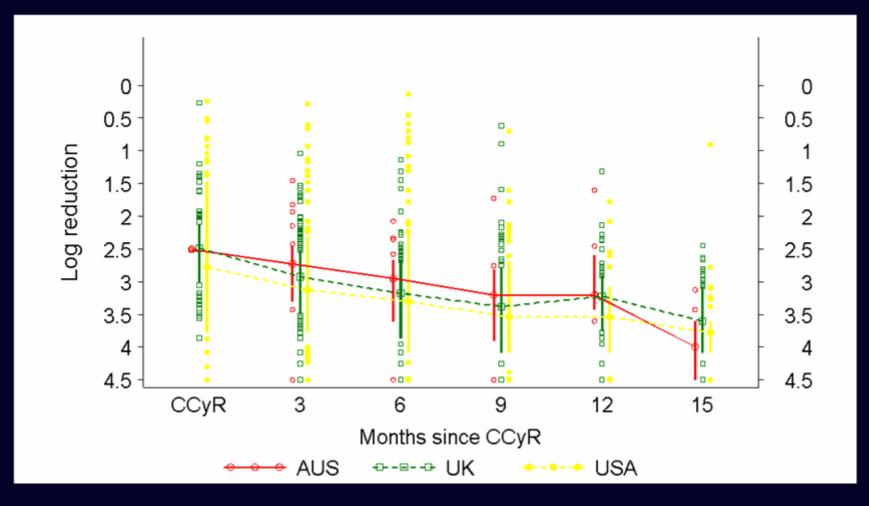
**Nick Cross** 

#### Wessex Regional Genetics Laboratory, Salisbury Human Genetics Division, University of Southampton

## **IRIS trial: variation in the three reference centres**



# IRIS trial: results following normalisation to 30 shared baseline samples

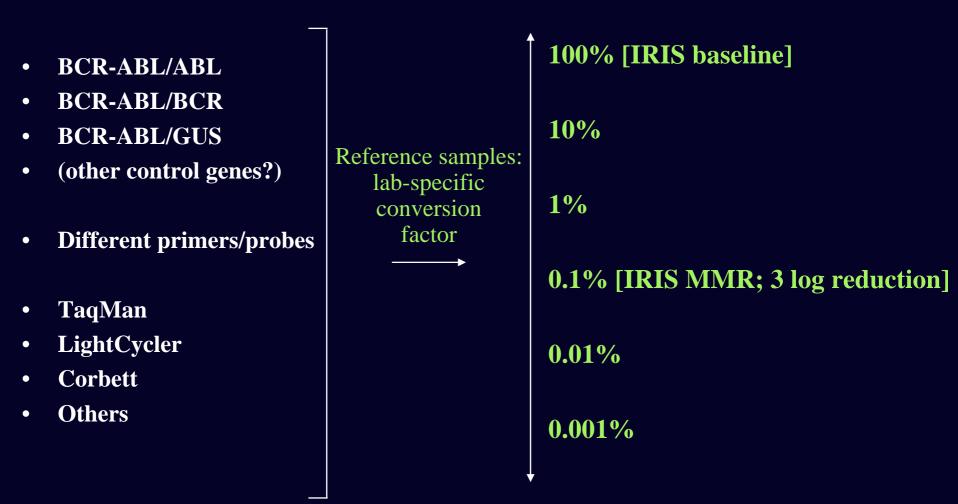


# Realising the international scale for BCR-ABL RQ-PCR

Monitoring CML patients responding to treatment with tyrosine kinase inhibitors: review and recommendations for harmonizing current methodology for detecting *BCR-ABL* transcripts and kinase domain mutations and for expressing results

Timothy Hughes, Michael Deininger, Andreas Hochhaus, Susan Branford, Jerald Radich, Jaspal Kaeda, Michele Baccarani, Jorge Cortes, Nicholas C. P. Cross, Brian J. Druker, Jean Gabert, David Grimwade, Rüdiger Hehlmann, Suzanne Kamel-Reid, Jeffrey H. Lipton, Janina Longtine, Giovanni Martinelli, Giuseppe Saglio, Simona Soverini, Wendy Stock, and John M. Goldman

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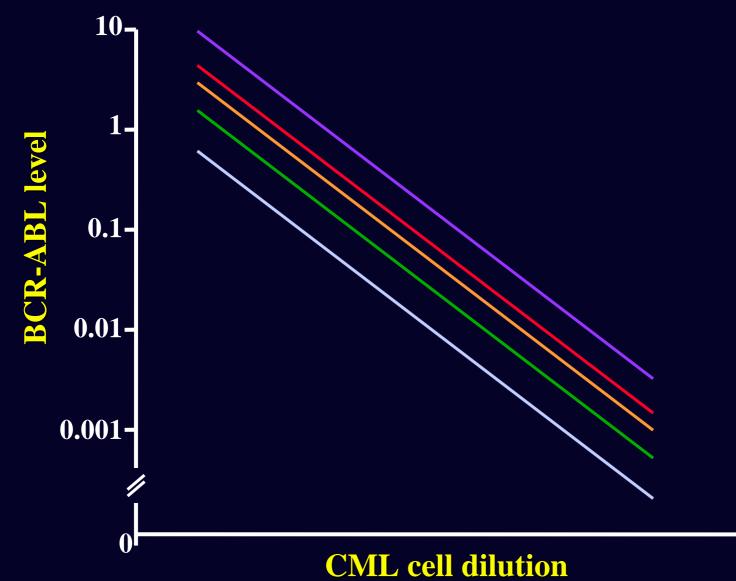


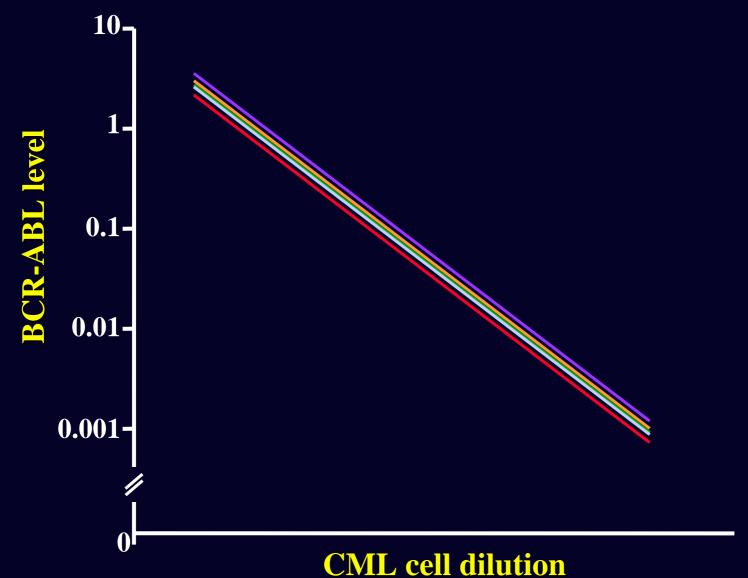
cf. International Normalized Ratio (INR) for Prothrombin time

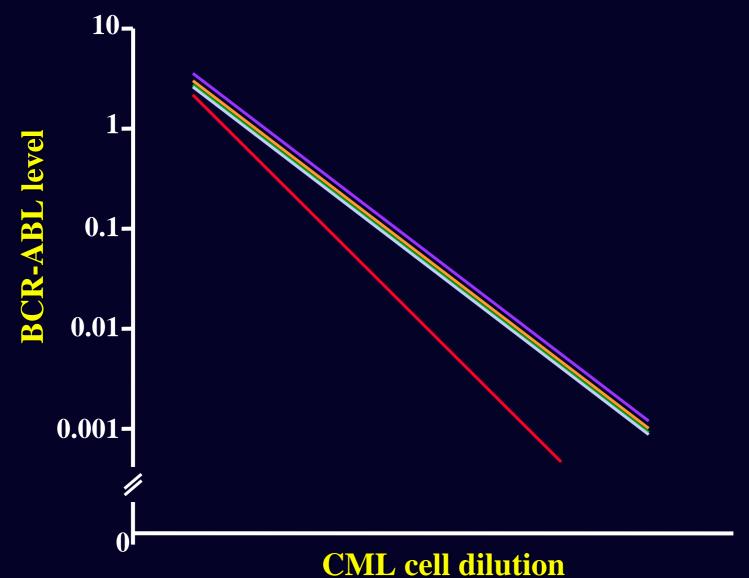
#### The Adelaide reference lab has maintained consistency of data since the MMR value was established for the IRIS trial in 2001- BCR-ABL/BCR 0.08%

#### Yearly quality control mean values

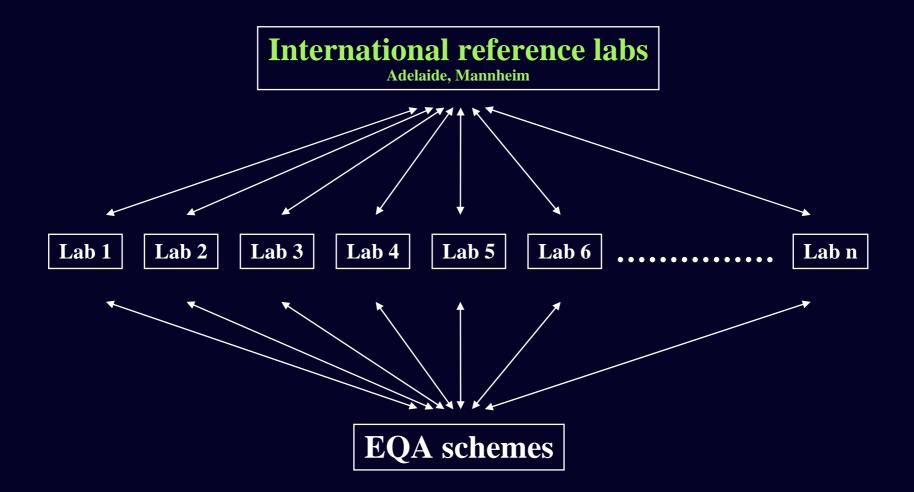
Control	Target mean	2001 (from July)	2002	2003	2004	2005	2006 (to March)
Low b3a2	0.07	0.07	0.06	0.08	0.06	0.08	0.09
High b3a2	85	82	77	93	69	94	93
Low b2a2 <	0.07	0.07	0.07	0.09	0.08	0.06	0.07
High b2a2	56	48	53	69	50	42	52
Number		36	111	147	156	217	34







### Conversion factors: current status



# **Derivation of conversion factors: current status**

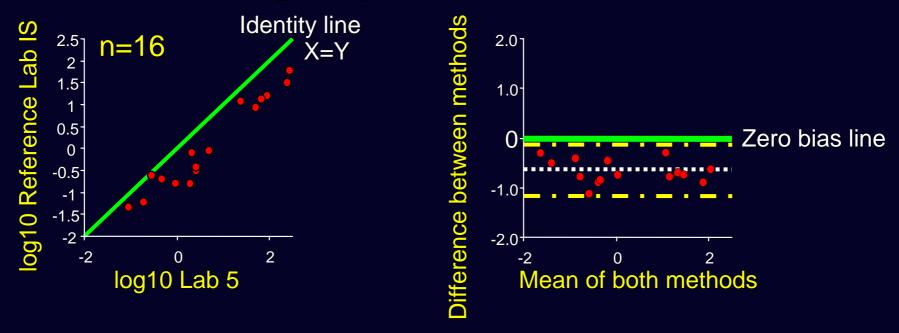
#### • Sample exchange:

- Either sends 20+ samples to each test lab; K562 or primary CML cells in normal leucocytes
- Or receives 20+ patient samples from test lab

#### • Derivation of conversion factor:

- Samples analysed in both centres; different operators, different days
- conversion factor calculated from Bland-Altman bias plots
- Validation:
  - Samples (20-30) from test lab covering minimum 3 log range sent back to reference lab for analysis and results compared.

## **Example of CF Calculation**

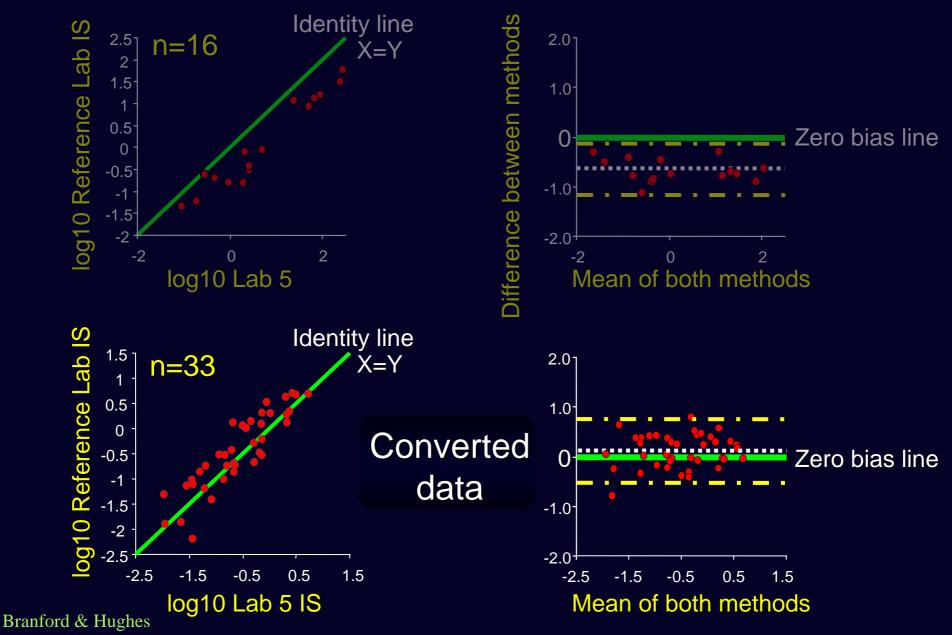


Lab 5 BCR-ABL/ABL% x 0.23 to convert to the international scale

The CF is the antilog of the Bias = 0.23

Branford & Hughes

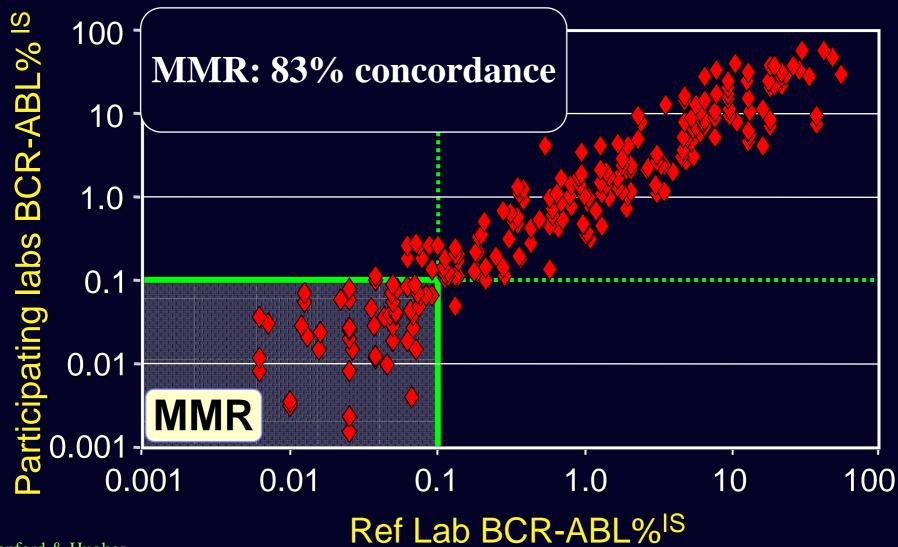
# Validation of Conversion Factor



# Concordance between Ref Lab and Lab 5



## What is achievable?

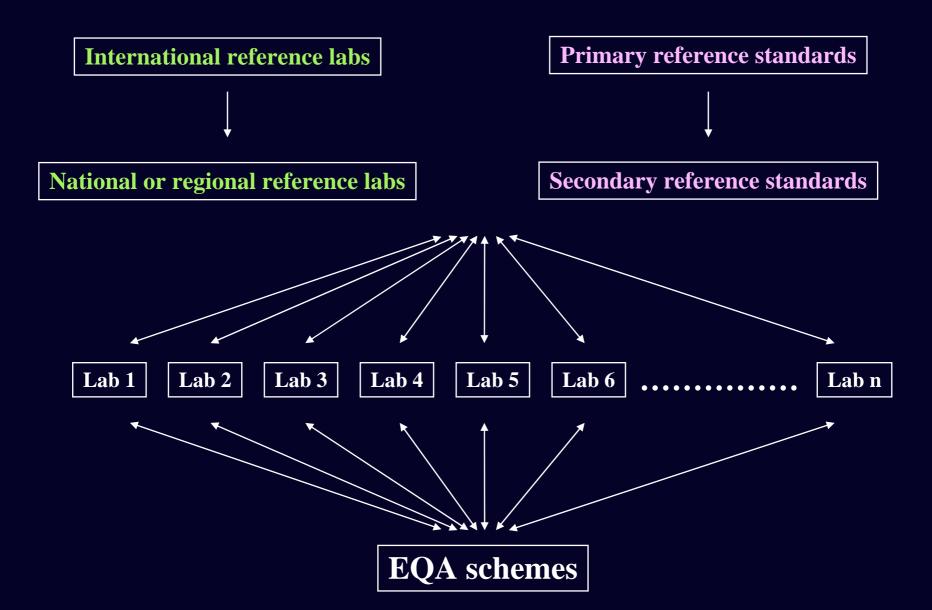


Branford & Hughes

## Conversion factors: current status

- Works well (for many labs), but very labour intensive
- Open to a limited number of labs at any time
- EUTOS programme: Europe including UK & Ireland
- Adelaide: Australasia/Asia
- USA: enthusiasm but not much action

# Conversion factors: future status



# **Primary reference standards**

- Ideally be as close as possible to real samples.
- Must be stable over several years (=freeze dried) and batches prepared that last several years.
- Must cover all or most existing methods (including RNA extraction).
- Prepared centrally and WHO accredited by NIBSC (but other routes possible).
- Depending on formulation may be very limited in supply
- Used by reference labs or companies to calibrate secondary reference standards

# Secondary reference standards

- Easily available: used by testing labs on every run? Every week?
- Prepared locally/nationally? Or by companies?
  - Asuragen, Molecular MD, (Ipsogen)
- Calibrated to primary reference reagents

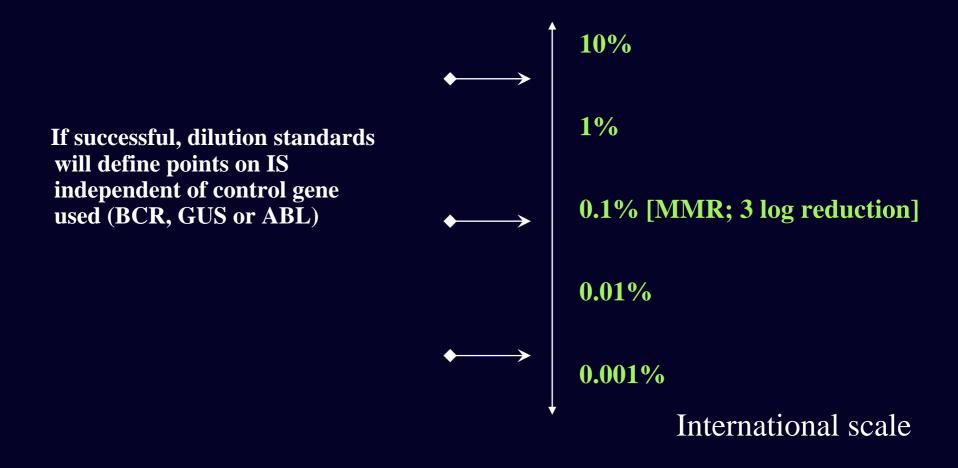
• Similar to real samples: function to monitor efficiency of RNA extraction and/or RT and assay drift but may also provide conversion factor

# Formulation for primary and/or secondary reagents

- [CML cells (primary or K562) diluted in normal leucocytes]
- Cell line mixtures
- Armored RNAs

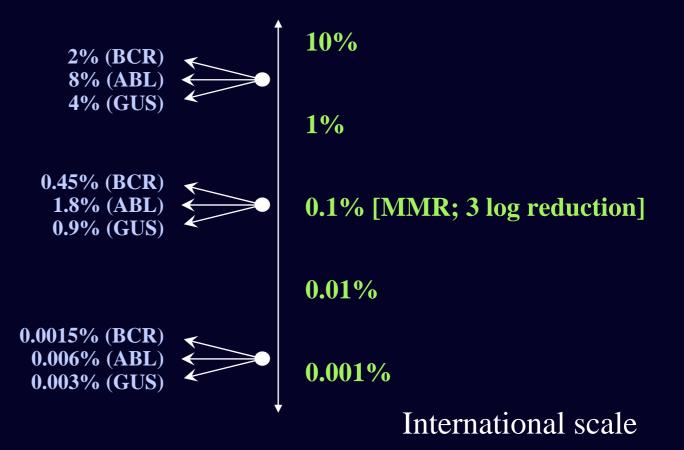
#### Cell lines: current status

Aim to find cell line or cell line mixture for which the ratio of BCR:ABL:GUS in the same as median of normal leucocytes



# What happens if we cannot find a line in which BCR:ABL:GUS is spot on?

Eg. if BCR:ABL:GUS in normal leucocytes is 1:2:4 cell line has a ratio 1:0.5:2



# **Evaluation of cell lines**

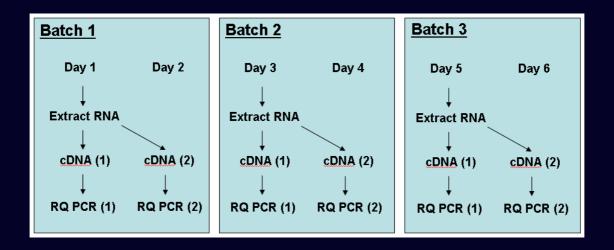
- K562 is fine for BCR-ABL
- Non BCR-ABL: Control genes (BCR, ABL and GUS) need to be expressed at levels comparable to normal leukocytes: KG1 and HL60 (at least the subclones we have tested).
- Pilot batch of freeze dried samples:
  - Cells grown and mixtures made in Salisbury
  - 4 dilutions of K562 in both KG1 and HL60; 10%-0.01%; 3x10<sup>6</sup> cells/vial (10<sup>9</sup> cells total)
  - Freeze dried at NIBSC (April 2007)
  - Initial tests at Salisbury, Mannheim and Marseilles
  - Full performance evaluation involving 14 labs worldwide July-October 2007

# **Field Trial Protocol**

- Freeze Dried Cells sent to 14 labs (4 control genes; 7 protocols; 9 platforms)
- Each lab sent 24 vials packaged into 3 batches
- Each batch contained 8 vials: HL60/K562 Levels 1-4

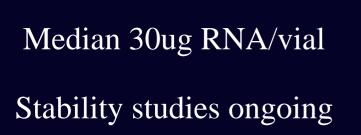
KG1/K562 Levels 1-4

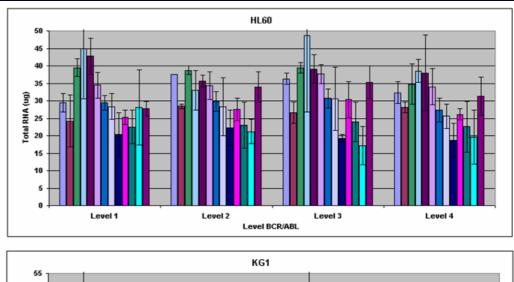
- Cells resuspended directly into 1ml Trizol (Invitrogen) or 600µl RLT Buffer (QIAGEN)
- Usual lab protocol for cDNA synthesis and RQ-PCR:

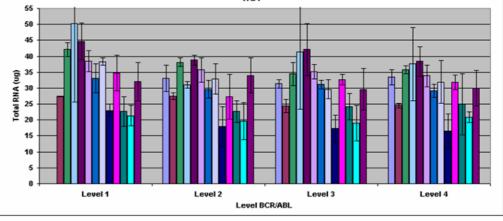


# Freeze dried cells: yields of RNA

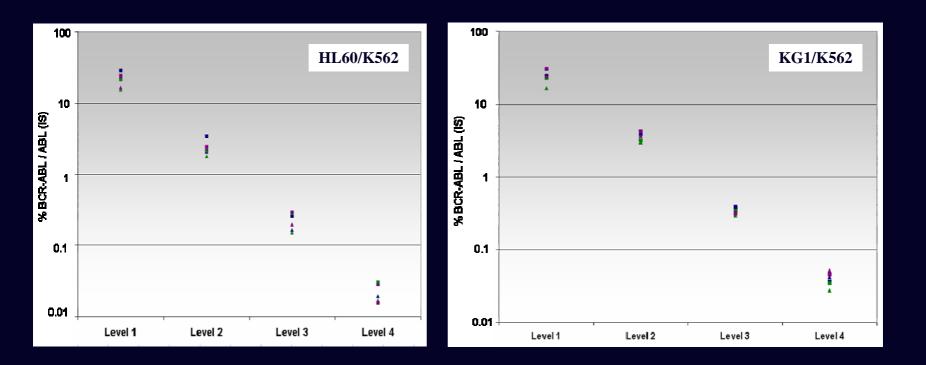








## Stability of freeze dried cells



- Pre Freeze Dry Original
- ▲ Pre Freeze Dry 2 months
- Trizol Resuspension Original
- ▲ Trizol resuspension 2 months
- PBS resuspension original
- ▲ PBS resuspension 2 months

# Trial summary

- Majority of labs obtained linear results with expected slope
- CVs were comparable to those described in other trials with patient samples
- Both HL60 and KG1 performed well

## Next steps

- Current trial: report to be circulated to participants for comments, then published/circulated more widely
- Selection of cell line combination and levels (HL60/K562?)
- Large scale grow ups and freeze drying
- Performance evaluation and certification
- 1.5x10<sup>6</sup> cells/vial
- 4 levels: Spanning range 10% 0.01%
- Values assigned to dilutions by reference labs

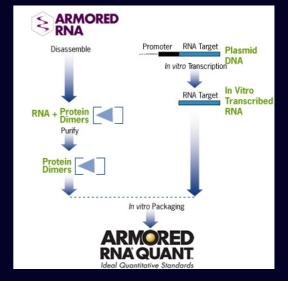
# How will vials be used?

- Vials should not be simply be available for any testing lab on demand: risk of depleting stock too quickly
- Essential to promote production and widespread availability of secondary reference reagents

# Armored RNAs

- Easily available in large quantities
- Stable
- Good track record for calibration of RNA virus detection assays
- Easy to adjust BCR:ABL:GUS ratio
- Flexible: can use directly for reverse transcription after heat lysis, put through RNA extraction or negative sample spike-ins.





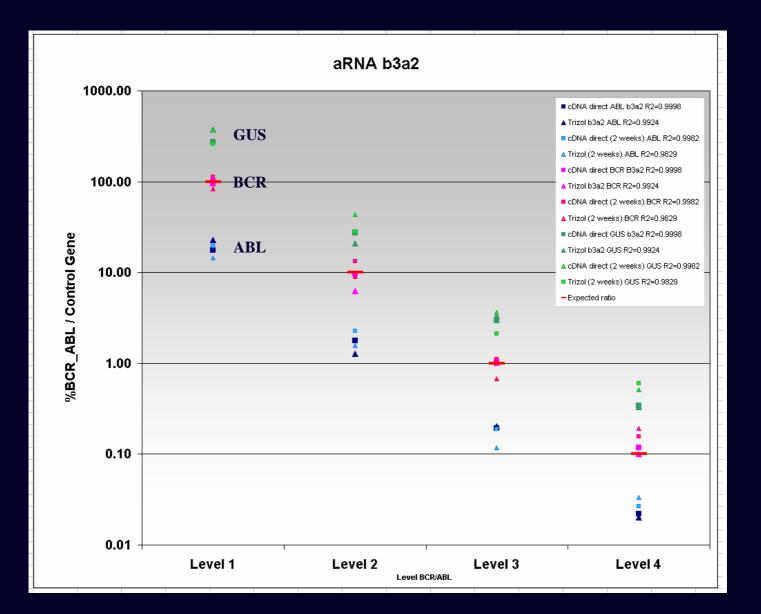


## Armored RNAs: current status

- Survey of primer sets performed July 2006
- Plasmids made (BCR, ABL, GUS, b2a2, b3a2) that cover the regions targeted by all members of the international group
- Sequence verified; sent (essentially gifted) to Asuragen Nov 2006
- Armored RNAs arrived in Salisbury May 2007
- Copy numbers estimated by NIST-traceable phosphate assay (except ABL)
  - Due to lack of sufficient ABL product yield, target was quantified through a standard OD260 conversion used for estimating copy numbers for Armored RNA nonquantitative products.
- Preliminary in house evaluation successful
- International evaluation round

**October – November 2007** 

#### Initial stability tests



# Armored RNA Field Trial

#### **Evaluation round**

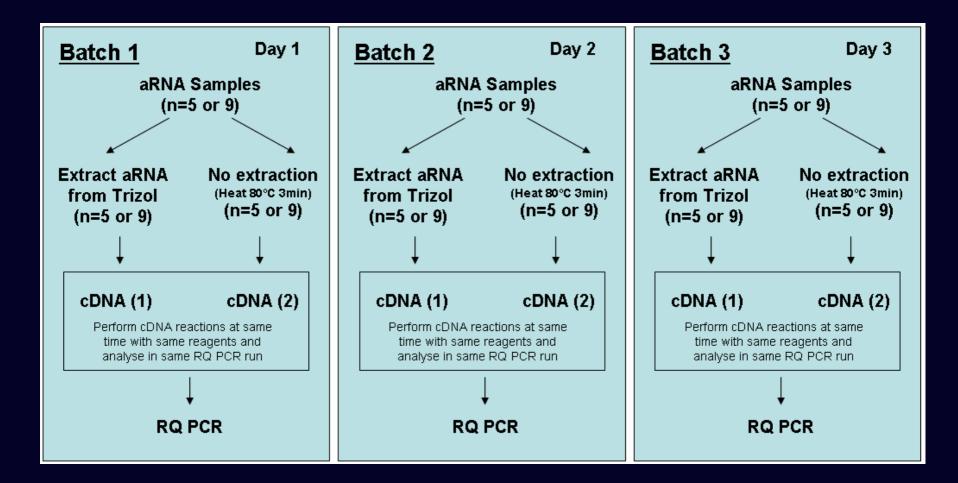
- aRNA mixes prepared in Salisbury
- 4 levels of BCR-ABL (in background of BCR+ ABL+ GUS) armored RNA mixtures tested before and after 'RNA' extraction protocol
- Samples sent at ambient temperature
- data return by 30th Nov 2007

#### aRNA sent to 29 labs (19 returned data so far: 18 ABL; 4 BCR; 11 GUS)

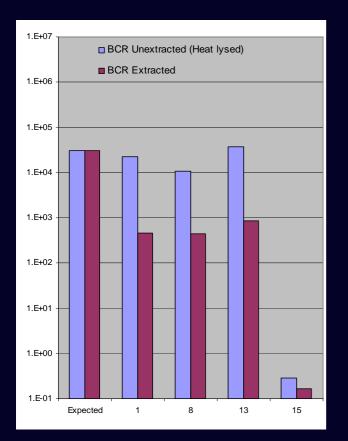
22 European: UK (5), France (4), Italy (3), Czech Republic (2), Germany (2), Spain (2), Austria (1), Finland (1), Greece (1), Sweden (1)

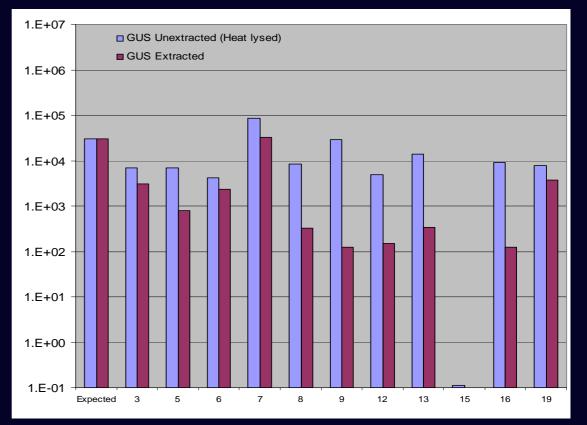
USA (3), Australia (1), Korea (1), Singapore (1), Japan (1)

## aRNA Field Trial Protocol



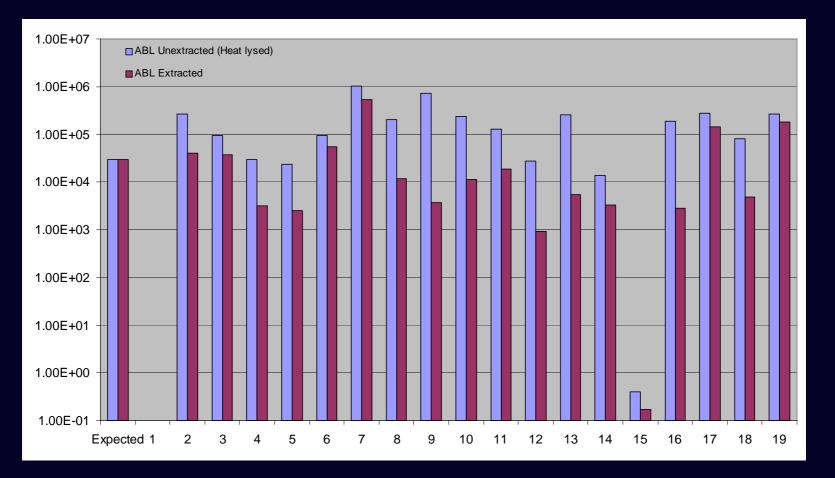
# aRNA results: BCR & GUS absolute copy numbers





Expected =  $3x10^4$ Median unextracted =  $1.6x10^4$ Median extracted =  $4.5x10^2$ n=4 Expected =  $3x10^4$ Median unextracted =  $7.8x10^3$ Median extracted =  $3.4x10^2$ n=10

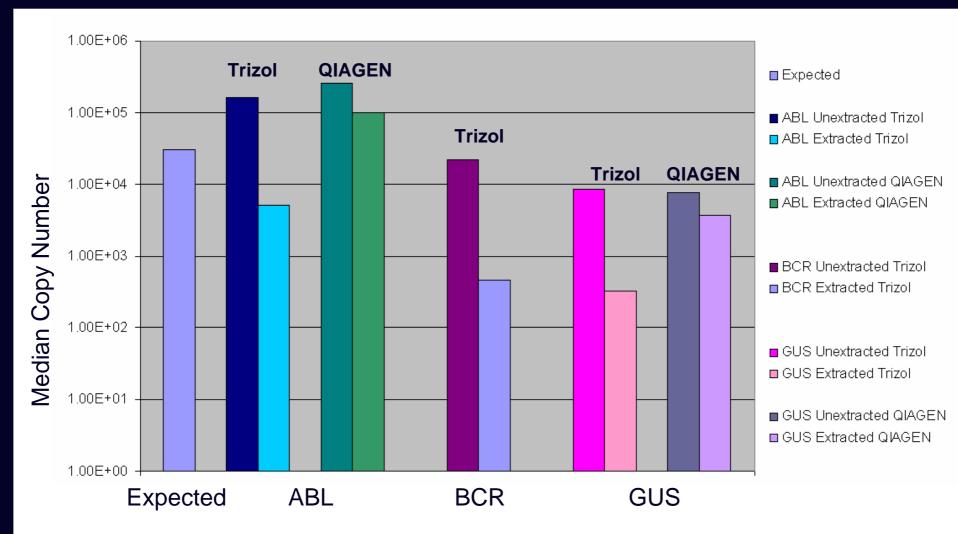
# aRNA results: ABL absolute copy number (corrected for protocol differences)



Expected =  $3x10^4$ ; Median unextracted =  $1.60x10^5$ ; median extracted =  $8.5x10^3$ ; n=18

Median ABL target values as formulated based on OD260 conversion and not NIST-traceable phosphate assay show an approximately 1 log difference from the expected value.

# aRNA losses during extraction: trizol vs Qiagen



## **Summary**

- aRNAs survived ambient temperature transportation
- aRNAs worked very well when heat lysed and directly converted to cDNA
- Variable losses on RNA extraction (particularly Trizol)
- Comparability between normalised values was good
- Next steps
  - Need to explore addition of carrier to improve Trizol extractions
  - Need to send out more concentrated aRNAs
  - Further trial?
    - » Adjustment of BCR/GUS/ABL ratios to those seen in leucocytes
    - » Provide test samples (and plasmid dilutions)
    - » Test ability of aRNAs to provide comparable results for test samples
    - » Test utility of spiking BCR-ABL RNAs into normal leucocytes

# **Acknowledgements**

- Adelaide
  - Tim Hughes, Sue Branford
- Mannheim
  - Andreas Hochhaus, Martin Muller
- Salisbury
  - Helen White
- NIBSC
- Asuragen
- International BCR-ABL Standardization Group







