Quality and Quantity Assessment of Nucleic Acids and Proteins

Ebru Serinsöz Linke, MD/PhD

- Quantification of nucleic acids
- Quality assessment of nucleic acids
- Quantification of proteins

• The amount and quality of molecules (nucleic

acids or proteins):

- Reproducibility
- Accuracy
- Efficiency

Quantification of Nucleic Acids

- Three quantification methods in common use:
 - Spectrophotometric measurement

(UV spectrometry)

- Fluorescent dye (Fluorometry) based measurement
- Real-time amplification (Absolute quantification)

- Measurement of light intensity at different wavelenghts
- *Transmittance*: the amount of light that passes completely through the sample
- *Absorbance*: measurement of light that is absorbed by the sample



- Bases in RNA/DNA absorb UV at 250~265nm
- Heterocyclic rings
- Measurements at A260nm, A280nm, A230nm
- Concentration estimation

- A260*nm*:
- Lambert-Beer Law: $C_{\mu g/\mu l} = A \times dilution factor \times \varepsilon$
- ε: molar extinction coefficient
 - physical constant
 - Unique
 - Amount of absorbance at 260nm of 1M nucleic acid solution measured in a 1cm path-length cuvette.

- A230 and A280 readings
 - A260/A280
 - A260/A230
- A260: DNA/RNA, Guanidine isothiocyanate
- A270: Phenol, TRIzol
- A280: Proteins
- A230: Phenol, TRIzol, Guanidine HCL

A260/A280: ~1.8 for DNA, ~2.0 for RNA

Low A260/A280 ratio

- Residual phenol or other reagent associated with the extraction protocol
- A very low concentration
- (> 10 ng/ul) of nucleic acid

High 260/280 ratio

• RNA/DNA contamination

• A260/A230*nm* ratio 2.0~2.2

Low A260/A230 ratio :

- Carbohydrate carryover (often a problem with plants).
- Residual phenol from nucleic acid extraction.
- Residual guanidine

(often used in column based kits)

• Glycogen used for precipitation.

High A260/A230 ratio

Making a Blank measurement

on a dirty pedestal

• Using an inappropriate

solution for the Blank

measurement.

Factors Affecting Absorbance

- A260/A280 ratio:
 - -рН
 - -ionic strength
- Water often has an acidic pH
- Buffered solution (Tris~EDTA at pH 8.0)

Conventional Spectrophotometers

• Conventional

spectrophotometers:

- Requires sample dilution
- Low sensitivity (lower
 limit 0.5~1µg nucleic
 acid)





- Miniaturization of UV spectrophotometers:
 - -Rapid
 - Direct quantification of nucleic acids in microvolumes



NANOSPECTROPHOTOMETRY~ NanoDrop

- Sample retension system
- Inherent surface tension of liquids
- Microvolume samples (0.5~2µl)
- Liquid column → Vertical optical path





- Vertical path length
 - Automatically changed
 - Shorter path length \rightarrow higher concentration of sample

- Benefits:
 - 1. Small sample volume at $0.5-2\mu l$
 - 2. Large dynamic range $(2ng/\mu l 3700ng/\mu l)$
 - 3. Cuvette free operation
 - 4. Short measurement time
 - 5. High accuracy and good reproducibility.





- DNA/RNA intercalating dyes
- Measurement of fluorescence

• \sim 1000 times more sensitive than UV absorbance

- Ethidium Bromide (EtBr)
- SYBR Green
- Hoechst 33258
- PicoGreen
- RiboGreen

EtBr:

- PCR products, gDNA
- Band intensity calculation
- Comparison to known
 reference
- Agarose and PAGE
- Not precise, relative

SYBR Green I:

- Highly sensitive
 - 25~100times more than

EtBr

- ssDNA, dsDNA
- Agarose and PAGE
- Less mutagenic

• Fluorometer

- Hoechst 33258 (DNA):
 - Binds to A-T bp in dsDNA
 - Emission Max. At 460nm

Emission max. at 530nm

- PicoGreen (DNA)
- RiboGreen (RNA)

TBS-380 Fluorometer 3800: 2mL or 50µl with minicell adapter



Aquaflour: 2mL





NanoDrop 3300 Fluorospectrometer : 2µl









The Qubit® 2.0 Fluorometer displays the standard curve after completion of the calibration.



Quantification by Real-Time PCR

- DNA, RNA (cDNA)
- Absolute quantification
- Serially diluted standards
 → Standard curve
- Determination of concentration of unknowns based on Ct (Threshold cycle) values



Quality Assessment of Nucleic Acids

Quality Assessment of Nucleic Acids

- Level of degradation
- Assay amenability (FFPE tissues)
- Method: nature of nucleic acid

Methods of Quality Assessment of Nucleic Acids

DNA

- 1. Agarose gel electrophoresis
- 2. PCR amplification of fragments with increasing length

RNA

- 1. Denaturing gel electrophoresis
- 2. RT-PCR amplification of mRNA fragments of increasing length

DNA and RNA

1. Agilent 2100 Bioanalyzer

Agarose gel electrophoresis (DNA)



PCR Amplification of Fragments with Increasing Length (DNA)

- Archival tissues
- Spesmen control size ladder
- Multiplex PCR



Denaturing Agarose Gel Electrophoresis (RNA)

- Most common method of integrity assessment :
- Secondary structure of RNA \rightarrow altered migration pattern
 - Electrophoresis buffer :Formaldehyde and MOPs (3-[N-Morpholino]-propanesulfonic acid)
 - Loading buffer: Glyoxal



RT-PCR Amplification of mRNA Fragments of Increasing Length

- mRNA integrity assessment
- Housekeeping gene
- cDNA subjected to amplification
- Fragments with increasing length
- Independent from rRNA integrity

Quality Assessment with Capillary Microchip Electrophoresis:

- LabChip systems (Caliper)
- MCE-202 MultiNA microchip electrophoresis system (Shimadzu)
- P/ACE MDQ (Beckman Coulter)
- **2100 BioAnalyzer (Agilent):** microfluidics, capillary electrophoresis, fluorescent detection

Agilent 2100 BioAnalyzer











Agilent 2100 BioAnalyzer





The RNA Integrity Number (RIN)

- 28S/18S rRNA ratio is reasonable but not ideal!!!
- Nondenaturing conditions
- Software algorithm for the the entire electrophoretic trace
- Estimation of the integrity of total RNA samples.
- Numbering system 1~10

RIN



RIN 1 : most degraded RIN 10: most intact



• Downstream application

-RT~PCR vs Microarray

- RINs > 7-8 work well for most experiments.
- RINs < 7 require extra validation studies

Quantification of Proteins

- Different methods:
 - Accuracy required
 - Amount and purity of protein

- Spectrophotometric assays:
 - UV Absorbance methods
 - Colorimetric and fluorescent-based detection

Quantification of Proteins

- Assay selection criteria:
 - 1. Sample volume (microplate assay vs cuvette-based)
 - 2. Sample recovery (UV spectroscopy)
 - 3. Throughput (microplate compatible rapid assay)
 - 4. Robustness (repeatability)
 - 5. Chemical modifications (Covalent modifications!)
 - 6. Protein aggregation (solubility of the protein)

Quantification of Proteins

- UV absorbance: quantitation of **purified** protein
 - Proteins that contain Trp, Tyr residues
 - Cys-Cys disulphide bonds
- Colorimetric assays: uncharacterized protein solutions and cell lysate
 - Bradford
 - BCA
 - Lowry

Quantification of Proteins: UV Absorption Spectroscopy

- UV Absorption at 280nm
 - Range 20~3000µg
 - Aromatic aa (tyrosine and tryptophan)
 - Molar extinction coefficient
 - Beer-Lambert Law: $A = a_m \ge C \ge 1$
 - Protein standard

Quantification of Proteins: UV Absorption Spectroscopy

- UV Absorption at 205nm
 - Range 1~100µg
 - Arsorption of photons by peptide bonds
 - Molar extinction coefficient at A205
 - Protein Standard

UV Absorption Spectroscopy

- NanoDrop 1000, NanoDrop 8000 A280 modules
- Concentration of **purified** protein samples
- 1µl sample

Dye-Based Assays: Bradford Assay

Bradford (Coomassie Blue) Assay (1~50µg):

• Binding of Coomassie Brillant Blue mainly to Tryptophan and tyrosine residues at acidic pH

• Shift in the absorbance of acidic CB solution from 465nm to 595nm

Bradford Assay







Dye-Based Assays: Lowry Assay

- Lowry (Alkaline Copper Reduction) Assay (5~100µg):
 - Two-step procedure
 - Reduction of Cu by proteins in alkaline solutions
 - Reduction of Folin reagent (a mixture of phosphotungstic acid and phosphomolybdic acid)
 - A blue color is formed with absorbance max. at 750nm

Dye-Based Assays: Bicinchoninic Assay

Bicinchoninic (BCA) Assay (0.2~50µg)

- Bicinchoninic acid (replacement of Folin's reagent)
- Improved sensitivity
- Tolerance to interfering substances
- Intense purple complex (562nm)

Protein analysis with the Agilent 2100 Bioanalyzer

- Microvolume analysis
- Different assays
 - For protein analysis in the low molecular weight range
 - General protein analysis up to 230 kDa
 - Picogram sensitivity

Quality Assessment of Proteins

- 1. Composition-Based and activity-based analysis
- 2. Electrophoretic methods (SDS gel electrophoresis)
- 3. Chromatographic methods
 - I. Gel filtration Chromatography

II. Reversed phase HPLC

- 4. Sedimentation velocity methods
- 5. Mass spectrometry methods
- 6. Light scattering methods