

# **ABMGG Molecular Genetics and Genomics CertLink**

## **Blueprint and Content Codes**

### **Blueprint**

<b>ID Number</b>	<b>Exam Content Topic</b>	<b>Approximate Percentage</b>
I	Normal structure and function	5%
III	Pathogenesis	35%
III	Application and interpretation	35%
IV	Laboratory management	10%
V	Nomenclature	5%
VI	Practice Guidelines	5%
VII	Emerging Applications	5%

## Content

- I. Normal structure and function
  - a) Chromosome/gene structure and function
    - i) RNA and DNA structures
    - ii) Elements/features of genes and RNA
    - iii) Transcription/translation/regulatory elements
    - iv) RNA processing
    - v) X-inactivation
- II. Pathogenesis
  - a) Mosaicism/pseudomosaicism
  - b) Mitochondrial mutations/heteroplasmy
  - c) Repeat expansion disorders
  - d) Point mutations
  - e) Imprinting/methylation
  - f) Polymorphisms/normal variants
  - g) Insertions/deletions/duplications
  - h) LOH/AOH/LCSH/UPD
  - i) Microsatellite instability
  - j) Monitoring minimal residual disease
  - k) Neoplasia
- III. Applications and interpretation
  - a) Core techniques, principles and practice
    - i) Copy number detection
      - (1) Chromosomal microarray
      - (2) MLPA
    - ii) Nucleic acid isolation
    - iii) PCR
      - (1) Qualitative
      - (2) Quantitative
    - iv) Allele discrimination
    - v) Sequencing
      - (1) Dideoxy sequencing
      - (2) NGS
    - vi) Methylation analysis
    - vii) Nucleic acid hybridization
    - viii) Targeted mutation analysis
    - ix) SNP analysis (molecular)
  - b) Diagnosis of symptomatic individuals
  - c) Predictive testing
    - i) Hereditary disease (presymptomatic)
    - ii) Pharmacogenetic
  - d) Carrier testing
  - e) Screening

- i) Population based
  - ii) Preconception/prenatal (including NIPT)
  - f) Prenatal diagnosis
  - g) Identity testing
    - i) Maternal cell contamination of fetal samples
    - ii) Paternity testing
  - h) Preanalytic variables
  - i) Analytic variables
  - j) Postanalytic variables
    - i) Interpretation of results
    - ii) Genotype:phenotype correlation
    - iii) Use of genomic databases
    - iv) Calculation of residual, recurrent risks
    - v) Test limitations and recommendations
    - vi) Variant classification
    - vii) Variants of uncertain significance
    - viii) Surveillance/clinical follow up, additional testing
    - ix) Bioinformatics
- IV. Laboratory management
- a) QA/QC
    - i) Professional and regulatory guidelines
    - ii) Proficiency testing
    - iii) Instrument validation and calibration
    - iv) Assay validation
    - v) Quality control/quality assurance
    - vi) Accuracy, precision, sensitivity, specificity
    - vii) Personnel competency assessment
  - b) Ethical/legal
    - i) HIPAA, GINA
    - ii) Patient consent/assent
    - iii) Genetic counseling
  - c) Lab safety
    - i) Chemical safety, fire safety
    - ii) Universal precautions, bloodborne pathogens
- V. Nomenclature
- a) Molecular nomenclature
- VI. Practice Guidelines
- VII. Emerging Applications