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Preparing for the American Board  
of Pathology (ABPath)  
Examination of Fundamental  
Knowledge and Skills

# Molecular Genetic Pathology

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*Content Specifications*

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## Overview:

### Molecular Genetic Pathology Content Specifications

This guide outlines the content related to molecular pathology that may appear on the American Board of Pathology Anatomic and Clinical Pathology exam, as well as the Molecular Pathology Subspecialty exam. It provides a framework based on the knowledge and skills that the trainee is expected to know or be able to perform ranging from the early learner through the Fellow-level.

#### Key to Designations:

C = Core/Foundational Knowledge

AR = Advanced Resident Knowledge

F = Fellow/Advanced Practitioner Knowledge

The exam assesses the knowledge, judgment, skills, and abilities necessary to identify specific entities, properly process specimens, and diagnose and monitor diseases using molecular methods. The specific diseases, tests, and concepts listed in this document are important for candidates to know, but it is not possible to create a fully comprehensive list of all the material needed for certification and effective practice. Candidates should use this guide as a reference for preparing for certification and professional practice.

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42 **1. Normal Structure and Function**

43	a. Chromosomes	C
44	b. Genes	C
45	c. DNA	
46	i. Exons, Introns, Non-Coding DNA	C
47	ii. Repetitive Elements (e.g., STRs, Microsatellite)	AR
48	d. RNA	
49	i. mRNA and tRNA	C
50	ii. miRNA and lncRNA	AR
51	e. Transcription, Translation, and Post-Translational Modification	C
52	f. Mitosis	C
53	g. Meiosis	C
54	h. Nomenclature	
55	i. Gene Nomenclature	C
56	ii. Protein Nomenclature	C
57	iii. Variant Nomenclature	AR

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59 **2. Molecular Genetic Principles**

60	a. Mechanisms and Types of Genetic Alterations/Defects	
61	i. Ploidy	C
62	ii. Copy Number Variants (CNV)	C
63	iii. Translocation and Other Rearrangements, including	
64	1. Deletions, Duplications, Inversions	C
65	iv. Single Nucleotide Polymorphisms (SNPs)	C
66	v. Methylation, Epigenetics	C
67	vi. Trinucleotide Repeats	C
68	1. Stabilizing Interruptions	F
69	vii. Multifactorial Events	C
70	viii. Mismatch Repair	C
71	ix. Point Mutations	C
72	x. Mosaicism	C
73	b. Inherited Defects	

74	i. Mendelian Inheritance	C
75	ii. Non-Mendelian Inheritance	C
76	iii. Oncogenes	C
77	iv. Tumor Suppressor Genes	C
78	v. Risk Calculations	C
79	vi. Hardy Weinberg Principle	AR
80	vii. Founder Mutations	F
81	viii. Bayes Analysis	F
82	c. Somatic Defects	
83	i. Oncogenes	C
84	ii. Tumor Suppressor Genes	C
85	iii. Loss of Heterozygosity (LOH)	AR
86	iv. Microsatellite Instability (MSI)	AR
87	v. Clonality	AR
88	vi. Genomic Instability	AR
89	vii. Chromothripsis	F

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### 91 3. Techniques and Methods

92	a. Cytogenetics	C
93	b. PCR, RT-PCR, and other NAAT	C
94	c. FISH	AR
95	d. Nucleic Acid Isolation & Quantitation	AR
96	e. Restriction Enzyme Digestion	AR
97	f. Fragment Analysis (i.e., Gels, Capillary Electrophoresis)	AR
98	g. Quantitative PCR and RT-PCR	AR
99	h. Nucleic Acid Sequencing (i.e., Sanger & Pyrosequencing)	AR
100	i. Next Generation Sequencing	AR
101	j. Constitutional Arrays (aCGH, SNP, and Oligo Arrays)	AR
102	k. Somatic Arrays (aCGH, SNP, and Oligo Arrays)	AR
103	l. Melt Curve Analysis	AR
104	m. Tumor Mutational Burden	AR
105	n. MLPA and Other Mutation Scanning Methods	F
106	o. Methylation Analysis	F
107	p. Mass Spectrometry	F
108	q. Proteomics	F

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### 110 4. Assay Performance and Validation

111	a. Proficiency Testing	AR
112	b. Validation versus Verification	AR
113	c. Preanalytical Considerations	AR
114	d. Analytical Considerations	AR
115	i. Stability	AR

116	ii. Specimen Selection	AR
117	iii. Specimen Collection	AR
118	iv. Anticoagulant	AR
119	v. Fixation	AR
120	e. Postanalytical Consideration, including	
121	i. Results, Interpretation, & Follow-up Testing	AR
122	f. Variant Classification	AR
123	g. Reporting	AR
124		
125	<b>5. Quality</b>	
126	a. Quality Assurance	AR
127	b. Quality Control	AR
128	c. Controls	
129	i. Internal Controls	AR
130	ii. Quantitative Controls	AR
131		
132	<b>6. Ethical, Legal, and Regulatory Issues</b>	
133	a. Ethical	
134	i. IRB	C
135	ii. Consent	C
136	iii. Secondary & Incidental Findings (e.g., ACMG List)	F
137	b. Legal	
138	i. HIPAA	C
139	ii. Genetic Information Non-Discrimination Act (GINA)	C
140	iii. Gene Patent	AR
141	c. Regulatory	
142	i. CLIA	AR
143	ii. CAP	AR
144	iii. CMS	AR
145	iv. FDA: LDT/LDP, IUO, RUO	AR
146	d. Coding and Reimbursement	
147	i. CPT	AR
148	ii. ICD	AR
149	iii. Laboratory Utilization	AR
150		
151	<b>7. Indications for Testing</b>	
152	a. Presymptomatic / Predictive	C
153	b. Diagnostic	C
154	c. Prenatal Diagnosis, including	
155	i. preimplantation genetic diagnosis (PGD)	C
156	d. Screening	

157	i.	Carrier Screening	C
158	ii.	Newborn Screening	C
159	iii.	Molecular Cancer Early-Detection Screening	F
160	e.	Cancer Monitoring (e.g., minimal residual disease, resistance)	F
161			
162			
163	<b>8.</b>	<b>Inherited Diseases</b>	
164	a.	Coagulation Disorders	
165	i.	Factor V Leiden (FVL)	C
166	ii.	Prothrombin	C
167	iii.	Factor VIII	C
168	iv.	Protein C	AR
169	b.	Non-Neoplastic Hematology	
170	i.	Thalassemia	C
171	1.	Methods of Detection	F
172	ii.	Hemophilia	C
173	iii.	Sickle Cell Disease	C
174	iv.	Hemochromatosis	C
175	1.	Penetrance	F
176	2.	Less Common Disease Producing Alleles	F
177	c.	Cardiovascular	
178	i.	Coronary Artery Disease; Hyperlipidemia	C
179	ii.	Heritable Cardiomyopathy	AR
180	iii.	Heritable Arrhythmias (e.g., Long QT)	AR
181	d.	Renal	
182	i.	Renal Cystic Disease	C
183	e.	Endocrine	
184	i.	Thyroid	C
185	ii.	Parathyroid	C
186	f.	Gastrointestinal	
187	i.	Hirschsprung Disease	AR
188	ii.	Celiac Disease	F
189	g.	Pulmonary	
190	i.	Cystic Fibrosis	C
191	1.	Variant Associated Therapy	F
192	2.	Genotype-Phenotype Correlation	F
193	3.	Residual Risk Based on Testing Method & Ethnicity	F
194	ii.	Alpha-1-Antitrypsin Deficiency	C
195	h.	Immunologic	
196	i.	Severe Combined Immunodeficiency (SCID)	C
197	i.	Biochemical Genetics	
198	i.	Metabolic Disorders	C
199		(e.g., Phenoketonuria (PKU), galactosemia, and	
200		fatty acid oxidation disorders)	

201	ii.	Storage Disorders	C
202		(e.g., Gaucher, Tay-Sachs, glycogen storage disorders)	
203	iii.	Leukodystrophies	C
204		(e.g., Krabbe, Canavan)	
205	j.	Mitochondrial Disorders	
206	i.	Inheritance Patterns	C
207		1. Nuclear versus Mitochondrial Genes	F
208	ii.	Heteroplasmy	C
209	k.	Neuromuscular Disorders/Dementia	
210	i.	Huntington	C
211	ii.	Alzheimer	C
212	iii.	Parkinson	C
213	iv.	Muscular Dystrophy	C
214	v.	Freidreich Ataxia	AR
215	vi.	Dystonia	F
216	vii.	Myotonic Dystrophia	F
217	viii.	Spinal Muscular Atrophy (SMA)	F
218	l.	Neurodevelopmental Variants	
219	i.	Down Syndrome	C
220	ii.	Fragile X	C
221	iii.	Autism Spectrum	F
222	m.	Cutaneous/Connective Tissue Disorders	
223	i.	Osteogenesis Imperfecta	C
224	ii.	Marfan Syndrome	C
225	iii.	Ehlers-Danlos Syndromes	AR
226	iv.	Epidermolysis Bullosa	AR
227	n.	Imprinting Disorders and Congenital Growth Disorders	
228	i.	Angelman/Prader-Willi Syndromes	C
229	ii.	Beckwith-Wiedemann/Russel-Silver Syndromes	F
230	iii.	Proteus Syndrome	F

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## 9. Inherited Cancer Syndromes

233	a.	Gastrointestinal Tumor Syndromes	
234	i.	Lynch Syndrome	C
235	ii.	APC-Associated Polyposis Conditions (e.g., FAP, GAPPS)	C
236	iii.	MUTYH Polyposis	AR
237	iv.	Peutz-Jeghers	AR
238	v.	Juvenile Polyposis Syndrome	AR
239	vi.	Hereditary Diffuse Gastric Cancer	AR
240	vii.	EPCAM Deletions	F
241	b.	Breast Cancer	
242	i.	BRCA 1 & 2	C
243	ii.	HRD Pathway Genes	AR

244	iii.	Cowden Syndrome	AR
245	iv.	Hereditary Diffuse Gastric Cancer	AR
246	v.	Reversion Mutations, Germline versus Somatic	F
247	vi.	Therapeutic and Management Implications	F
248	c.	Prostate Cancer (e.g., HRD Pathway, MMR)	F
249	d.	Pancreatic Cancer (e.g., HRD Pathway)	F
250	e.	Renal Cell Carcinoma	C
251	i.	<i>SMARCB1</i> -Deficient Renal Medullary Carcinoma	AR
252	f.	Endocrine	
253	i.	von Hippel-Lindau	C
254	ii.	Multiple Endocrine Neoplasia (MEN) 1 & 2	C
255	g.	Soft Tissue & Bone	
256	i.	Neurofibromatosis	C
257	ii.	Li-Fraumeni Syndrome	C
258	iii.	Carney-Stratakis Syndrome	
259		AR	
260	h.	Hematologic	F
261	i.	Müllerian	
262	i.	<i>BRAC 1 &amp; 2</i>	C
263	ii.	Lynch Syndrome	C
264	iii.	<i>POLE &amp; POLD1</i>	F

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## 10. Solid Tumors, Sporadic

266	a.	Breast	
267	i.	HER2 Status	C
268	ii.	HER2 FISH Interpretations	AR
270	iii.	Prognostic Gene Expression Panels	AR
271	iv.	HER2 Detection by Other Methods	F
272		1. (e.g., <i>ESR1</i> , <i>PIK3CA</i> )	
273	b.	Uterine, Cervical	
274	i.	HPV Status	C
275	ii.	HPV Molecular Methods	F
276	c.	Müllerian	
277	i.	TP53 Diagnostic Classification	AR
278	ii.	MMR Status	AR
279	iii.	<i>POLE</i> Mutation	AR
280	iv.	HER2 Amplification	AR
281	v.	HRD Pathways, including BRCA 1 & 2	AR
282	vi.	<i>KRAS</i> Mutation	F
283	d.	Lung	
284	i.	EGFR	C
285	ii.	<i>KRAS</i> Mutation	C
286	iii.	Guideline Recommended Gene Targets	AR



287	iv.	On versus Off Resistance Mechanisms	AR
288	v.	Biomarkers for Adjuvant Therapy (e.g., EGFR)	F
289	vi.	DNA versus RNA for Biomarker Assessment (e.g., <i>MET</i> Exon 14)	F
290	e.	Gastrointestinal (GI)	
291	i.	Upper GI	
292		1. HER2	AR
293		2. MMR	F
294	ii.	Colorectal	
295		1. MMR	C
296		2. Sporadic MSI/MLH1	AR
297		3. <i>RAS/RAF</i> Mutation Testing	AR
298		4. HER2	F
299		5. <i>POLE</i>	F
300	iii.	Pancreas	
301		1. <i>RAS</i>	F
302		2. Fusion	F
303	f.	Genitourinary	
304	i.	Prostate	
305		1. <i>TMPRSS2::ERG</i> Fusion	AR
306		2. Homologous Recombination Deficiency (HRD)	F
307	ii.	Bladder	
308		1. <i>FGFR</i> Mutations	F
309		2. <i>TP53</i> Mutation Status	F
310	g.	Central Nervous System	
311	iii.	Molecular Classification	AR
312		1. (e.g., <i>TERT</i> Promoter Mutations, <i>IDH1</i> & 2, <i>BRAF</i> Fusion)	
313	iv.	<i>MGMT</i> Promoter Methylation	AR
314	v.	Methylation-Based Classifiers	F
315	h.	Endocrine	
316	i.	Thyroid	
317		1. <i>BRAF</i> Status	C
318		2. <i>RET</i> Fusion Status	C
319		3. <i>RAS</i> Family Variants	AR
320		4. Targetable Alterations	F
321		5. Multigene Classifiers (e.g., for indeterminate cytology)	F
322	ii.	Pheochromocytoma	F
323		1. <i>SDH</i> Gene Family Deficiency	F
324	iii.	Neuroblastoma	AR
325		1. N- <i>MYC</i>	AR
326	i.	Bone and Soft Tissue (including GIST)	
327	i.	Diagnostic Genetic Alterations	AR
328	j.	Cutaneous	
329	i.	Melanoma	C
330		1. <i>BRAF</i>	C

331	2. Site Specific Mutations (i.e., Cutaneous, Mucosal, Uveal)	AR
332	3. UV Mutational Signatures	F
333	4. Molecular Classifiers (e.g., Microarray, FISH, TERT)	F
334	ii. Merkel Cell Carcinoma	AR
335	iii. Squamous and Basal Cell Carcinomas	F
336	k. Pan-Tumor Markers (e. g., Tumor Mutational Burden (TMB), MSI, <i>NTRK</i> Fusions)	
337	F	
338		

## 339 11. Hematopathology

340	a. Clonality (e.g., IGH, IGK, TRB, TRG)	AR
341	b. Disease Monitoring	AR
342	c. Specific Disorders	
343	i. Lymphoid	
344	1. B-ALL	
345	a. <i>BCR::ABL1</i>	AR
346	b. <i>ETV6::RUNX1</i>	F
347	c. <i>TCF3::PBX1</i>	F
348	d. <i>KMT2A::AF4</i>	F
349	e. <i>Ph</i> -Like ALL (e.g., <i>CRLF2</i> -Rearranged)	F
350	2. Mature T cell	
351	a. <i>NPM1::ALK</i>	AR
352	b. <i>JAK-STAT</i> Pathway	F
353	3. Mature B cell	
354	a. Common Translocations	
355	i. (e.g., <i>BCL2</i> , <i>BCL6</i> , <i>CCND1</i> , <i>MYC</i> )	AR
356	b. Recurrent Mutations	
357	ii. (e.g., <i>BRAF</i> , <i>MYD88</i> , <i>TP53</i> )	
358	c. <i>JAK-STAT</i> Pathway	F
359	d. <i>IGH</i> Somatic Hypermutation	F
360	e. Predictive Biomarkers (e.g., <i>BTK</i> , <i>PLCG2</i> , <i>EZH2</i> )	F
361	4. Plasma cell	F
362	ii. Myeloid	
363	1. <i>MPN</i> -associated mutations (e.g., <i>JAK2</i> , <i>CALR</i> , <i>MPL</i> , <i>CAF3R</i> )	C
364	2. Clonal hematopoiesis (e.g., <i>DNMT3A</i> , <i>TET2</i> , <i>ASXL1</i> )	AR
365	3. <i>MDS</i> -associated mutations (e.g., <i>SF3B1</i> , <i>EZH2</i> , <i>TET2</i> )	AR
366	4. Kinase Fusions (e.g., <i>PDGFR A/B</i> , <i>FGFR1</i> , <i>JAK2</i> , <i>ABL1</i> )	AR
367	5. AML	
368	a. Associated mutations (e.g., <i>NPM1</i> , <i>FLT3</i> , <i>CEBPA</i> )	AR
369	b. Associated translocations	AR
370	(e.g., <i>PML::RARA</i> , <i>RUNX1::RUNX1T1</i> , <i>CBFB::MYH11</i> )	
371	6. CML	
372	a. Diagnostic studies	C
373	b. Monitoring studies	AR

374	c. Mastocytosis-associated mutations	AR
375	d. Resistance	F
376		

377 **12. Pharmacogenetics**

378	a. Cytochrome P450 (e.g., CYP2D6, CYP2C9, CYP2C19)	C
379	b. Other Drug Metabolizing Enzymes	F
380	(e.g., UGT1A1, Thymidine Kinase, DPYD, TPMT)	
381	c. HLA-associated Pharmacogenetics (e.g., abacavir, carbamazepine).	F
382		
383		

384 **13. Infectious Disease**

385	a. Detection	C
386	b. Differentiation/Subtyping	AR
387	c. Antimicrobial Resistance	AR
388	d. Sequencing	F
389	e. Molecular Epidemiology	F
390	f. Viruses	
391	i. Hepatitis	C
392	ii. HIV	C
393	iii. HPV	C
394	iv. Viral Infections of Immunocompromised Hosts	AR
395	v. CNS Viral Infections	AR
396	vi. Respiratory Viruses	AR
397	vii. Gastrointestinal Viruses	AR
398	g. Bacteria	
399	i. <i>Staphylococcus aureus</i> and MRSA	C
400	ii. <i>Clostridioides difficile</i>	C
401	iii. Respiratory Bacteria	AR
402	iv. Mycobacteria	AR
403	v. Gastrointestinal Pathogens	
404	1. <i>E. coli</i>	AR
405	2. <i>Campylobacter</i>	AR
406	vi. Sexually Transmitted Pathogens (e.g., GC/CT, <i>Trichomonas</i> )	AR
407	vii. Central Nervous System Pathogens	AR
408	viii. Microbiome / Metagenomics	F
409	h. Fungi	AR
410	i. Parasites	AR
411		

412 **14. Identity Testing**

413	a. Chimerism	AR
414	b. Maternal Cell Contamination Studies	AR
415	c. Sample Contamination	AR

416	d. Paternal/Relationship Testing	F
417	e. Forensic Applications	F
418	f. Gestational Disease (Molar Pregnancy)	F
419		
420		
421	<b>15. Histocompatibility</b>	F
422		
423	<b>16. Informatics</b>	
424	a. Genomic Databases (e.g., CoSMIC, gnomAD, ClinVar)	AR
425	b. NGS Bioinformatics Pipelines	F
426		
427	<b>17. Gene Therapy</b>	
428	a. Therapeutic mechanisms	
429	i. Gene Editing (e.g., CRISPR)	F
430	ii. Gene Transfer (e.g., Viral Constructs)	F
431	iii. Gene Modifying (e.g., CFTR, DMD, SMA)	F
432	b. Adverse Events	F
433	c. Ethical Considerations	F