

Certainly! Below is a **concise, genetics-focused summary** of each disorder as per your professor's requirements, covering:

1. Genetic Defect | **2. Pathophysiology** | **3. Inheritance Pattern**

1. Xeroderma Pigmentosum (XP)

- **Genetic Defect:**
 - Mutations in **XPA-XPG** genes (NER pathway).
 - **Types:** Missense, nonsense, or deletions disrupting DNA repair.
- **Pathophysiology:**
 - Defective NER → UV-induced pyrimidine dimers persist → DNA mutations → skin cancers.
- **Inheritance:** Autosomal recessive (25% recurrence risk if both parents are carriers).

2. Osteogenesis Imperfecta (OI)

- **Genetic Defect:**
 - Mutations in **COL1A1**/**COL1A2** (glycine substitutions common).
 - **Types:** Null alleles (mild) vs. structural mutations (severe).
- **Pathophysiology:**
 - Abnormal collagen fibrils → brittle bones, fractures, blue sclerae.
- **Inheritance:**
 - Autosomal dominant (Types I–V) or recessive (Types VI–VIII).

3. Cystic Fibrosis (CF)

- **Genetic Defect:**
 - **F508del** (3-bp deletion in **CFTR**) → misfolded protein (Class II).
 - Other classes: Nonsense (Class I), gating defects (Class III).
- **Pathophysiology:**
 - Loss of CFTR Cl⁻ channel → thick mucus → lung/pancreatic damage.
- **Inheritance:** Autosomal recessive (25% risk if both carriers).

4. Retinoblastoma

- **Genetic Defect:**
 - Biallelic loss of **RB1** (tumor suppressor).
 - **Types:** Germline (40%) or somatic (60%) mutations.
- **Pathophysiology:**
 - Dysfunctional pRb → uncontrolled cell cycle → retinal tumors.
- **Inheritance:** Autosomal dominant (familial cases; 50% transmission).

5. Huntington Disease (HD)

- **Genetic Defect:**
 - CAG repeat expansion (>36) in **HTT** (gain-of-function).
- **Pathophysiology:**
 - Toxic polyglutamine aggregates → neuronal death → chorea/dementia.
- **Inheritance:** Autosomal dominant (50% risk; anticipation if paternal transmission).

6. Marfan Syndrome

- **Genetic Defect:**

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- **Genetic Defect:**
 - Mutations in ***FBN1*** (fibrillin-1) → dominant-negative effect.
- **Pathophysiology:**
 - Dysregulated TGF- β signaling → aortic aneurysms, lens dislocation.
- **Inheritance:** Autosomal dominant (50% risk).

7. Hemophilia A

- **Genetic Defect:**
 - Mutations in ***F8*** (factor VIII): inversions (45%), deletions, nonsense.
- **Pathophysiology:**
 - Impaired clotting cascade → spontaneous bleeding (joints/muscles).
- **Inheritance:** X-linked recessive (50% risk to sons of carrier mothers).

8. Duchenne Muscular Dystrophy (DMD)

- **Genetic Defect:**
 - Deletions (65%) or duplications in ***DMD*** → frameshift → no dystrophin.
- **Pathophysiology:**
 - Loss of dystrophin → muscle membrane instability → degeneration.
- **Inheritance:** X-linked recessive (50% risk to sons of carriers).

9. Color Vision Deficiencies

- **Genetic Defect:**
 - Unequal crossover in ***OPN1LW*** (red) or ***OPN1MW*** (green) opsins.
- **Pathophysiology:**
 - Altered cone photopigments → red/green colorblindness.
- **Inheritance:** X-linked recessive (higher prevalence in males).

Key Patterns Summary Table

Disorder	**Gene**	**Genetic Defect**	**Inheritance**
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XP	*XPA-XPG*	NER pathway mutations	Autosomal recessive
CF	*CFTR*	F508del (Class II)	Autosomal recessive
HD	*HTT*	CAG repeat expansion	Autosomal dominant
DMD	*DMD*	Deletions (frameshift)	X-linked recessive
Hemophilia A	*F8*	Inversions/nonsense	X-linked recessive

10 Genetics-Focused MCQs

1. ***XP results from mutations in:***

- Mismatch repair genes
- *XPA-XPG* (NER pathway)**
- *BRCA1***

****Answer: b****

2. ***F508del in *CFTR* causes:***

- Hyperactive Cl⁻ channel
- Misfolded protein degradation
- Enhanced mRNA splicing

****Answer: b****

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1. **XP results from mutations in:**

- a) Mismatch repair genes
- b) *XPA-XPG* (NER pathway)
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Answer: b

2. **F508del in *CFTR* causes:**

- a) Hyperactive Cl⁻ channel
- b) Misfolded protein degradation
- c) Enhanced mRNA splicing

Answer: b

3. **HD's CAG repeats are in:**

- a) *HTT*
- b) *FMR1*
- c) *DMPK*

Answer: a

4. **DMD patients lack:**

- a) Myosin
- b) Dystrophin
- c) Actin

Answer: b

5. **Marfan Syndrome involves mutations in:**

- a) *COL1A1*
- b) *FBN1*
- c) *DMD*

Answer: b

This summary **strictly adheres** to your professor's focus: **genetic defect → pathophysiology → inheritance**. Let me know if you'd like to adjust the depth for any disorder!