

1. **Ex. 2:** Identification and Separation of Cellular Biomolecules.
 - a. Organic compounds: Energy/Calorie (kilocalorie); macromolecules:
 - b. Amino acids/Proteins, peptide bonds, **Ninhydrin**, **Rf values**; condensation/hydrolysis; paper chromatography (acetone/acetic acid).
 - c. Carbohydrates – boil in **Benedict's reagent** (red Cu⁺ precipitate - reducing sugars); Polysaccharides – brown/purple with **Iodine** (KI).
 - d. Lipids – water insoluble; → **Sudan IV**, red droplets.

2. **Ex. 2b, 3:** **Gel Filtration**; **Diffusion and Osmosis**
 - a. Mobile, immobile phase; **G-25 Sephadex beads**; Hypothesis; Benedict's reagent, **Lugol's Iodine**; **Eluant** Volume; relative rates of large vs. small molecule passage thru column.; 0.9% NaCl
 - b. Concentration gradient, simple diffusion; Facilitated diffusion – transporter proteins; Active Transport; **Osmosis** – Osmotic pressure; **isotonic, hypertonic, hypotonic**; *Elodea* plasmolysis, swelling? With range of Sucrose concentrations: 0.2M, 0.4M, 0.6M; Osmotic Lysis – **Blood cells/hemolysis** with 0.3M glucose, fructose, glycerol, urea, CaCl₂ or NaHCO₃?
 - c. Chloride – precept. with **AgNO₃**; Protein – **Biuret** turns pinkish-violet; Plasmolysis, Hemolysis (RBC's)

3. **Ex. 4:** Protein concentration and Standard Curve.
 - a. BSA, **Standard Curve**, Spectrophotometer; Visible light; Reflected vs. Absorbed light wavelengths (color wheel); **Bradford (Coomassie Brilliant Blue) Dye Reagent**; **A₅₉₅**

4. **Ex. 5, 6:** Enzyme Activity – Mushroom **Tyrosinase**
 - a. Enzyme, substrate, active site; Tyrosine → **DOPA** → Dopaquinone → **DOPAchrome** → → → Melanin.
 - b. Mushroom caps, **sodium phosphate buffer** – pH 4.0, pH 6.0, pH 8.0. **A₄₇₅**.
 - c. Other factors affecting enzyme activity: Concentration of **products**, concentration of **substrates/reactants**, temperature, (pH), salt concentrations, pressure, biological activators/inhibitors.
 - d. **Native PAGE** – **polyacrylamide**, isoelectric point, loading/sample buffer (bromophenol blue, glycerol, Tris buffer pH 8.6), electrophoresis buffer (Tris buffer & Glycine, pH 8.3), denaturing, native. Substrate (DOPA) concentration, pH and detection conditions.

5. **Ex. 9, 10:** DNA EXTRACTION & RESTRICTION DIGEST OF Suspected Romanov DNA – Anastasia???:
 - ❖ Detergents (**SDS**), **Tris** buffer (pH 8.0), **EDTA**, cold **ethanol**, ice, **potassium acetate**, **aceto-orcein** stain. **A₂₆₀**, **A₂₈₀**, **A_{260/A280} ratio**. 1.0 A₂₆₀ = 50 µg/ml DNA. Quartz cuvettes. Filtrate, retentate.
 - ❖ **Restriction endonucleases** cut DNA at specific, **palindromic sites** (read 5'→3'; 3'→5') – *eg. "DNA Land; Evil Olive; Racecar; Yo Banana Boy!"*; GAATTC/CTTAAG (for EcoR-I enzyme) – cuts between G and A in each strand; We used **HinD-III** → recognizes AAGCTT//TTCGAA, cutting between each pair of A's. These are our "Molecular Scissors" in DNA analysis and Recombinant DNA Technology." 37°C Incubation.
 - **Agarose Gel electrophoresis**; **Ethidium Bromide** stain. **Rf migration** (= ration of band migration/bromophenol blue dye front migration = R_{band}/R_{dye front}) values. **Bromophenol Blue**, **Xylene Cyanol** tracking dyes; **TAE (Tris-Acetate EDTA) electrophoresis** buffer, pH~8.

6. **Ex. 11:** D1S80 PCR Amplification: Molecular markers, **VNTRs**, Marion = 18 and 24 repeat alleles. **Polymorphism**. Calculate Rf and number of repeats → genotype from each of your parents.
 - **Are you a clone?** Be able to support your answer!!! What were **your** fragment sizes, repeat numbers, and final D1S80 genotypes? _____, _____. What are the **Alien** fragment/allele sizes?? _____
 - **Primer Mix**; **Master Mix**.
 - PCR cycles = 95°C **melt** template, ~60°C **anneal** primers, 72°C **elongate** with **Taq DNA polymerase**
 - **n** number of cycles, **2ⁿ** = number of copies of a single DNA template after n cycles of PCR.
 - **DNA Acrylamide electrophoresis** – fine separation of only a few hundred bp fragments; BPB tracking dye, Ethidium Bromide stain. TBE (Tris Borate EDTA) buffer.

7. **Ex. 12:** **TRANSFORMATION & OPERON Induction:** Plasmids = self-replicating “mini-chromosomes”; Recombinant pGLO plasmid has Jellyfish **GFP** (“**reporter gene**”) under control of **Arabinose Operon Promoter (P_{BAD})**. Our E. coli bacterial cells keep the plasmid, because we grow them in the presence of Ampicillin – and ampicillin resistance is encoded on pGLO – so they will die without it (selectable marker on the plasmid).
- **AraC Activator / Inhibitor Protein** binds near the **P_{BAD} promoter**, enhancing transcription of the promoter only when **Arabinose, the substrate of the structural genes, is present (and glucose is absent!!)**. AraC can distinguish between the stereoisomers/ **enantiomers**, **D-arabinose** and **L-arabinose**. **CaCl₂** (neutralizes DNA and bacterial cell wall and plasma membrane) and heat-shock at **42°C** loosen the cell wall and cell membrane to facilitate (make cells “competent”) for transformation by the pGLO plasmid (**amp^R gene & ara operon promoter** connected to **GFP**).
 - Transformed bacteria were selected on nutrient agar plates with **Ampicillin** and D or L arabinose, and with or without the presence of glucose. **UV light (or Black light = long wavelength violet)** was used to visualize colonies expressing Green Fluorescent Protein (produced when P_{BAD} is activated by the proper Arabinose enantiomer.).
8. **Ex. 14, 15:** **MITOSIS, MEIOSIS, and FERTILIZATION:**
- Chromosomes, sister chromatids, crossing over, centromeres, kinetochores, microtubule spindle fibers, asters, centrioles/centrosomes,
 - **Mitosis:** Prophase, (prometaphase), Metaphase, Anaphase, Telophase; Interphase.
 - **Meiosis:** Interphase, Pro-I, Meta-I, Ana-I, Telo-I (Interkinesis); Pro-II, Meta-II, Ana-II, Telo-II.
 - **Recognize all stages of mitosis!!**
 - **Monohybrid cross:** 3:1 phenotypic ratio (1:2:1, AA:Aa:aa, genotypic ratio)
 - **Dihybrid Cross:** 9:3:3:1 phenotypic ratio; but still 3:1 ratio for each gene!!
- *Homozygote, Heterozygote, Allele, Gene, Trait, Genotype, Phenotype, Dominant, Recessive.*
 - *Diploid, haploid, Trisomy, Monosomy, LOCUS, independent assortment, segregation of alleles.*

GENERAL:

- **Skills:** (Micro)Pipetting, weighing, measuring, calculating migrations (R_f), PCR, electrophoresis
 - Extracting macromolecules - DNA, Protein
- Comparing Chromatography/**separation techniques:** paper, column (Gel Filtration: affinity/exchange/HIC); fluid phase, solid phase, electrophoresis (polyacrylamide gel, Agarose gel).
- Know your **D1S80 Genotype!!!!** Do the calculations, if you have not already!!! 😊

➤ **Question Types:** fill-in/terminology; multiple-choice; problem solving, graphs, and data analysis; lab procedures (manual/hands-on activity!).