Smooth muscle	CARDIAC MUSCLE	SKELETAL MUSCLE
INTERNAL ORGANS	HEART	LEG
Anesthesia Structure/Function of Skeletal Muscle Muscle:		
Light I band Dark A band	Mitochondrion Sarcolemma Myofibril	
Sarcoplasmic Sarco reticulum Thin (actin) filament Z disc H z		

Muscle fiber: large multinucleate cells that are fused myblasts \rightarrow contractile proteins = myofibril, striation due to arrangement of myofibrils

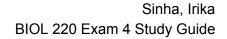
Sarcomere: Thin **actin** filaments with regulatory **troponin** and **tropomyosin** anchored to z-disk. Thin **myosin** filaments which consist of two globular heads and a long tail.

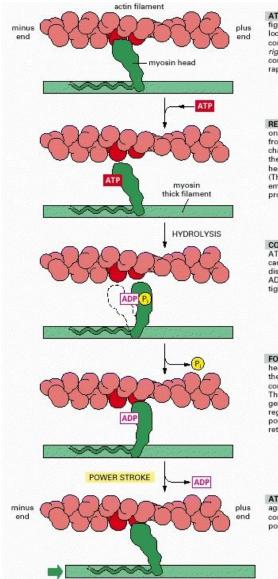
Calcium ²⁺ release:

 Ca^{2+} stored in the sarcoplasmic reticulum \rightarrow action potential reaches muscle cells \rightarrow spreads down T-tubules \rightarrow conformational change in the voltage sensitive DHP receptor \rightarrow RYR receptor (Ca²⁺ channel) opens \rightarrow Ca²⁺ release causes muscle contraction

Cross-Bridge Cycling:

- Hydrolysis of 1 ATP molecule required \rightarrow rigor mortis without ATP
- Myosin cross-bridge attached to actin → ATP binding leads to dissociation of myosin from actin → shape changed to mysoin head is "cocked" after hydrolysis although ADP and Pi bound → myosin binds to actin + weak initial binding releases P → powerstroke occurs and ADP releases; myosin head returns to low energy conformatino → generates force and pulls actin towards the center of the sarcomere
 - **Tension** (amount of force produced by muscle) depends on proportion of active cross-bridges





ATTACHED At the start of the cycle shown in this figure, a myosin head lacking a bound nucleotide is locked tightly onto an actin filament in a *rigor* configuration (so named because it is responsible for *rigor mortis*, the rigidity of death). In an actively contracting muscle, this state is very short-lived, being rapidly terminated by the binding of a molecule of ATP.

RELEASED A molecule of ATP binds to the large cleft on the "back" of the head (that is, on the side furthest from the actin filament) and immediately causes a slight change in the conformation of the domains that make up the actin-binding site. This reduces the affinity of the head for actin and allows it to move along the filament. (The space drawn here between the head and actin emphasizes this change, although in reality the head probably remains very close to the actin.)

COCKED The cleft closes like a clam shell around the ATP molecule, triggering a large shape change that causes the head to be displaced along the filament by a distance of about 5 nm. Hydrolysis of ATP occurs, but the ADP and inorganic phosphate (P_i) produced remain tightly bound to the protein.

FORCE-GENERATING A weak binding of the myosin head to a new site on the actin filament causes release of the inorganic phosphate produced by ATP hydrolysis, concomitantly with the tight binding of the head to actin. This release triggers the power stroke—the forcegenerating change in shape during which the head regains its original conformation. In the course of the power stroke, the head loses its bound ADP, thereby returning to the start of a new cycle.

ATTACHED At the end of the cycle, the myosin head is again locked tightly to the actin filament in a rigor configuration. Note that the head has moved to a new position on the actin filament.

Calcium regulation of contraction

Ca²⁺ released from the sarcoplasmic reticulum and binds to **troponin**. Troponin holds **tropomyosin** in lace and blocks myosin binding sites

Tension is a function of Ca^{2+} concentration \rightarrow relaxation due to CaATPase pumping Ca out of cytoplasm

Excitation-Contraction Coupling:

T-tubules - invaginations of sarcolemma in muscle fiber \rightarrow AP in muscle fiber conducted into interior of muscle cell along T-tubles \rightarrow lumen continuous w/ extracellular fluid

DHP is voltage gated Ca2+ channel and directly connected to RYR on SR \rightarrow pulls open Action potentials determine rate of Ca²⁺ leaving SR

Gradient is a Ca²⁺ EC gradient, resistance is number of open channels

Neuromuscular Junction:

Motor neuron to muscle cell synapse \rightarrow Ach released \rightarrow binds to muscle cell \rightarrow causes "end plate" graded potential = always action potential in healthy muscle cell

Acetylcholine binds to AchR, a cation channel that moves both Na⁺ and K⁺ \rightarrow more Na⁺ in that potassium out \rightarrow depolarization

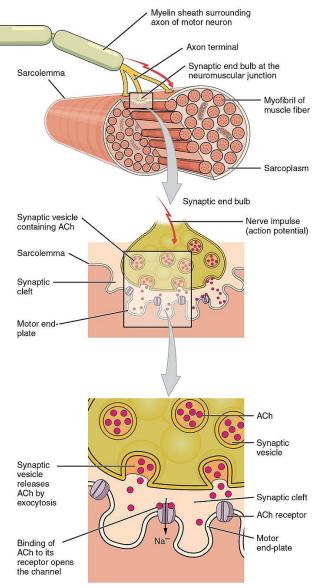
Acetylcholinesterase breaks down acetylcholine

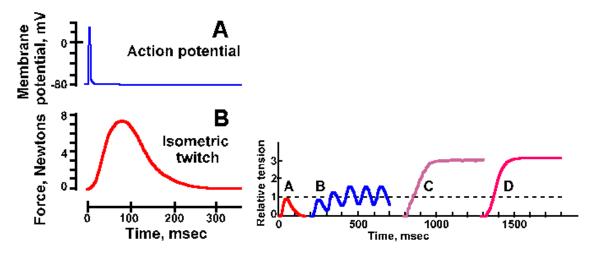
End plate potential: depolarizations of skeletal muscle fibers caused by neurotransmitters binding to the postsynaptic membrane in the neuromuscular junction ---> no AP

- Each muscle cell has one neuromuscular junction = signals from one somatic efferent neuron
- Action potential in neuron releases ACh which binds to DHP and Na⁺ enters the cell, causing depolarizing EPSPs above threshold and triggering AP
 - In muscle cell EPSP always above threshold so always triggers AP → degree that EPSP exceeds threshold = safety factor
- Motor Unit: A somatic efferent neuron and all the fbers it innervates

Skeletal muscle twitch: single contraction and

relaxation cycle produced by an action potential within the muscle fiber itself; doesn't occur in healthy individual





Tetany: Summation of action potentials = tentany \rightarrow repeated simulation \rightarrow steady state of tension

Control Systems general model of physiology

Sense + Respond to internal/external environment

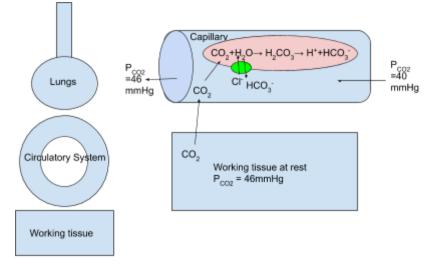
- 1. Stimulus
- 2. Receptor
- 3. Input
- 4. Integrator
- 5. Output
- 6. Target
- 7. Response

CO_2 and O_2 in the Blood

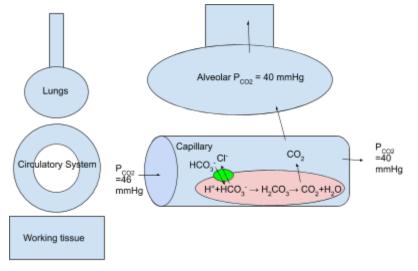
 $\rm CO_2$ enters capillary then blood cell, converted to bicarb in blood cell through set of reactions Carbonic anhydrase catalyzes carbon dioxide and water to carbonic acid

CI- moved in for charge balance

CO2 enters blood and stored as bicarb



CO2 out of bicarb then out of blood at lungs



 $\text{Bicarb} \rightarrow \text{RBC} \rightarrow \text{H2CO3} \rightarrow \text{CO2} \rightarrow \text{plasma} \rightarrow \text{aveoli}$

Chemosensors: surface of medulla, sensitive to PCO2 and pH of cerebrospinal fluid

Chemosensors on large blood vessels leaving heart sensitive to decreased O2 availability

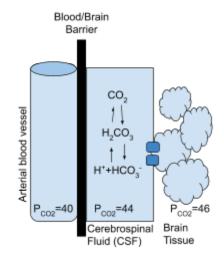
Bodies sense high arterial pCO2 + increase ventilation

• Stimulus: low pH, low CO2

- Receptor: H+ gated sodium channel on central chemoreceptor
- Input: AP from central chemoreceptor (sensory neuron)
- Integrator: respiratory center
- Output: phrenic nerve
- Target: diaphragm
- Response: Increase ventilation

Involuntary Ventillation Regulation: Central chemoreceptors

Amount of CO2 in CSF determined by amount in from tissue and amount out into artery by diffusion (mass balance)



SA Node cells

F-Na+ depolarizes cell automatically

Norepinephrine (NE): increases permeability to Na+ in F-Na+ channels \rightarrow increases heart rate

Acetylcholine: slows V-K closing (stronger hyperpolarization) = lower heart rate (opens K+ channels)

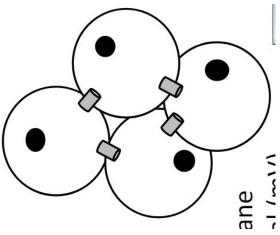
Blood pressure: measurement of force applied to artery walls

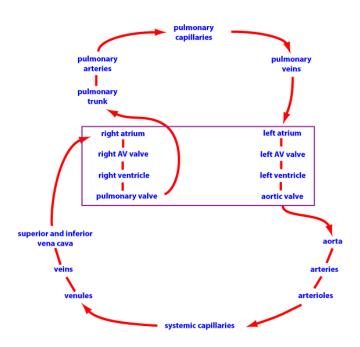
- peripheral resistance, blood volume, and cardiac output
- PR= Blood vessel diameter, blood viscosity, and total vessel length

Mean Arterial Pressure: $CO \times TPR = MAP$ $CO = HR \times SV$

CO = cardiac output HR = heart rate SV = stroke volume TPR = total peripheral resistance

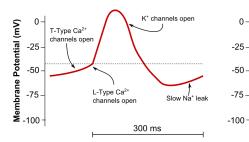
SV changes due to end-diastolic volume (EDV) = how much total volume heart has before contraction/end of filling



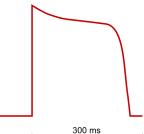


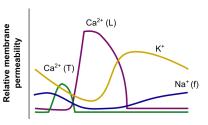
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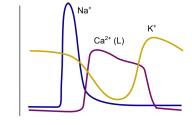
Pacemaker AP











Frank-Starling: stroke volume of the heart increases in response to an increase in the volume of blood in the ventricles

Cardiac muscle cells = different to skeletal muscle but still striated due to sarcomeres **Function:** pump blood \rightarrow intercalated disks, 1 or 3 nuclei, branched

Intercalated disks connect cardiac muscle cells and are critical for the flow of electrical signals from cell to cell and, thus, coordination of heartbeat

Gap junctions = electrical coupling **Desmosomes** = provide strength

Heartbeat: SA node initiates signal \rightarrow AV node contracts \rightarrow bundle of His \rightarrow perkinje fibers \rightarrow ventricular cells contract

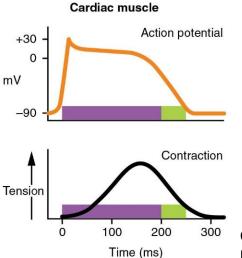
Cardiac muscle cell excitation:

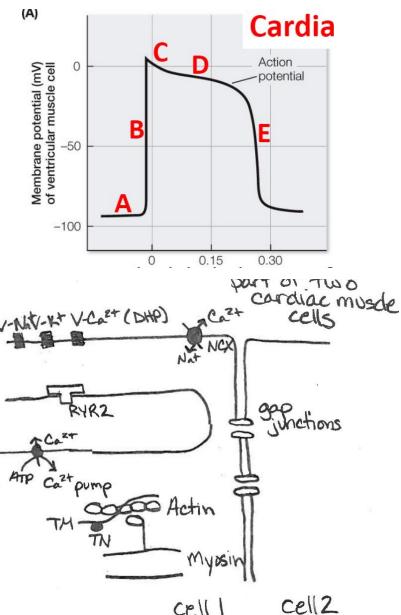
- 1. V-Na+ opens
- 2. Na+ enters, increases mV
- 3. V-Na+ closes, V-K+ opens
- 4. V-Ca++ opens, V-K+ still open
- 5. V-Ca++, V-K+ closes

Systole: Contraction while blood pumped into circulation

Diastole: Relaxation as chambers fill with blood

Cardiac muscle cannot reach tetany.





Ca++ enters cytoplasm through V-Ca++ and RYR Exits through NCX and ATP-Ca++

NE increases stroke volume \rightarrow binds to NER \rightarrow RYR stays open longer = more Ca++ in cell = more crosbridge cycling

Atrial cells receive signal for AP from gap junctions & SA node **Ventricular muscles** receive AP signal from purkinje fibers

Main steps of sketletal muscle contraction:

- 1. Excitation
- 2. Ca++ release
- 3. Contraction/crossbridge
- 4. Relaxation

Main steps of cardiac muscle contraction:

- 1. Excitation
- 2. Ca++ release and crossbridges
- 3. V-Ca++ Ca in
- 4. NCX and CaATPase remove Ca++
- 5. Relaxation

$MAP = CO \times TPR$ $CO = HR \times SV$

HR = SA node, SV = EDV

Aka driven by radius of vessel

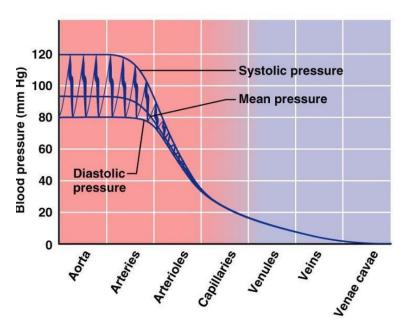
 \rightarrow arteries have highest resistance (largest drop)

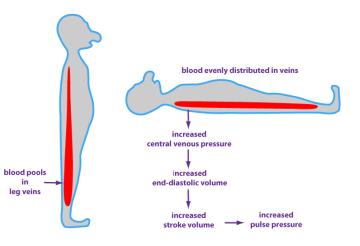
 \rightarrow arterial pressure increases as resistance increases aka vasoconstriction

Use mass balance for this

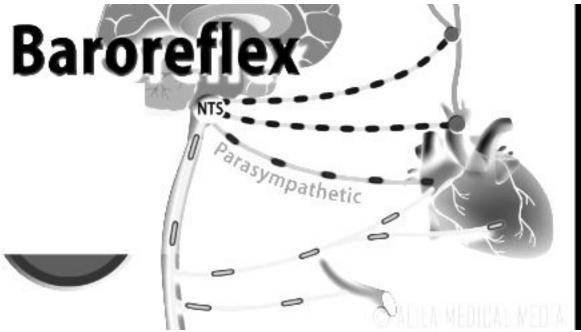
Blood pressure regulated by neurons

- 1. Stimulus: BP
- 2. Receptor: Baroreceptor
- 3. Input: baroreceptor sensory neuron
- 4. Integrator: Cardiovascular center of branstem
- 5. Output: Motor neuron
- 6. Target: cardiac cells
- 7. Response: Change SV/force of contraction





Baroreceptor: functions as a mechanically gated Na+ channel \rightarrow more stretch = more AP in aorta



Decrease in blood pressure:

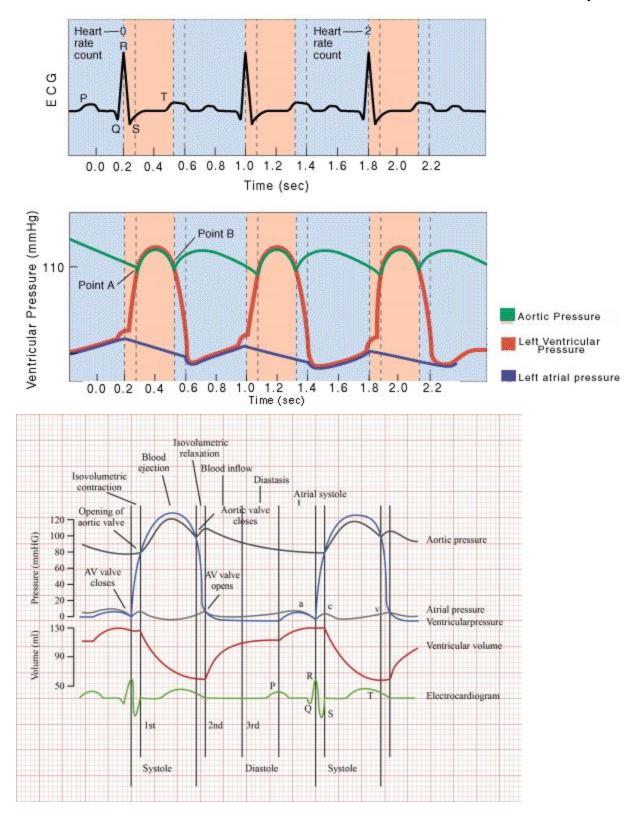
SA node: less ACh \rightarrow less slowing of HR, more NE \rightarrow increased HR **Cardiac muscle**: NE increases, NER binding increases, more Ca2+, stronger contraction, increase SV

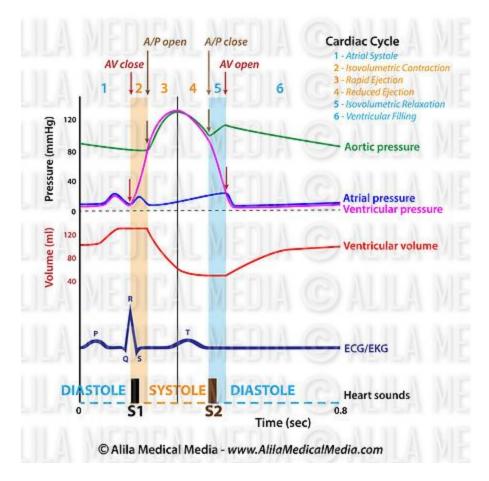
Arterioles: NE increases \rightarrow more Ca++, vasoconstriction \rightarrow more TPR

Parasympathetic: ACh \rightarrow only affects SA node **Sympathetic**: NE

Decreased stretch \rightarrow decrease in graded potentials \rightarrow less baroreceptor AP \rightarrow increased sympathetic AP

Sinha, Irika BIOL 220 Exam 4 Study Guide





Lazy Pandas

Pandas are in the class carnivora but diet = mostly bamboo \rightarrow low nutritional quality, specialized digestive system needed

Hypothesis: statement about world views **Prediction**: what will be measured/quantified

FALSE morphological changes in digestive tract (long herbivore, short carnivore)

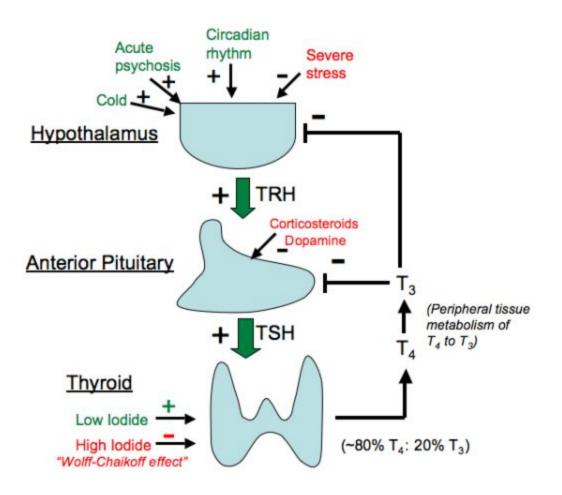
Pandas have shortish \rightarrow digestive tract not specialized to extract maximum energy from plant material therefore energy in is not greater in rate

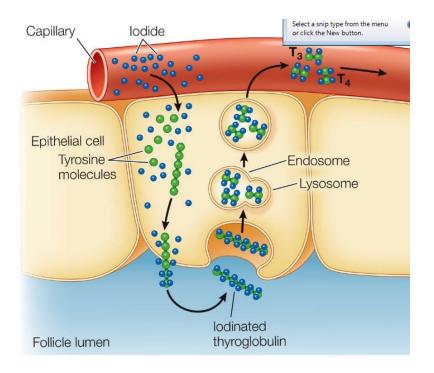
Pandas use less energy relative to other bears/ mammals \rightarrow lower daily energy expenditure compared to other mammals of same size

How?

Changes in behavior \rightarrow less activity in wild

Better insultation to reduce lost body heat = less surface temperature Smaller energy hog organs \rightarrow brain, liver, kidney.... Heart is bigger than others of similar size Change in hormone levels (thyroid) = metabolic \rightarrow low t3/t4





Goiter: low T4, above normal + goiter \rightarrow lots of thyroglobulin in lumen

