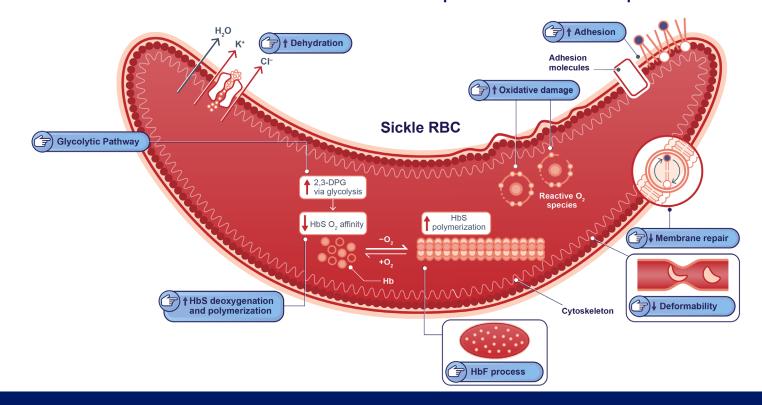


Red Blood Cell Health

Elements of SCD Pathophysiology

Overview: Elements of SCD Pathophysiology

- Sickled RBCs have higher levels of 2,3-DPG and lower levels of ATP than healthy RBCs^{1,2}
- Elevated levels of HbF have been associated with improved RBC lifespan in SCD³

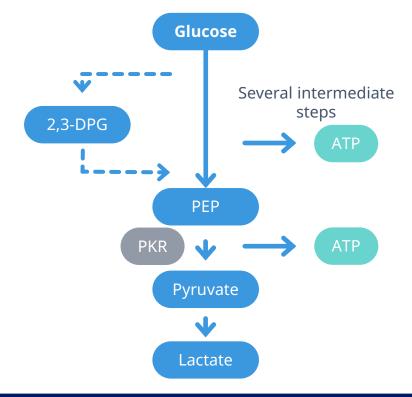


Impaired RBC health in SCD leads to hemolysis, vaso-occlusion, and reduced RBC lifespan^{1,4}

HbS polymerization results in sickled RBCs¹

- SCD is a genetic disorder characterized by abnormal hemoglobin (HbS) that polymerizes under conditions of low oxygen tension¹
- 2,3-diphosphoglycerate (2,3-DPG) is a byproduct of glycolysis and is a negative regulator of hemoglobinoxygen affinity^{2,3}
- HbS exists in a relaxed state (R-state) or a tense state (T-state), which impacts oxygen affinity^{1,4}
 - In the R-state, Hb has a high affinity for oxygen and is associated with low levels of 2,3-DPG⁴
 - In the T-state, Hb has a low affinity for oxygen and is associated with higher levels of 2,3-DPG⁴

Glycolysis pathway²



The glycolytic pathway is a source of ATP generation in RBCs for membrane repair, hydration, and deformability^{2,5}

ATP, adenosine triphosphate; Hb, hemoglobin; HbS, sickle hemoglobin; PEP, phosphoenolpyruvate; PKR, pyruvate kinase; RBC, red blood cell; SCD, sickle cell disease.

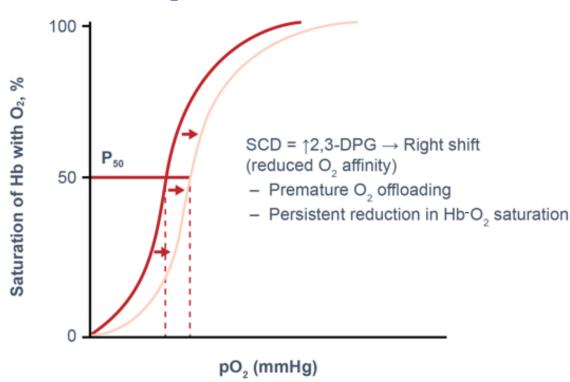
1. Kato GJ, et al. *Nat Rev Dis Primers*. 2018;4:18010. 2. van Wijk R, van Solinge WW. *Blood*. 2005;106(13):4034-4042. 3. Eaton WA, Bunn HF. *Blood*. 2017;129(20):2719-2726. 4. Oder E, et al. *Br J Haematol*. 2016;175(1):24-30. 5. Huisjes R, et al. *Front Physiol*. 2018;9:656.





2,3-DPG is the major allosteric effector of Hb-O₂ affinity¹

O₂-Hb dissociation curve²



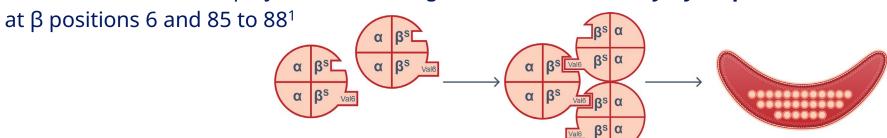
- 2,3-DPG regulates the ability of RBCs to carry and deliver O_2^2
- Under conditions of low O_2 tension, 2,3-DPG binds to the cleft between β subunits, stabilizing HbS in the tense state¹
- Increased 2,3-DPG
 - Stabilizes HbS polymers by decreasing solubility¹
 - Decreases intracellular pH, which reduces HbS solubility and O₂ affinity (the Bohr effect)^{1,2}
- Ultimately, increased 2,3-DPG potentiates HbS polymerization¹

Fetal Hb (HbF) does not interact with 2,3-DPG, resulting in a high affinity for O_2^3

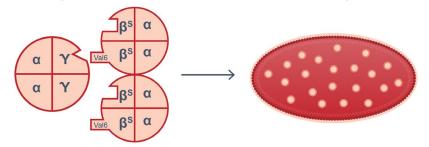


HbF prevents polymerization of HbS

• In SCD, HbS tetramers polymerize into long strands, **stabilized by hydrophobic interactions** between strands



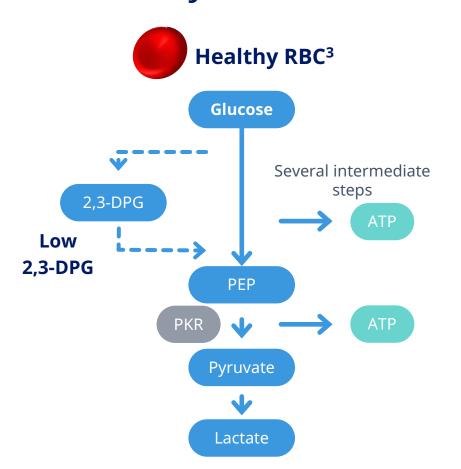
- HbF has a γ-globin with Gln at position 87 (vs Thr), which weakens the hydrophobic interaction
 between strands, blocking polymerization and preventing the formation of long polymer strands found in SCD¹
- Additionally, HbF has a decreased affinity for 2,3-DPG, which further prevents HbS polymerization^{2,3}

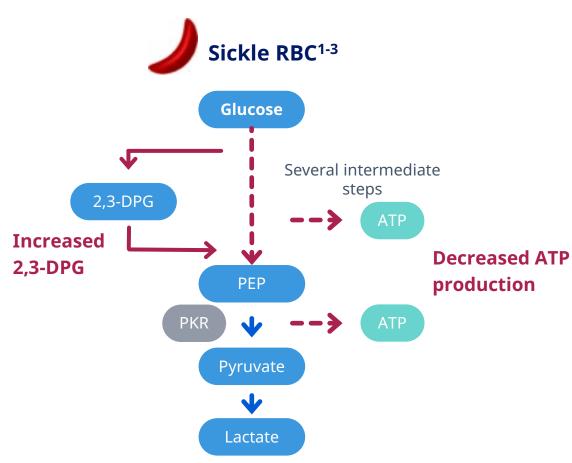


Even small increases in HbF have shown to improve RBC health and clinical outcomes⁴



Sickle RBCs have higher levels of 2,3-DPG and lower levels of ATP than healthy RBCs^{1,2}

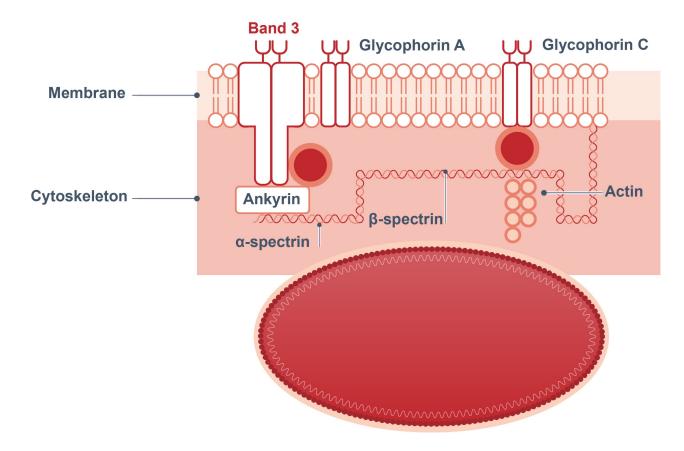






Sickle RBCs have impaired deformability¹

Healthy RBCs need to deform to traverse vessels under shear stress without rupture.1

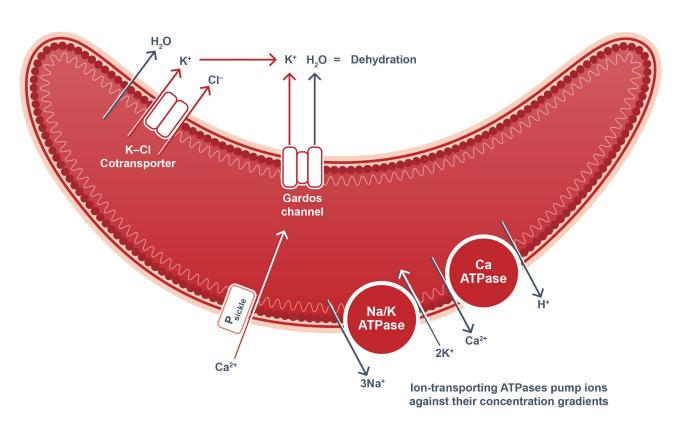


- Sickle RBCs have impaired deformability and become stiff due to polymerization¹
- RBC deformability is regulated and maintained by interactions between membrane and cytoskeletal proteins¹
- Low levels of ATP in sickle RBCs, along with HbS polymerization, result in impaired deformability^{1,2}



Impaired ion hemostasis causes sickle RBC dehydration¹⁻³

Healthy RBCs maintain significant ion gradients via the action of ATP-dependent cation pumps.²



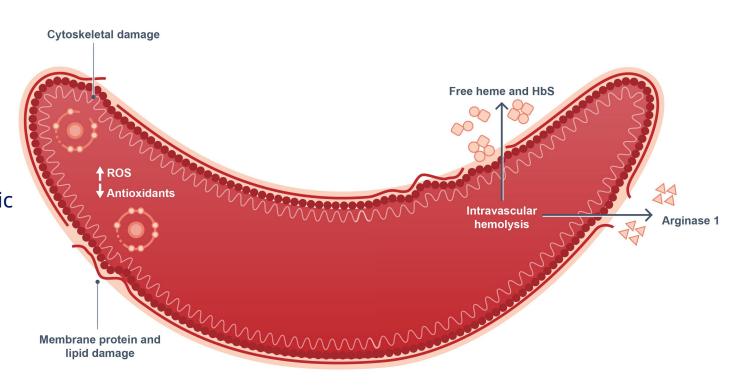
- Sickle RBCs leak cations as a result of sickling¹
- Divalent cation permeability is particularly important as increased intracellular Ca²⁺ and decreased Mg²⁺ stimulate K⁺ efflux^{2,3}
 - As intracellular K⁺ concentration decreases, water flows out of the RBC^{2,3}
- Dehydrated RBCs sickle more rapidly due to the higher concentration of HbS⁴
- Homeostasis can be restored/maintained only by the expenditure of ATP to pump ions against their concentration gradients²



Sickle RBCs have decreased antioxidant capacity¹

Healthy RBCs contain helpful enzymes that serve as antioxidants to scavenge ROS.²

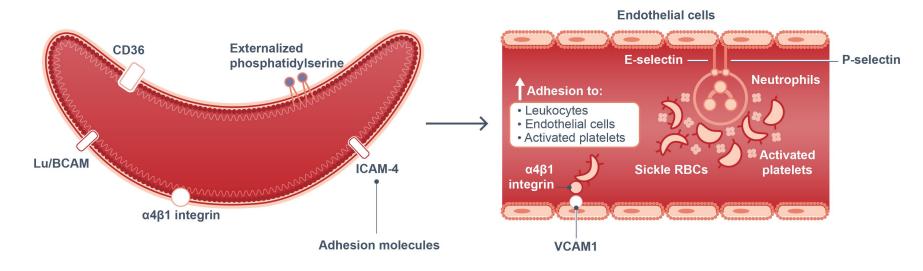
- ROS are harmful to membrane proteins and can cause lipid damage and cytoskeletal defects¹
- Antioxidant enzymes, including glutathione reductase and superoxide dismutase, are ATP-dependent¹
- Lack of ATP results in decreased enzymatic activity that causes increased oxidative damage to the membrane and cytoskeleton, leading to increased intravascular hemolysis^{1,3}





Sickle RBCs exhibit increased adhesion¹

Healthy RBCs express amino phospholipids, such as phosphatidylserine (PS), on the inner leaflet of the membrane and do not aggregate.²



- Sickle RBCs exhibit increased adhesion and overexpress adhesion molecules, such as $\alpha 4\beta 1$ integrin and Lu/BCAM. This may lead to endothelial activation, thereby causing vaso-occlusion^{1,3,4}
- Sickling induces reorientation of PS from the inner to the outer surface of RBCs, making RBCs stickier and leading to premature clearance of sickled RBCs by macrophages and vaso-occlusion⁵
- PS can be reoriented to the inner leaflet by an ATP-dependent translocase^{2,4}

ATP, adenosine triphosphate; CD, cluster of differentiation; ICAM-4, intracellular adhesion molecule 4; Lu/BCAM, Lutheran/basal cell adhesion molecule; RBC, red blood cell; VCAM1, vascular cell adhesion molecule 1.



^{1.} Kato GJ, et al. Nat Rev Dis Primers. 2018;4:18010. 2. Weiss E, et al. Anemia. 2011;2011:379894. 3. McMahon TJ. Front Physiol. 2019;10:1417. 4. Hannemann A, et al. Br J Haematol. 2018;182(4):567-578. 5. Setty BN, et al. Blood. 2002;99(5):1564-1571.

Low levels of ATP result in impaired membrane repair¹

Healthy RBCs maintain functional integrity of key cytoskeletal and membrane components, which are dependent on ATP.²

- Translocase (also known as *flippase*) reorients PS to the inner leaflet¹
- Flippase requires ATP to repair membranes in various hemoglobinopathies, including SCD³
- Inhibition of flippase can be triggered by oxidant stress, increased intracellular Ca²⁺, and the activity of protein kinases³

