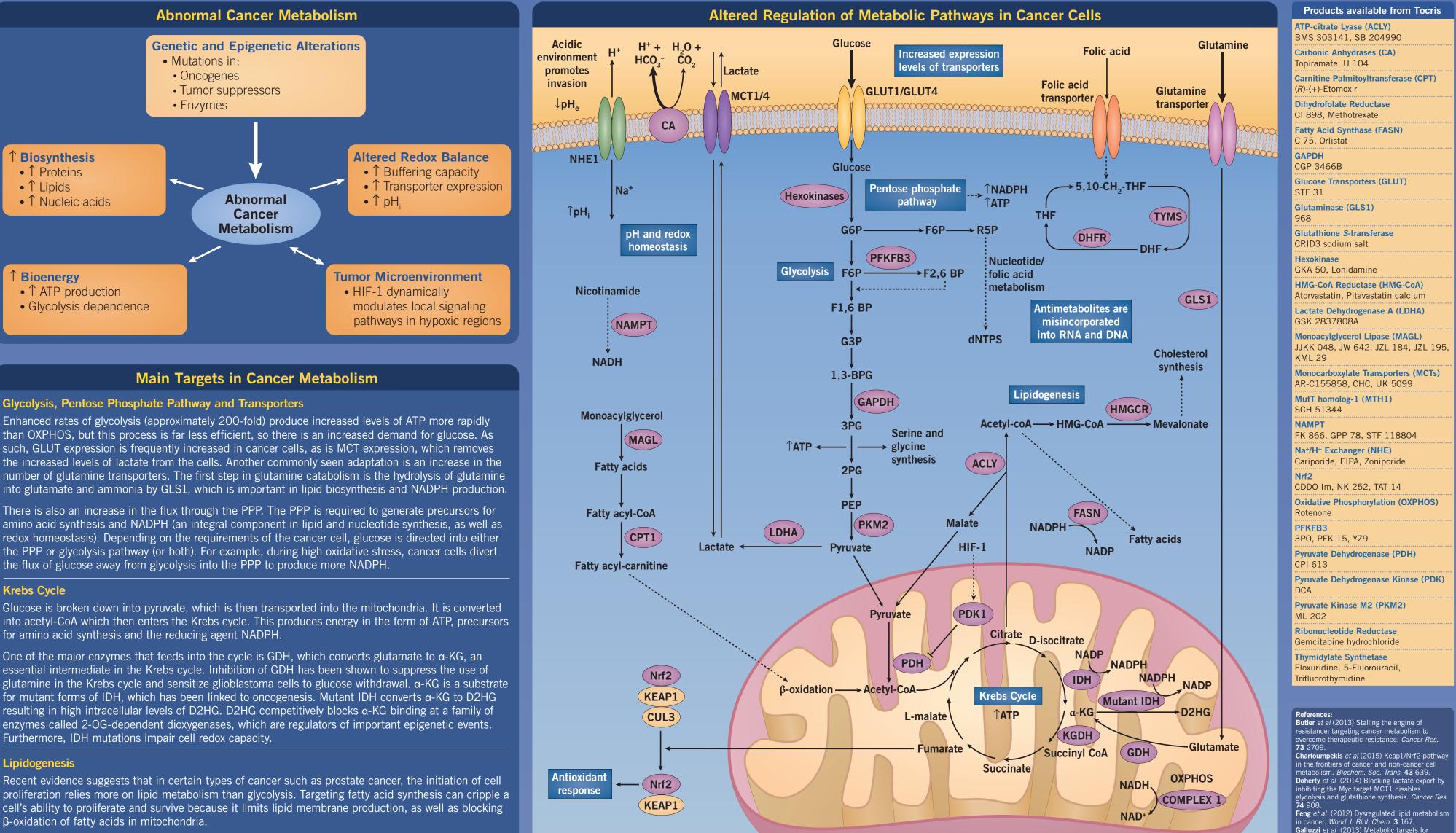
Cancer Metabolism

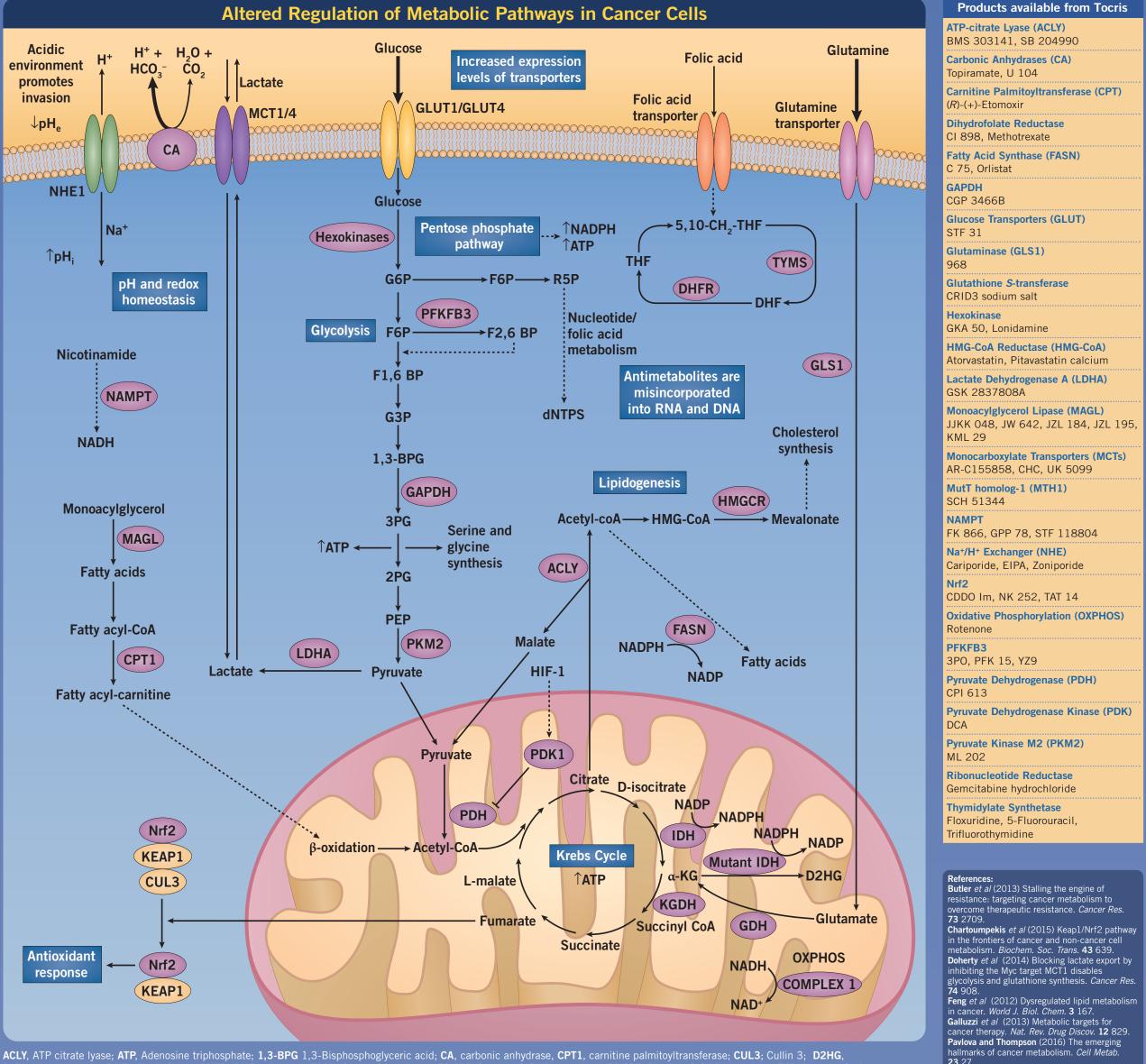
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In 1924 Otto Warburg first discovered that cancer cells generated a large proportion of their ATP by metabolizing glucose via aerobic glycolysis (as opposed to mostly through oxidative phosphorylation (OXPHOS) in normal cells). Initially it was thought that this Warburg effect was a cause of cancer, but it was later established that this shift to glycolytic metabolism was an effect of cancer cell transformation. Genetic changes and epigenetic modifications in cancer cells alter the regulation of cellular metabolic pathways. These distinct metabolic circuits could provide viable cancer therapeutic targets.



pH and Redox Balance

Cancer cells are able to survive in their hostile microenvironments because of increased expression of proton pumps and ion transporters. Aberrant regulation of hydrogen ions leads to a reversal of the pH gradient across tumor cell membranes, resulting in a more basic intracellular pH (pH_i) and a more acidic extracellular pH (pH_a). It is critical to cancer cell survival that the intracellular environment does not become acidified because this could induce apoptosis.





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D-2-hydroxyglutarate; DHF, dihydrofolate; DHFR, DHF reductase; FASN, fatty acid synthase; F1,6BP, fructose-1,6-bisphosphate; F2,6BP, fructose-2,6-bisphosphate; F6P, fructose 6-phosphate; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; GDH, glutamate dehydrogenase; GLUT, glucose transporter; GLS1, glutaminase; G3P, Glyceraldehyde-3-phosphate; G6P, glucose-6-phosphate; HIF-1, Hypoxia-inducible factor 1; HMGCR, HMG-CoA reductase; IDH, isocitrate dehydrogenase; α-KG, α-ketoglutarate; KGDH, α-ketoglutarate dehydrogenase; LDHA, lactate dehydrogenase A; MAGL, monoacylglycerol lipase; MCT, monocarboxylate transporter; NAD+/NAD+; Nicotinamide adenine dinucleotide (oxidised/reduced forms respectively); NADPH, Nicotinamide adenine dinucleotide phosphate; **NAMPT**, nicotinamide phosphoribosyltransferase; **NRF2**, Nuclear factor (erythroid-derived 2)-like 2; **OXPHOS**, oxidative phosphorylation; **PDH**, pyruvate dehydrogenase; **PDK**, pyruvate dehydrogenase kinase; **PEP**, phosphoenolpyruvate; **PFKFB3**, 6-phosphofructo-2-kinase/fructose- 2,6-bisphosphatase; **2PG**, 2-phosphoglycerate; **3PG**, 3-phosphoglycerate; **PKM2**, pyruvate kinase M2 isoform; **PPP**, Pentose Phosphate Pathway; **ROS**, reactive oxygen species; **R5P**, ribose-5-phosphate; **5,10-CH**₂-**THF**, 5,10-methylene tetrahydrofolate; **THF**, tetrahydrofolate; **TYMS**, thymidylate synthase

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