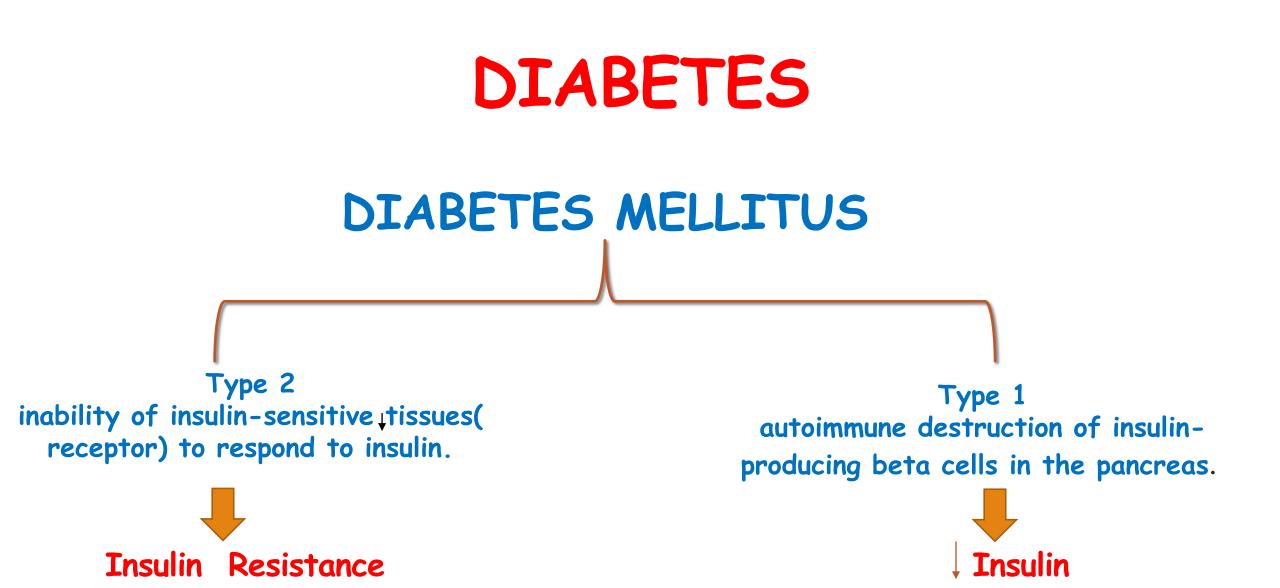


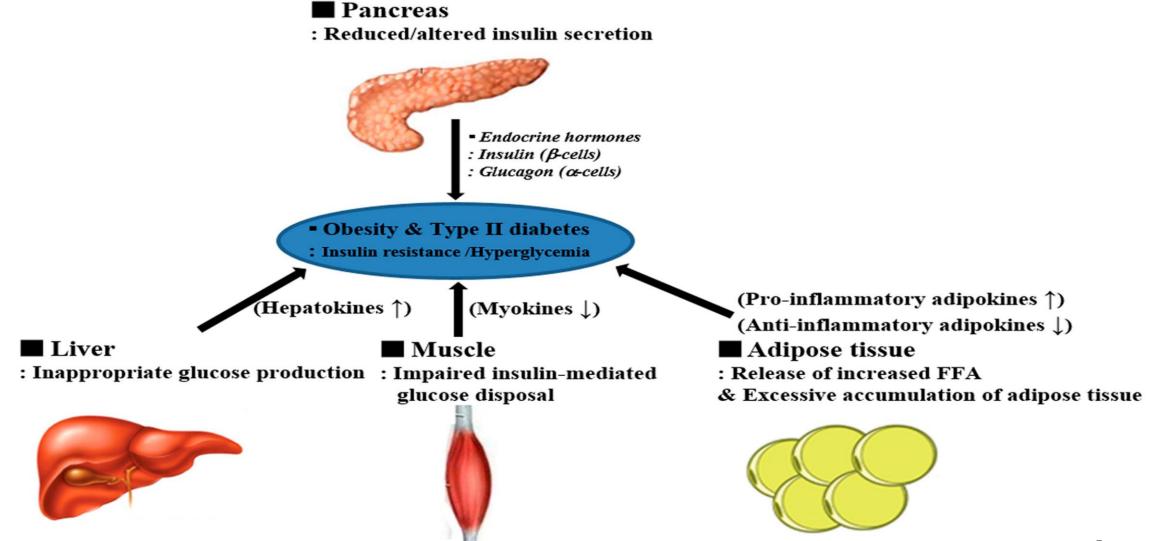


Aquaporin's in insulin resistance and diabetes: More than channels!

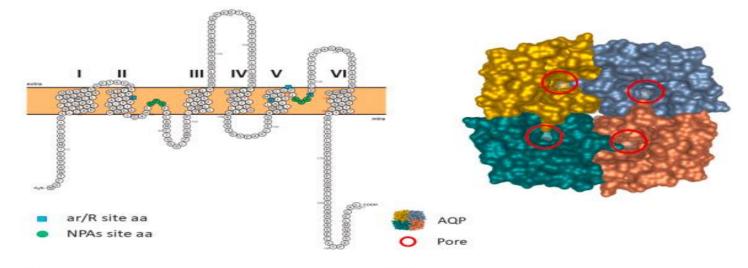
Nepton Soltani



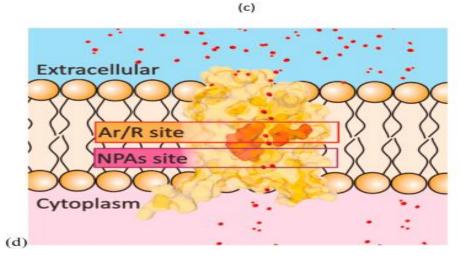
Insulin resistance



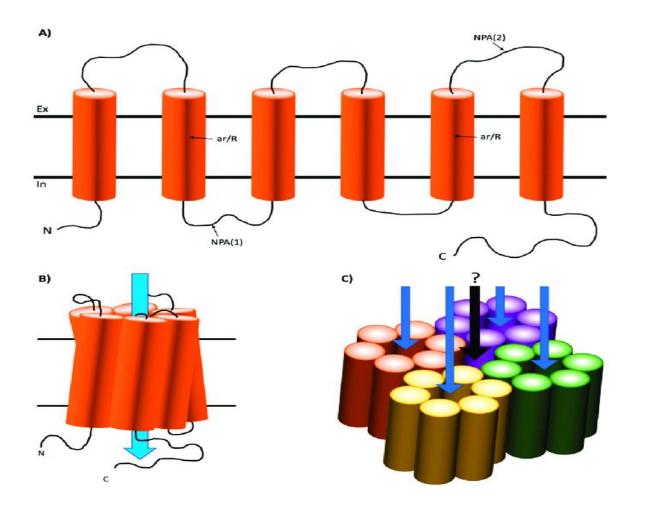
Aquaporin's Structure



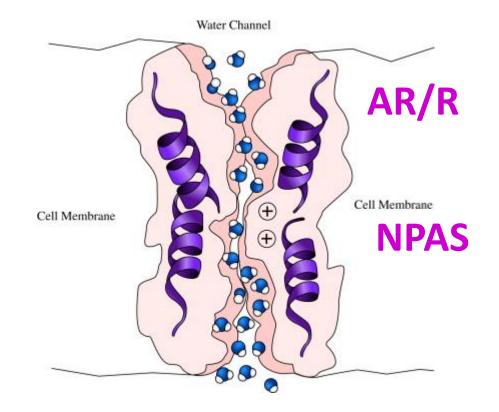
(b)



Aquaporin's Structure



Aquaporin's Structure



Different kinds of aquaporin's

So far, 13 AQPs have been identified in human.
AQP3, 7, 9, and 10 are subcategorized as aquaglyceroporins which permeabilize glycerol as well as water.
Many investigators have demonstrated that AQPs play a

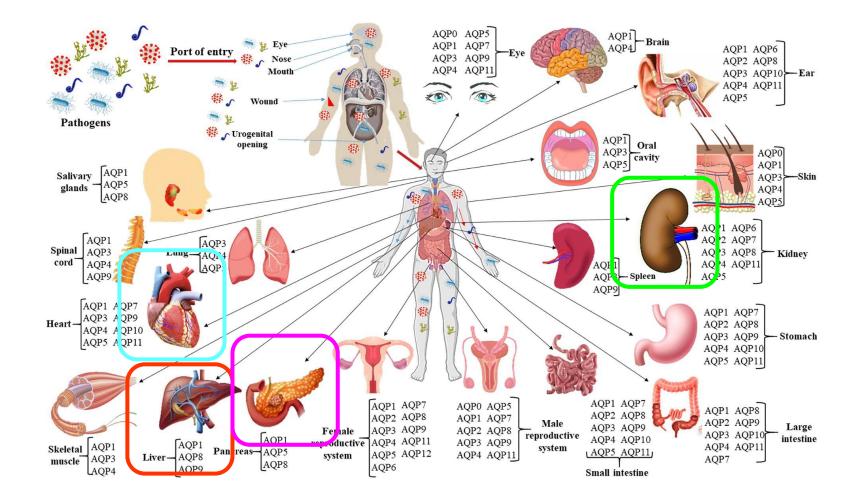
AOP0, AOP1, AOP2,

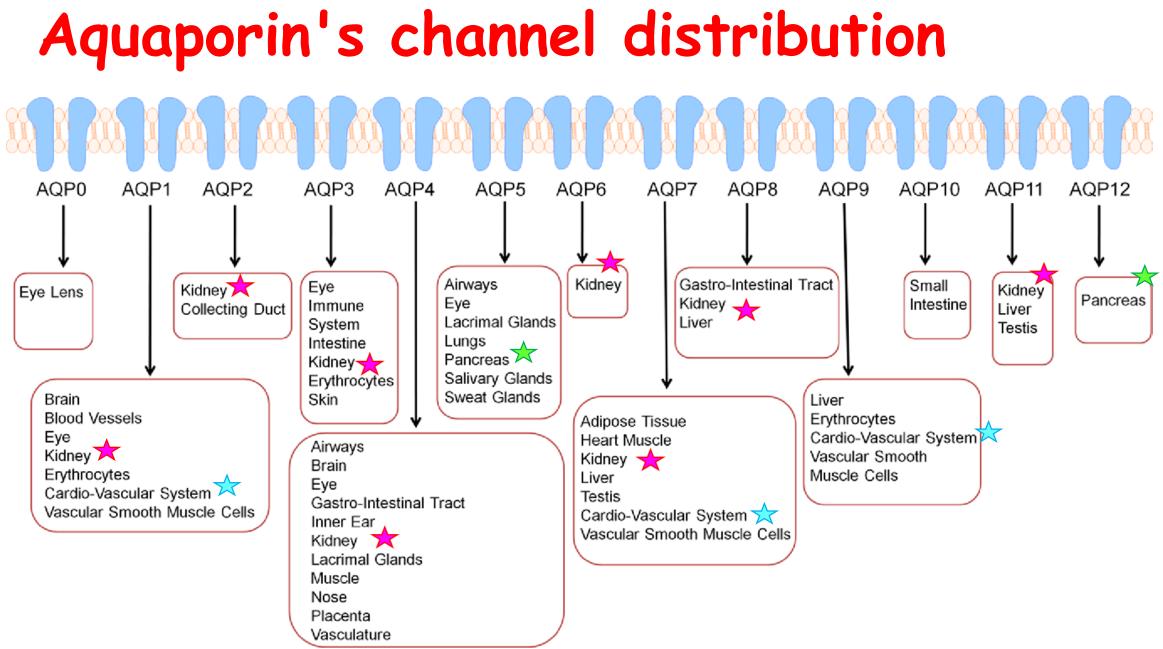
- Many investigators have demonstrated that AQPs play a crucial role in maintaining water homeostasis, but the physiological significance of some AQPs as a glycerol channel is not fully understood.
- Adipose tissue is a major source of glycerol and glycerol is one of substrates for gluconeogenesis.

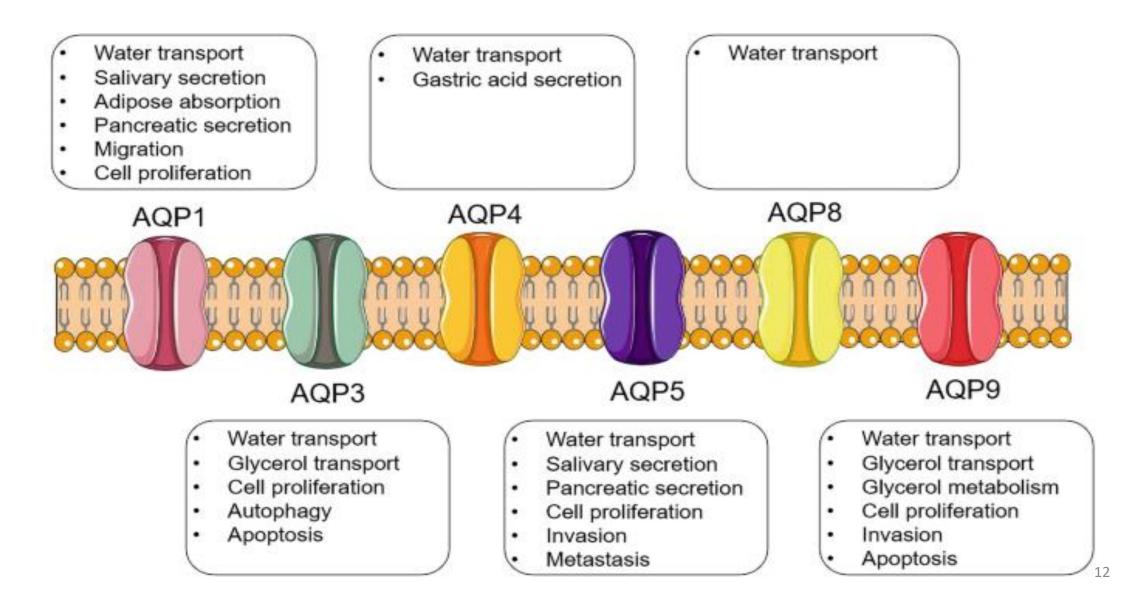
Intracel	llul	lar
muace	nu	lai

Tetrameric Aquaporin

Aquaporin's channel distribution

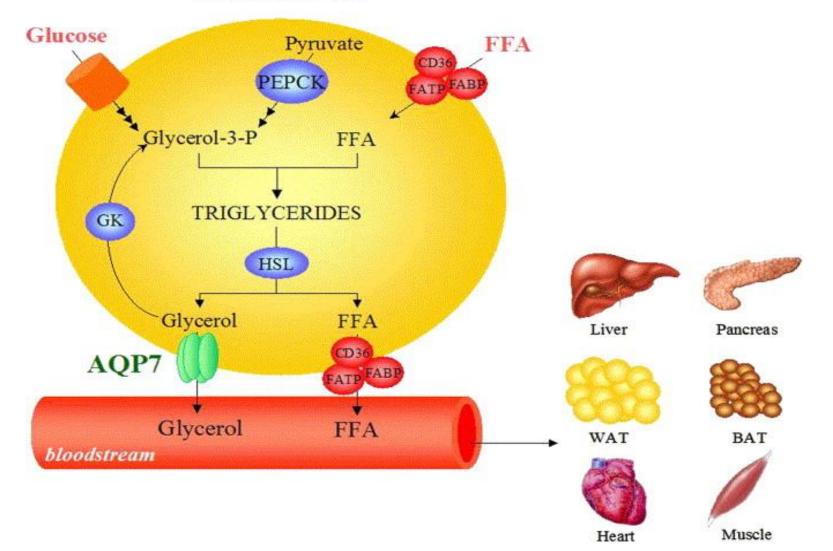


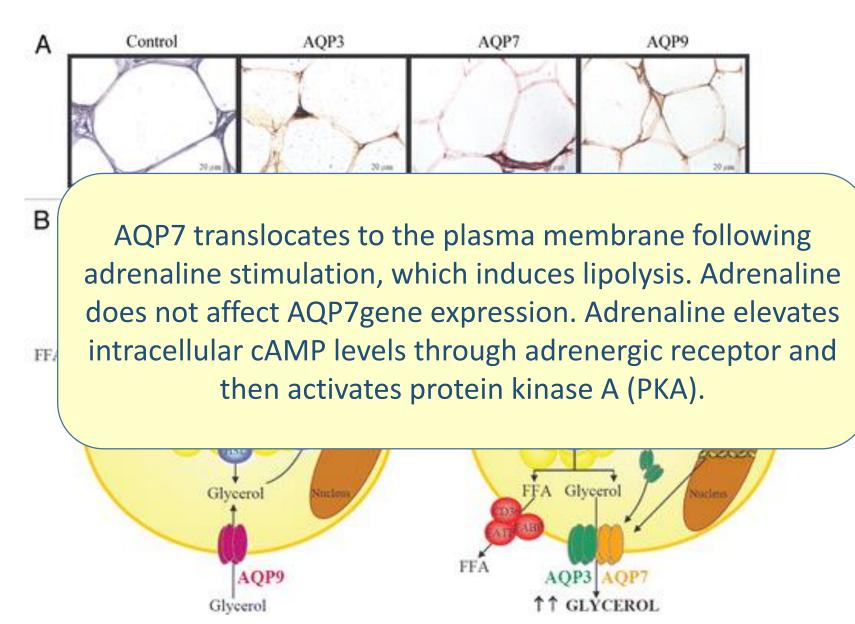




Aquaporins	Metabolic Function	Fertility Function	
AQP3	Adipocyte glycerol efflux Upregulated by insulin Upregulated by leptin	Sperm osmoadaptation	
AQP7	Adipocyte glycerol efflux Upregulated by insulin Downregulated by leptin Increases insulin sensitivity	Sperm motility Spermiogenesis	
	Adipocyte glycerol influx	Spermatogenesis	
AQP9	Hepatocyte glycerol influx Upregulated by insulin Downregulated by leptin	Regulation of epididymal osmolarity Lactate secretion for germ cells Upregulated by sex hormones	

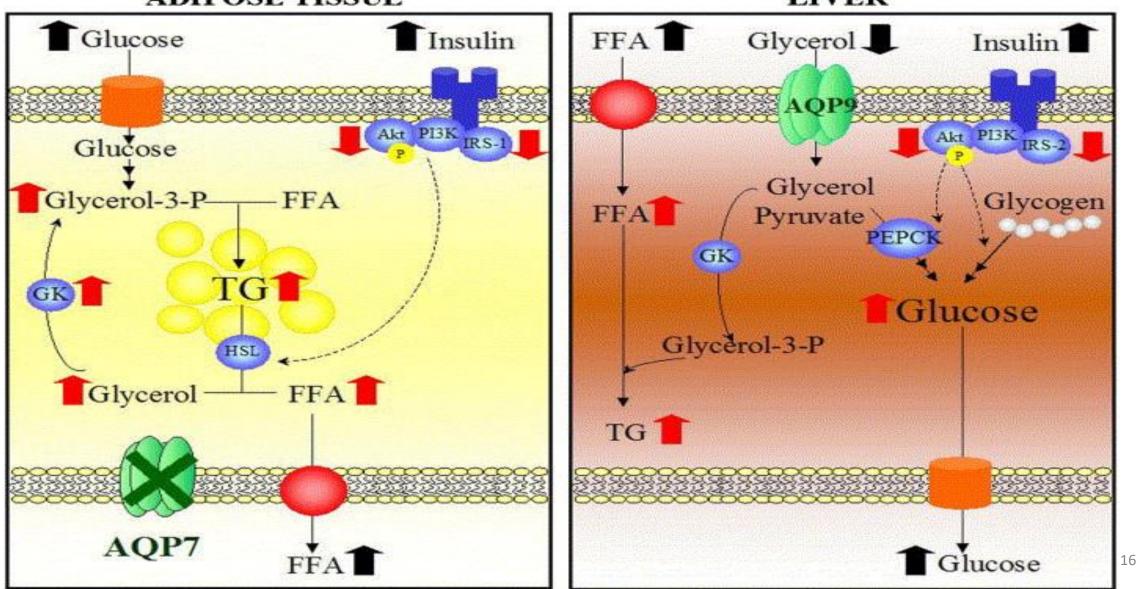
ADIPOCYTE

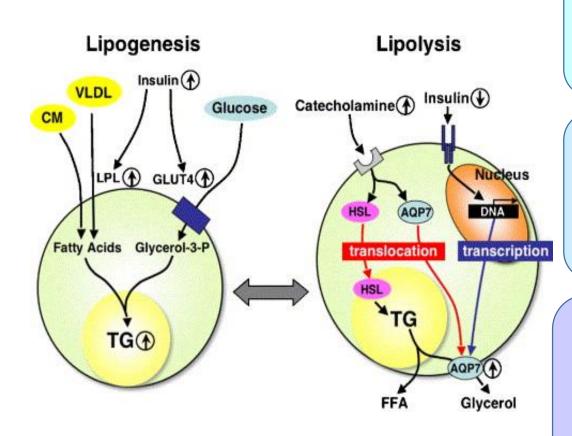




ADIPOSE TISSUE

LIVER





Moreover, AQP7 mRNA levels are elevated by the decrease of insulin signaling cascade. Thus, long-term regulation of AQP7 is under the control of insulin while short-term regulation is under catecholamines.

AQP7 is highly expressed in white adipose tissue (WAT), brown adipose tissue (BAT), and testis. Furthermore, a weak expression of AQP7 is also observed in the heart, skeletal muscles, and kidneys.

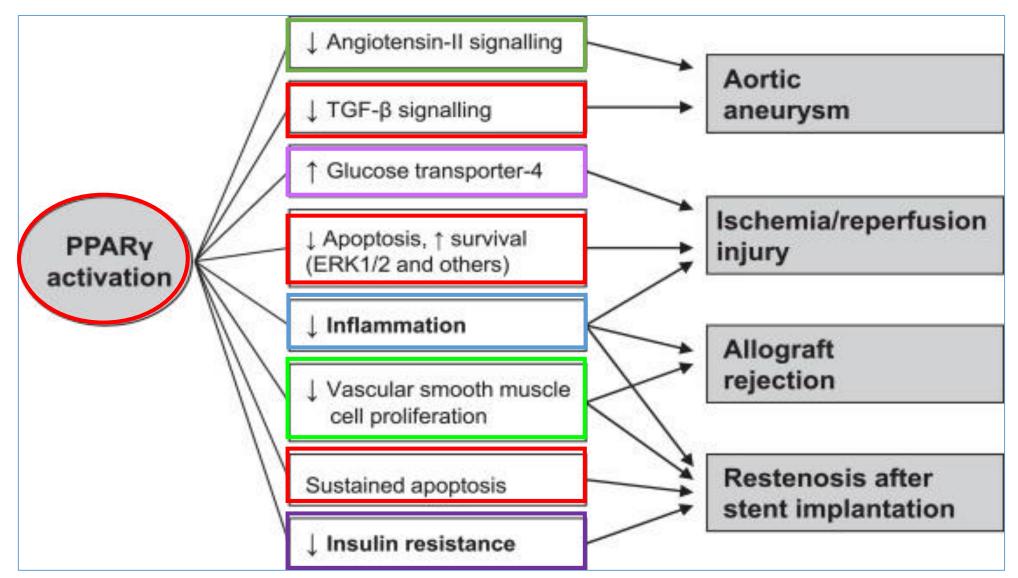
Peroxisome proliferator activated receptor γ (PPAR γ) is a master regulator of adipocytes differentiation and regulates several adipose-specific genes at the transcriptional level. PPAR γ forms a heterodimer with **retinoic acid X receptor** α (RXR α), and binds to peroxisome proliferator response element (PPRE) site. The PPRE site is identified in the promoter region of AQP7 gene

PPARy actions

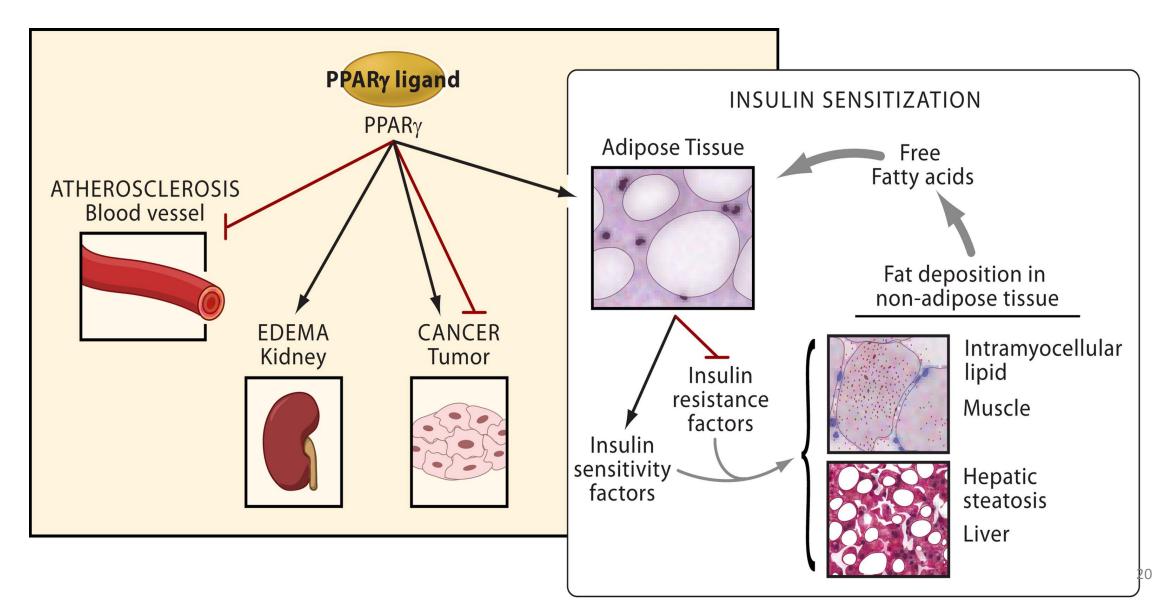
	Diet,	Lipoxygenases, Cyclooxigenases	
	OFurthermor	e, administration of thiazolidinediones	
	(TZDs), wh	ch are insulin-sensitizing agents and	
	exogenous f	PARy ligands, increase AQP7 mRNA level	S
	in adipocyte	s and in adipose tissues of mice. The	
	precise med	hanism of TZD in ameliorating insulin	
	resistance i	s not fully understood.	
	OTo clarify w	whether the TZD-induced adipose AQP7 i	is
	related to t	he amelioration of insulin resistance,	
	requires fu	ther studies in the future.	
05			

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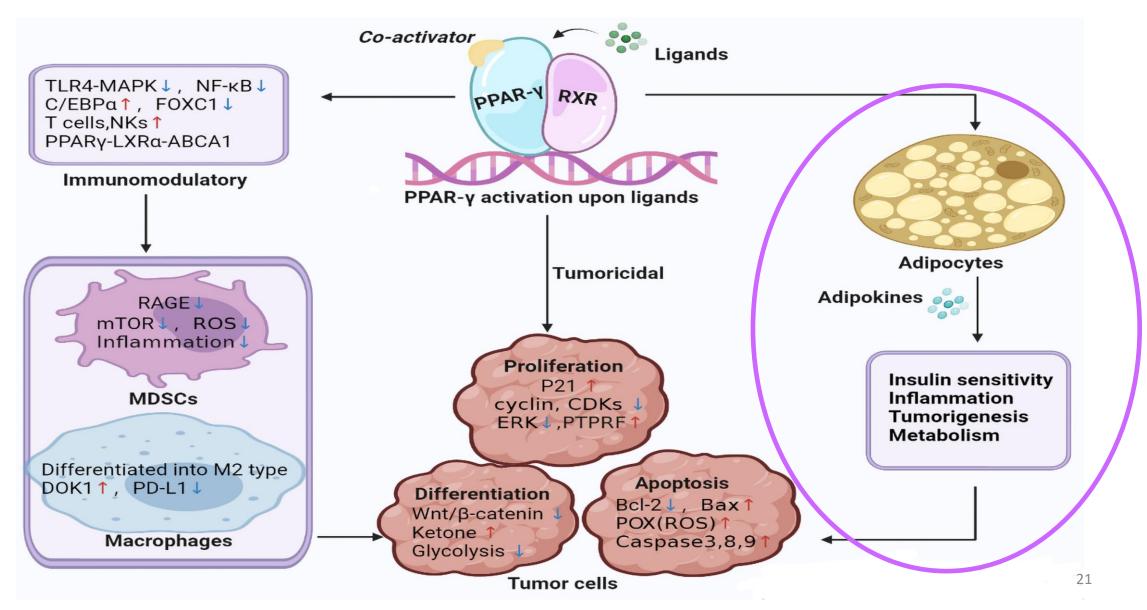
PPARy actions



PPARy actions







Effect of insulin on aquaporin 7

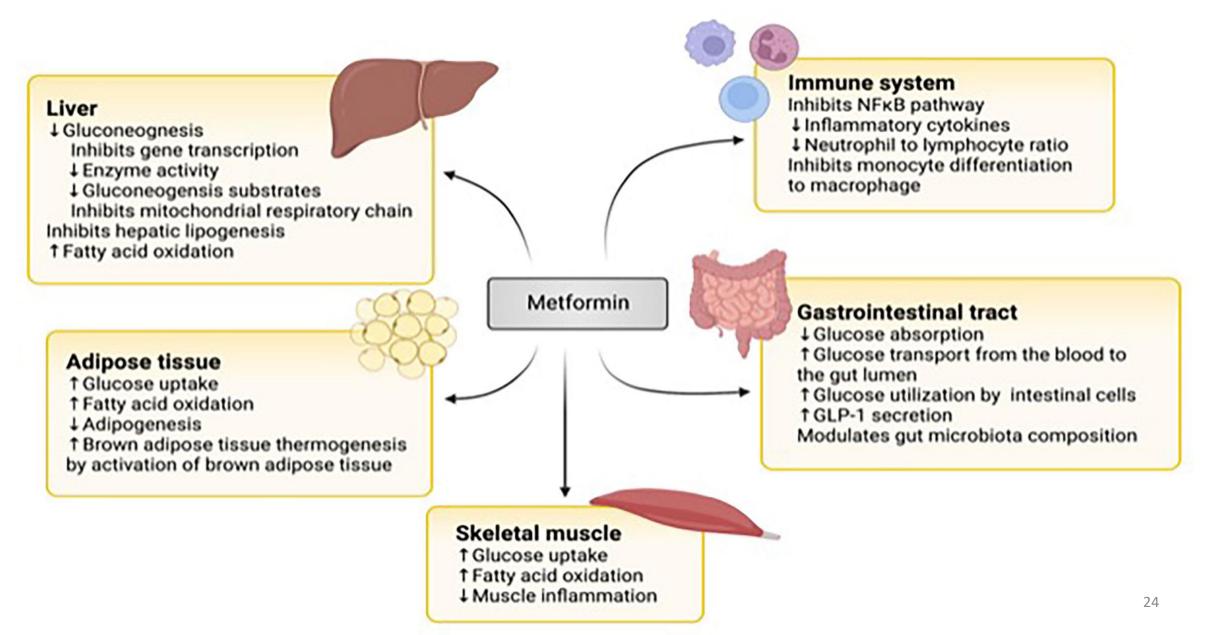
• The insulin negative response element (IRE) is identified in the promoter region of AQP7 gene.

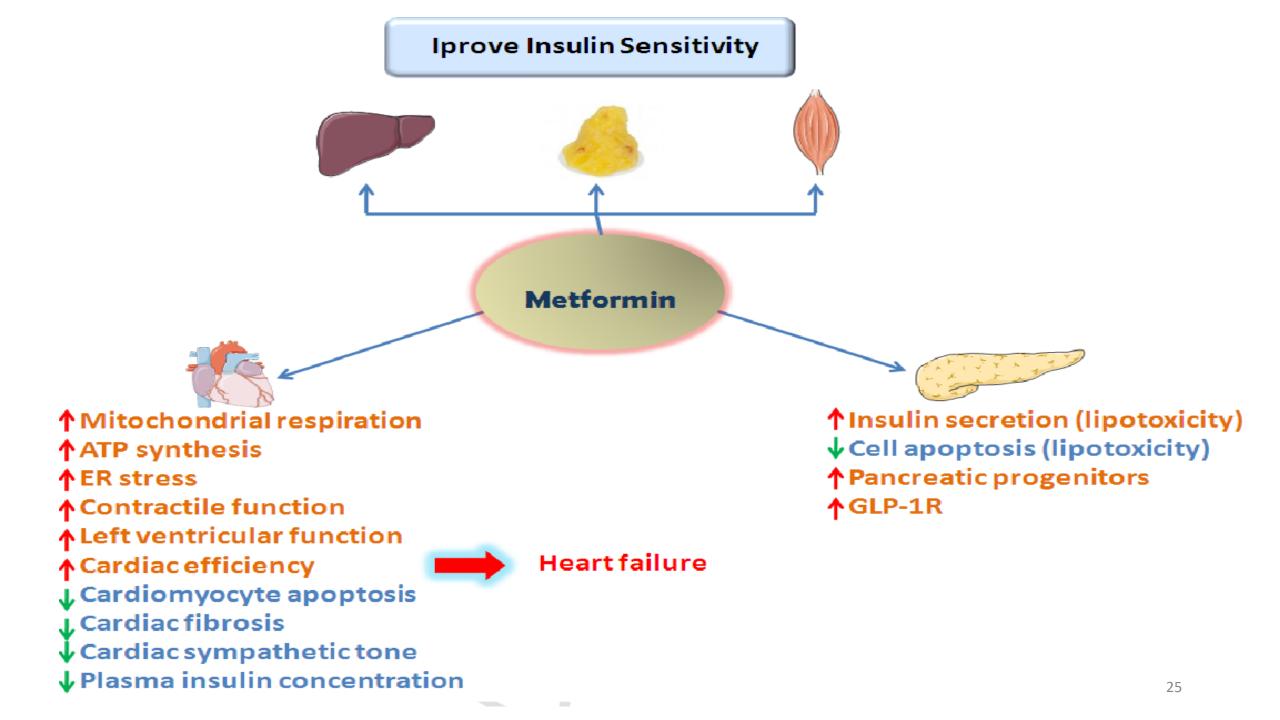
- This result indicates that AQP7 mRNA expression is closely regulated by insulin at the transcriptional level. Furthermore, glucose-6-phosphatase (G6Pase) and phosphoenolpyruvate carboxykinase (PEPCK), which are key enzymes of gluconeogenesis, also contain IRE in their promoter regions.
- Insulin also suppresses the mRNA levels of G6Pase and PEPCK. Taken together, plasma glycerol levels are partly determined by insulin through adipose AQP7, and suggests that adipose AQP7 may be associated with glucose metabolism.

Metformin

•Metformin is a widely used antidiabetic drug for treating type 2 diabetes mellitus (T2DM) that enhances insulin regulation of glucose, promotes weight loss, and reduces appetite. It reduces insulin resistance and decreases blood glucose concentration by inhibiting gluconeogenesis and suppressing hepatic glucose production.

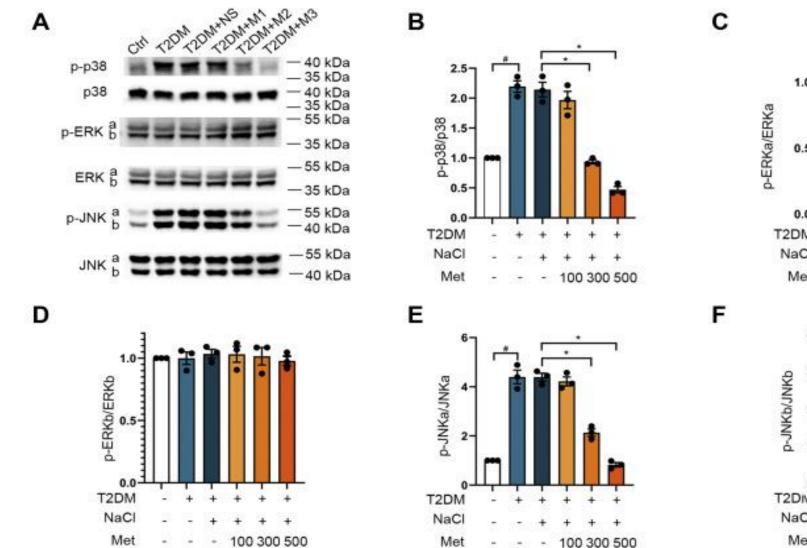
Metformin actions



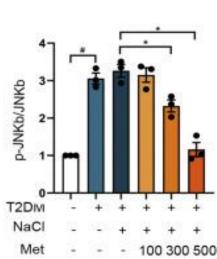


OPancreatic β-cell injury and insulin resistance appear to be partially triggered by inflammatory, oxidative, and endoplasmic reticulum stress-induced pathways, including the mitogen-activated protein kinases (MAPK) signaling cascade. However, whether MAPK modulations affect AQP7 expression in pancreatic tissue of T2DM treated with metformin remains poorly understood.

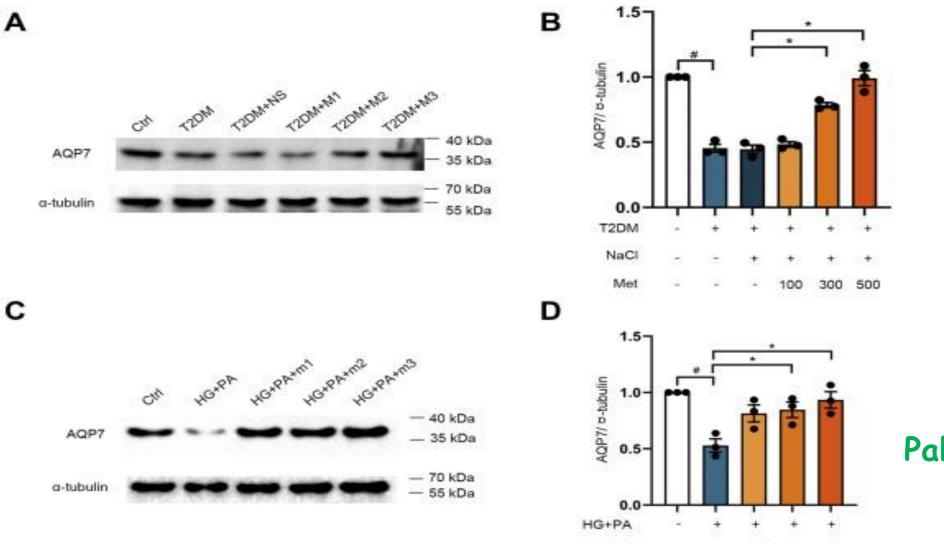
Effect of Metformin on JNK pathway



1.0 0.5 0.0 T2DM - + + + + + NaCl - - + + + + + Met - - - 100 300 500



Aquaporin's channel gene expression by Metformin



C

Palmitate

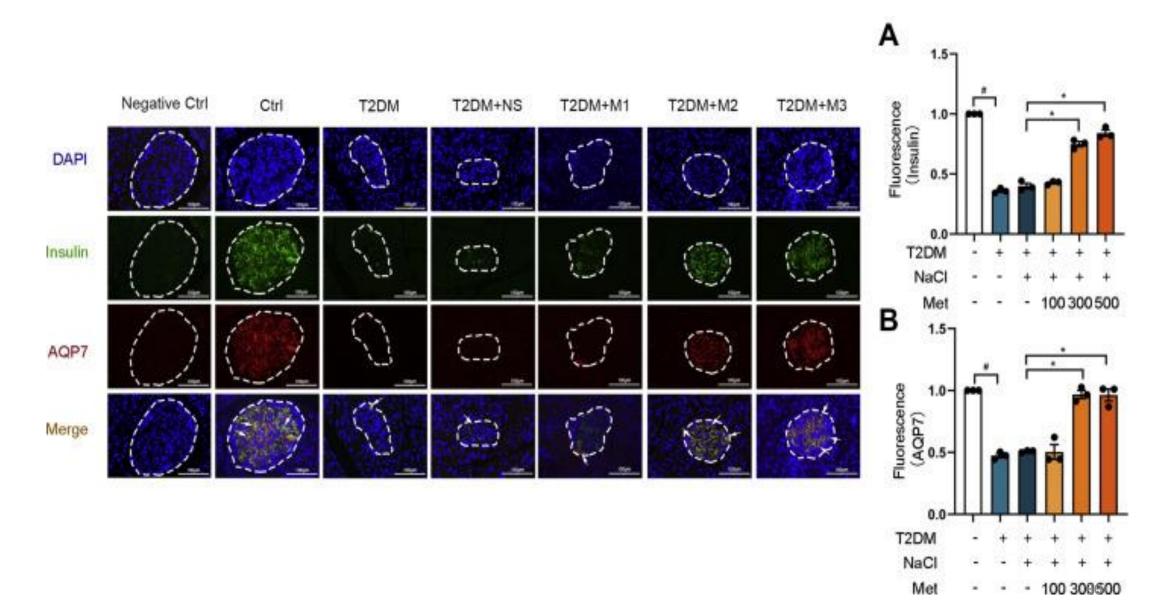
0.5

1

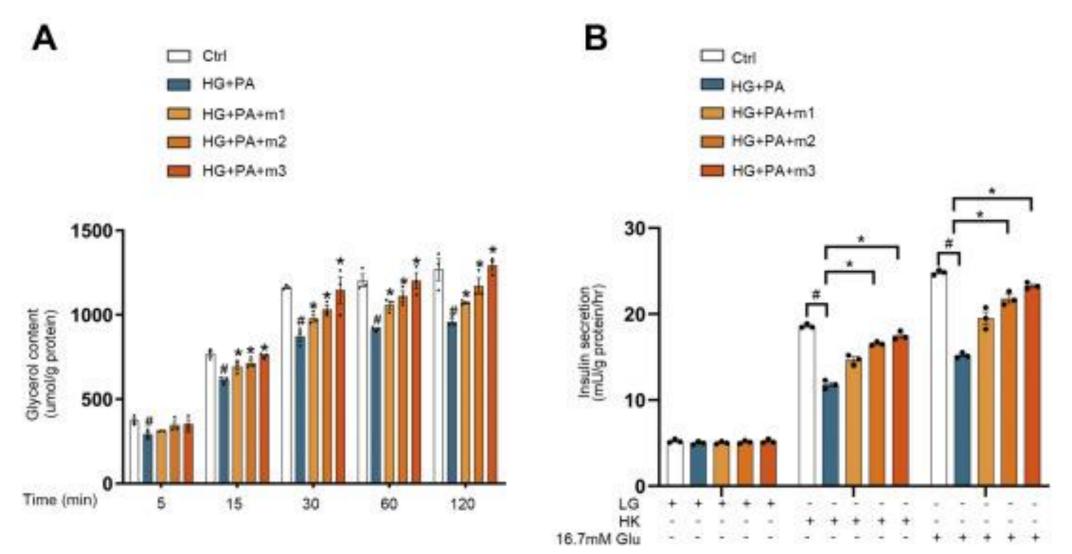
Met

2

Aquaporin's channel gene expression



Aquaporin's channel and insulin secretion





كداميك درحال تقلب است؟

AQP3, AQP8 (in PM) regulation of H₂O₂ cellular uptake

amplification/diminishment of downstream redox signaling cascades (role in cell migration, immune responses, wound healing, proliferation)

 H_2O_2

changes of AQPs expression profiles influence on cell susceptibility to external H₂O₂ signaling?

Ν

i to tal e

P mtAQP8 knockdown oxidative stress LO XO due to ROS accumulation mtAQP8 (in IMM) H₂O₂ release H_2O_2 \blacksquare O_2^- М regulation from ETC

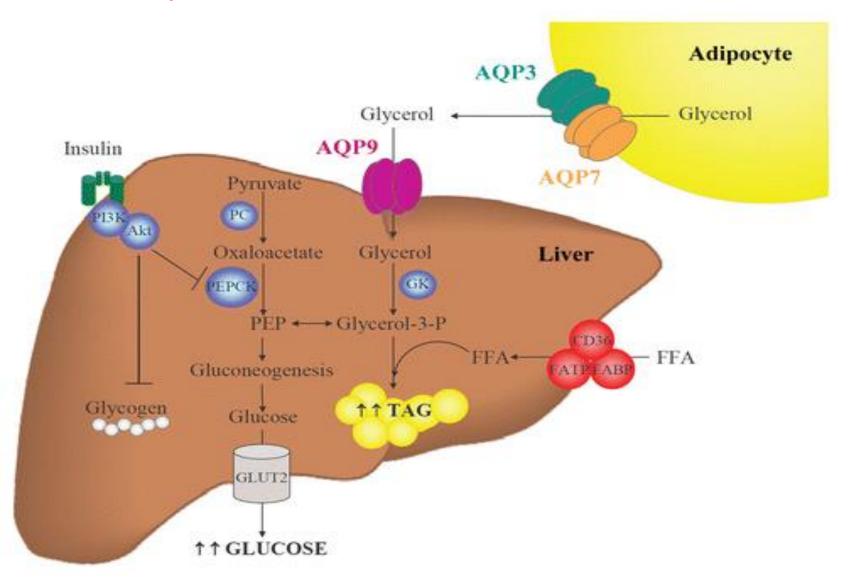
 O_2

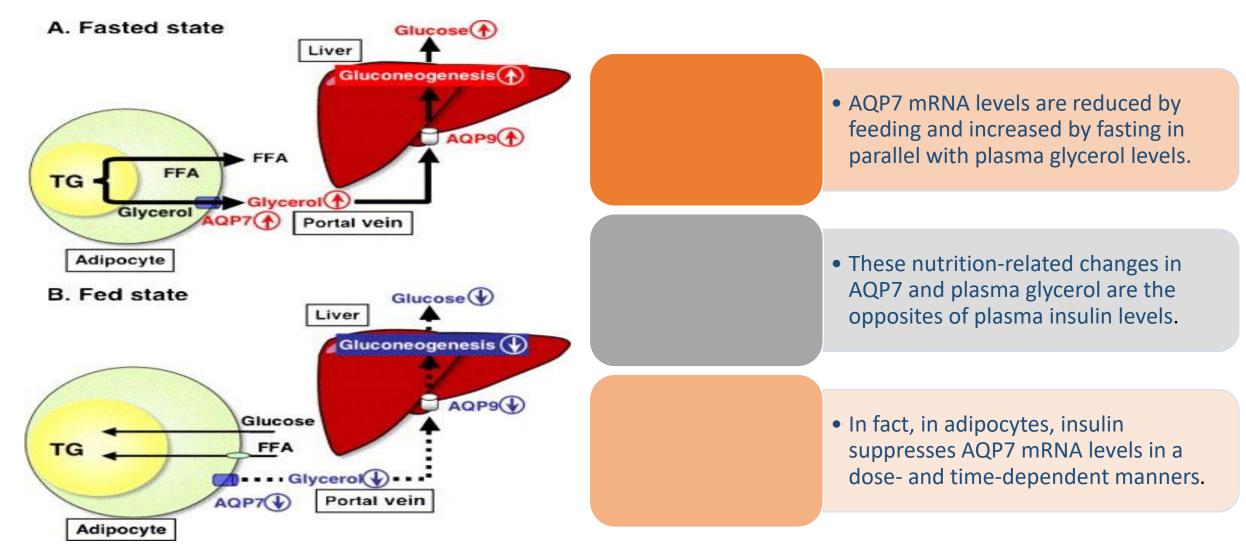
NADP

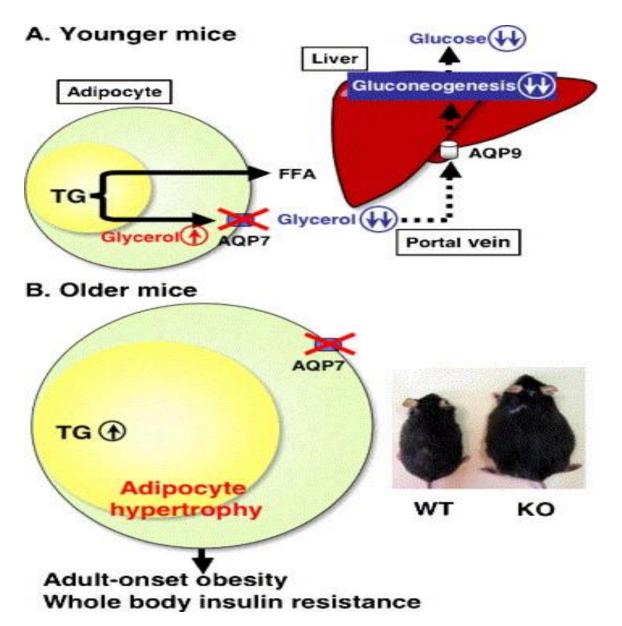
NOX

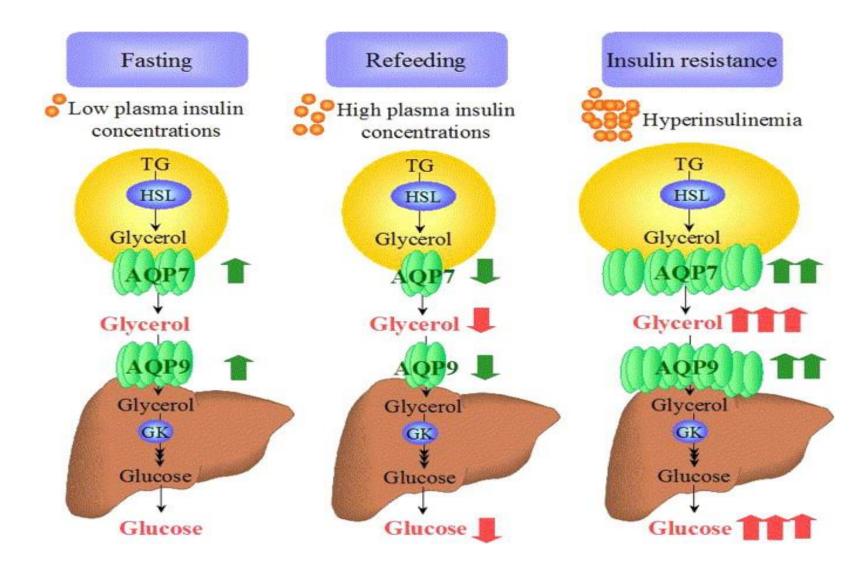
NADPH

 O_2^{-}

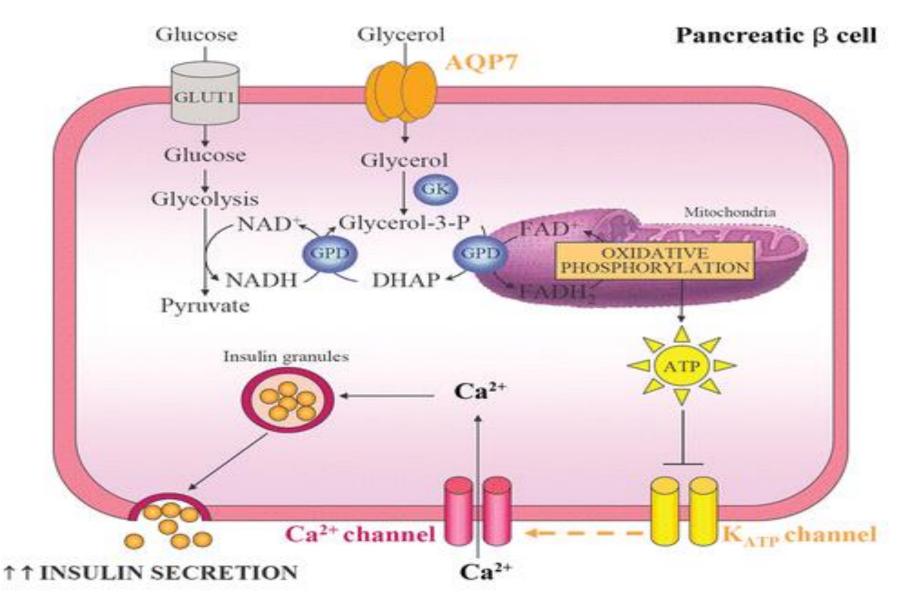








Aquaporin's channel action



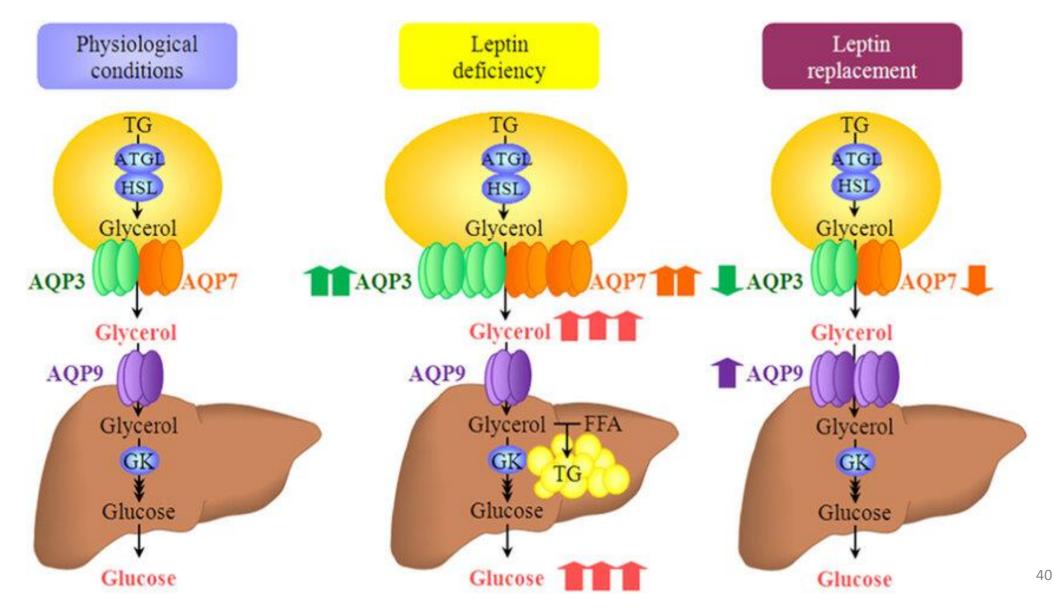
Endocrine pancreas

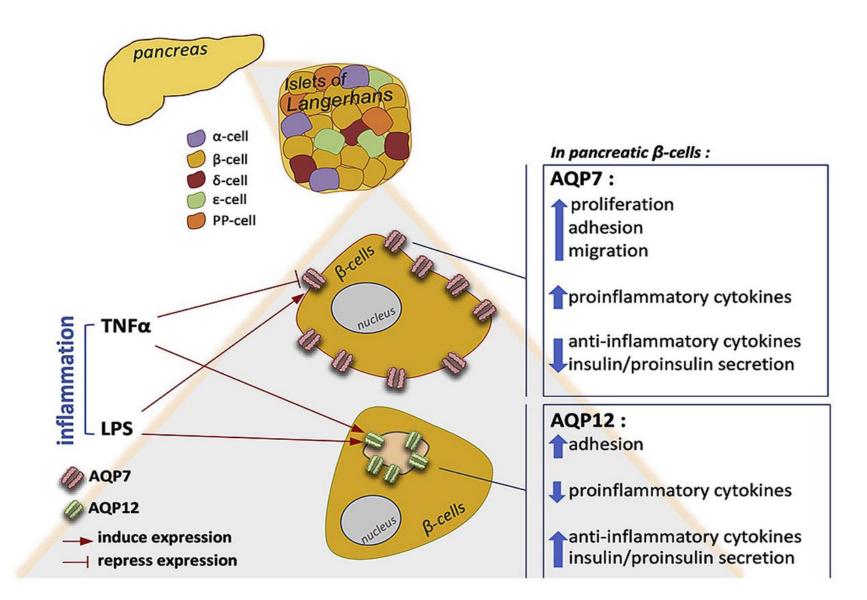
Pan

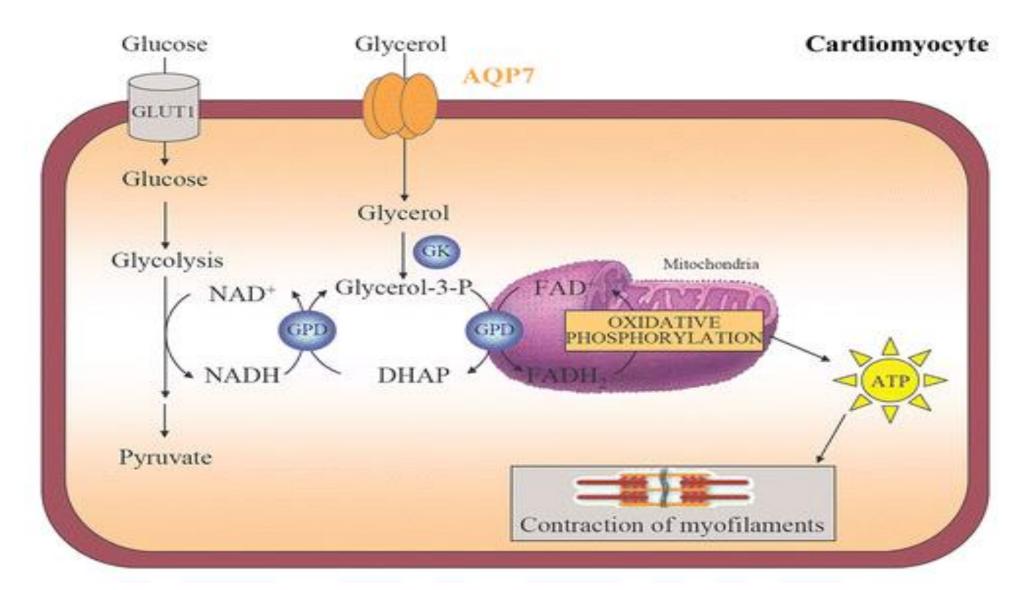
Sleeve gastrectomy was associated with an upregulation of AQP7 together with a normalization of the increased AQP12 levels in the rat pancreas. Interestingly, ghrelin and GLP-1 repressed AQP7 and AQP12 expression in RIN-m5F β -cells. AQP7 protein was negatively correlated with intracellular lipid accumulation in acylated ghrelin-treated cells and with insulin release in GLP-1-stimulated β -cells.

ulin etion

Effect of leptin on Aquaporin's channel







Effect of insulin on aquaporin 9

 Insulin suppresses AQP9 mRNA levels in time- and dose-dependent manners in H4IIE hepatocytes.

- Promoter analysis demonstrates that insulin reduces AQP9 mRNA via IRE locating at -496/-502 promoter region.
- Administration of STZ results in increased AQP9 mRNA and protein levels in insulin-insufficient mice.

