

C.79-Hepatitis C virus and type 2 diabetes: Molecular homology between HCV E2 envelope and glucagon

Written by GMK Tran

Friday, 12 April 2013 16:48 - Last Updated Friday, 14 June 2013 12:56

We found a homology between human glucagon, especially miniglucagon (19-29),

and HCV E2 protein chimera; the glucagon motif H.....RR..DFVQ W LM

was conserved in HCV E2 motif H... ..RR..DF .QGW.PI,

including the complete receptor binding motif located at the COOH-terminus and containing the residues D15, A19, F22, W25, L26, M27, T29 (M27 replaced by I; and *linearly* D15 is missing, A19 replaced by V):

Receptor binding residues * * * * *

Numbering 15 19 22 25262729

Glucagon 15-**DSRRAQDFVQ-W-LMNT**-29

HCV E2 chimera 459-**CRRVE DFVQGWGL I N**-473

HCV E2 variant **T**-474

17-**RR**-18 is a protease cleavage site.

The 2 gaps for glycines G (=Gly) are also encountered in a phylogenetically related glucagon in insects, the Adipokinetic Hormone pQVNFTPGWGTG (Gade G, 1997; Tran GMK, 2011), an 11 residues long peptide, which possesses the GWG motif of HCV E2.

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Adipokinetic Hormone

pQVNFTPGWGT

(pQ = pyroGlu)

HCV 2b E2

E- DFRI GWGT

Mini-glucagon (19-29) AQDFVQWLMNT is 1000 fold more powerful than glucagon itself (Dalle S, 2002). It is a very potent (ID50 close to 0.1 pmol/l) inhibitor of insulin release from beta-cell.

The glucagon (1-21) and des-(22-26) glucagon have potencies of