

Virtual Three Minute Thesis



Investigating localization and function of transcription factor FOXO4 in Basal breast cancer (BBC) and Glioblastoma multiforme (GBM) cell lines

Presented by Millat Jahan

Background

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- FOXO transcription factors
- Glioblastoma Multiforme
- Basal Breast Cancer

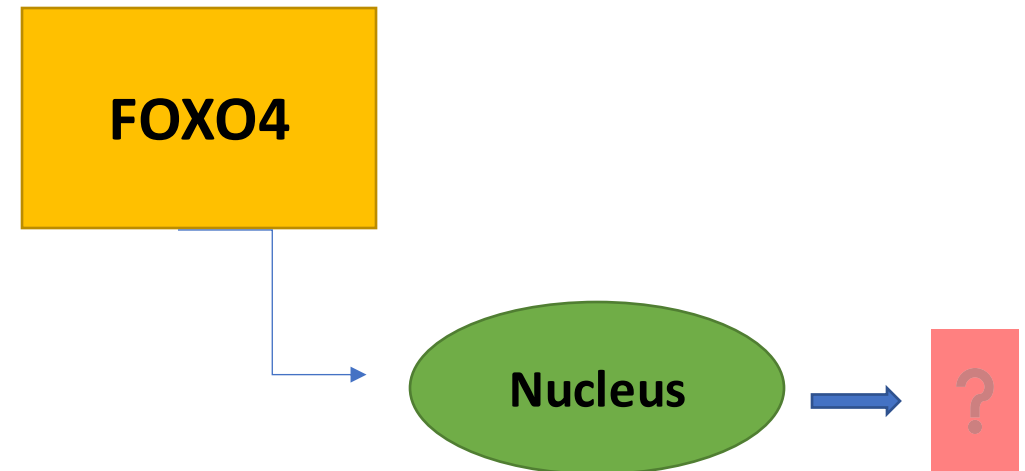


Introduction cont..

- FOXO4 was extensively identified as a key tumor suppressor by regulating its target genes associated with antioxidative stress, cell cycle arrest, and apoptosis (Wang et al.,2016)

Hypothesis

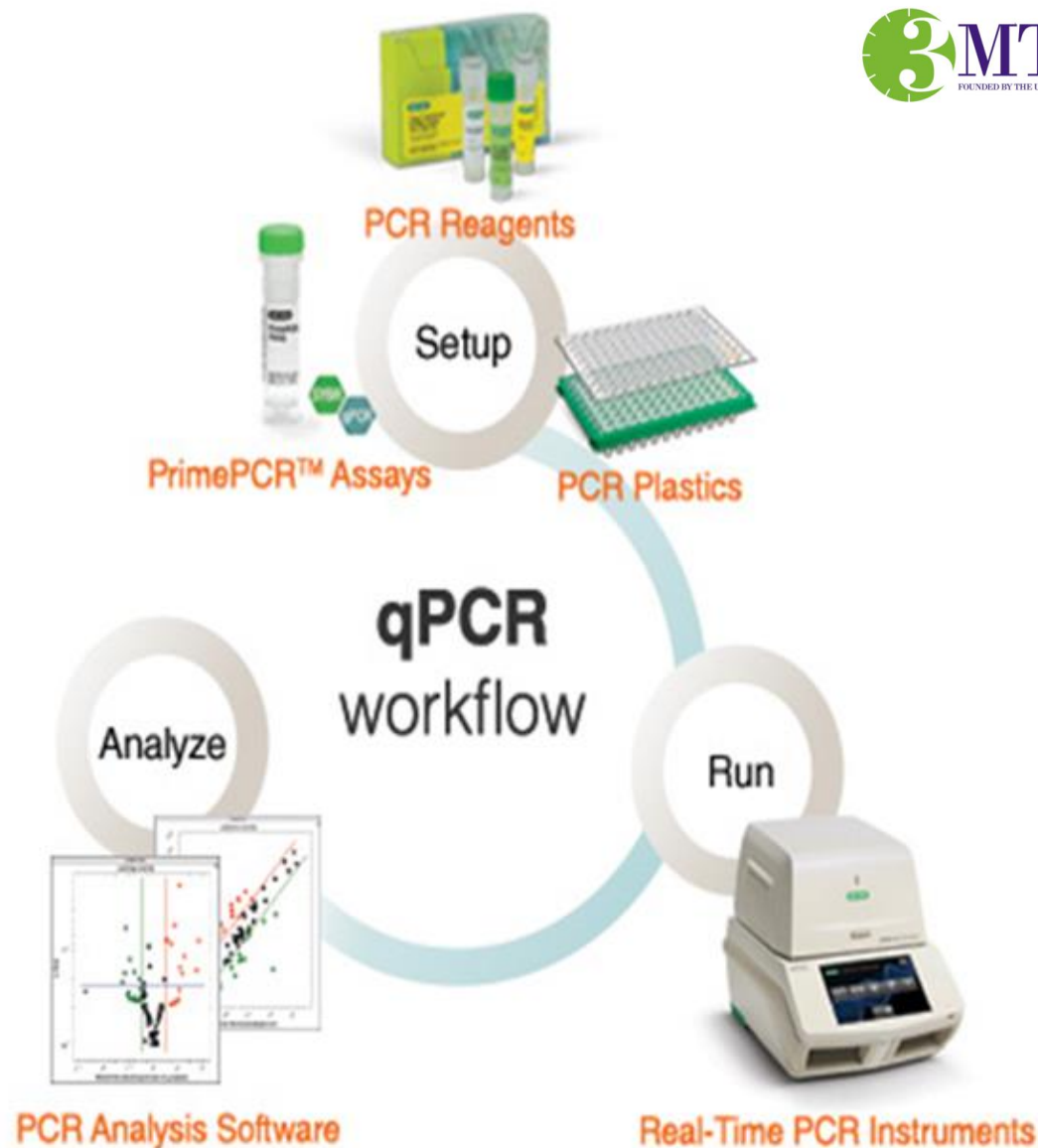
- For some reason FOXO4 is always in the nucleus.
- I hypothesize that this factor rewires cancers and enable nuclear FOXO4 factor to directly promote the expression of stem cell genes



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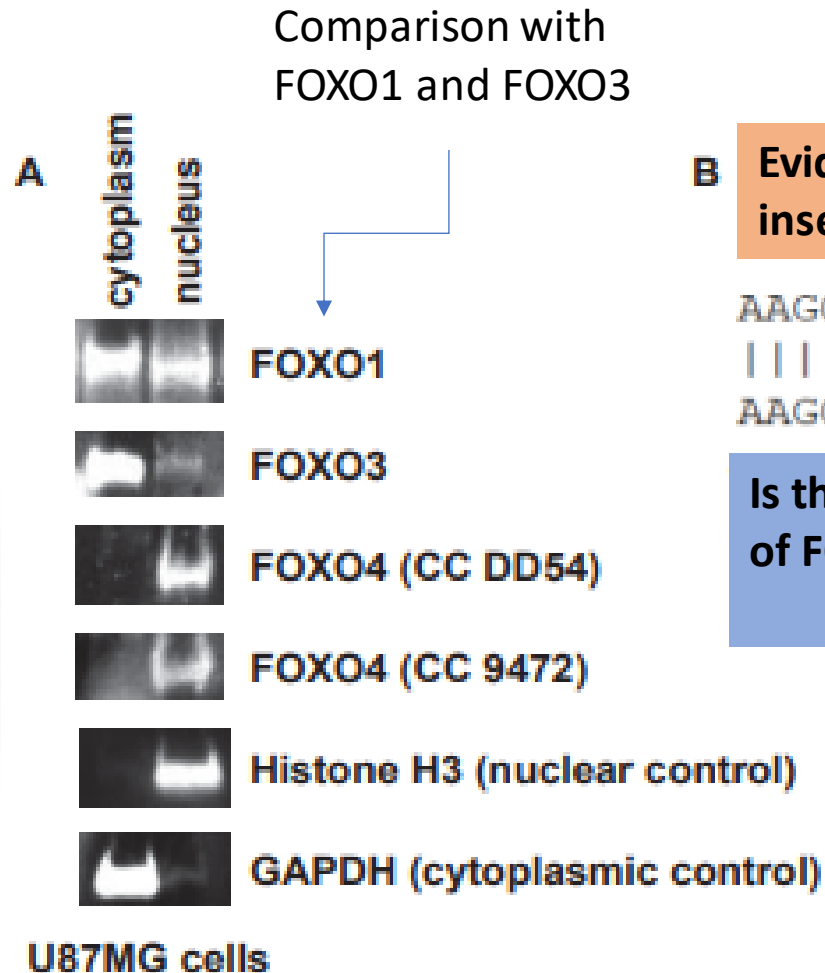
Methods

- Cell culture
- Immunofluorescence experiment
- qPCR to measure changes in gene expression
- Western blot
- RNA seq looking for specific interactions



Preliminary experimental outcome

- Here Cell line is U87MG
- Western blot analysis was done
- FOXO1 and FOXO3 was in cytoplasm and nucleus, but FOXO4 was always in nucleus



B Evidence of in frame 6 nucleotide insertion in the DNA binding domain

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AAGGGTGACAGCGACAGC Wild-type FOXO4
|||||                |||||
AAGGGT-----GACAGC U87MG FOXO4
  
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Is this the reason of nuclear localization of FOXO4?

Fig. Western blot analysis showing nuclear localization of FOXO4

Expected outcomes

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- Identifying novel mechanism that determines localization of FOXO4 which could be pharmacologically targeted as innovative avenues for therapeutic interventions
- Delineation of mechanism employed by FOXO4 factors that can regulate stem cell genes in aggressive cancers



Conclusions

- Repeating experiments can be done to confirm over expression of FOXO4 lead to stem cell characteristics
- Investigating how FOXO4 transcription factors promote cancer via regulation of stem cell genes



References

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Thank you!