



SDI Review Form 1.6

Journal Name:	Ophthalmology Research: An International Journal
Manuscript Number:	Ms_OR_38734
Title of the Manuscript:	ASSOCIATION OF TRANSCRIPTION FACTOR 7 LIKE 2 (TCF7L2) POLYMORPHISMS WITH DIABETIC RETINOPATHY
Type of the Article	Review Article

General guideline for Peer Review process:

This journal's peer review policy states that **NO** manuscript should be rejected only on the basis of '**lack of Novelty**', provided the manuscript is scientifically robust and technically sound. To know the complete guideline for Peer Review process, reviewers are requested to visit this link:

(<http://www.sciencedomain.org/page.php?id=sdi-general-editorial-policy#Peer-Review-Guideline>)

PART 1: Review Comments

	Reviewer's comment	Author's comment (if agreed with reviewer, correct the manuscript and highlight that part in the manuscript. It is mandatory that authors should write his/her feedback here)
Compulsory REVISION comments	<ul style="list-style-type: none"> The title of the paper has to be modified because many documented reports already show that TCF7L2 is involved in RD. So this same title can not stand for this paper. The title should reflect the relationship between TCF7L2, VEGFα and DR. The abstract says nothing about TCF7L2, VEGFα. This must be included to make a strong abstract without which nothing is new in the abstract. No data shows that if TCF7L2 & VEGF α are implicated in DR then it must be due to TCF7L2 polymorphism. This must be demonstrated. And even if it is due to polymorphism then which polymorphism is involved?. The background does not seem to be sufficiently rich. TCF7L2 factor needs to be located in the Wnt Signalling pathway. This figure says nothing about TCF7L2 & VEGF. Again nothing is new in the conclusion. These are already existing in published data 	<ul style="list-style-type: none"> Title correction noted and Modified VEGF incorporated into abstract We tried to explain the association of TCF7L2 polymorphism with upregulation of VEGF using Luo et al. 2014 as ref. TCF7L2 is a transcription factor in wnt signalling pathway and VEGF is a wnt targeted gene. We have shown that TCF7L2 is a member of the TCF family which is shown in Fig 2 Conclusion is modified
Minor REVISION comments	<p>The English needs to be revised.</p> <p>It will be necessary to indicate the sources of Fig 1, 2 & 3.</p>	<ul style="list-style-type: none"> English proofread done. All Figs were referenced appropriately.
Optional/General comments		