

Shikshak Bahuuddeshiya Shikshan Mandal Wadi Nagpur
VIMALTAI MAHILA MAHAVIDYALAYA

(Affiliated by Gondwana University Gadchiroli)
Saoli, Tah. Saoli, Dist- Chandrapur - 441225

Phone No: 07174-269656

Email: principalvmmsaoli@gmail.com

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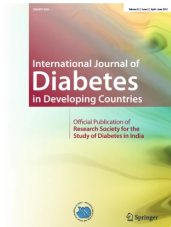
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Submissions are sourced globally, particularly from countries affected by the diabetes epidemic.

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All published materials support their claims with transparent data and comply with ethical standards.

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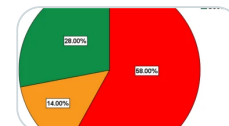
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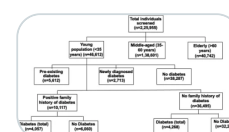
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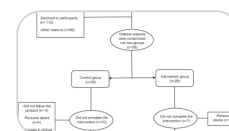
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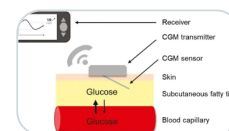
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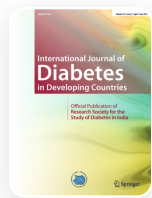
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SNPs in the *catalase* promoter: a study based on Indian diabetic individuals

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Abstract

Background

It is known that anti-oxidant defense is compromised in diabetic individuals. In our study, we indeed found decreased catalase activity in diabetic individuals compared to non-diabetic controls. Therefore, we analyzed the single nucleotide polymorphisms (SNPs) in the promoter region of catalase in diabetic individuals. To investigate catalase promoter SNP's contribution (if any) to the low catalase activity and assess their functional significance by reporter assay.

Methods



The activity of catalase was quantitated from 109 non-diabetic and 138 diabetic individuals. Genomic DNA isolated from these individuals was screened for SNPs in the *catalase* promoter by direct sequence analysis of PCR products. The functional effect of these polymorphisms was checked by reporter assay.

Results

We found six reported SNPs of which, three were polymorphic in our study groups at -330, -89, and -20 position viz. rs1001179: C>T, rs7943316: A>T, and rs1049982: T>C. Out of the three SNPs, only rs1001179: C>T showed a positive association with the occurrence of low activity of catalase in diabetic individuals ($p<0.05$). However, reporter assay confirmed that the presence of these SNPs does not significantly affect the transcriptional activity of the gene ($p>0.05$). Furthermore, our in silico analysis revealed that the presence of these SNPs did not significantly affect the binding of transcription factors except at -330 position.

Conclusion

Our reporter assay suggesting no functional relevance of promoter SNPs in reduced catalase activity observed in Indian diabetic individuals is a novel finding.

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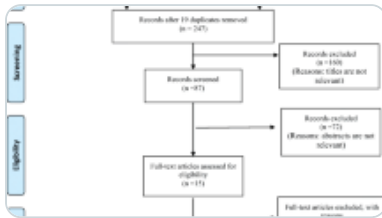
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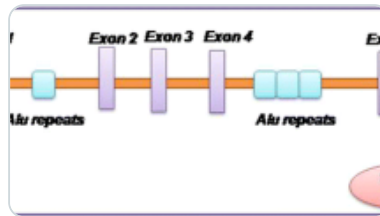
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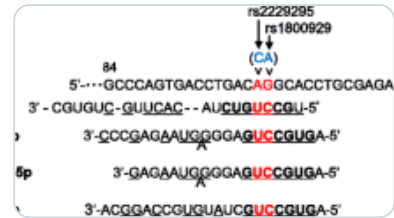
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Data availability

The data that supports the findings of this study are available in Appendix, table-1 of this article.

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Author information

Authors and Affiliations

Department of Zoology, Center of Advanced Studies, Savitribai Phule Pune University, Pune, 411007, India

Dipak A. Kadam, Saurabh D. Kalamkar, Amit Saraf, Imran Pathan, Jhankar Acharya, Komal Pekhale, Saroj Ghaskadbi & Richa Ashma

Dr. B. N. Purandare Arts and Smt. S.G. Gupta Commerce & Smt. S. A. Mithaiwala Science College Lonavala, Pune, 410403, India

Dipak A. Kadam

Microbial Culture Collection, National Centre for Cell Science, Pune, 411007, India

Yogesh Shouche

National Institute of Virology, 20 –A, Dr. Ambedkar Road, P.B. No. 11, Pune, 411007, India

Kavita Lole

Contributions

Dipak Kadam: formal analysis; methodology; resources; writing—review and editing. Saurabh D. Kalamkar: Resources. Amit Saraf: methodology. Imran Pathan: investigation, methodology, writing—original draft preparation. Jhankar Acharya: formal analysis; writing—review and editing. Komal Pekhale: formal analysis; methodology. Yogesh Shouche: resources. Kavita Lole: resources. Richa Ashma: conceptualization; formal

analysis; supervision; validation; writing—original draft preparation; writing—review and editing. Saroj Ghaskadbi: conceptualization; funding acquisition; supervision; validation; writing—review and editing.

Corresponding authors

Correspondence to [Saroj Ghaskadbi](#) or [Richa Ashma](#).

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The authors declare no competing interests.

Informed consent

Informed consent was obtained from all individual participants included in the study.

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