IDENTIFICATION OF ENDOGENOUS OVEREXPRESSION OF CATALASE, AS A INHIBITOR OF EMT SIGNALLING THE PROGRESSION OF BREAST CANCER.

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ABSTRACT:

KEYWORDS: Breast tumor; Exercise; Catalase; Vimentine; Beta-catenin; Ecadherin; innovative method.

BACKGROUND: Tumor is an abnormal cell growth that spreads to other organs. Breast tumor develops in the tissues of the breastRegular exercise can help to lower the chance of developing breast tumor. For the year 2020, the expected incidence of tumor patients in India was 679,421 (94.1 per 100,000) for males and 712,758 (103.6 per 100,000) for females.. **AIM:** To analyse the endogenous over the expression of catalase as an inhibitor of EMT signaling in breast tumor through molecular docking.

MATERIALS AND METHODS:

The molecular docking analysis is a bio informatic study conducted in a private dental college. The endogenous substance catalase which is secreted after post exercise is used as our target protein. The interaction of catalase with the proteins relevant to breast tumor namely Vimentin, Beta-catenin, Ecadherin are included for docking analysis. The protein structure is retrieved using protein data bank, Protein protein docking done using patch Dock server followed by visualisation of protein-protein interaction using pymol.

RESULT:

The surface representation of catalase with Vimentin, Beta-catenin and Ecadherin showed good shape complementarity. The results showed that Catalase forms strong interaction with Vimetine, Beta catenin, Ecatherine proteins in terms of hydrogen bond interaction, hydrophobic and non bonded interaction. Through this interaction these proteins might control the overexpression of catalase activity in breast tumor.

CONCLUSION:

From the obtained result it can be concluded that catalase may have a protective role against breast tumor through its interaction with Vimentin, E-Cadherin, Beta-Catenin. The present study has suggested a possible mechanism of catalase in the inhibitor of EMT signaling in breast tumor.

KEYWORDS: Breast tumor; Exercise; Catalase; Vimentine; Beta-catenin; Ecadherin; innovative method.

INTRODUCTION:

The catalase CC genotype is associated with 17% reduction in breast tumor and has a therapeutic role and antioxidant enzyme that decomposes hydrogen peroxide, damages the DNA and protein and builds up toxic levels. Using mitochondrial tumour cells and tumour stromal cells to target catalase inhibits ROS-driven tumour development and metastasis. As a result, enhancing the mitochondrial compartment's antioxidant capacity could be a sensible therapeutic strategy for invasive breast tumor.¹

Using mitochondrial tumour cells and tumour stromal cells to target catalase inhibits ROS-driven tumour development and metastasis. As a result, enhancing the mitochondrial compartment's antioxidant capacity could be a sensible therapeutic strategy for invasive breast tumor.³. The trials from previous studies⁴ ^{5,67,85,65910111210,121314 15} have led us to concentrate on the study.

Important characteristics of basal breast tumor, such as its proclivity for poor overall survival, imply that it could be a promising pharmaceutical target for this aggressive breast tumor subtype16. Ecadherin is a calcium-dependent cell adhesion protein that is one of the most investigated tumour suppressors in breast tumor. In various cases, signalling molecules and transcription factors that regulate Ecadherin expression have been identified to promote EMT ¹⁷. Evidence suggests that EMT may play a role in the course of human breast tumor, as well as the prognostic value of vimentin expression.¹⁸. Studies at molecular levels were performed by our team of researches which insisted us to proceed this study ^{19–26},²⁷,²⁸,²⁹,^{30,31},³²,³³,^{34–38}Thus the present study attempts to analyse the role of endogenous over the expression of catalase as an inhibitor of EMT signalling the progression of breast tumor.

MATERIALS AND METHODS:

The molecular docking analysis is a bio informatic work conducted in a private dental college. The endogenous substance catalase which is secreted post exercise is used as our target protein. The interaction of catalase with the proteins relevant to breast tumor namely Vimentin, Beta-catenin, Ecadherin are included for docking analysis.

Procedure:

Retrieval of Target proteins structures from Protein data bank

In order to study the mechanism of interaction between Catalase with Vimetine, Beta catenin, Ecatherine proteins, the three dimensional structures were downloaded from Protein Data Bank using the respective ids(Pdb ids: Catalase- 1DGF; Vimetine -1GK4; Beta catenin -4DJS; Ecatherine - 2O72)³⁹

Protein-Protein Docking.

Patch Dock (http://bioinfo3d.cs.tau.ac.il/PatchDock) is a geometry-based molecular docking technique⁴⁰ used to study the interaction between Catalase with Vimetine, Beta catenin, Ecatherine proteins. The Patch Dock service calculates docked transformations that result in strong molecular shape complementarity. The Connolly dot surface representation of the molecules is divided into concave, convex, and flat patches using the algorithm. In order to produce various transformations, the patches were paired according to their complementarity. For clustering, a default value of 4 Å was used and redundant solutions were discarded by RMSD clustering. The geometric score, desolvation energy, interface area scale, and the actual rigid transformation of the solutions are created by the Patch Dock output. For each complex, twenty solutions were created, from which one complex was selected for further analysis based on the scoring geometric shape for both complexes.

Visualization of Protein – Protein interactions.

Using the academic version of the Pymol, the residual interactions between docked complexes were viewed. The colour intensity for interactions was clearly visible here, and the findings were exported. ⁴¹ Pdbsum was used to determine the types of interactions that Catalase has with the proteins Vimetine, Beta catenin, and Ecatherine.

RESULTS:

The results suggests that Catalase forms strong interaction with Vimetine, Beta catenin, Ecatherine proteins (Figure 1,2,3) in terms of hydrogen bond interaction, hydrophobic and non bonded interaction. Once the docking process is completed, it will send the web link to the mail which we gave during the docking process. The output of results contains 20 top ranked complexes. Based on the, the interface area size the best complex was selected for each protein and further analyzed using a pymol visualization tool. Some the amino acids from catalase like ALA-8, SER-9, LYS-77,THR-115, ALA-117, GLY-118,jjj GLY-121, SER-122, ALA-123, VAL-126, ARG-127,ASP-128, GLN-168, LYS-19, THR-174, LEU-176, ASP-178 and VAL-182 alternatively involved in the formation of complex with the Vimetine, Beta catenin, Ecatherine. Through this interaction these proteins might control the overexpression of catalase activity in breast tumor. The present study has suggested a possible mechanism of catalase in the inhibition of EMT signaling in breast tumor.(table 1)

S.No	Protein Name	Score	ACE (Atomic
			contact energy)
1	Vimetine	15794	- 389.49
2	Beta catenine	22658	-490.24
3	Ecatherine	19246	-273.28

Table 1: Molecular docking results of Catalase with Vimetine, Beta catenine, Ecatherine proteins

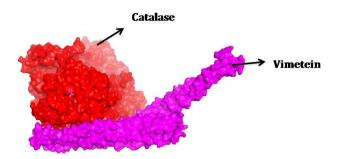


Figure 1-Protein-Protein interaction between Catalase-Vimetein

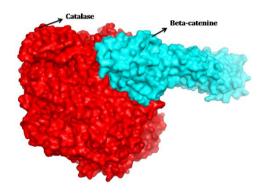


Figure 2- Protein-Protein interaction between Catalase-Beta catenin

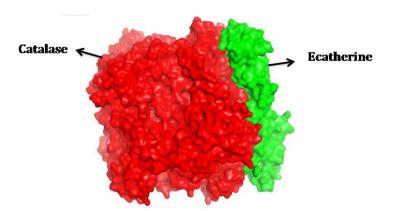


Figure 3-Protein-Protein interaction between Catalase-Ecadherin

DISCUSSION:

The docking study's findings could add to the growing body of data indicating exercise-induced endogenous catalase is a protective molecule against breast tumor cell growth. It was possible to design a more selective and efficient inhibitor by understanding these types of interactions. We found that catalase, which is created after exercise, functions as a regulator for EMT signalling in breast tumor in our study..

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Protein-protein interactions (PPI) are important in cellular biological processes, immunological response, and cellular organisation. As a result, PPI analysis is extremely important and can aid in the identification of pharmacological targets as well as treatment design. High-throughput technologies like yeast two-hybrid screenings (Y2H) and mass spectrometric protein complex identification (MS-PCI) have yielded a lot of data, but they're also expensive and time-consuming. Furthermore, these approaches may not be relevant to proteins from all species, resulting in false-positive results. Its goal is to find docking changes that result in substantial molecular shape complementarity. When applied, such enhancements result in extensive interface areas as well as minor amounts of steric conflicts. Patch dock takes two molecules in PDB format as input, which we can upload or have the server obtain from the pdb using their respective pdb ids. Regular exercise has an influence on acute and chronic symptoms of breast carcinoma. ⁴².

EMT allows epithelial tumour cells to acquire mesenchymal characteristics that aid in metastasis. ⁴³. During the epithelial to mesenchymal transition during early metastasis, the impact of Vimentin on tissue factor expression is limited by negative regulation of TF in mRNA. ⁴⁴.. Beta-catenin is also known as Catenin-1. It has a dual function that is regulation and coordination. Cell-Cell adhesion, gene transcription takes place. It is located in the membrane at the side of the cytoplasm. ^{45,46,47}

Ecadherin determines whether the tumour is ductal or lobular. It regulates the formation and stability of ECadherin. It plays an important role in morphogenesis and homeostasis. The EGF receptor was shown to be overexpressed in tumors. Ecadherin expressions in menopausal status, hormone receptor status, and age were found to have no significant association.⁴⁸. Ecadherin expression is controlled by a number of signalling molecules and transcription factors. Ecadherin loss has been linked to EMT in a number of situations. Because of the frequent changes in their expression, it's possible that the Ecadherin pathway's loss of function is important in the development of breast tumor.⁴⁹. Understanding the basic development of tumours and increasing early diagnosis to expediting the process of turning prospective therapeutic targets into clinical features are all things that we're working on. ⁵⁰. For further understanding the study can be carried out in vitro and in a large population.

CONCLUSION:

The study concludes that exercise induced endogenous over the expression of catalase may act as an inhibitor of EMT signaling in preventing breast tumor initiation and progression. From this we analyse the role of endogenous over the expression of catalase as an inhibitor of EMT signalling in the progression

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CONFLICT OF INTEREST

All the authors declare that there was no conflict of interest in the present study.

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