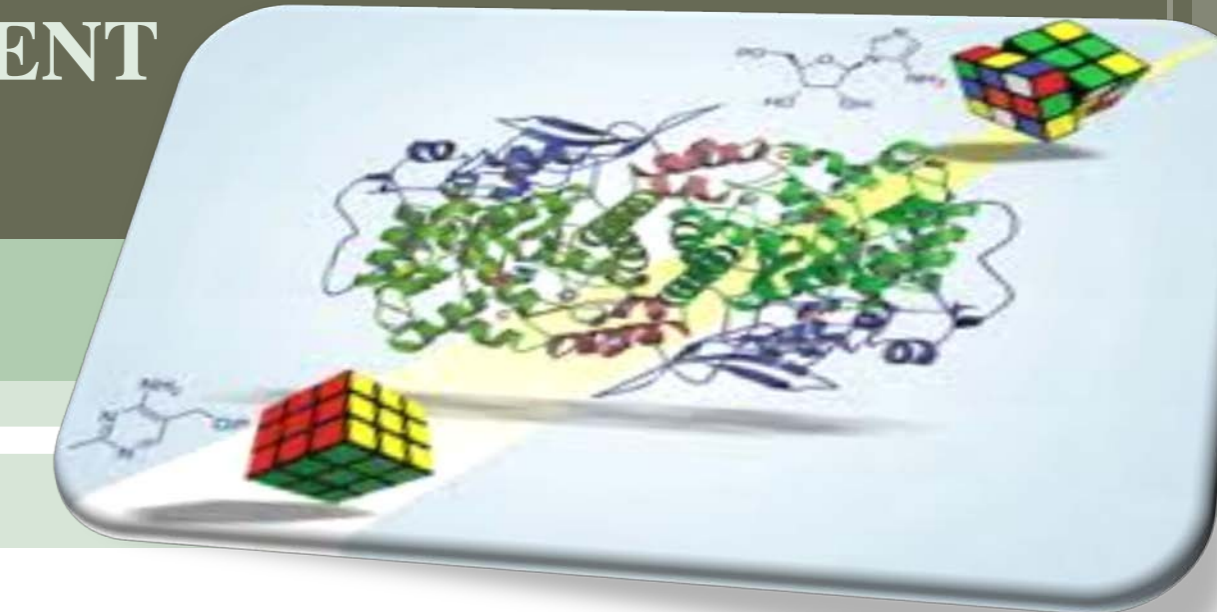


PROTEIN ENGINEERING OF GLUTATHIONE TRANSFERASE FOR THE DEVELOPMENT OF OPTICAL BIOSENSOR TO DETECT XENOBIOTICS



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Abstract

Glutathione transferases (GSTs, EC. 2.5.1.18) are inducible enzymes that play essential role in detoxification and degradation of toxic compounds, including pesticides. The purpose of the present study is the development of an optical enzyme biosensor based on GSTs, for the detection and determination of pesticides in environmental samples. Protein engineering was used for the creation of a GST variant with higher selectivity towards pesticides. cDNA libraries were created from *Phaseolus vulgaris* and *Glycine max* stressed plants using degenerated primers and reverse transcription-PCR. Large diversity in GST genes was accomplished employing directed evolution through DNA shuffling of a mixture of GST genes from *P. vulgaris* and *G. max* stressed plants. The shuffled library of chimaeric GST genes was cloned in *E. coli* expression plasmid. Screening of the library led to the isolation of a novel GST enzyme that displays both glutathione transferase and glutathione peroxidase activities. The enzyme was purified by affinity chromatography and characterized by kinetic analysis towards 20 different substrates and 66 different pesticides. The results showed that the organochlorine insecticides and strobilurins (fungicides) are strong inhibitors of the enzyme. The specificity of the enzyme towards pesticides was further improved using site-saturation mutagenesis at position Phe117. The mutant Phe117Ile displays 5-fold higher catalytic efficiency and selectivity towards organochlorine insecticides. Therefore, the mutant GSTPhe117Ile was used for the development of an optical biosensor. The enzyme was immobilized in alkoxysilane (TEOS/PTMOS) sol-gel system in the presence of the pH indicators bromocresol purple (acidic) and phenol red (basic). The bioactive material exhibits linearity in the range of 0.625-30 μM α -endosulfan (pH=4-7) at 562 nm and was used for the development of an analytical method for the determination of α -endosulfan in environmental samples.

Results

Stressed *Phaseolus vulgaris* and *Glycine max* plants were used for the creation of cDNA libraries using degenerated primers and reverse transcription-PCR. Large diversity in GST genes was accomplished employing directed evolution through DNA shuffling of a mixture of GST genes from *P. vulgaris* and *G. max* stressed plants. The shuffled library of chimaeric GST genes was cloned in *E. coli* expression plasmid. Screening of the library led to the isolation of a novel GST enzyme *PvGmGSTUG* that displays both glutathione transferase and glutathione peroxidase activity. This novel GST enzyme belongs to *tau* class and is strongly inhibited by organochlorines (insecticides) and strobilurins (fungicides).

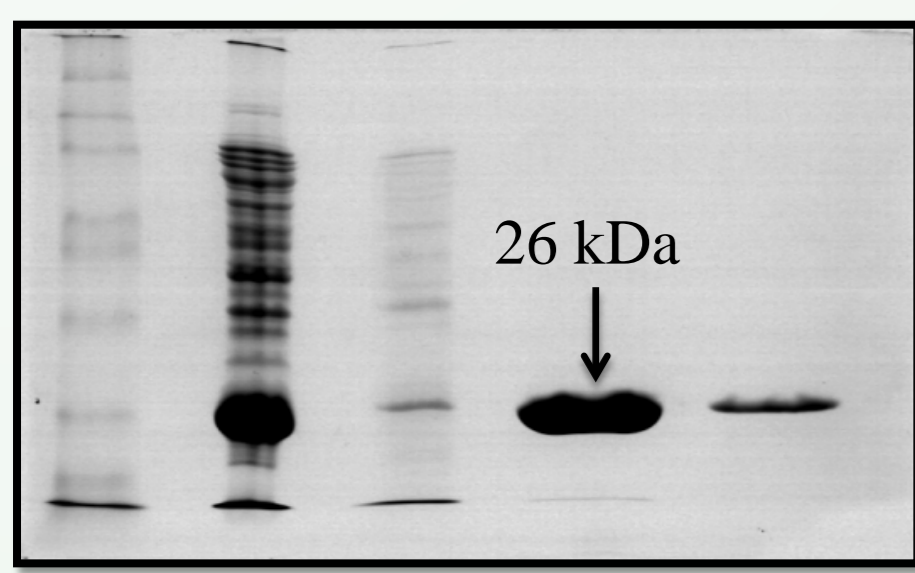


Fig.1. SDS-PAGE electrophoresis of *PvGmGSTUG* purification by affinity chromatography.

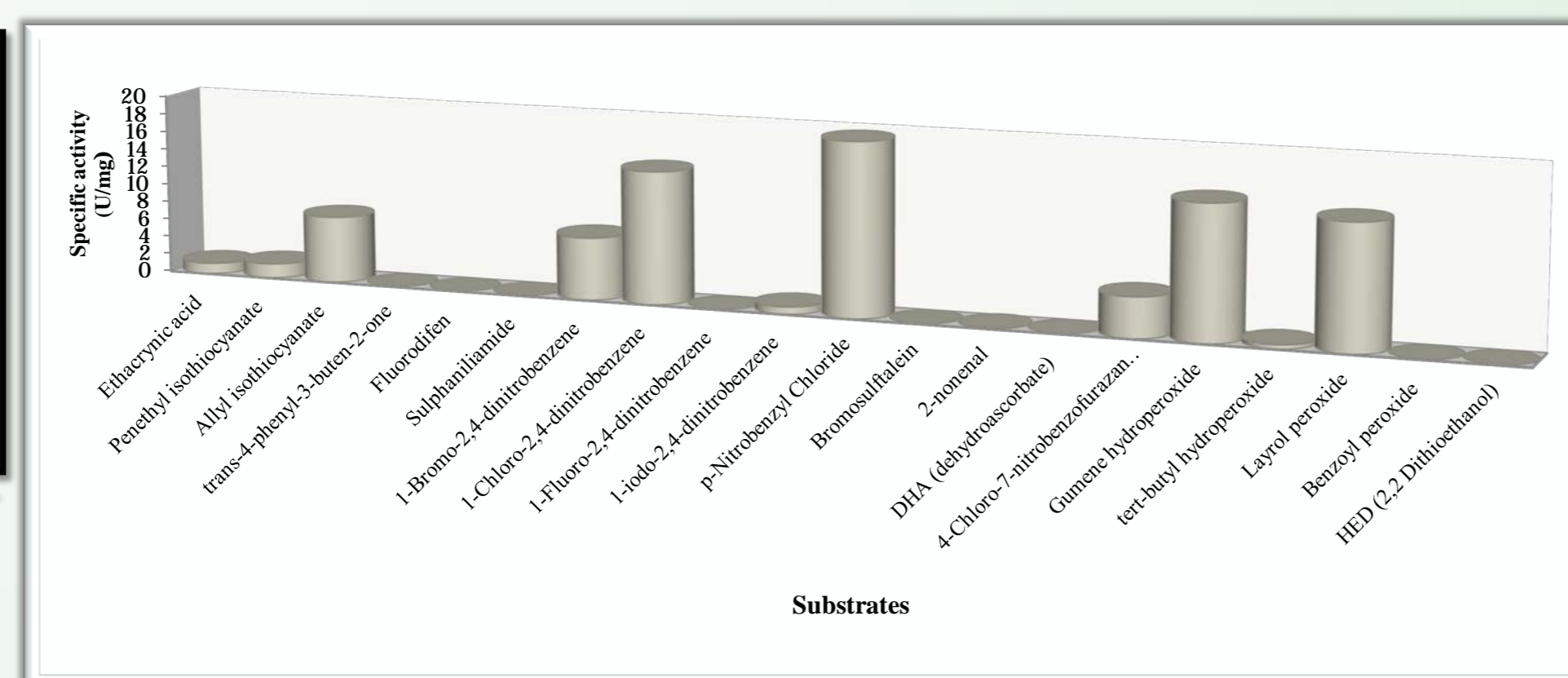


Fig.2. Specific activity of *PvGmGSTUG* towards 20 different substrates

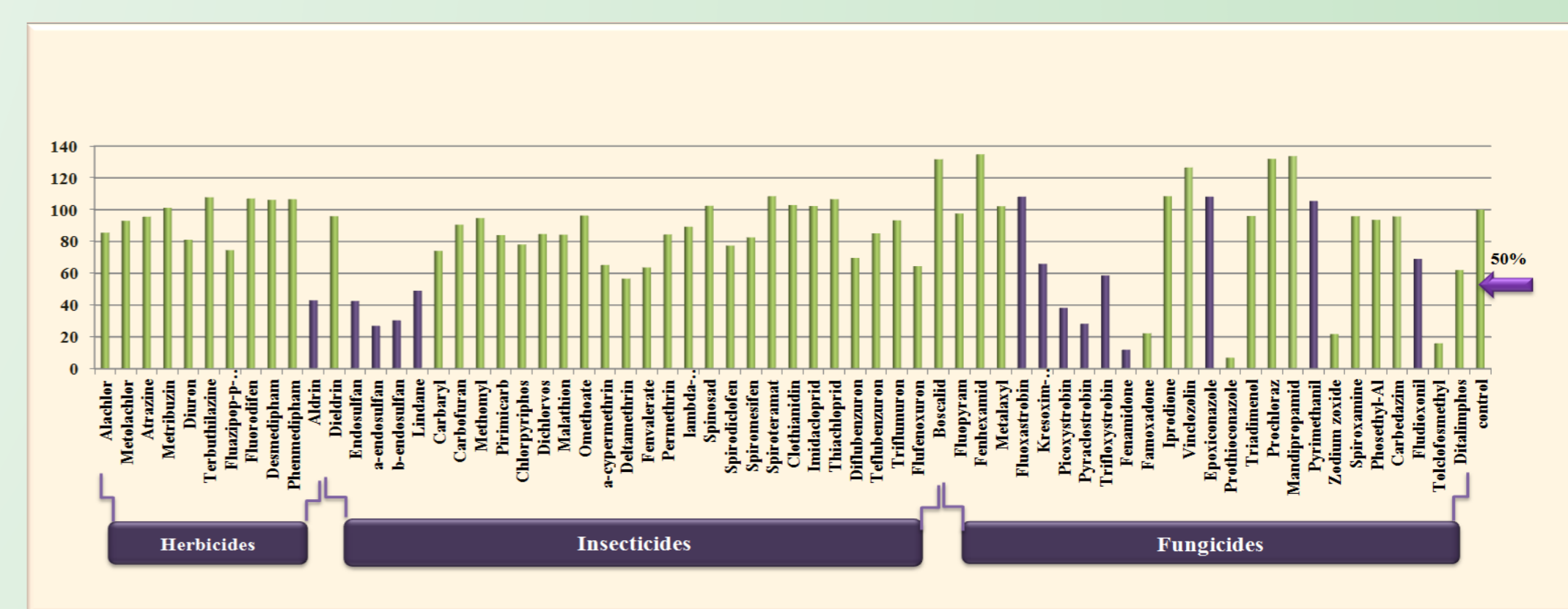
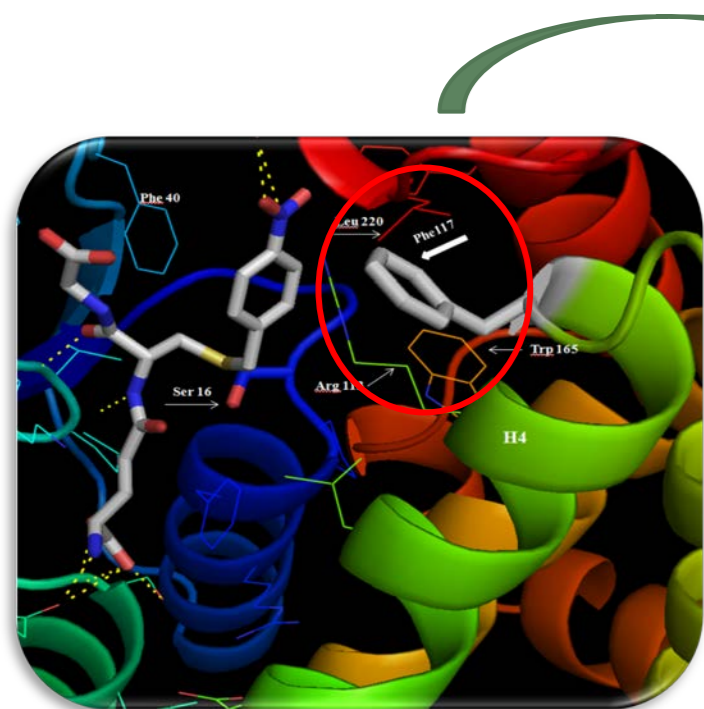


Fig.3. Inhibition (% residual activity) of *PvGmGSTUG* by 66 different pesticides

For the enhancement of *PvGmGSTUG* pesticide selectivity, site-saturation mutagenesis was carried out at position Phe117. Phe117 is located at the entrance of the substrate binding-site. The enhanced selectivity of each mutant is shown in Fig.5. The mutant Phe117Ile exhibits strong inhibition by organochlorine insecticides. Therefore, the mutant F117I was selected for the development of the biosensor. This mutant also shows 5-fold higher k_{cat}/K_m than wild type enzyme. The mutant F117I was immobilized in sol-gel composed by alkoxysilanes (TEOS and PTMOS), with aging in TEOS. Fig 6. depicts kinetic analysis of the immobilized mutant enzyme.



- F117I
- F117R
- F117W
- F117S
- F117T
- F117G
- F117H

Fig.4. Site-saturation mutagenesis in the position 117

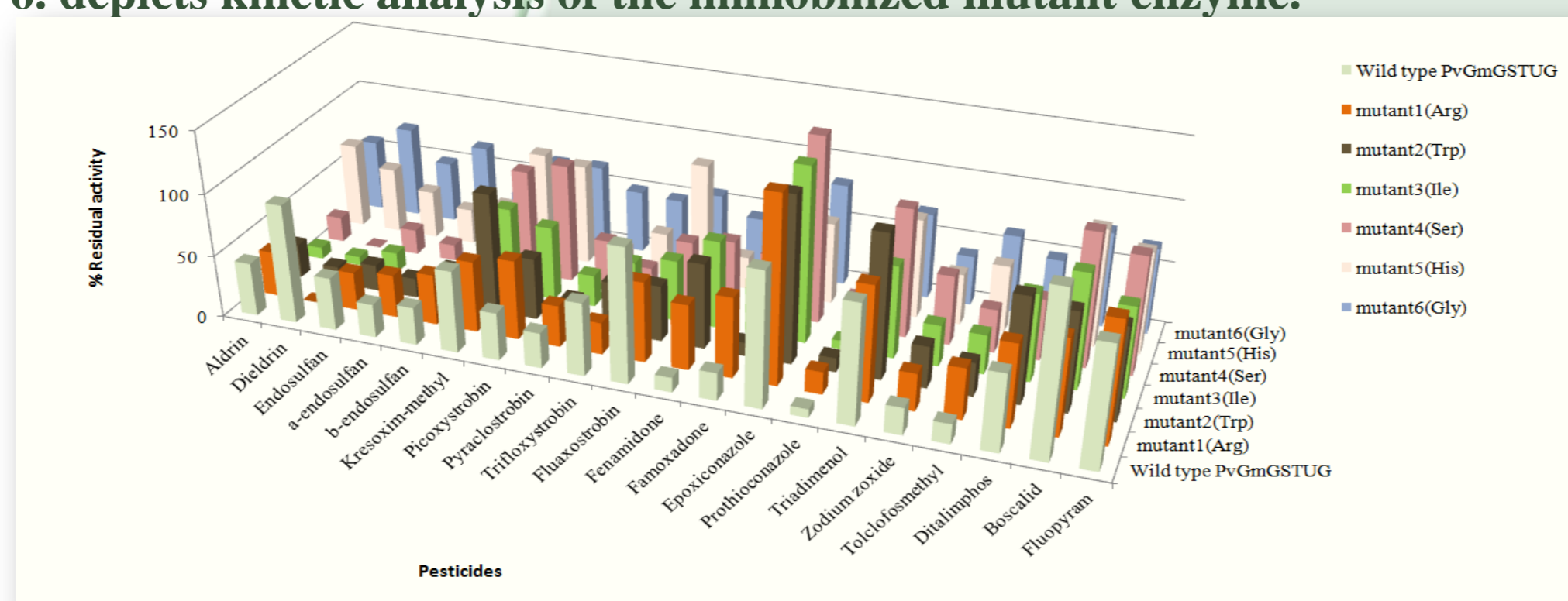


Fig.5. Screening of the mutants towards pesticides

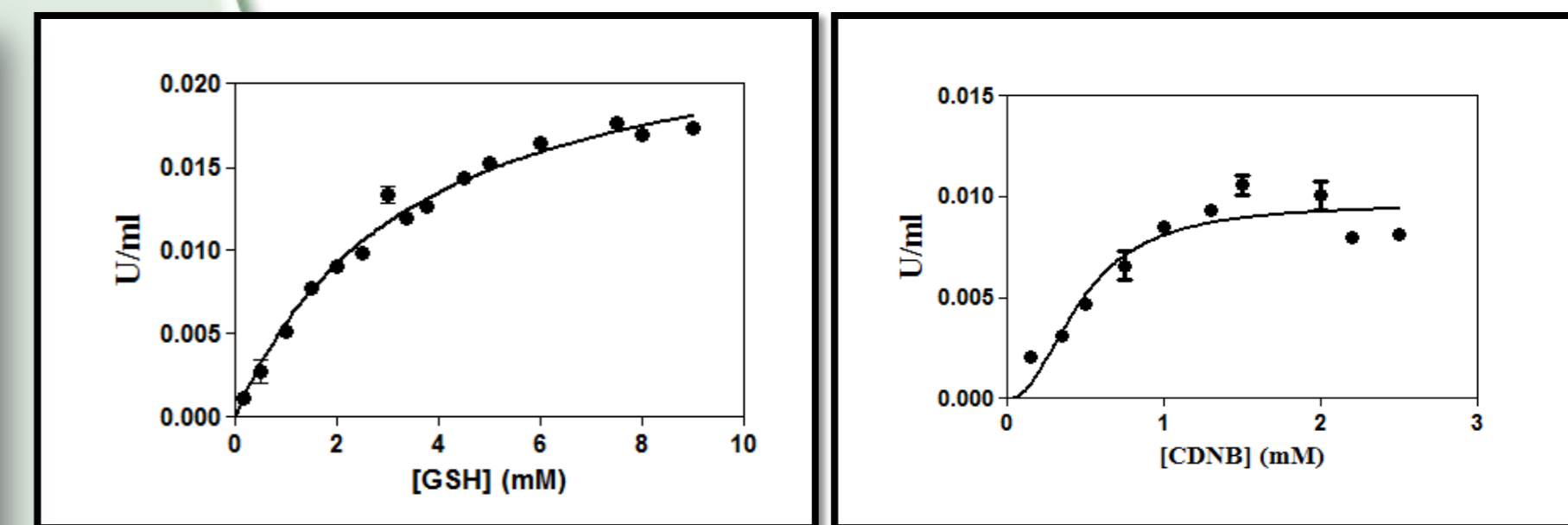


Fig.6. Kinetic analysis of the immobilized mutant F117I

The analytical method was based on the enzyme inhibition by α -endosulfan. The bioactive material consists of two different sol-gels that led to the immobilization of F117I mutant and the pH indicators (bromocresol purple and phenol red).

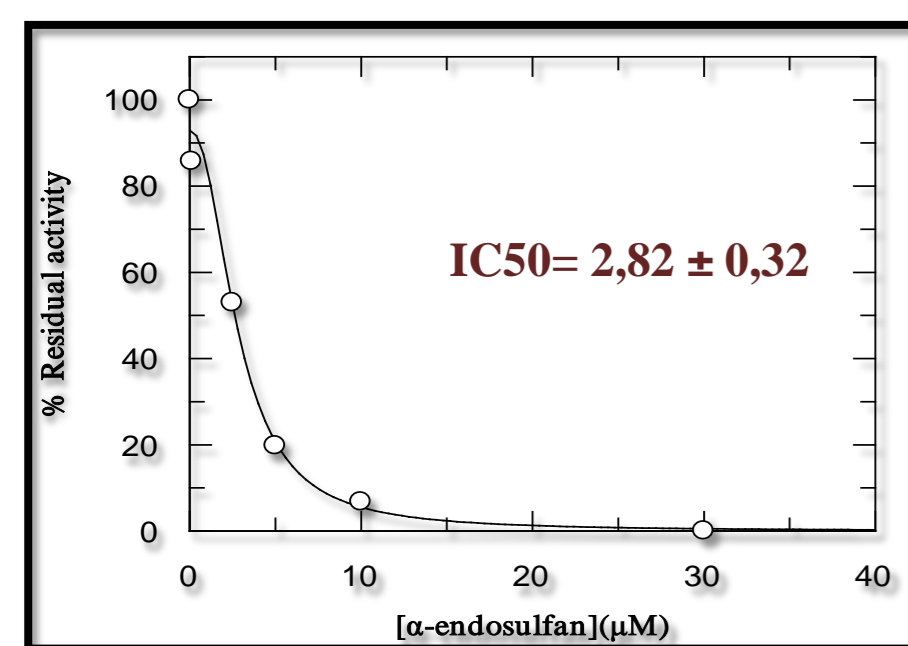


Fig.7. Dose response curve of α -endosulfan/immobilized F117I mutant enzyme.

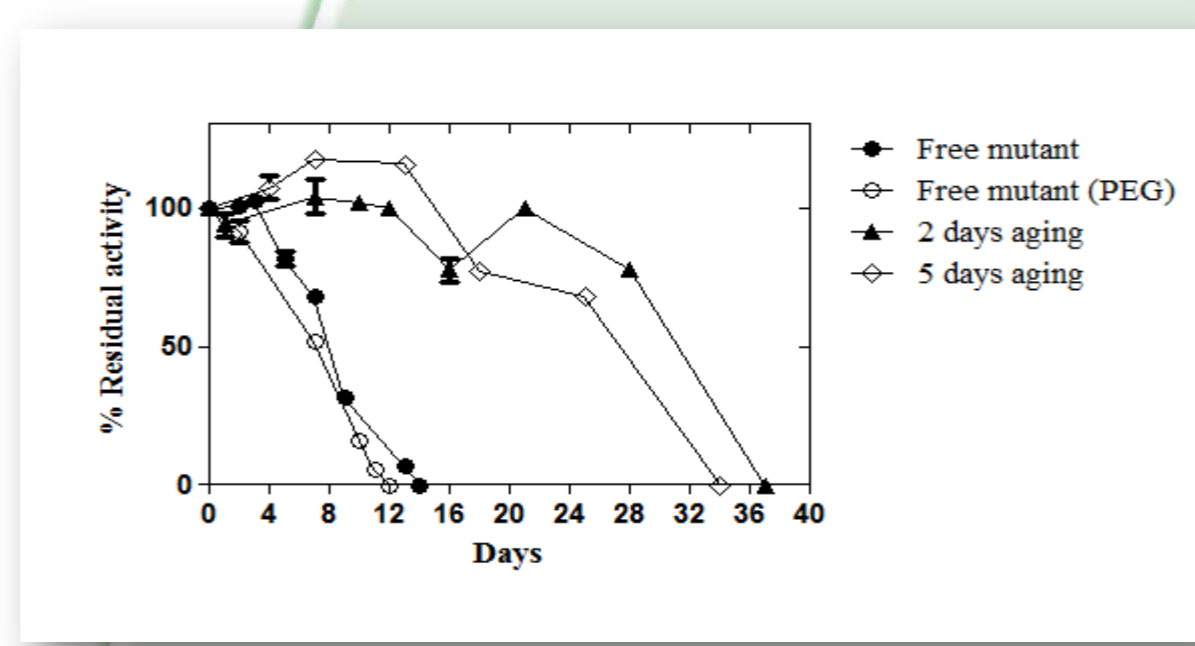


Fig.8. Stability of the free and immobilized F117I mutant enzyme.

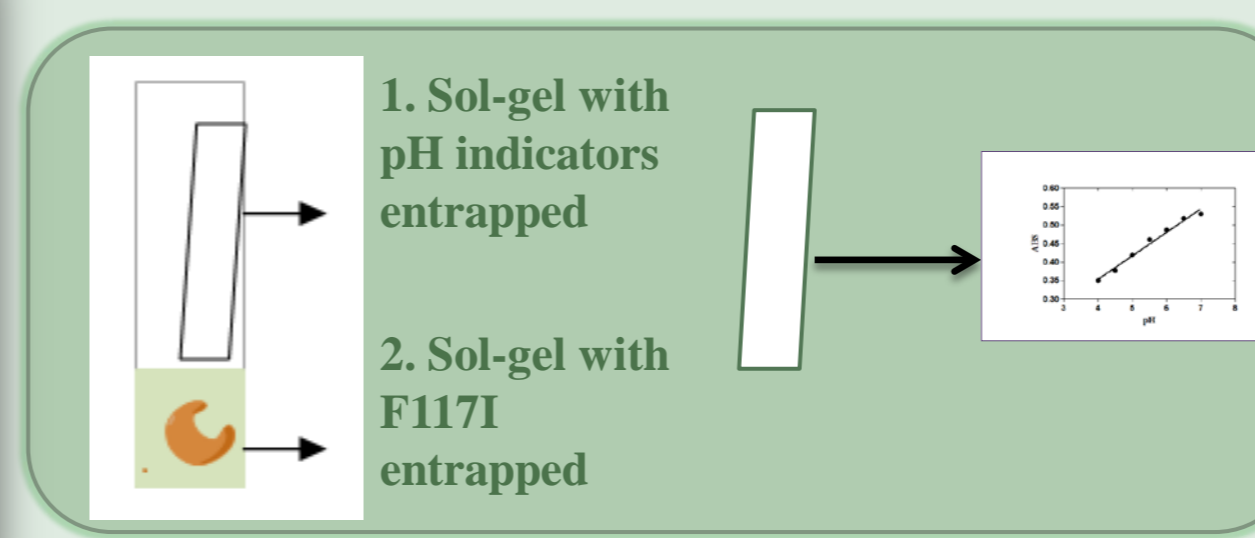


Fig.9. The biosensor used for the development of the analytical method for α -endosulfan determination

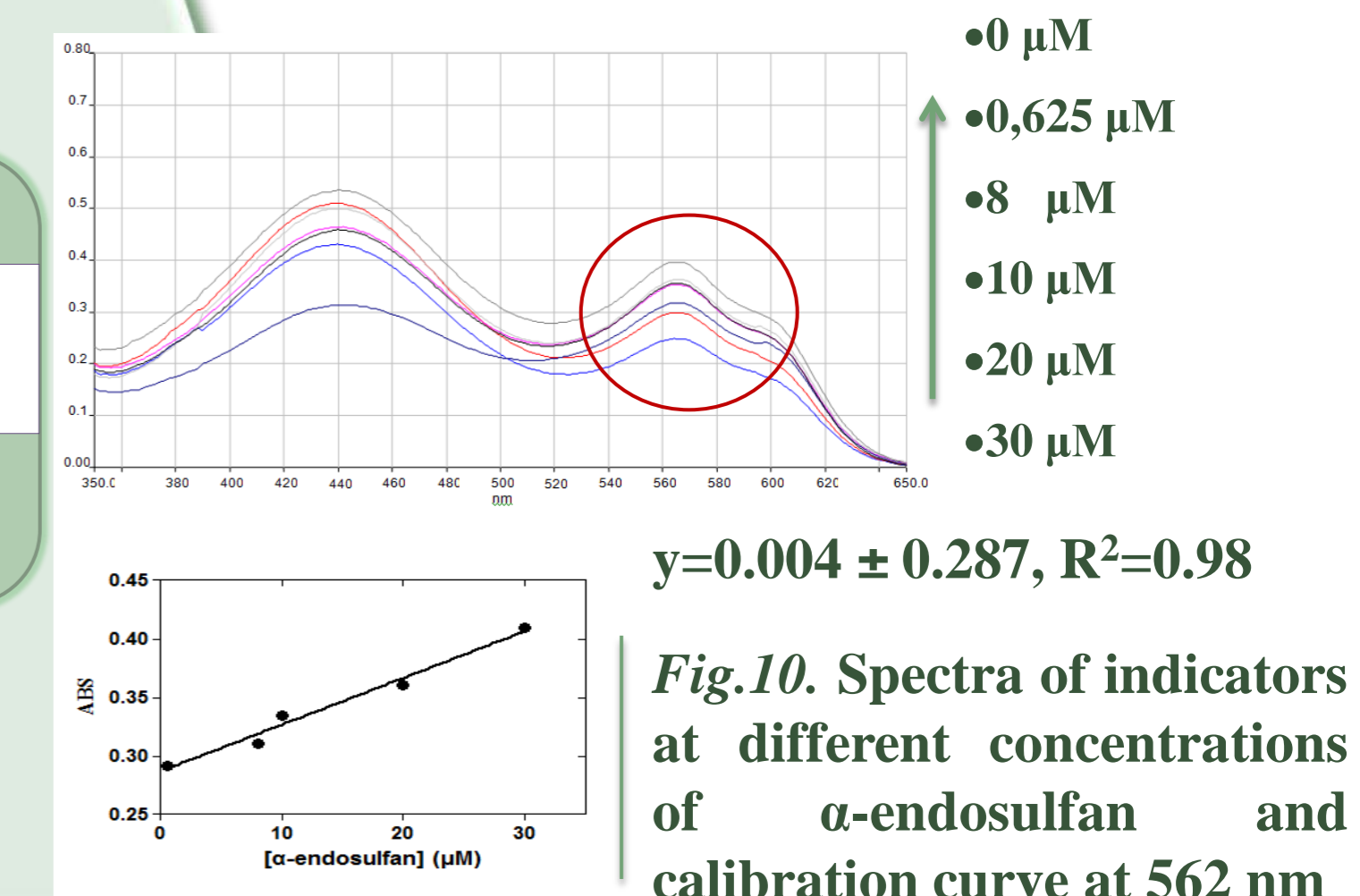


Fig.10. Spectra of indicators at different concentrations of α -endosulfan and calibration curve at 562 nm

Conclusion

Sol-gel encapsulated enzymes have been widely employed in the construction of biosensors using different detection methods such as electrochemical and optical methods. In the present work a novel GST was created using DNA shuffling and site-saturation mutagenesis. The mutant F117I displayed higher selectivity towards organochlorine insecticides. The enzyme was immobilized in alkoxysilane (TEOS/PTMOS) sol-gel system in the presence of the pH indicators (bromocresol purple and phenol red). The bioactive material exhibits linearity in the range of 0.625-30 μM α -endosulfan at 562 nm and high operational stability.

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