

N°1139 / PC

TOPIC(s) : Enzyme discovery and engineering / Industrial biocatalysis

Discovery of a novel ultra-thermostable Carbonic Anhydrase for enzymatic CO₂ sequestration using high-throughput metagenomic analysis

AUTHORS

KONSTANTINOS RIGKOS / NATIONAL HELLENIC RESEARCH FOUNDATION (N.H.R.F), VASILEOS KONSTANTINOU 48, ATHENS

GEORGIOS FILIS / NATIONAL HELLENIC RESEARCH FOUNDATION (N.H.R.F), VASILEOS KONSTANTINOU 48, ATHENS

PAVLOS SARIDIS / NATIONAL HELLENIC RESEARCH FOUNDATION (N.H.R.F), VASILEOS KONSTANTINOU 48, ATHENS

DIMITRA ZARAFETA / NATIONAL HELLENIC RESEARCH FOUNDATION (N.H.R.F), VASILEOS KONSTANTINOU 48, ATHENS

GEORGIOS SKRETAS / BIOMEDICAL SCIENCES RESEARCH CENTRE "ALEXANDER, FLEMING", FLEMINGK 34, VARI

Corresponding author : DIMITRA ZARAFETA / dzarafeta@eie.gr

PURPOSE OF THE ABSTRACT

Anthropogenic CO₂ emissions have been dramatically increasing for the past 30 years, an outcome that has led to an enhanced greenhouse effect, causing the Earth's temperature to rise at an accelerated rate. This phenomenon, known as Global Warming, is expected to have devastating effects on humanity if not addressed. This is well reflected in the commitment the European Commission made in 2019 to achieve net-zero emissions by 2050. In this landscape, the need for novel, low cost and sustainable CO₂ Capture and Storage Technologies (CCS) is urgent. Today, the go-to technology relies on the use of amines for the chemical separation of CO₂ from flue gas. Unfortunately, amines have a negative environmental impact due to high toxicity and high-energy consumption related to their regeneration process. Consequently, more eco-friendly technologies to mitigate the negative impact of CO₂ rising emissions are pursued by researchers worldwide. An alternative green technology for CO₂ sequestration involves the use of carbonic anhydrase (CA), an enzyme that catalyzes the reversible hydration of CO₂ to HCO₃⁻ and H⁺ ions. This catalytic activity is vital for many biological processes, including respiration and photosynthesis and because of this, CAs are found in all living organisms. Despite the fact that CAs are abundant in nature and can speed up CO₂ absorption kinetics in aqueous media, the harsh conditions present in industrial setups (high temperatures, alkaline pH and high salinity) compromise the stability of common enzymes. A promising approach to overcome this issue is the identification of thermostable enzymes which belong to the protein repertoire of extremophilic organisms. A bottleneck to this strategy is the fact that the protein space of such organisms is not accessible through classic microbiology, as 99% of them cannot be cultured. Metagenomic analysis can overcome this limitation by giving access to this huge, unexplored protein space, and driving the discovery of extremozymes of interest, such as thermostable CAs. In the present study, the discovery and characterization of a novel carbonic anhydrase is presented. High-throughput bioinformatic analysis of large, metagenomic data sets found in NIH's Sequence Read Archive (SRA) and in the Joint Genome Institute (JGI) open-access databases, resulted in the identification of an ultra-thermostable CA that was produced recombinantly in *E. coli* and studied. The novel CA-KR1 carbonic anhydrase, shares low similarity to known enzymes and exhibits very high thermostability, which, to our knowledge ranks it amongst the most thermostable CAs reported to date. This trait renders CA-KR1 a very strong candidate biocatalyst for enzymatic industrial CO₂ sequestration.

FIGURES

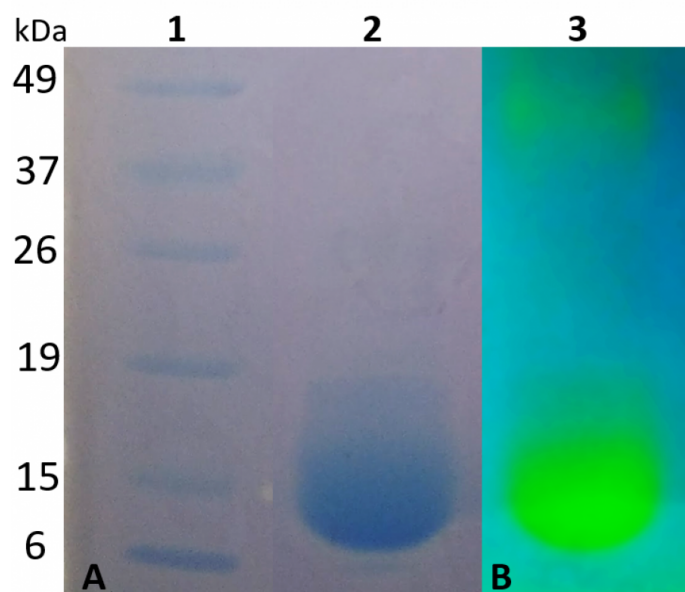


FIGURE 1

SDS-PAGE & Protonography analysis of CA-KR1 protein.

(A) Unboiled sample of purified CA-KR1 protein (Lane 2) and protein standard (Lane 1). (B) Protonography analysis of CA-KR1 (Lane 3). The color shift from blue to yellow indicates CA activity.

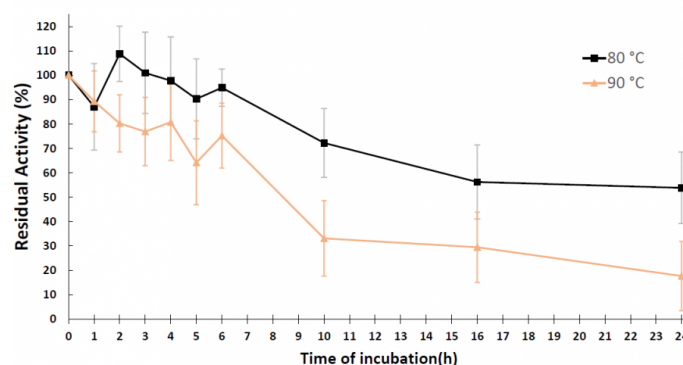


FIGURE 2

CA-KR1 Thermostability

The residual activity of CA-KR1 was measured following incubation of the enzyme at 80oC and 90oC for up to 24 h. The reported values correspond to the mean value from five independent experiments performed at least in triplicate +/- SD.

KEYWORDS

Enzyme discovery | CO₂ Capture | Carbonic Anhydrase | Bioinformatic Analysis

BIBLIOGRAPHY

- [1] Ozensoy Guler O, Capasso C, Supuran CT. J Enzyme Inhib Med Chem. 2016, 31(5):689-94.
- [2] Angeli A, Supuran CT. J Enzyme Inhib Med Chem. 2023, 38(1):2166503.
- [3] Hanifa M, Agarwal R, Sharma U, Thapliyal PC, Singh LP. J CO₂ Util. 2023, 67:102292.
- [4] Olabi AG, Wilberforce T, Sayed ET, Shehata N, Alami AH, Maghrabie HM, et al. Energies. 2022, 15(22):8639.
- [5] Dutcher B, Fan M, Russell AG. ACS Appl Mater Interfaces. 2015, 7(4):2137-48.
- [6] Anderson C, Harkin T, Ho M, Mumford K, Qader A, Stevens G, et al. Energy Procedia. 2013, 37:225-32.
- [7] Barzagli F, Giorgi C, Mani F, Peruzzini M. J CO₂ Util. 2017, 22:346-54.
- [8] Talekar S, Jo BH, Dordick JS, Kim J. Curr Opin Biotechnol. 2022, 74:230-40.