

## **2-oxoglutarate modulates the affinity of FurA for the** *ntcA* **promoter in** Anabaena sp. PCC 7120



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In cyanobacteria, 2-oxoglutarate (2-OG) is a metabolite whose function is to provide carbon skeletons that allow the incorporation of ammonium through the GS-GOGAT cycle, connecting both carbon and nitrogen metabolism. It has also been described as a signal molecule that reflects the cellular carbon/nitrogen balance, modulating the DNA-binding activity of the key regulator of nitrogen metabolism NtcA. Since FurA (Ferric uptake regulator) from the cyanobacterium Anabaena sp. PCC7120 is a global regulator that controls several cellular processes, including nitrogen metabolism, we sought to investigate whether this transcriptional regulator could also be sensing carbon/nitrogen balance via 2-OG.



Firstly, in order to **predict** if **2-OG** was able to **bind to FurA**, a model of FurA three-dimensional structure was built and docked with 2-OG.





**Figure 1.** Prediction of the 2-OG binding sites in the modelled structure of FurA from *Anabaena* sp. PCC7120. A. Cartoon representation of FurA (green) in complex with 2-OG (cyan). Residues involved in 2-OG binding are represented as sticks and by atom type (C, green; N, blue). B. Surface representation of FurA in complex with 2-OG (cyan).

These simulations **predicted** that **2-OG could bind to FurA** and revealed the existence of two potential 2-OG binding sites (Fig. 1). One of this sites involved Arg24 and Arg63 and the other involved Arg70, residues that were predicted to interact with 2-OG by **hydrogen bonds**.

## **3. 2-OG enhances DNA binding activity of FurA to** *ntcA* **promoter**

The effects of 2-OG on DNA binding activity of FurA were studied by

Figure 2. Analysis of the interaction between FurA from Anabaena sp. PCC7120 and 2-OG by ITC. **A.** Titration of FurA with 2-OG. **B.** Titration of FurA with 2-OG in the presence of 100 μM Mn<sup>2+</sup> and 1 mM DTT. **C.** Titration of FurA with glutamate in the presence of 100 µM Mn<sup>2+</sup> and 1 mM DTT.

Results showed that **FurA** was able to **interact** *in vitro* with **2-OG** but only in the **presence** of both **Mn<sup>2+</sup>** and **DTT**, the same conditions that FurA requires to bind to DNA (Fig 2A and 2B). Besides, this interaction was proven to be **specific**, since **FurA was not able to bind glutamate**, a molecule which a similar structure to 2-OG (Fig 2C).

## 4. Arg70 is involved in 2-OG binding

As docking simulations suggested two putative binding sites, we constructed two variants of FurA in which the residues that were predicted to interact with 2-OG were replaced by alanines.



**Figure 5.** EMSA assays showing the effect of 2-OG on FurA binding to the *isiB* promoter. The *pkn22* internal fragment was used as control for unspecific binding.



Figure 6. A. Analysis of the interaction between R24R63FurA and 2-OG by ITC. **B.** EMSA assays showing that R24R63FurA was unable to bind to the *ntcA* promoter. The pkn22 internal fragment was used as control for unspecific binding.

Figure 7. A. Analysis of the interaction between R70FurA and 2-OG by ITC. B. EMSA assays showing the effect of 2-OG on FurA binding to the *ntcA* promoter. The *pkn22* internal fragment was used as control for unspecific binding.

ITC and EMSA assays showed that FurA variant lacking both Arg24 and Arg63 (R24R63) was able to bind to 2-OG but unable to bind to DNA (Fig. 6). On the contrary, FurA variant lacking Arg70 (R70) was unable to **interact with 2-OG** and this metabolite **had no effects on its DNA binding** activity (Fig 7), revealing that Arg70 was involved in 2-OG binding.

As it is showed in Fig 4, 2-OG was able to **mildly enhance** the **binding** of FurA to the ntcA promoter. However no effects were observed when this experiment was carried out with the flavodoxin (*isiB*) promoter region (Fig 5).



>A 2-OG binding site in FurA from Anabaena sp. PCC7120 has been identified by combining **bioinformatic predictions** and **experimental procedures**.

> FurA from Anabaena sp. PCC 7120 probably acts as a sensor of carbon/nitrogen **balance**, since it **binds 2-OG** and this interaction influences its **DNA binding activity** 





If you have any questions or are interested in our work please do not hesitate to contact me: jguio@unizar.es • Follow us on Twitter! @cyanofur Funding for this study was from 'Ministerio de Ciencia, Innovación y Universidades' (PID2019-104889GB-I00 to M.F.F. and grant FPU2018/03619 to J.G.)