

UvrA and UvrC subunits of the *Mycobacterium tuberculosis* UvrABC excinuclease interact independently of UvrB and DNA

Manoj Thakur, Sugith Badugu and Kalappa Muniyappa 

Department of Biochemistry, Indian Institute of Science, Bangalore, India

Correspondence

K. Muniyappa, Department of Biochemistry,
Indian Institute of Science, Bangalore
560012, India
Tel: +91 80 2293 2235/2360 0278
E-mail: kmbc@iisc.ac.in

(Received 26 August 2019, revised 31
October 2019, accepted 31 October 2019)

doi:10.1002/1873-3468.13671

Edited by Peter Brzezinski

The UvrABC excinuclease plays a vital role in bacterial nucleotide excision repair. While UvrA and UvrB subunits associate to form a UvrA₂B₂ complex, interaction between UvrA and UvrC has not been demonstrated or quantified in any bacterial species. Here, using *Mycobacterium tuberculosis* UvrA (MtUvrA), UvrB (MtUvrB) and UvrC (MtUvrC) subunits, we show that MtUvrA binds to MtUvrB and equally well to MtUvrC with submicromolar affinity. Furthermore, MtUvrA forms a complex with MtUvrC both *in vivo* and *in vitro*, independently of DNA and UvrB. Collectively, these findings reveal new insights into the pairwise relationships between the subunits of the UvrABC incision complex.

Keywords: *M. tuberculosis*; nucleotide excision repair; protein-protein interaction; UvrABC excinuclease; UvrA-UvrC complex

In bacteria, nucleotide excision repair (NER) is a vital DNA repair pathway that acts on a wide variety of bulky DNA adducts such as cyclobutane pyrimidine dimers, alkylation adducts and cisplatin interstrand cross-links [1,2]. In the well-studied *Escherichia coli* paradigm, the multistep NER pathway begins with the binding of UvrA₂B₂ heterotrimer to the damaged region [3–11]. Subsequently, the dissociation of UvrA₂ from UvrB₂ enables the latter to perform damage verification, recruit UvrC and assemble the pre-incision complex [3,11,12]. The ATP hydrolysis by UvrA₂ facilitates the release of UvrA dimer from DNA [13,14]. After incision at either side flanking the DNA lesion by UvrC within the UvrB-UvrC-DNA complex, the postincision complex comprising of 12- to 13-mer oligonucleotide and UvrB-UvrC is removed by UvrD and Pol I [15–20]. The resulting gap is filled in by DNA polymerase I,

and the broken strands are sealed by DNA ligase [21].

Although NER is one of the most extensively studied DNA repair processes in *E. coli*, much remains unknown about the formation and disassembly of NER repair machinery [1,2,22]. Additionally, the mechanisms underlying the coordination between the various proteins/enzymes involved in the NER pathway are not fully understood [1,2]. The translation of *in vitro* models of the *E. coli* NER pathway to an actual *in vivo* context remains challenging [10,11]. Furthermore, to what extent the findings from the *E. coli* NER pathway can be extrapolated and applied to other strains and species, and during different physiological conditions, remains to be shown. While the enzymes of the NER pathway exist in *Mycobacterium tuberculosis* [23], very little is known about their catalytic and regulatory mechanisms. Notwithstanding,

Abbreviations

MtUvrA, *M. tuberculosis* UvrA; MtUvrB, *M. tuberculosis* UvrB; MtUvrC, *M. tuberculosis* UvrC; NER, nucleotide excision repair; RU, response units; SEC, size-exclusion chromatography; SPR, surface plasmon resonance; ssDNA, single-stranded DNA; Y2H, yeast two-hybrid system.