



STRUCTURAL BIOLOGY  
COMMUNICATIONS

**Volume 72 (2016)**

**Supporting information for article:**

**Crystal structure of FhuD at 1.6 Å resolution: a ferrichrome-binding protein from the animal and human pathogen *Staphylococcus pseudintermedius***

**Francesca Abate, Roberta Cozzi, Martina Maritan, Paola Lo Surdo, Domenico Maione, Enrico Malito and Matthew James Bottomley**

## S1. Materials and Methods

### S1.1. Analytical size-exclusion chromatography

Purified FhuD samples at concentrations of 0.5 mg/mL, 0.75 mg/mL and 1.5 mg/mL (i.e. up to approximately 50  $\mu$ M concentration) were loaded as 20  $\mu$ l samples onto a Superdex-75 resin, 5/150 analytical SEC column (GE Healthcare) equilibrated at room temperature (18-26  $^{\circ}$ C) in buffer containing 20 mM Tris-HCl, 150 mM NaCl, pH 8.0, with flow rate 0.15 mL/min. At all three concentrations tested, the elution profile revealed a single symmetrical peak, with a maximum at 11.1 min. The elution time was used to obtain an apparent molecular weight by running standard proteins (Bio-Rad markers ranging from 1.35-670 kDa) on the same column and under the same conditions as used for FhuD.

### S1.2. Differential scanning calorimetry

The thermal stability of FhuD was assessed by differential scanning calorimetry (DSC) using a MicroCal VP-Capillary DSC instrument (GE Healthcare). Purified FhuD was prepared at 10  $\mu$ M concentration in phosphate buffered saline pH 7.4 (PBS). DSC experiments were performed in the presence or absence of 200  $\mu$ M ferrichrome ( $C_{27}H_{42}O_{12}N_9Fe$ ; MW 740; obtained from EMC microcollections GmbH). The DSC temperature scan ranged from 10  $^{\circ}$ C to 110  $^{\circ}$ C, with a thermal ramping rate of 200  $^{\circ}$ C per hour and a 4 second filter period. Data were analyzed by subtraction of the reference data for a sample containing only buffer (e.g. PBS only, or PBS + 200  $\mu$ M ferrichrome), using the Origin 7 software.

### S1.3. Surface plasmon resonance

The binding of ferrichrome to FhuD was assessed by surface plasmon resonance (SPR) using a Biacore T200 instrument equilibrated at 25  $^{\circ}$ C in HBS-EP running buffer (GE Healthcare). Experiments were performed using a CM5 sensor chip with covalent immobilization via amine coupling of FhuD (prepared at 50  $\mu$ g/mL in 10 mM sodium acetate buffer pH 4.5, reaching a surface density of approximately 2000 RU), essentially as described previously ([Mariotti \*et al.\*, 2013](#)). The analytes were tested using injections at 1  $\mu$ M concentration.