

# Structure Factor Check

## 2FBW

Title: AVIAN RESPIRATORY COMPLEX II WITH CARBOXIN BOUND  
Date: 10-DEC-05  
PDB code: 2FBW

### Crystal

Cell parameters:

a: 118.70 A b: 200.75 A c: 67.63 A  
 $\alpha$ : 90.00  $\beta$ : 90.06  $\gamma$ : 90.00

Space group: P 1 21 1

### Structure Factors

#### Input

Nominal resolution range: 64.1 – 2.05 A  
Reflections in file: 166892  
Unique reflections above 0: 166892  
above 1 $\sigma$ : 166666  
above 3 $\sigma$ : 153945

#### SFCHECK

Nominal resolution range: 64.1 – 2.10 A  
05max. from input data, min. from author05  
Used reflections: 162211  
Reflections out of resolution: 4681  
Completeness: 88.3 %  
R\_stand(F) =  $\langle \sigma(F) \rangle / \langle F \rangle$  : 0.035  
Anisotropic distribution of Structure Factors  
ratio of eigen values: 0.7481 1.0000 0.6648  
B\_overall (by Patterson): 30.A<sup>2</sup>  
Optical resolution: 1.63 A  
Expected opt. resol. for complete data set: 1.63 A  
Estimated minimal error: 0.019 A

### Model

19494 atoms (2000 water molecules)  
Number of chains: 51  
Volume not occupied by model: 46.8 %  
<B> (for atomic model): 30.4 A<sup>2</sup>  
 $\sigma(B)$ : 14.95 A<sup>2</sup>  
Matthews coefficient: 2.95  
Corresponding solvent % : 58.01

### Model vs. Structure Factors

R-factor for all reflections: 0.193  
Correlation factor: 0.935  
R-factor: 0.198  
for F > 2.0  $\sigma$   
nom. resolution range: 64.09 – 2.10A  
reflections used: 162012  
Rfree: 0.235  
Nfree: 8000  
R-factor without free-refl.: 0.197  
Non free-reflections: 154012  
<u> (error in coords by Luzzati plot): 0.232 A  
Estimated maximal error: 0.111 A  
DPI: 0.166 A

### Refinement

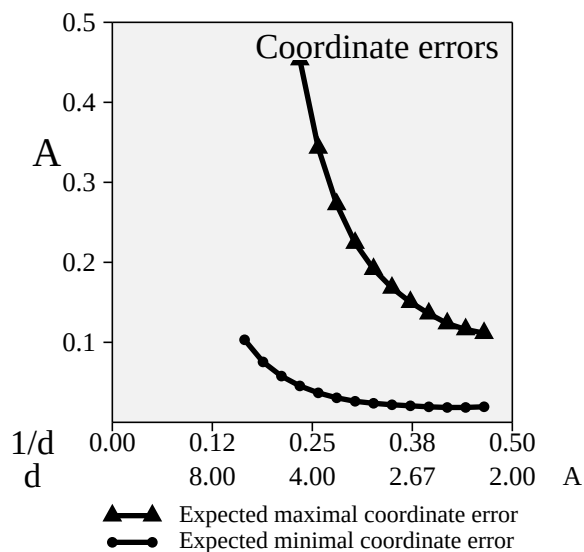
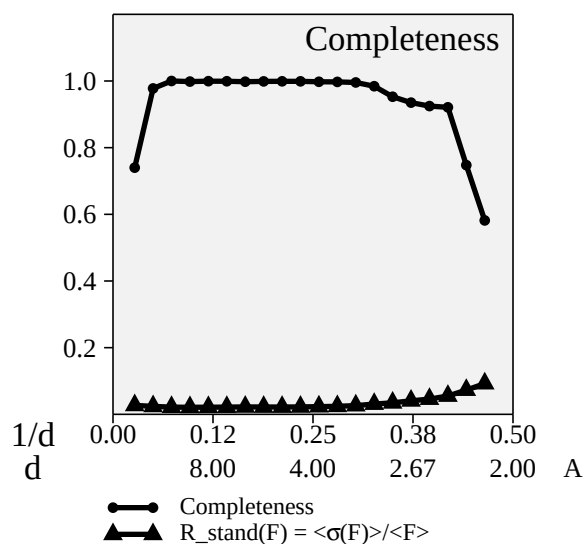
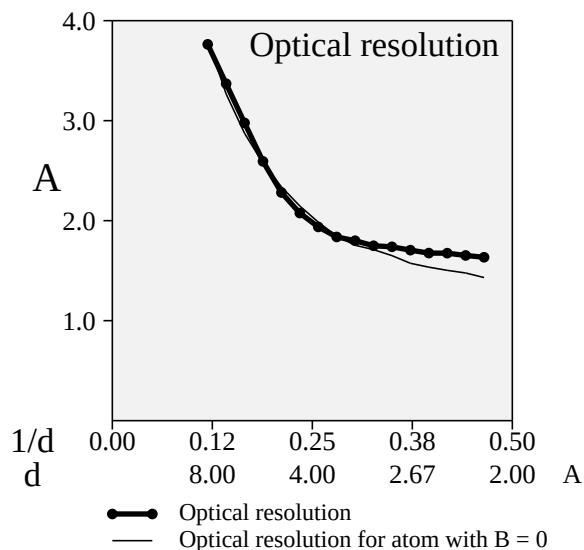
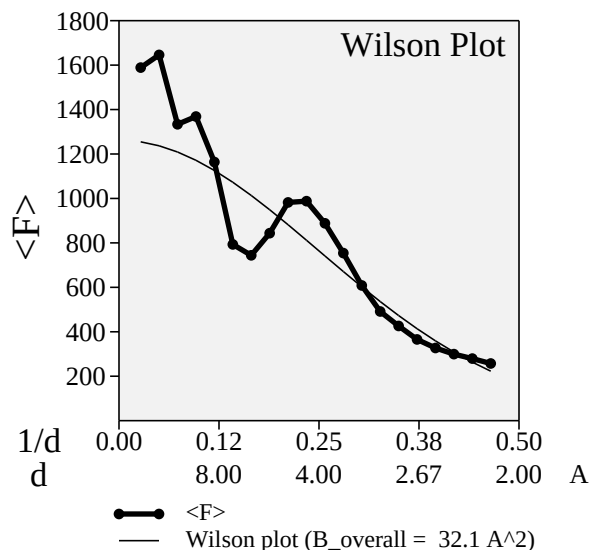
Program: CNS 1.1  
Nominal resolution range: 64.1 – 2.10 A  
Reported R-factor: 0.187  
Number of reflections used: 162208  
Reported Rfree: 0.23  
Sigma cut-off: N.A.

#### Scaling

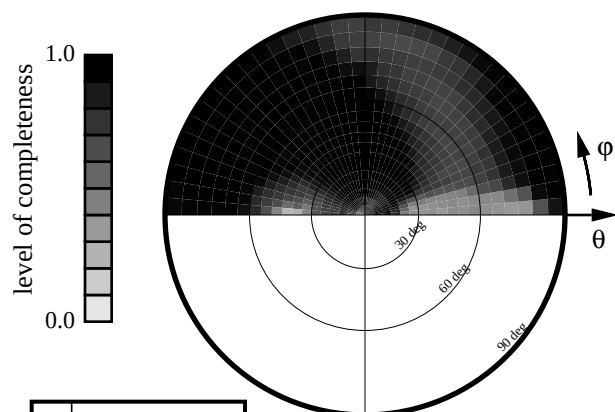
Scale: 1.056  
Bdiff: -0.67  
Anisothermal Scaling (Beta):  
-0.3776 2.1252 -0.5333 0.0000 -0.2013 0.0000  
Solvent correction – Ks,Bs: 0.900 492.305

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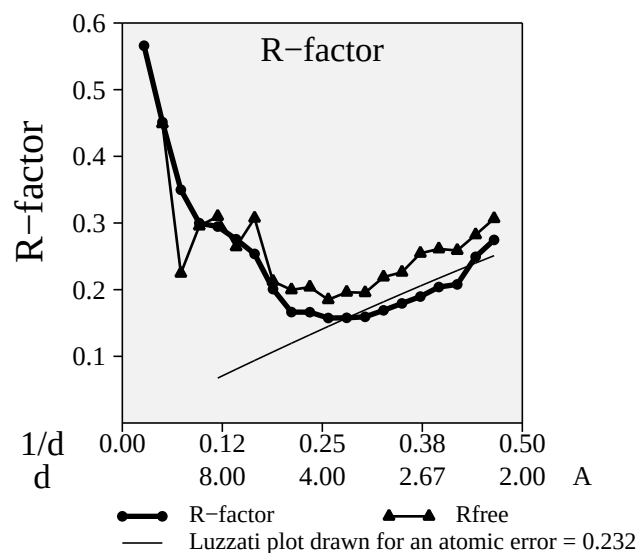


Stereographic projection of the averaged radial completeness



|   | $\theta$ | $\phi$ |
|---|----------|--------|
| h | 89.94    | 0.00   |
| k | 90.00    | 90.00  |
| l | 0.00     | 0.00   |

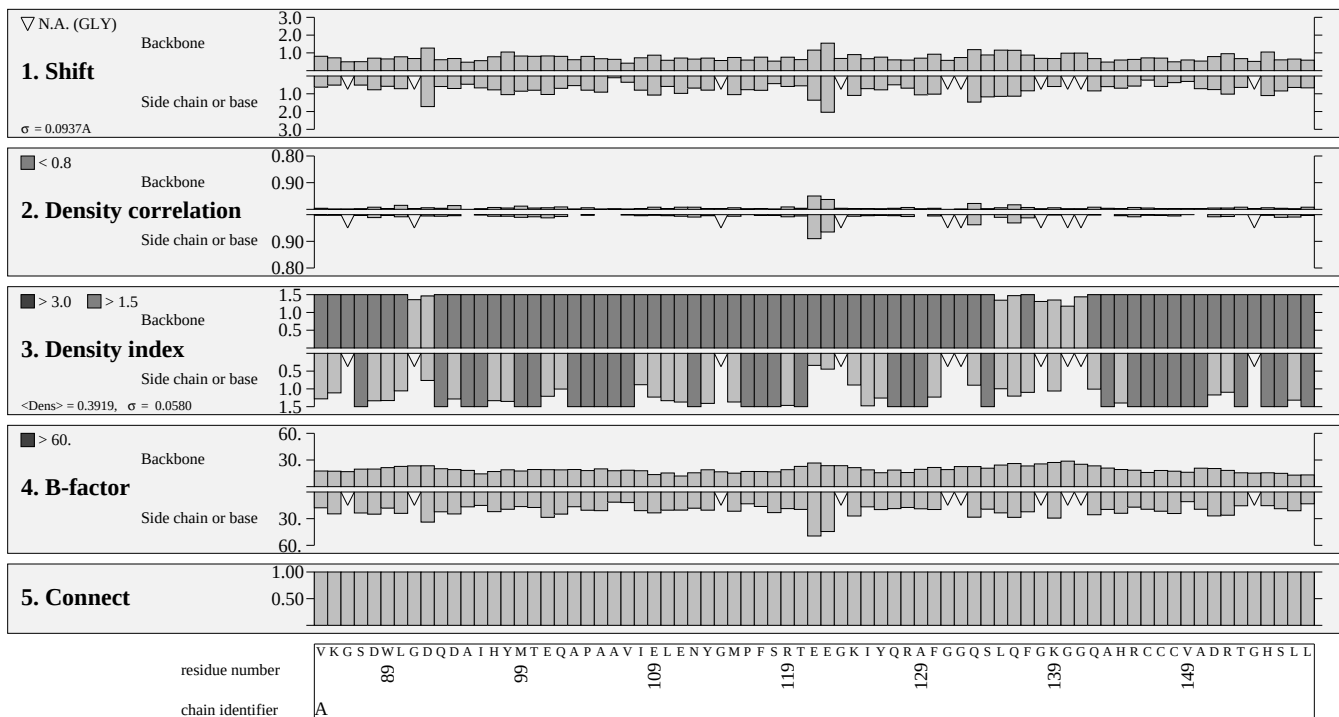
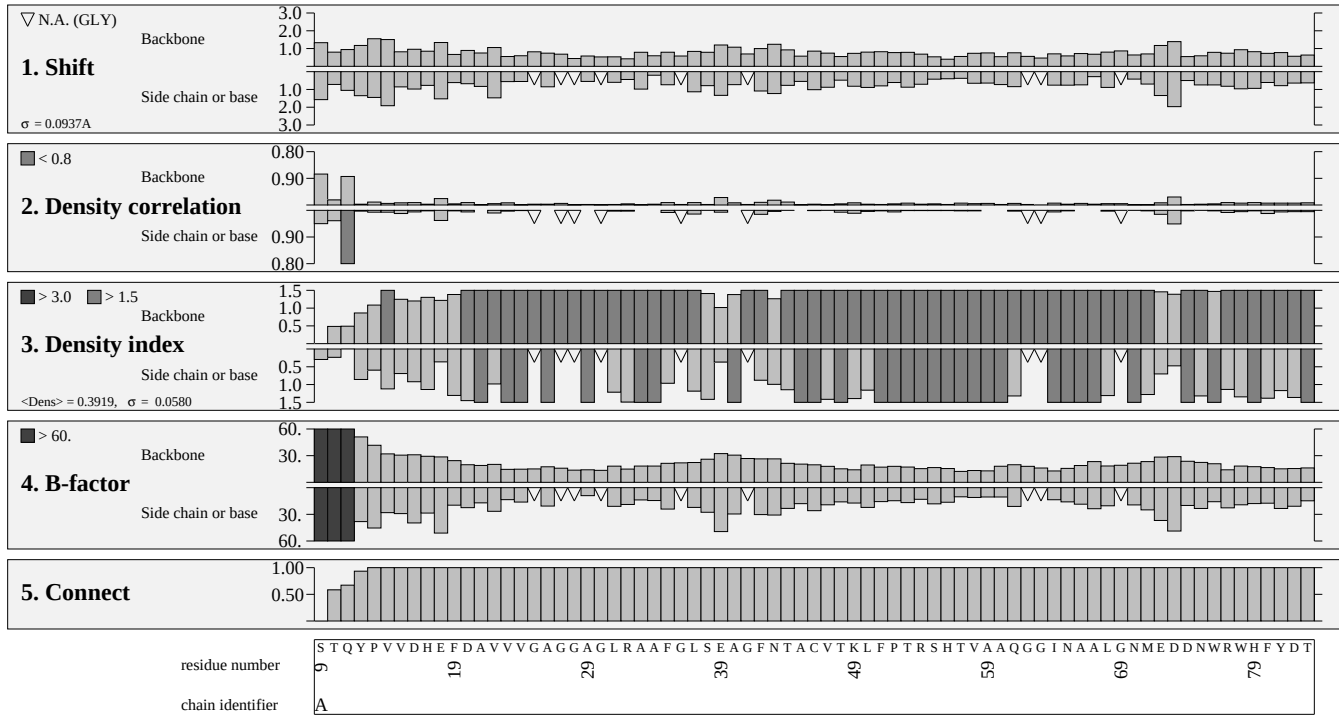
polar coordinates of the crystallographical axes



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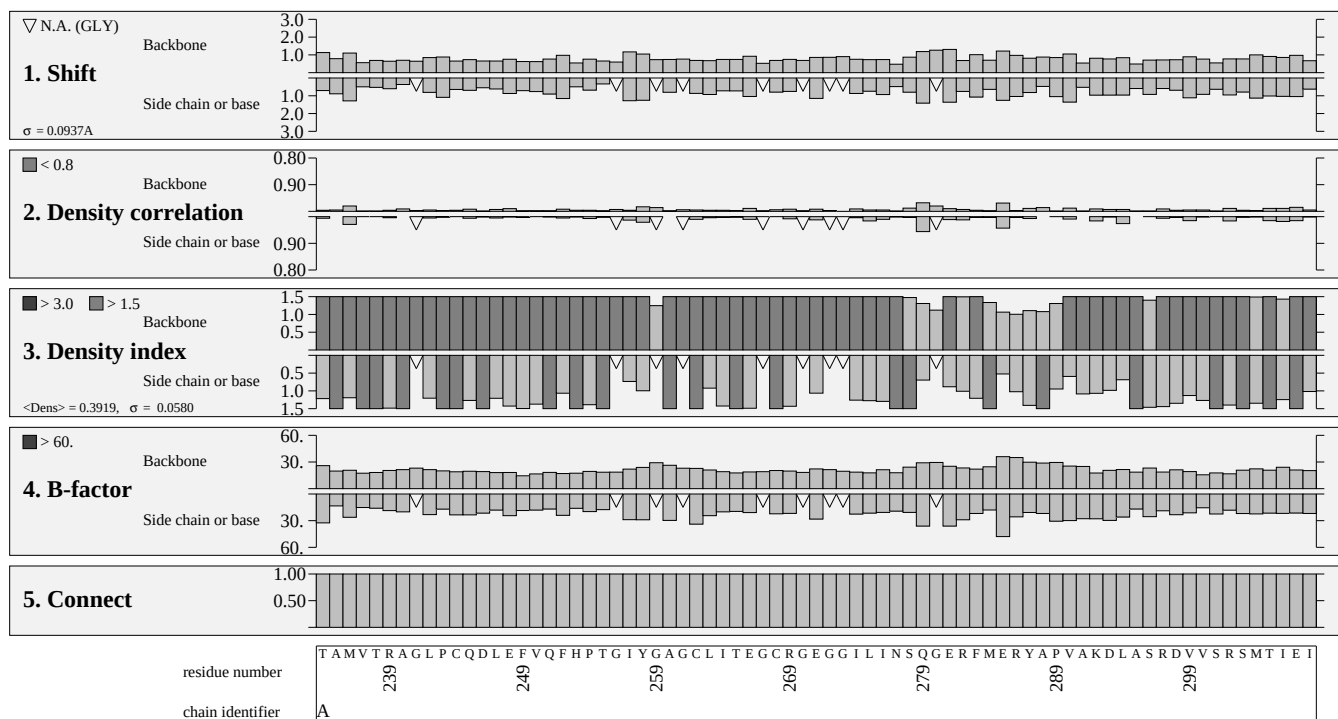
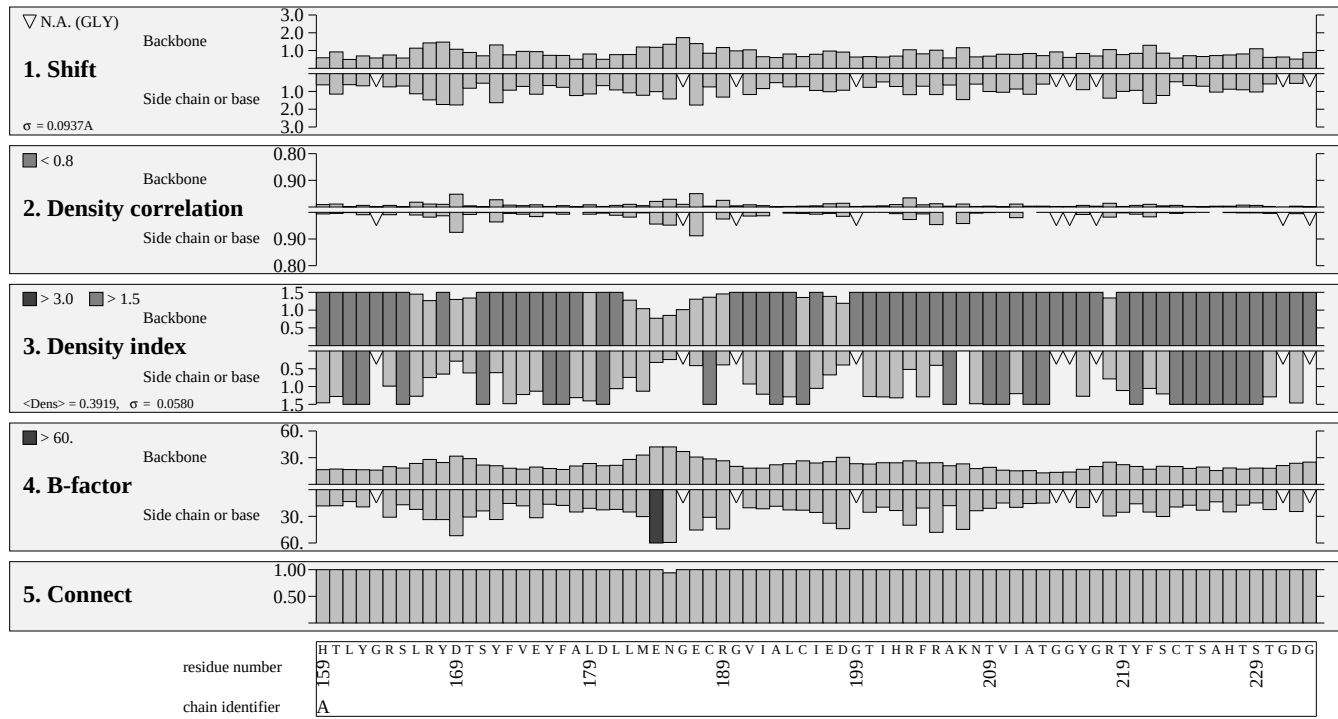
### Local estimation



# Structure Factor Check

## 2FBW

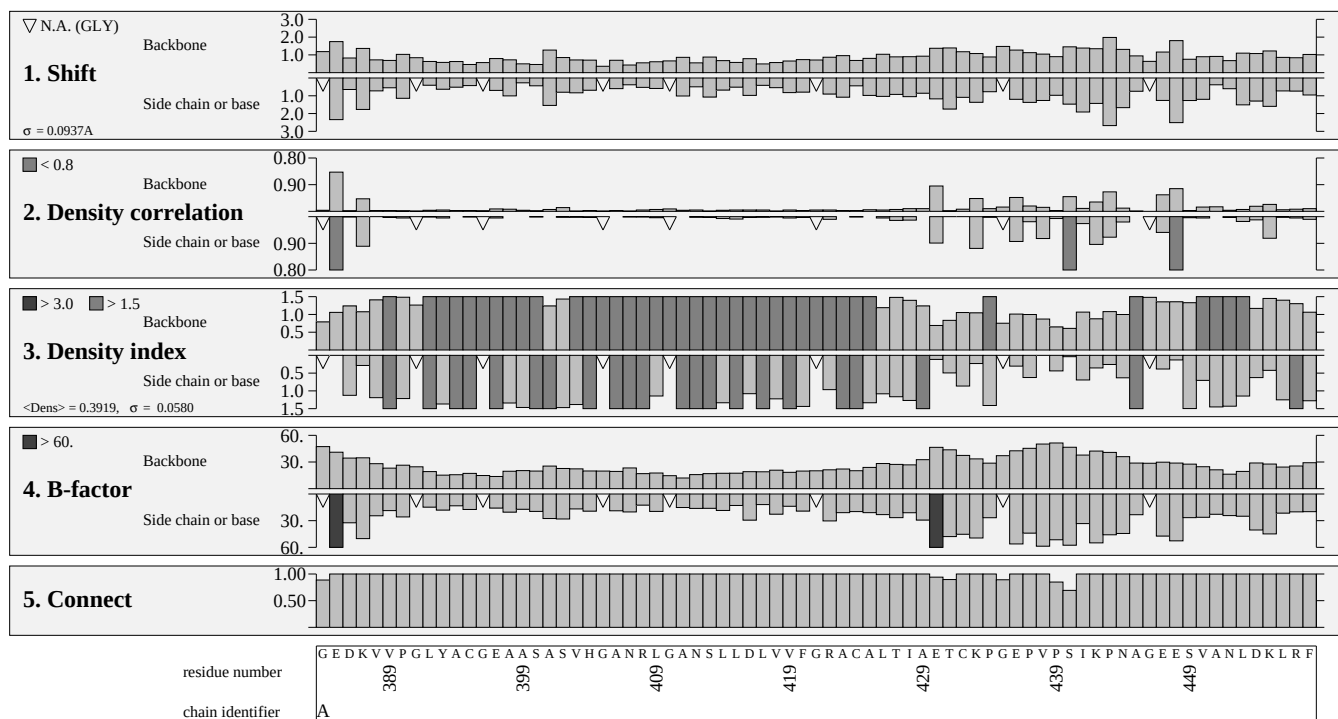
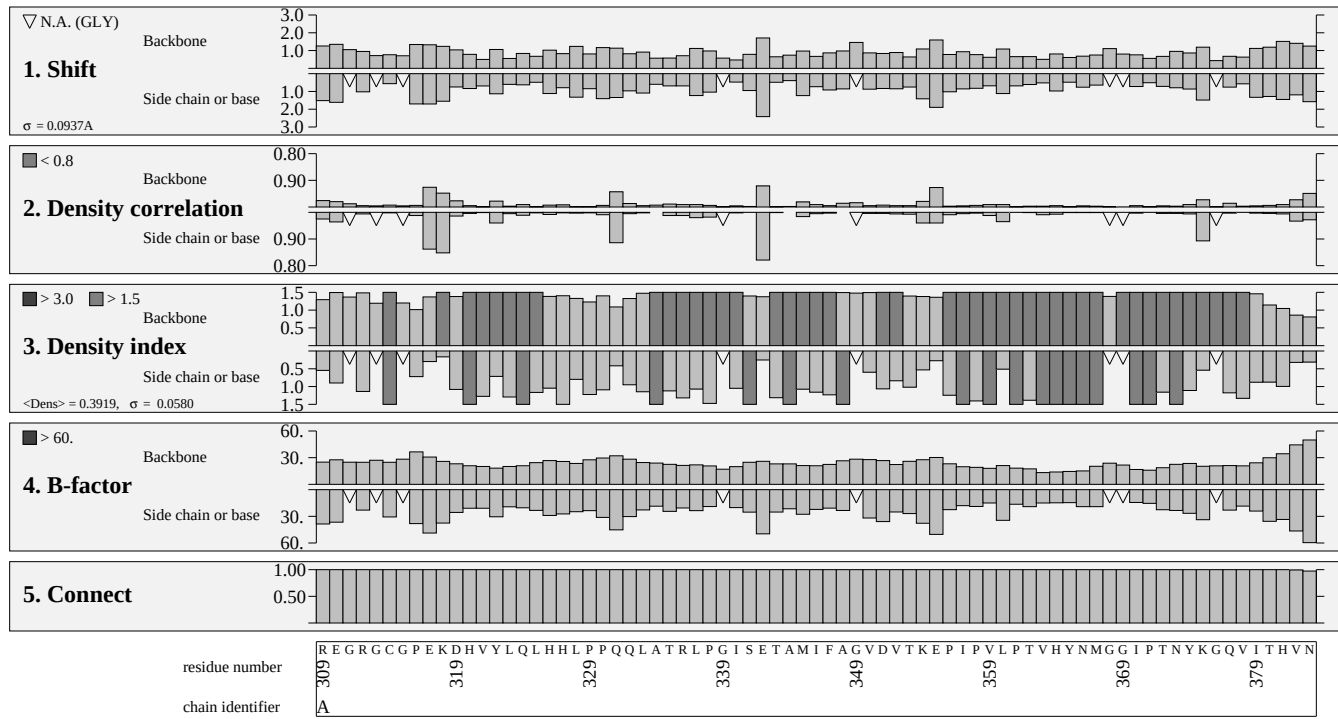
### Local estimation (2)



# Structure Factor Check

## 2FBW

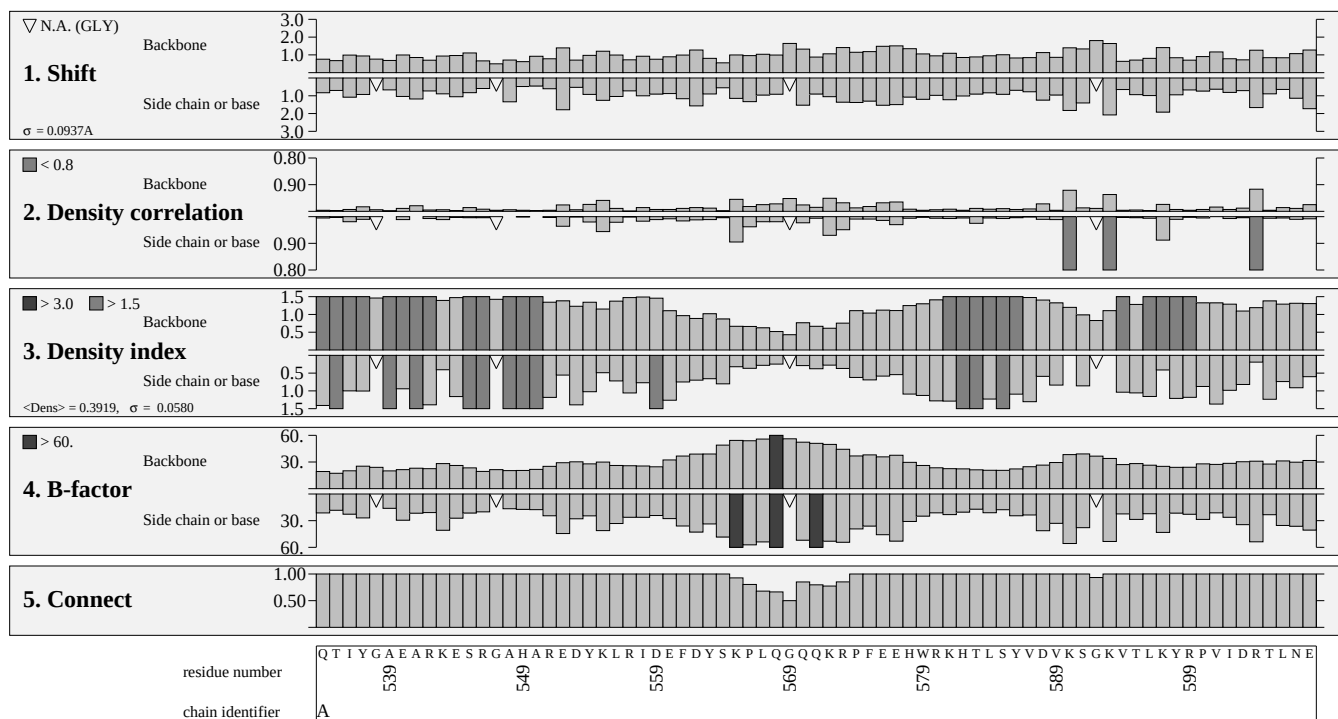
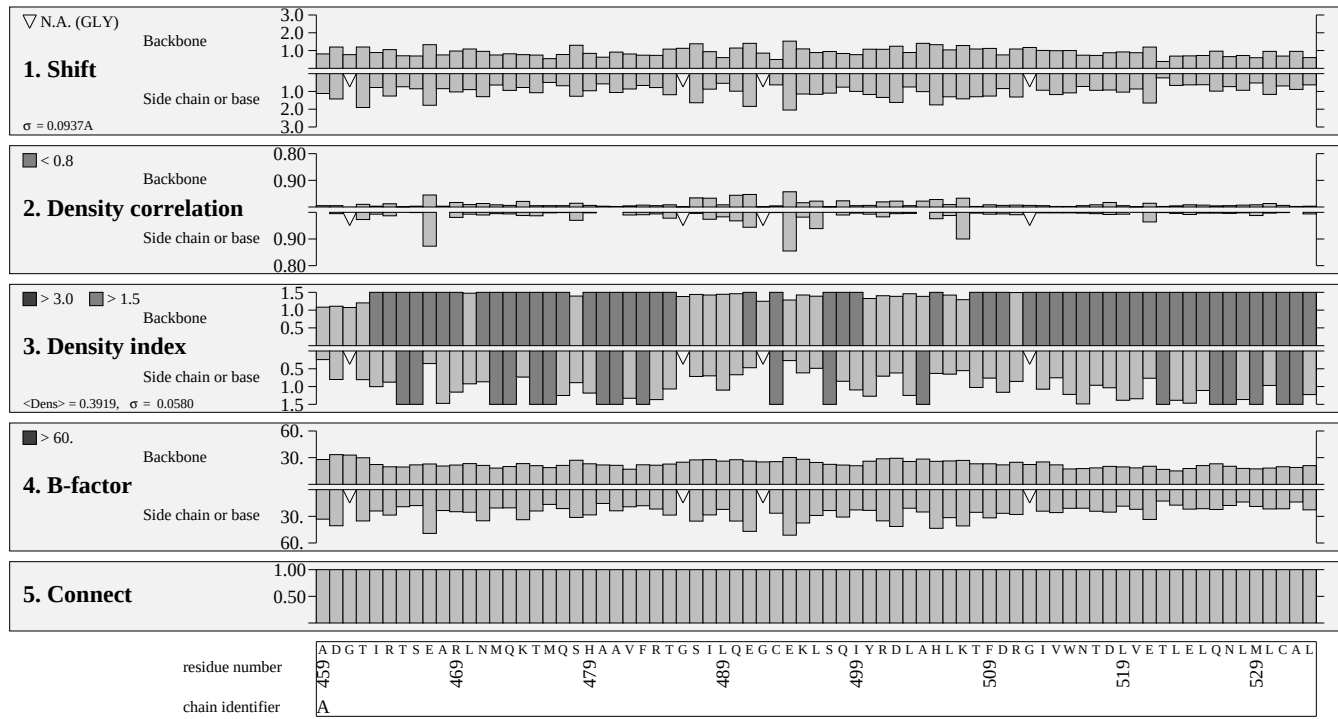
### Local estimation (3)



# Structure Factor Check

## 2FBW

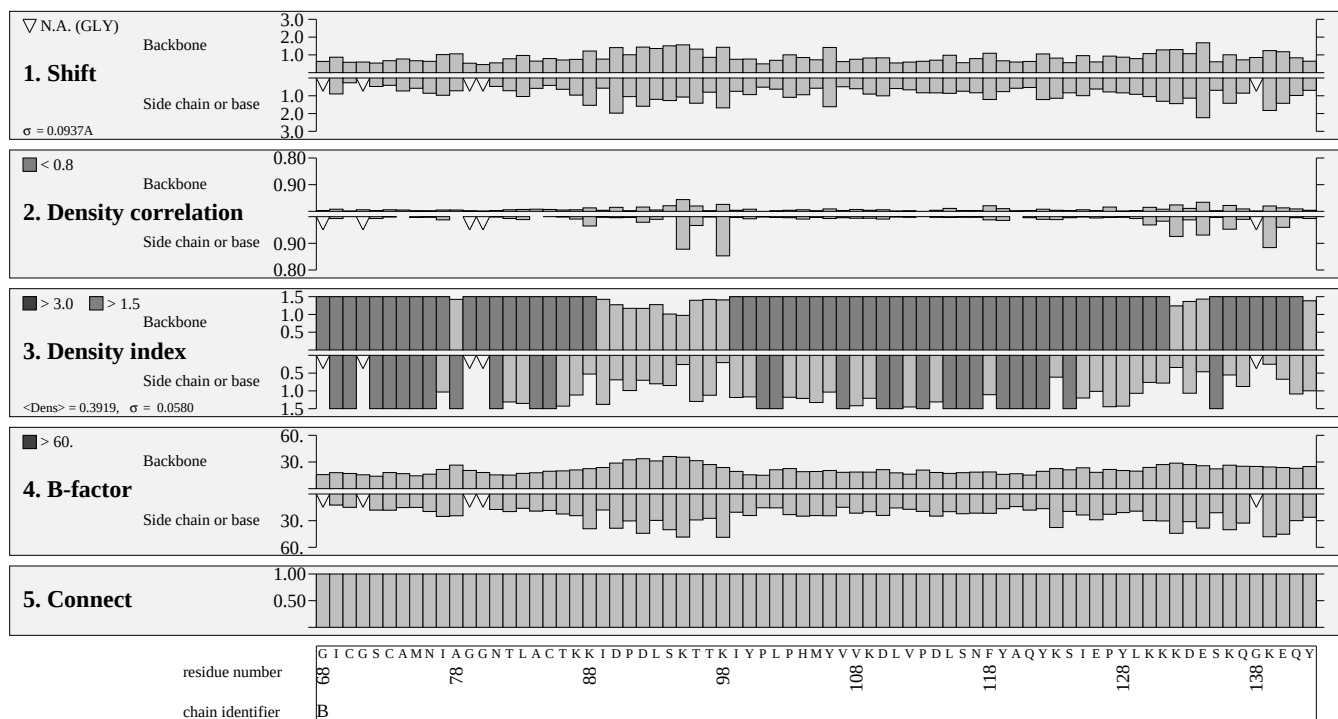
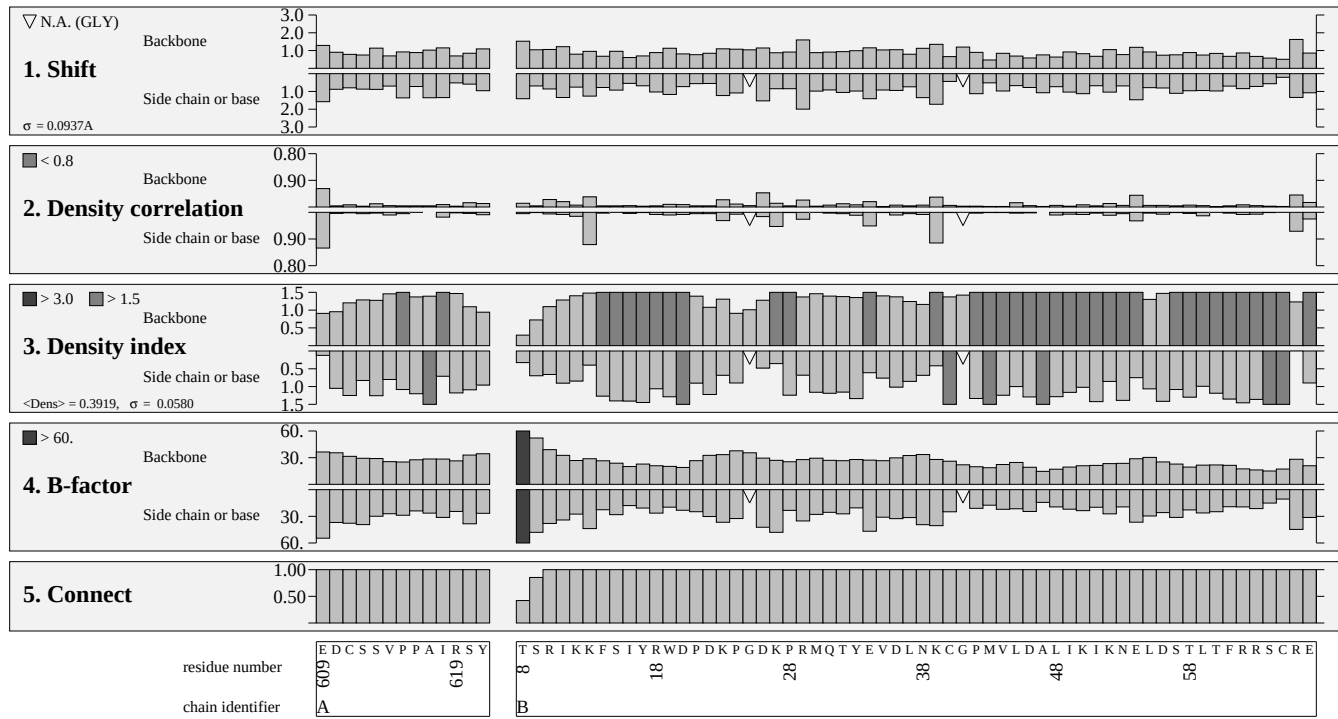
### Local estimation (4)



# Structure Factor Check

## 2FBW

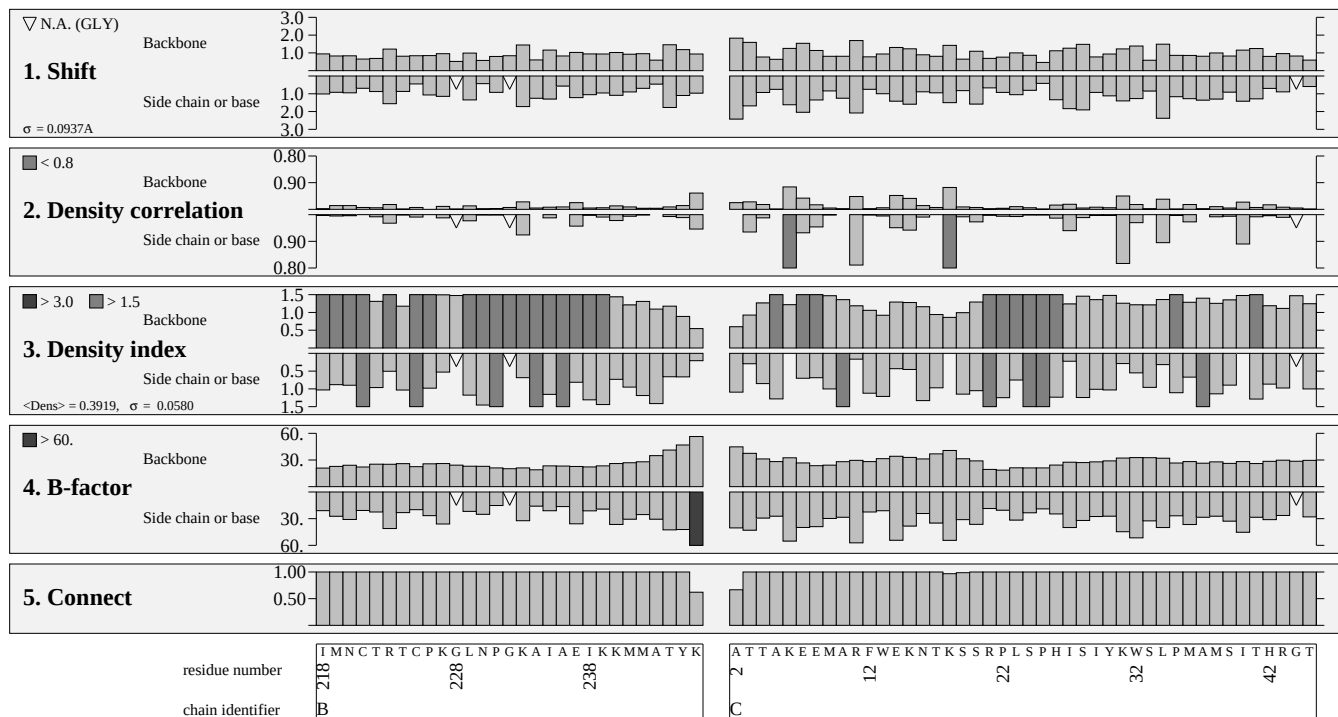
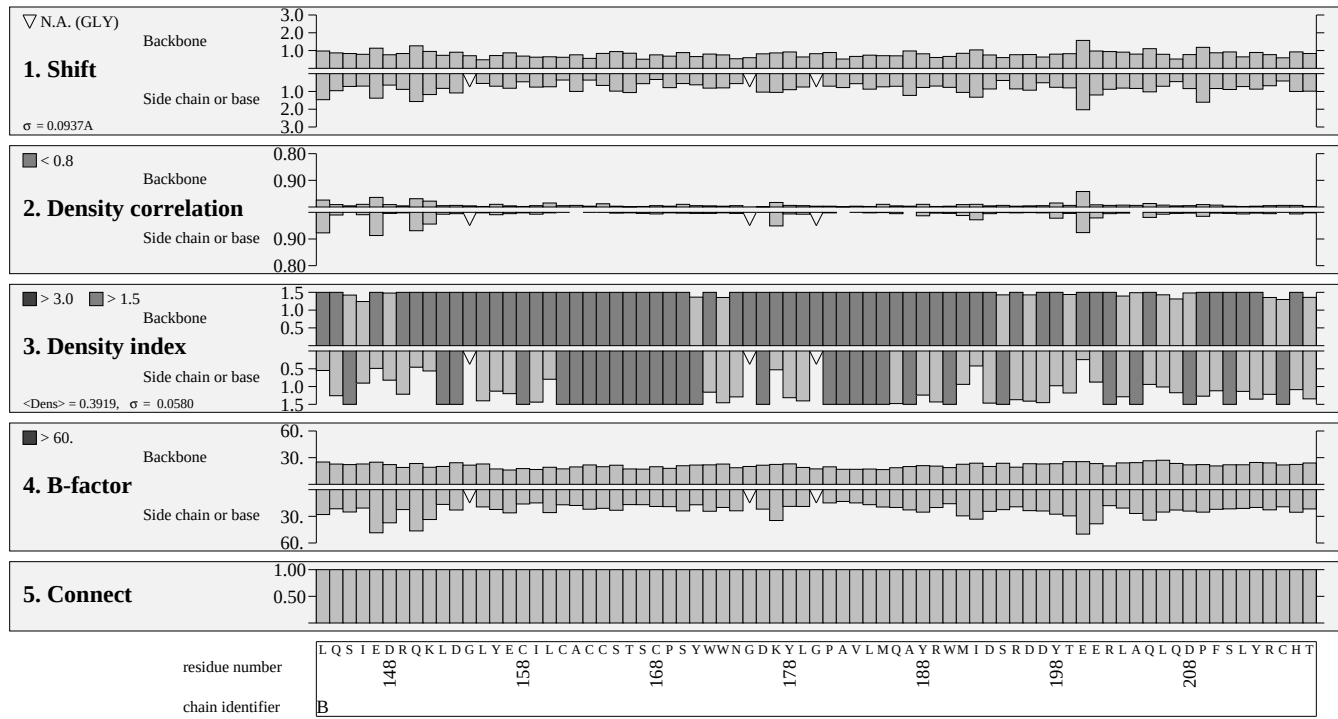
### Local estimation (5)



# Structure Factor Check

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### Local estimation (6)

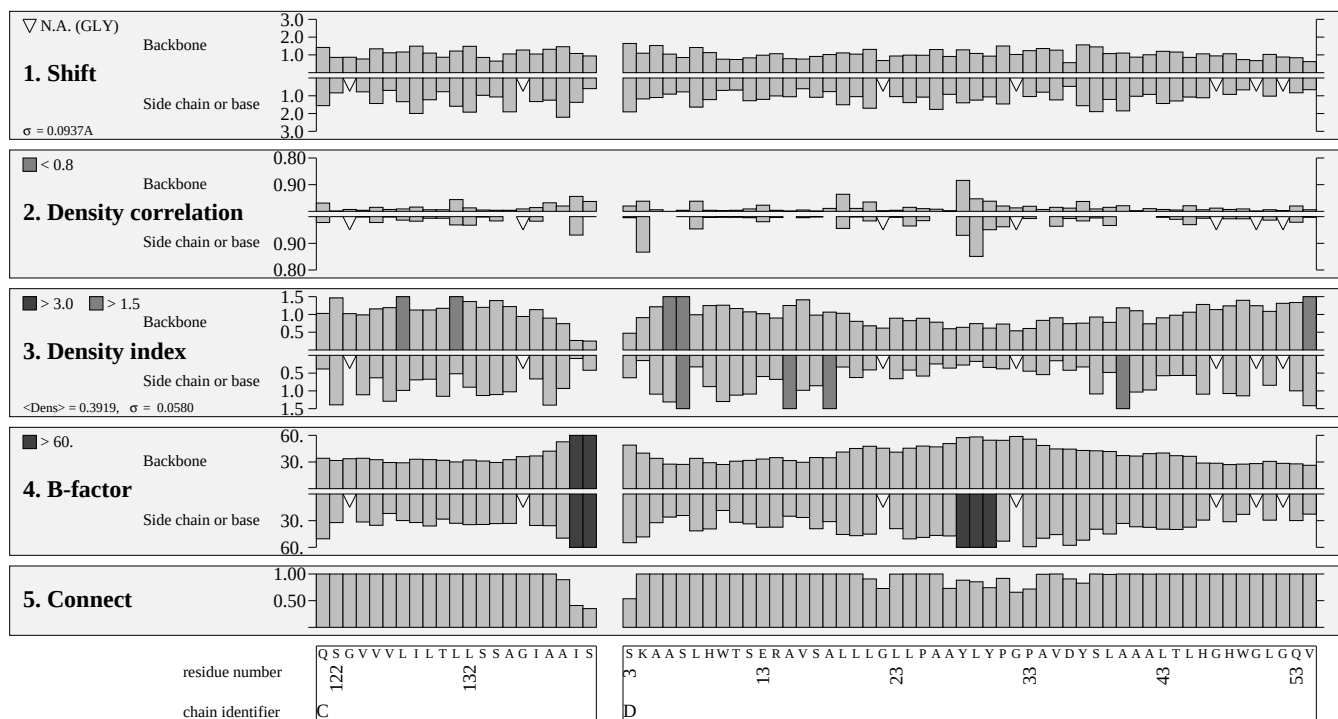
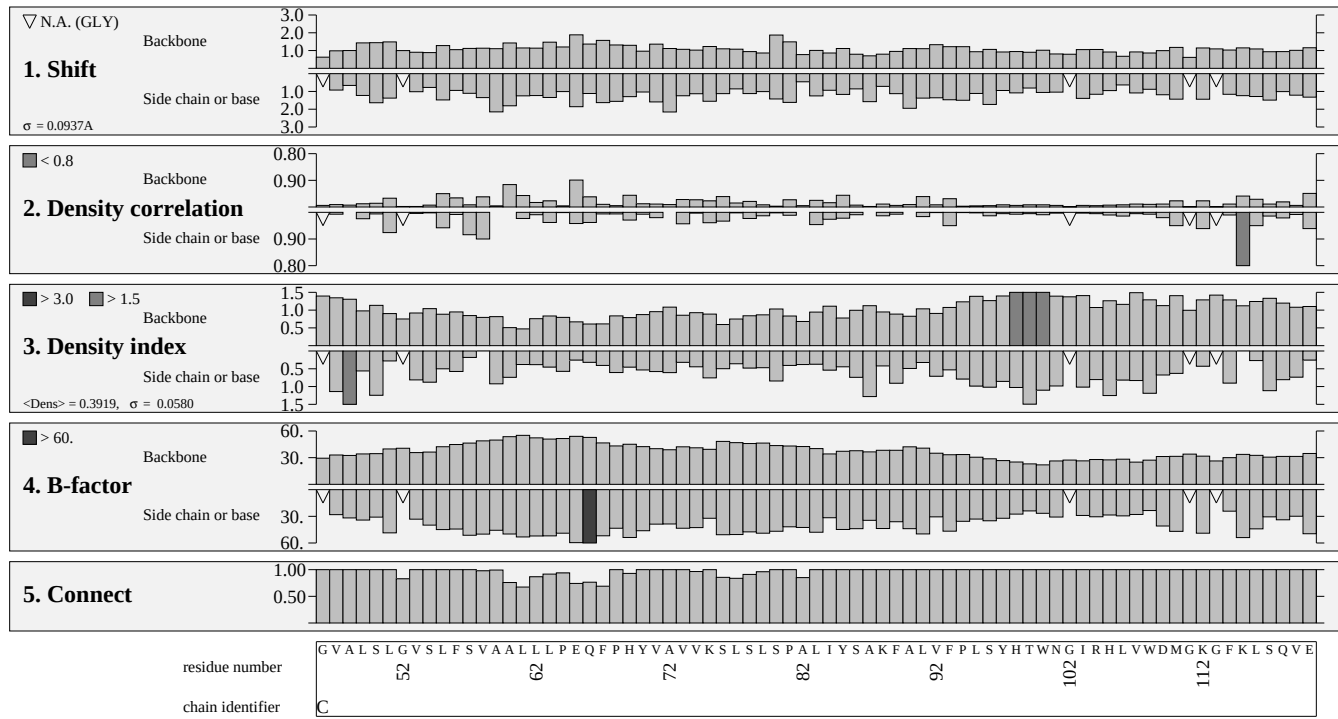




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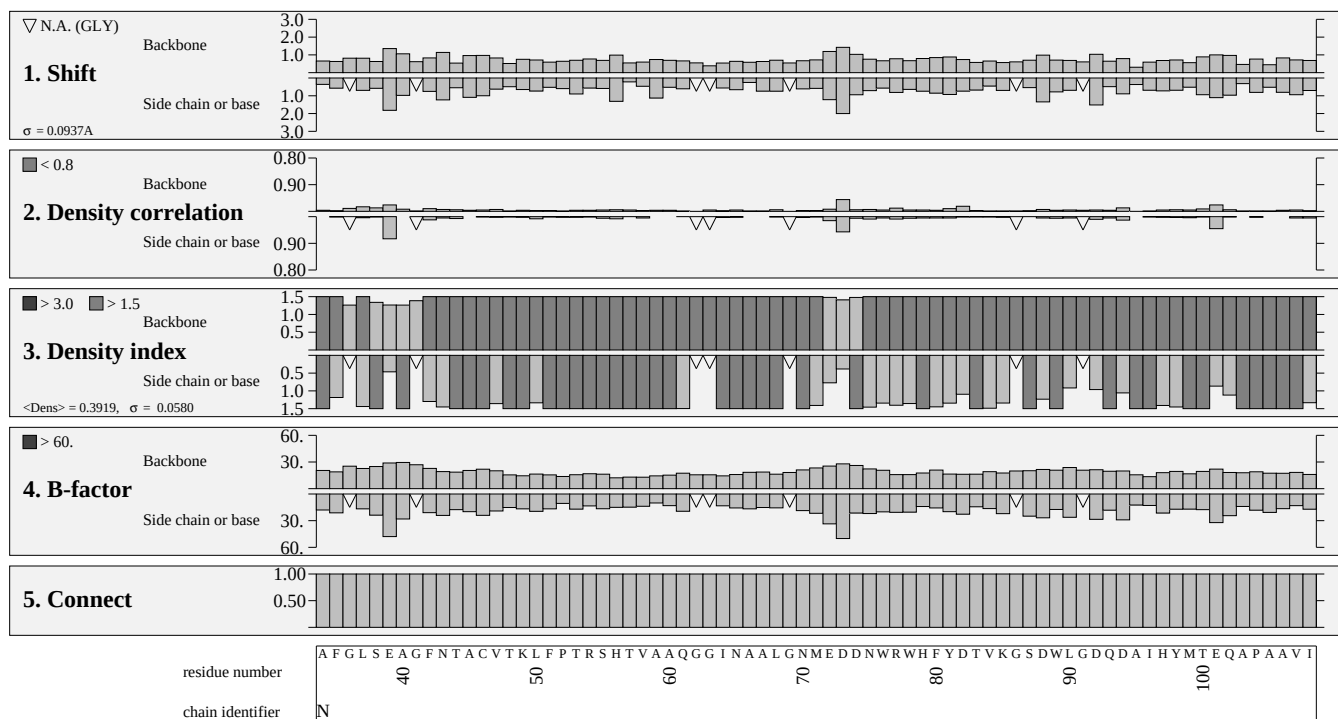
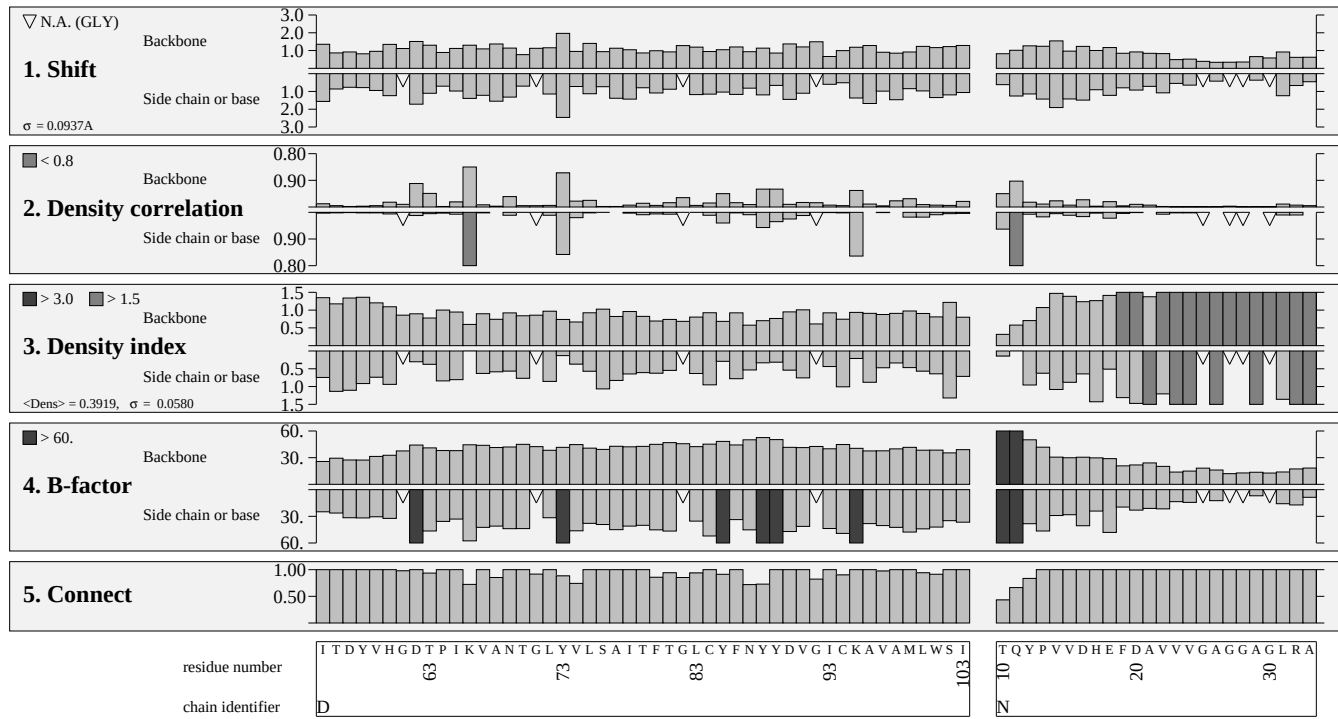
### Local estimation (7)



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## 2FBW

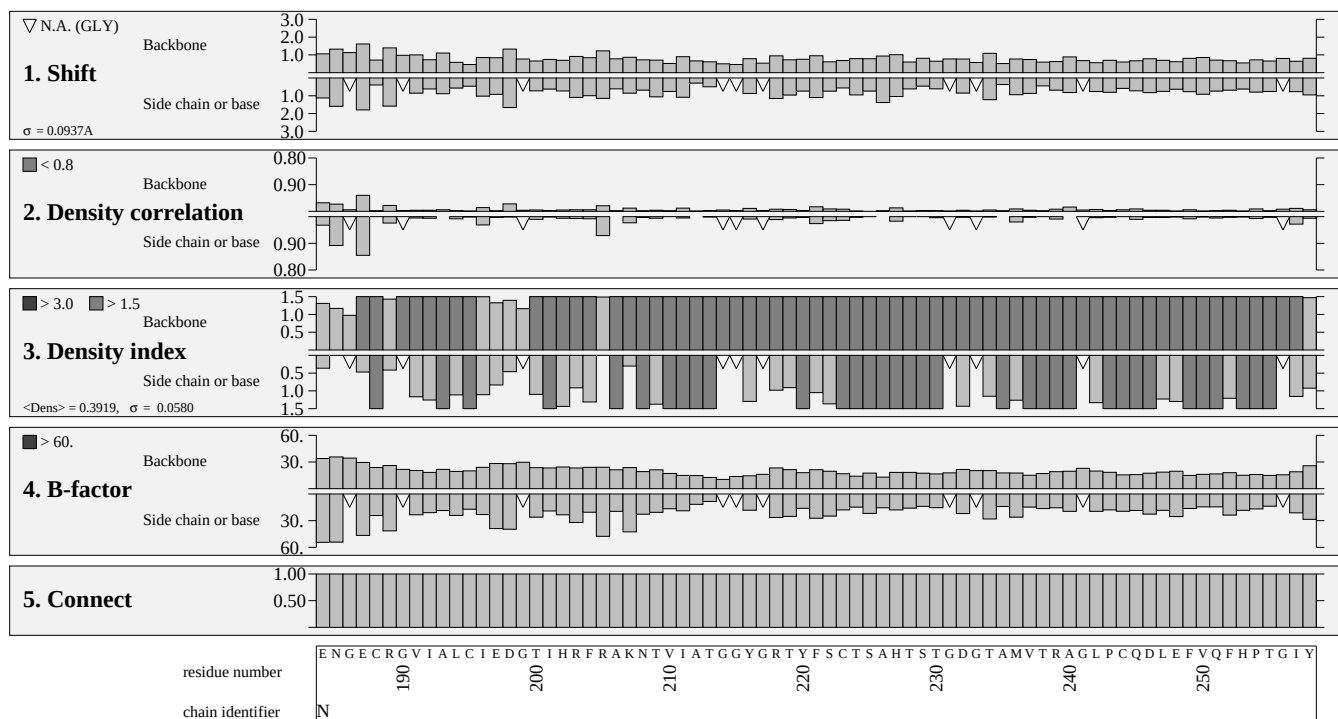
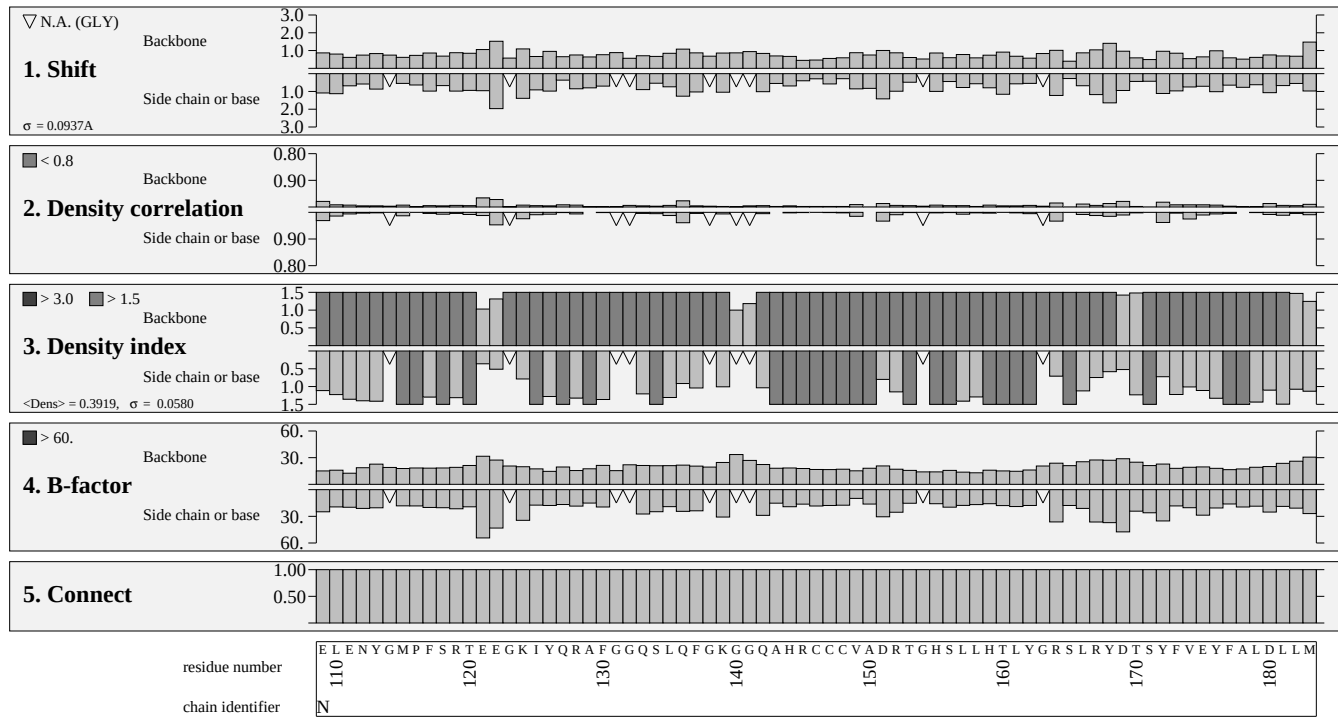
### Local estimation (8)



# Structure Factor Check

## 2FBW

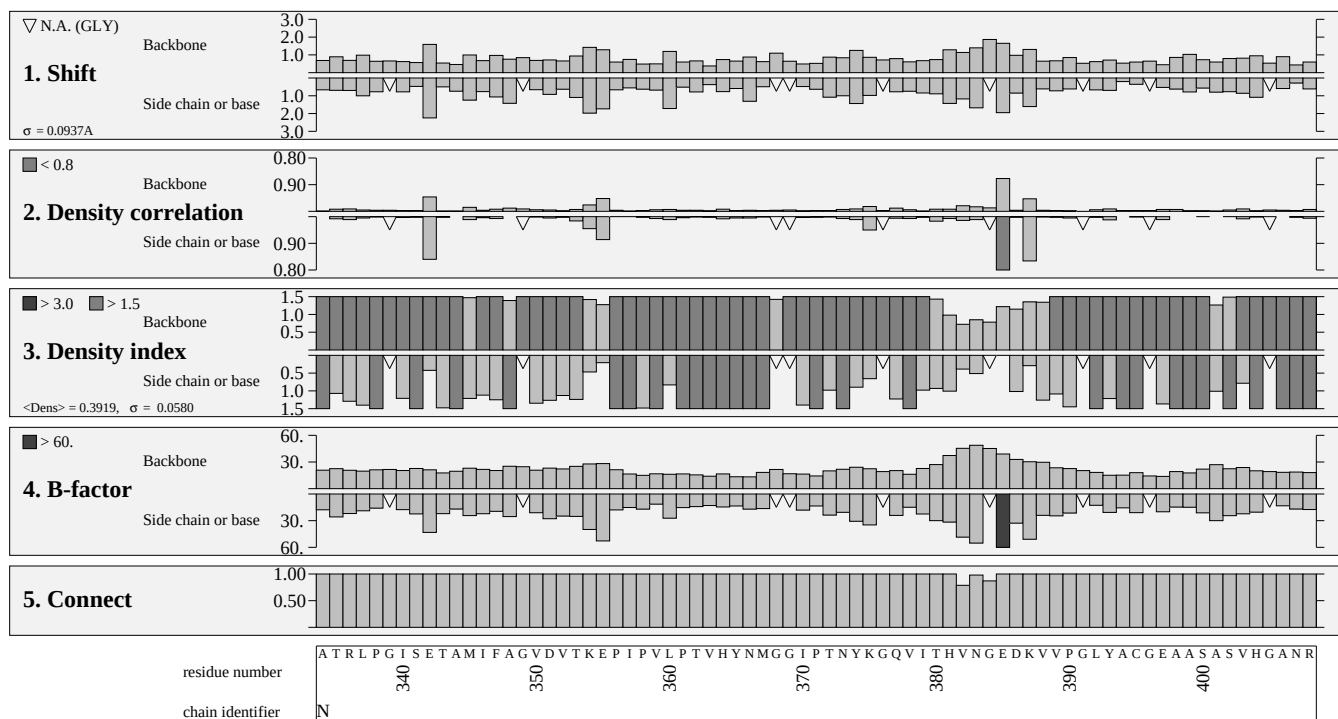
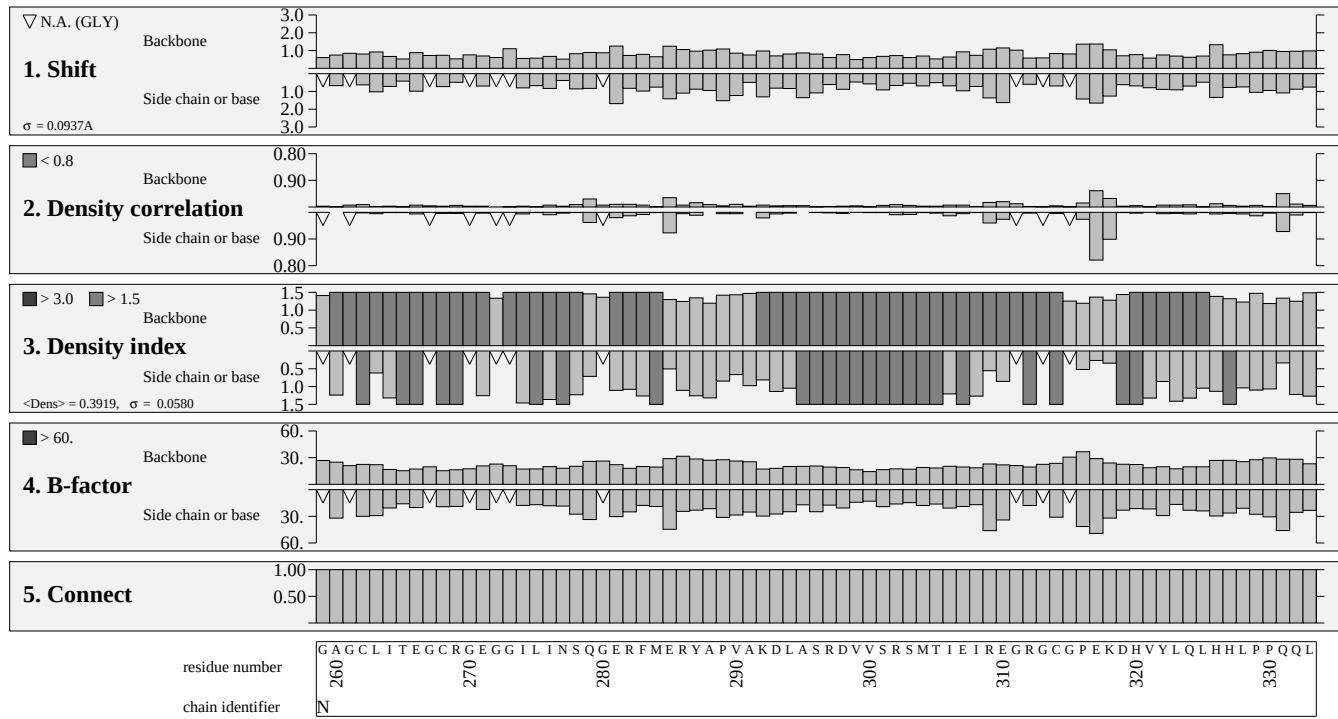
### Local estimation (9)



# Structure Factor Check

## 2FBW

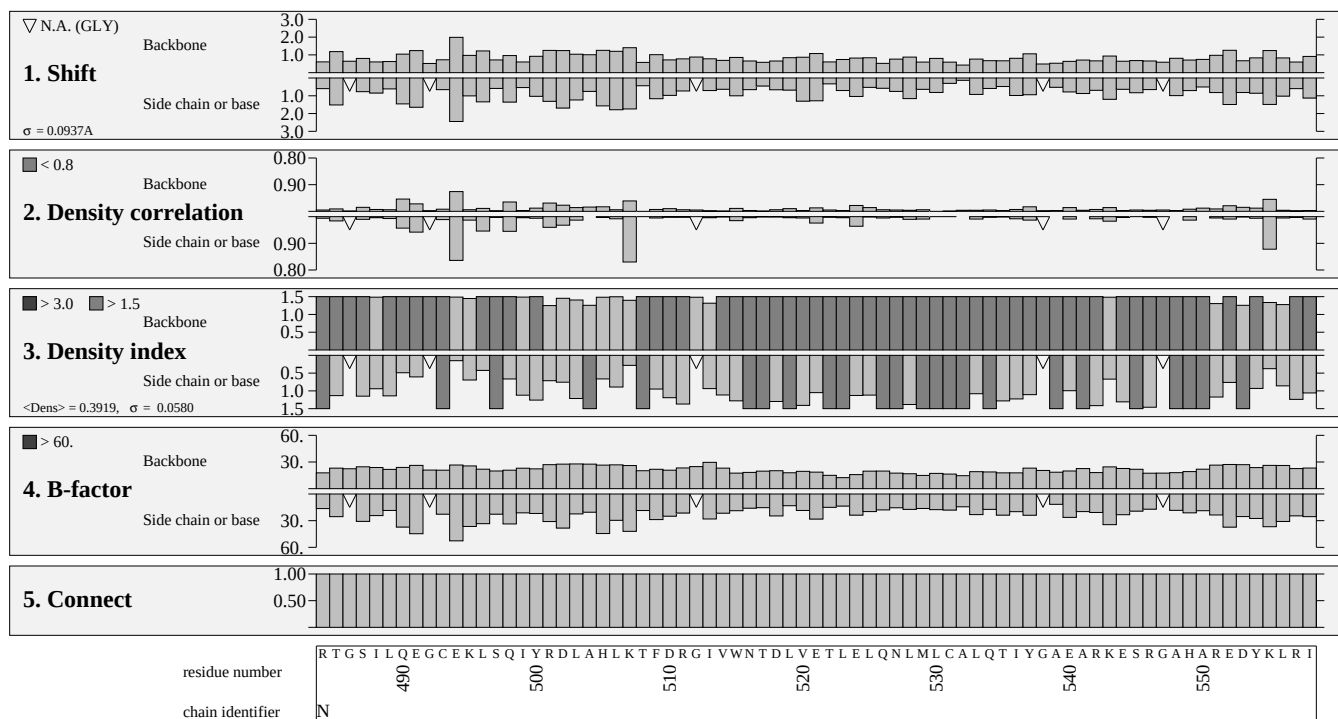
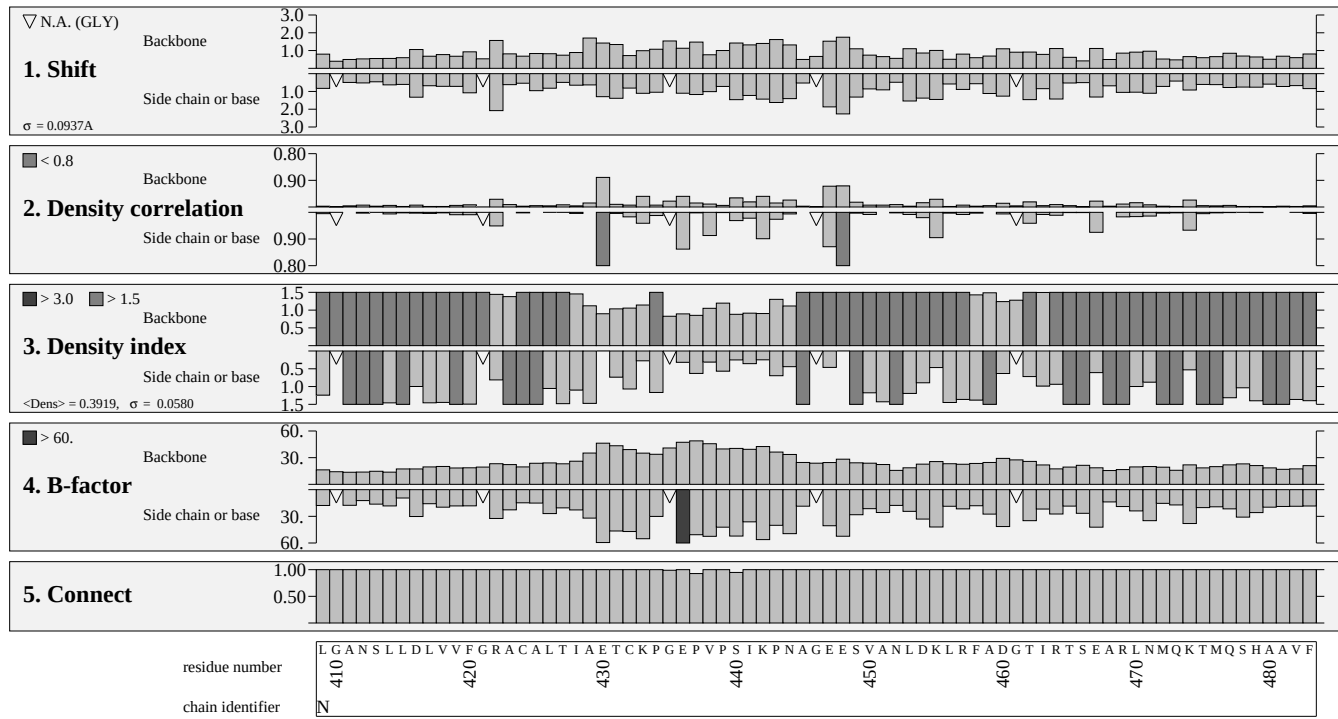
### Local estimation (10)



# Structure Factor Check

## 2FBW

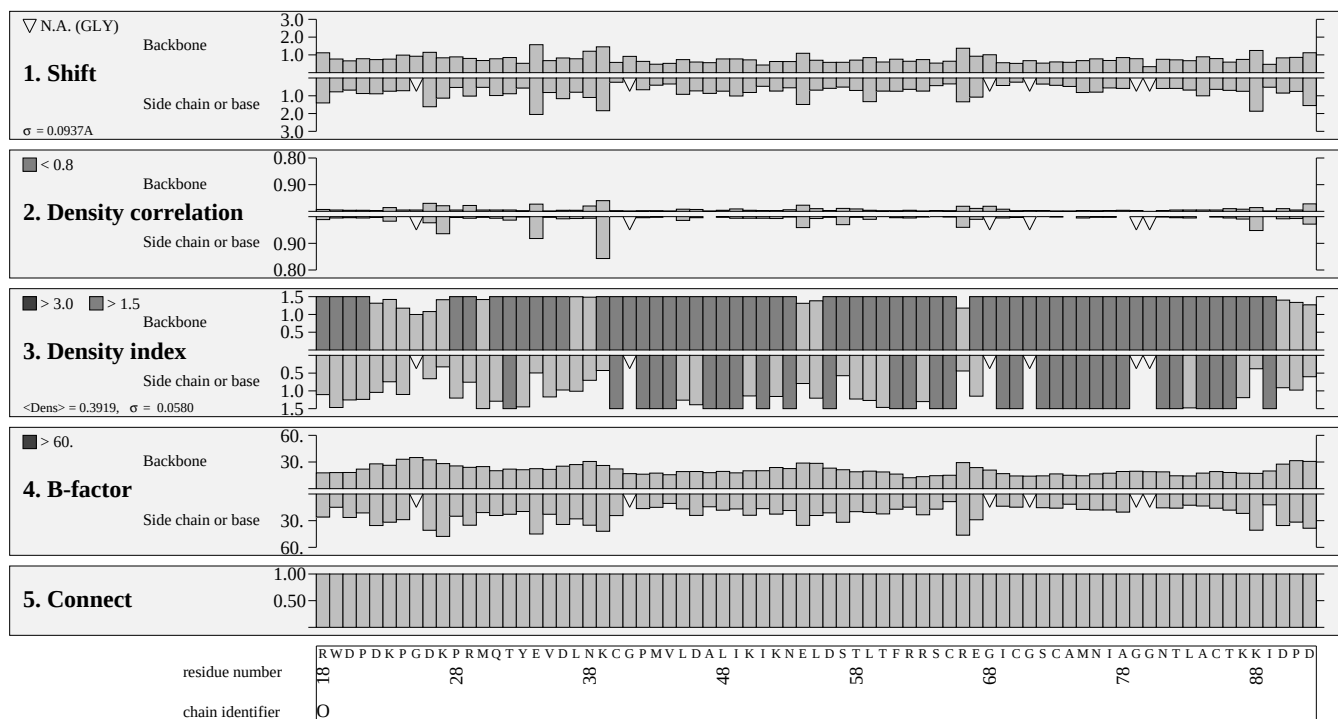
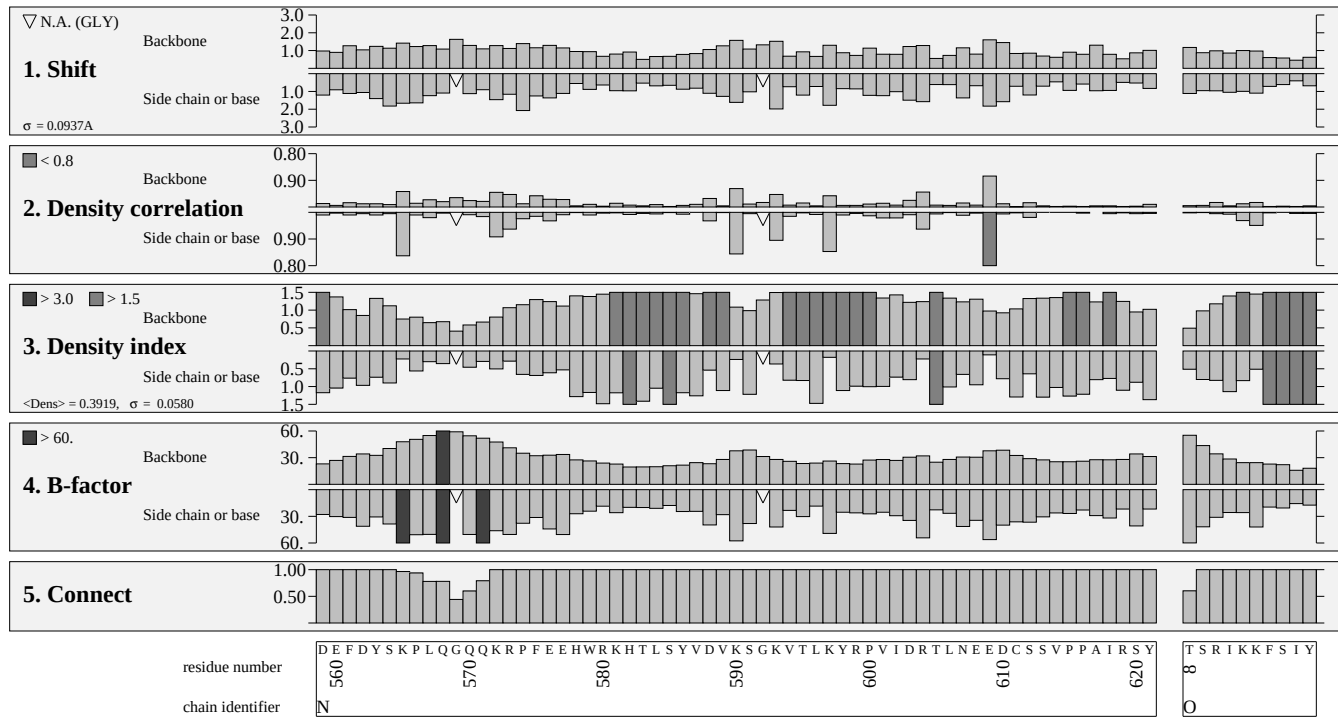
### Local estimation (11)



# Structure Factor Check

## 2FBW

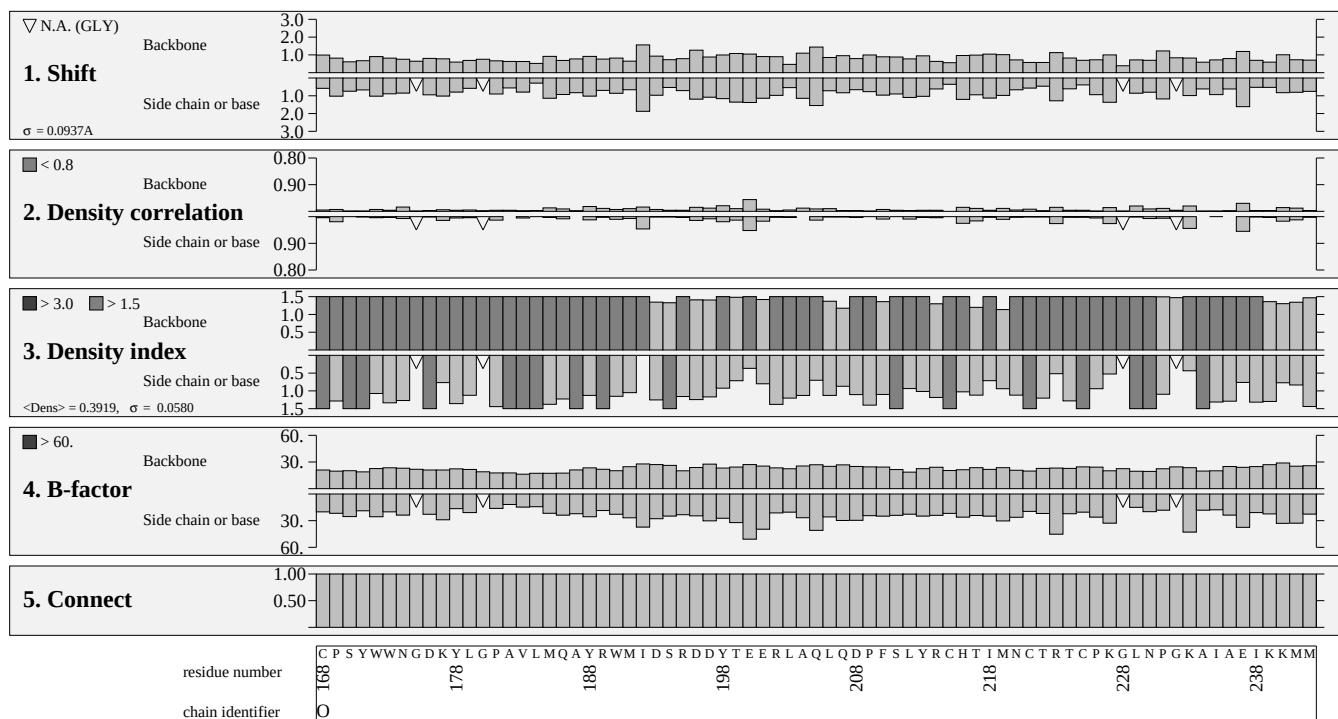
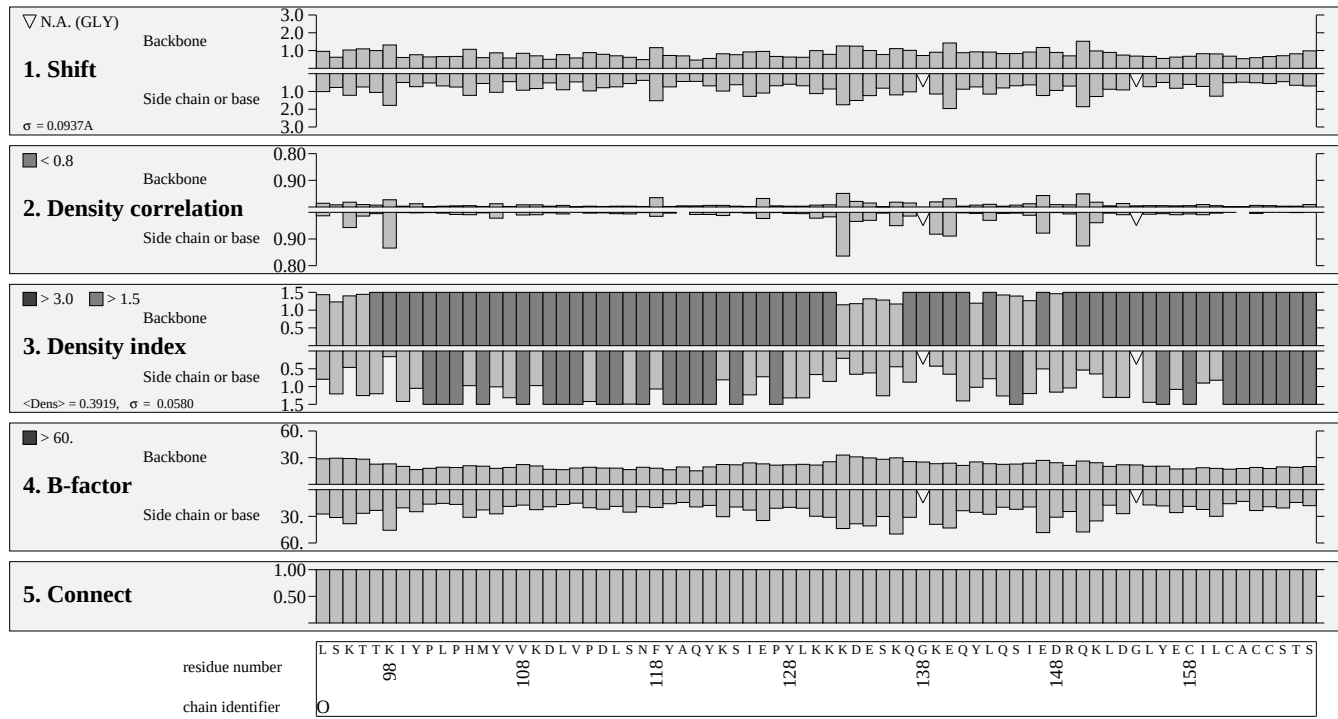
### Local estimation (12)



# Structure Factor Check

## 2FBW

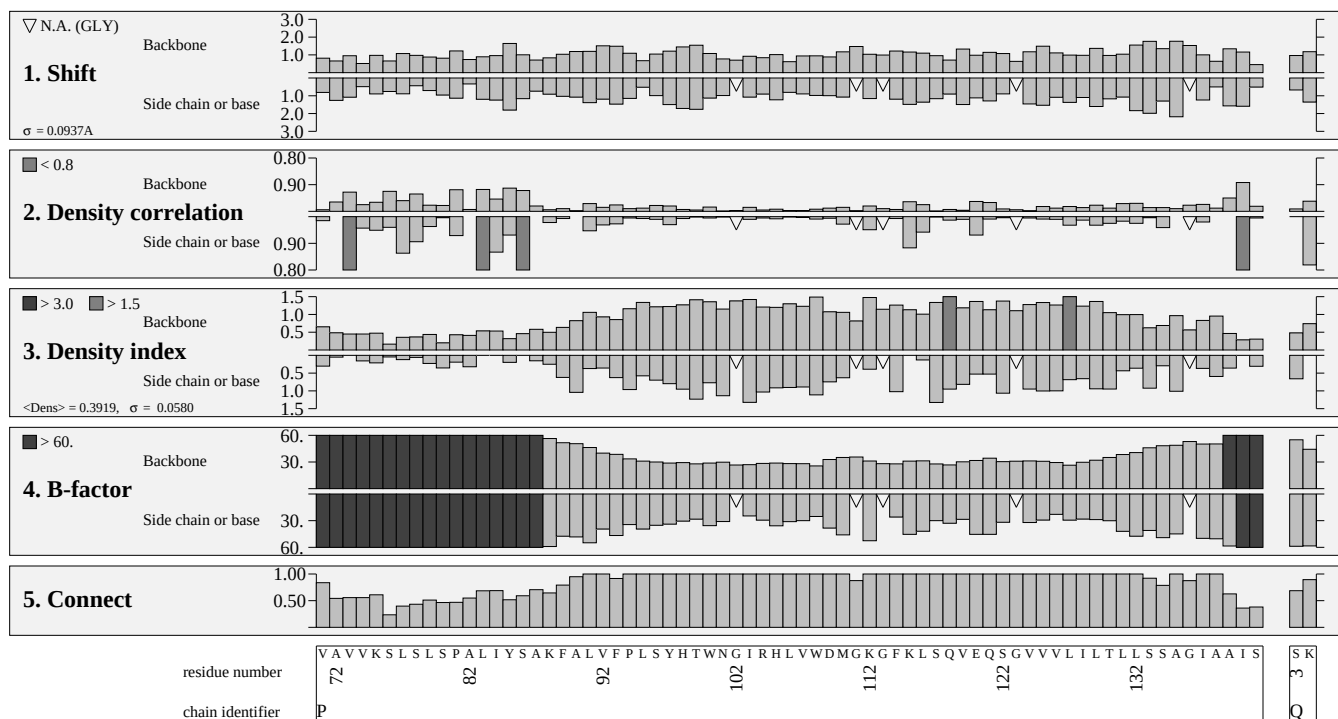
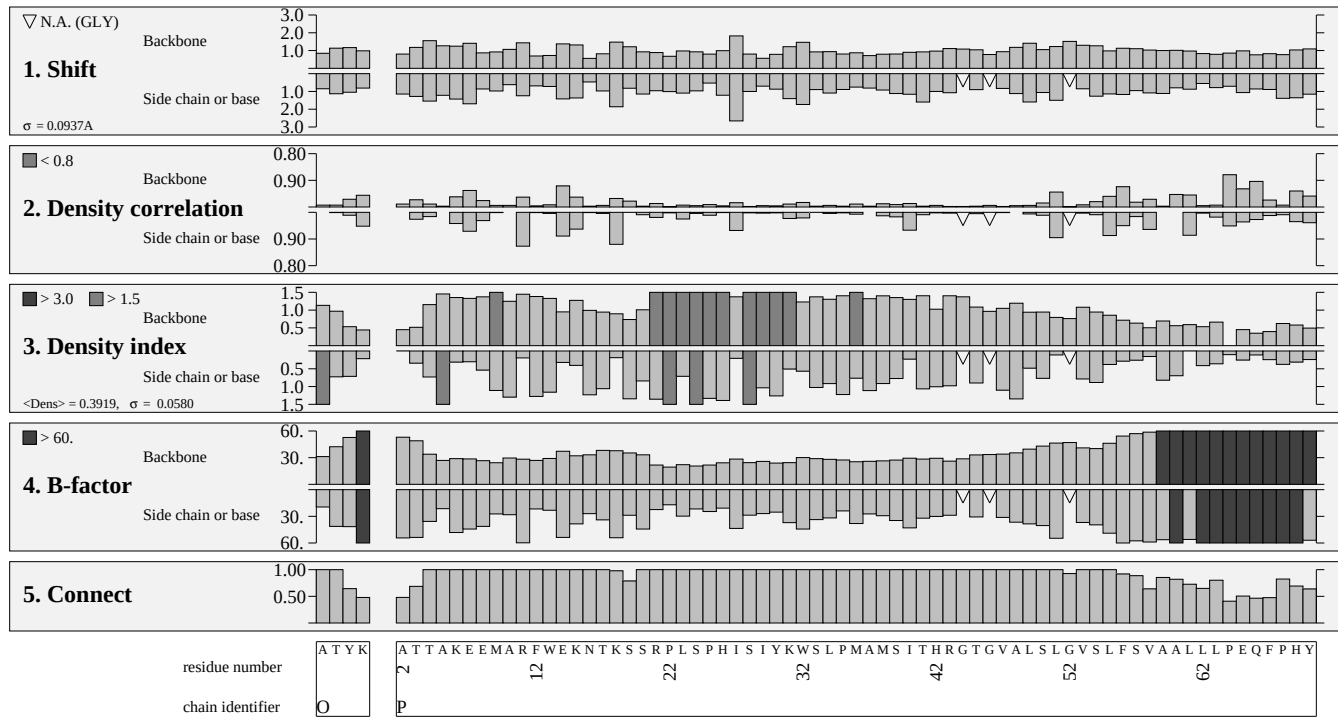
### Local estimation (13)



# Structure Factor Check

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### Local estimation (14)

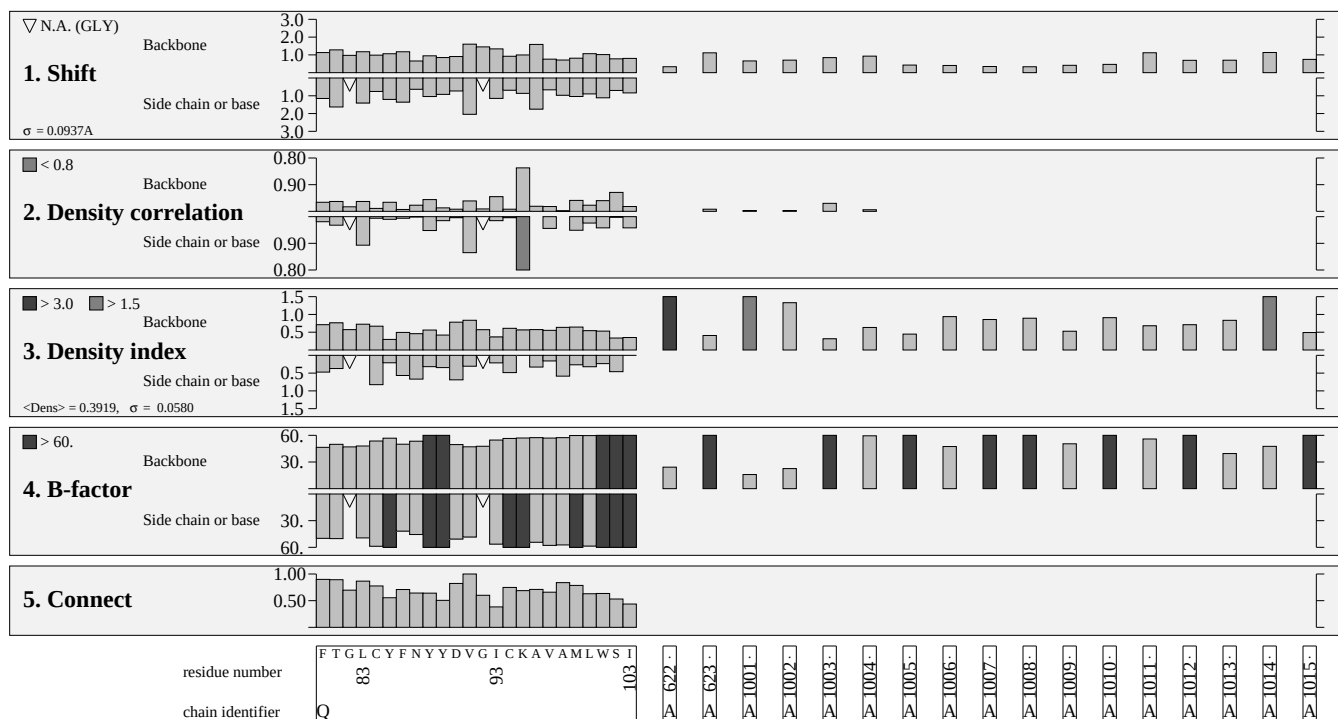
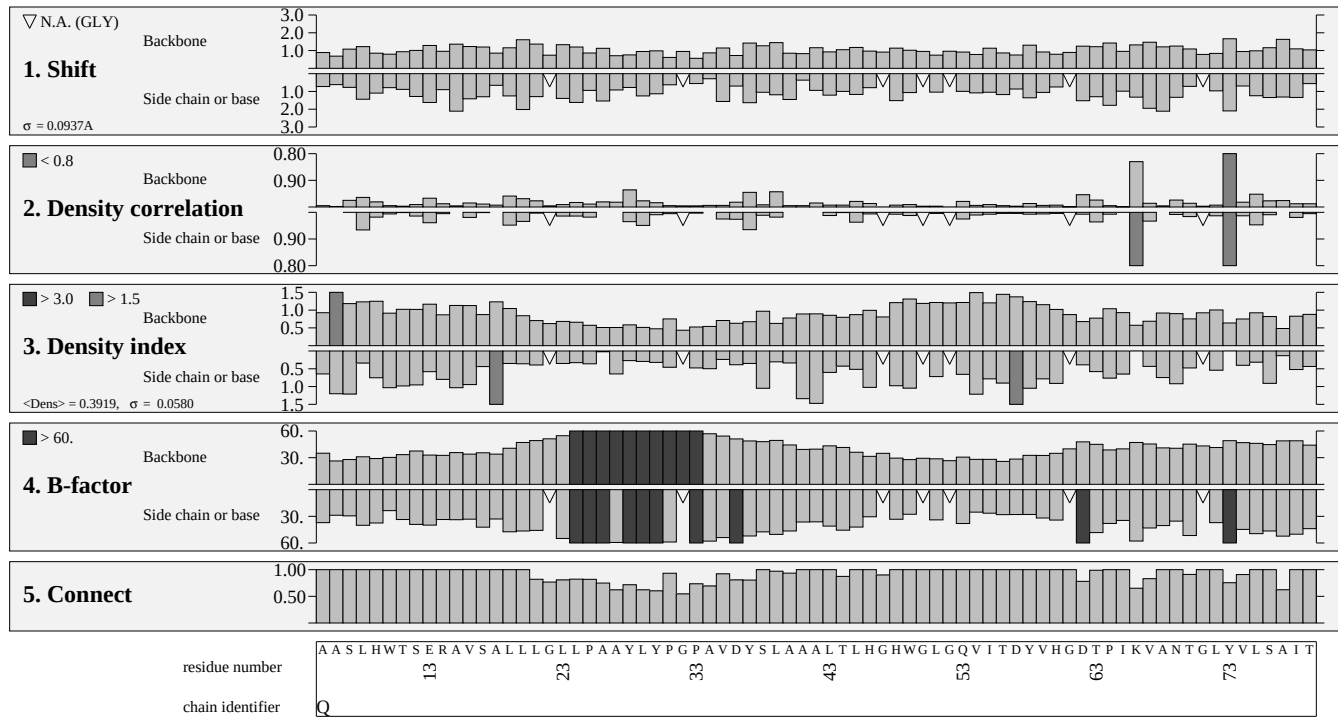




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## 2FBW

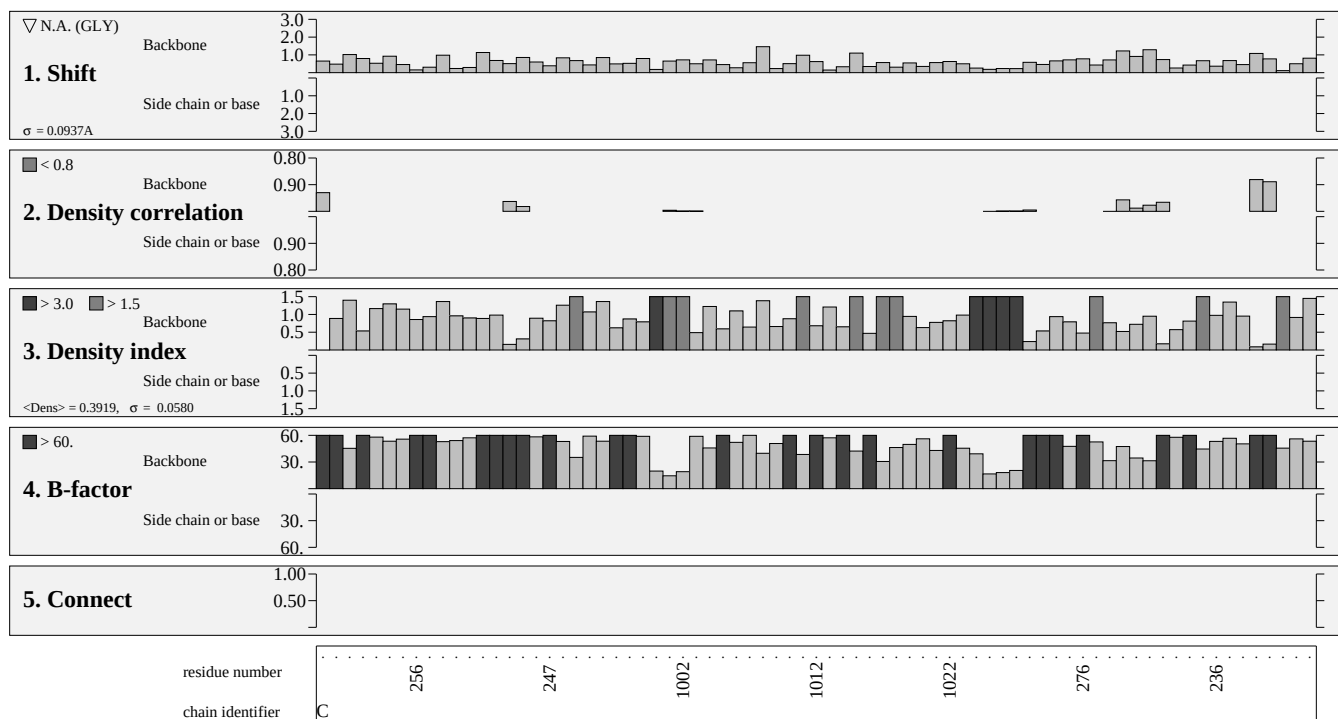
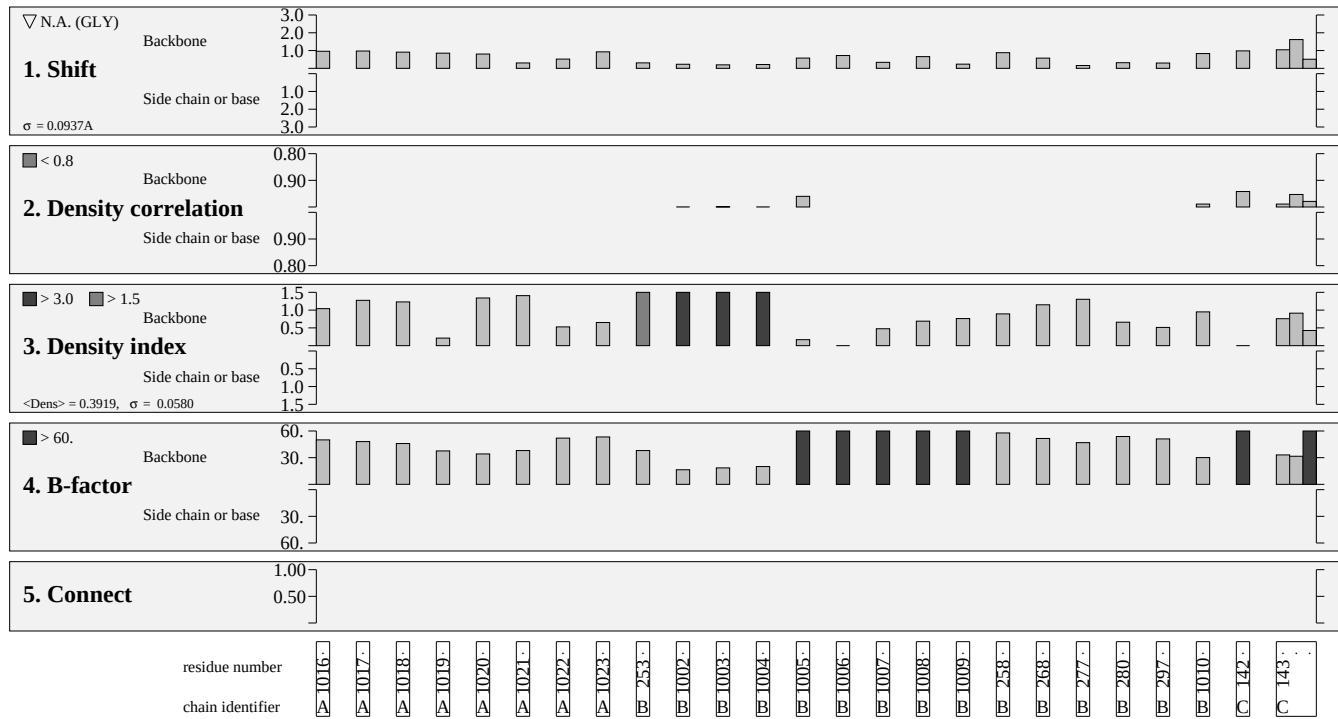
### Local estimation (15)



# Structure Factor Check

## 2FBW

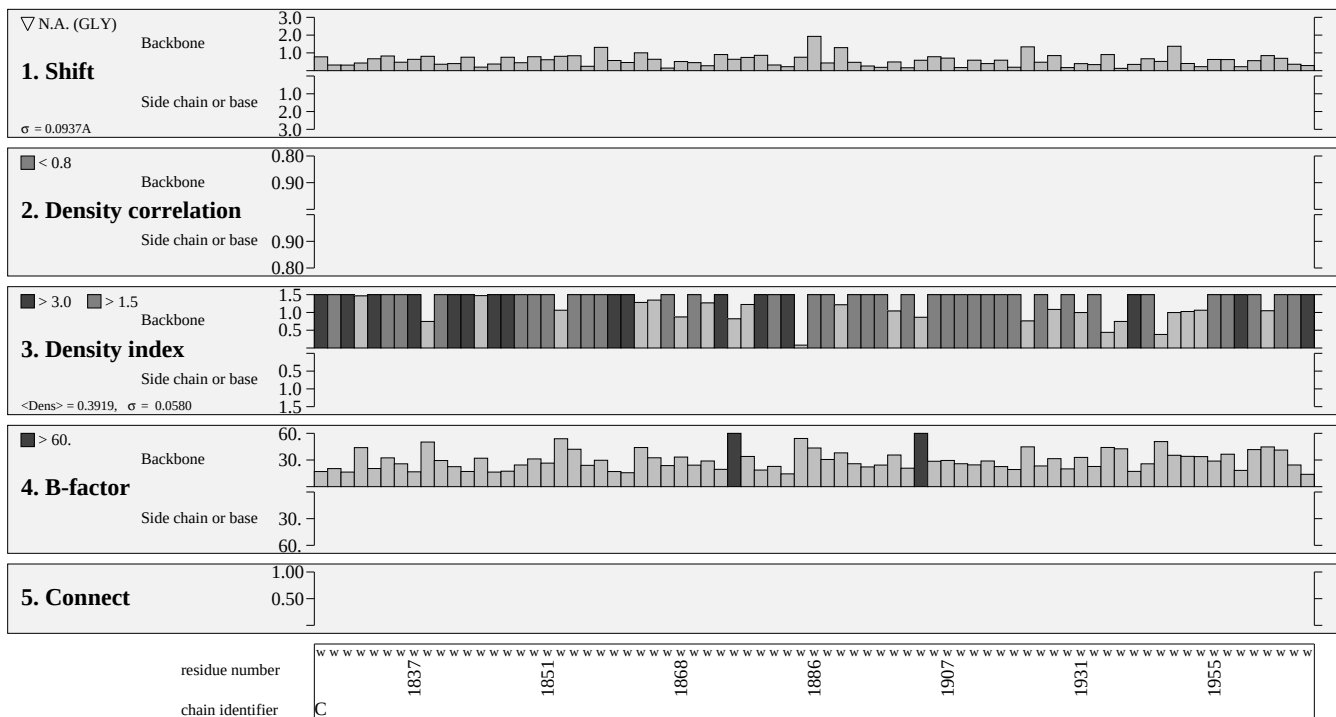
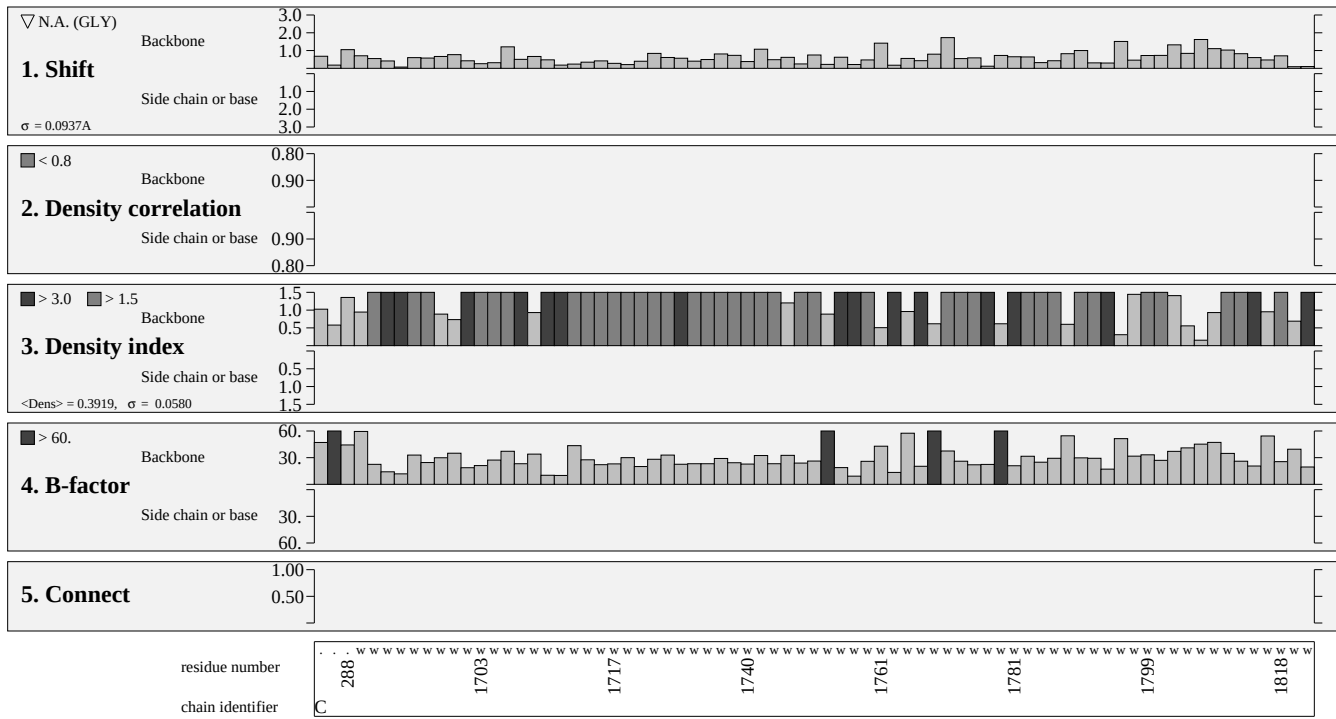
### Local estimation (16)



# Structure Factor Check

## 2FBW

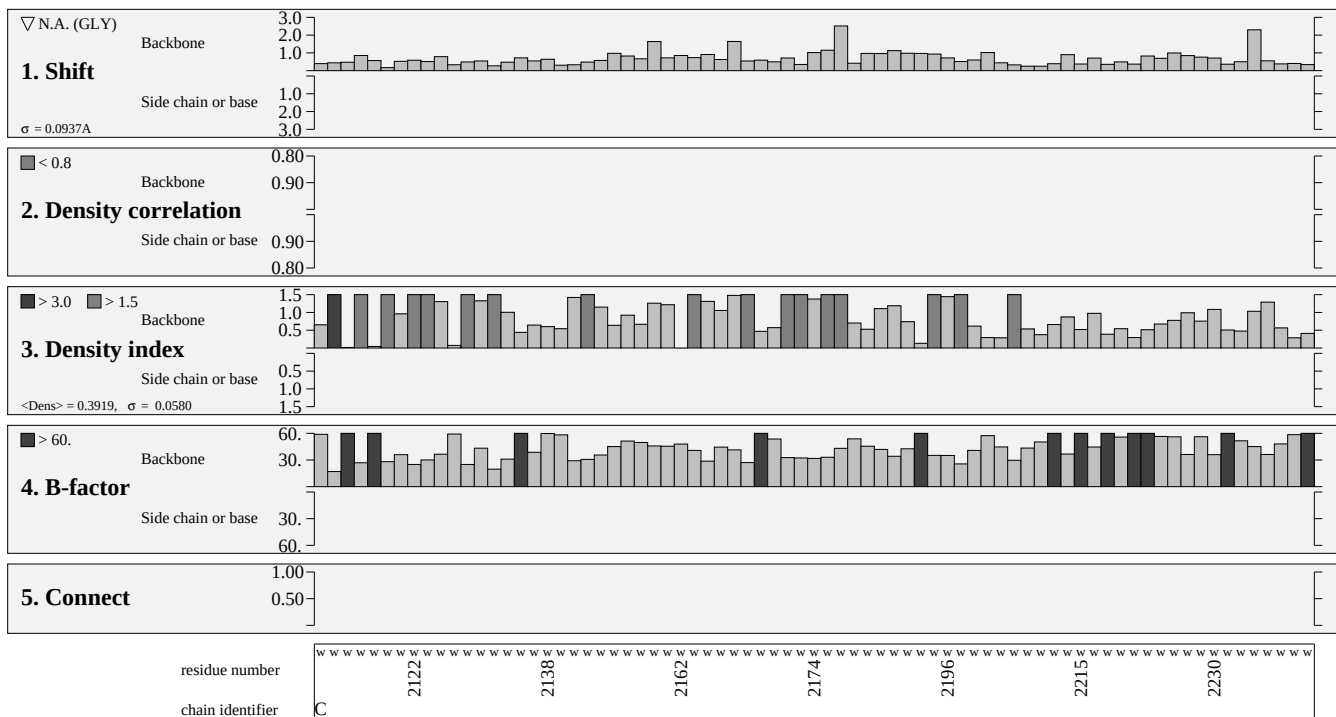
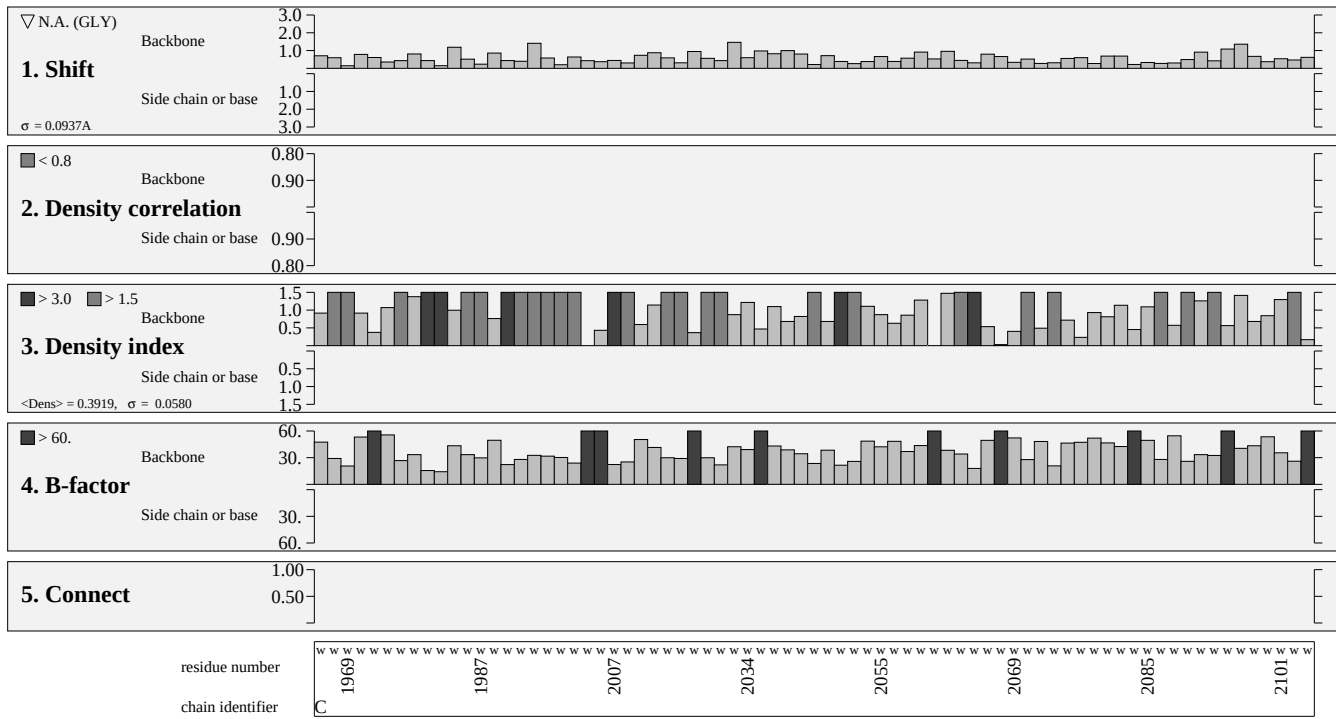
### Local estimation (17)



# Structure Factor Check

## 2FBW

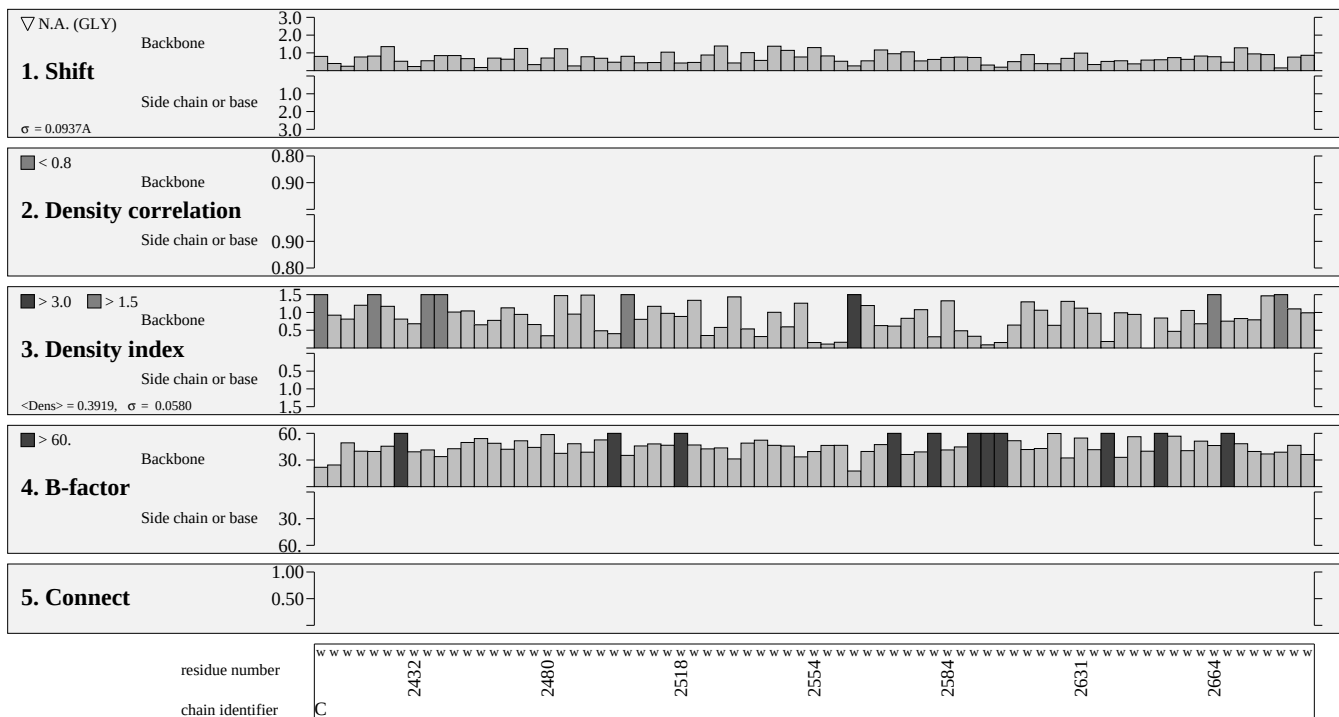
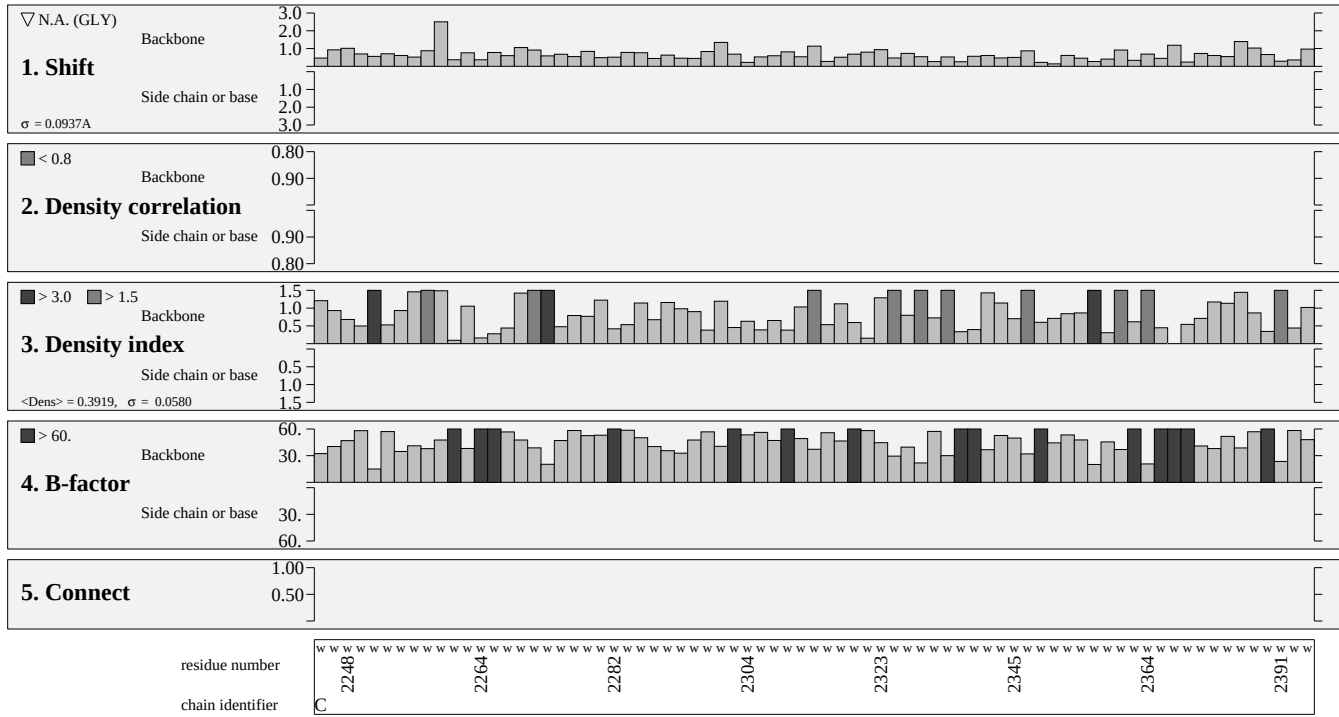
### Local estimation (18)



# Structure Factor Check

## 2FBW

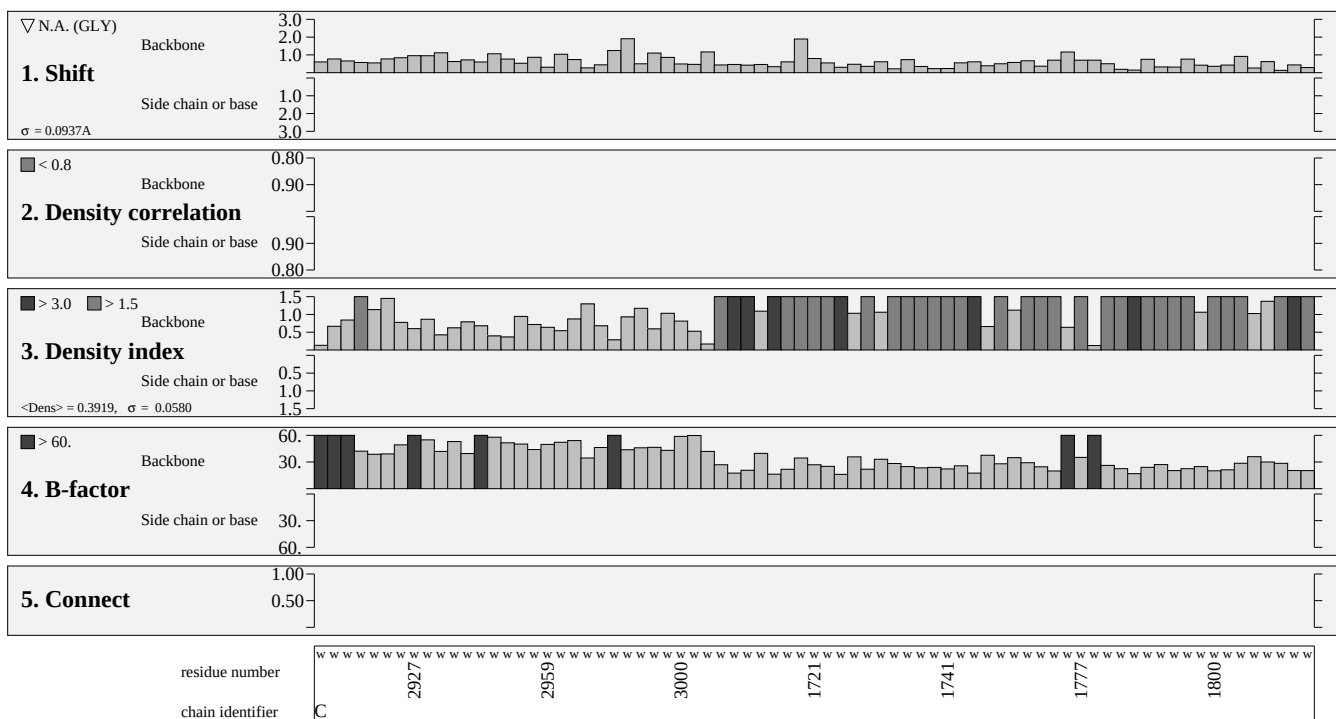
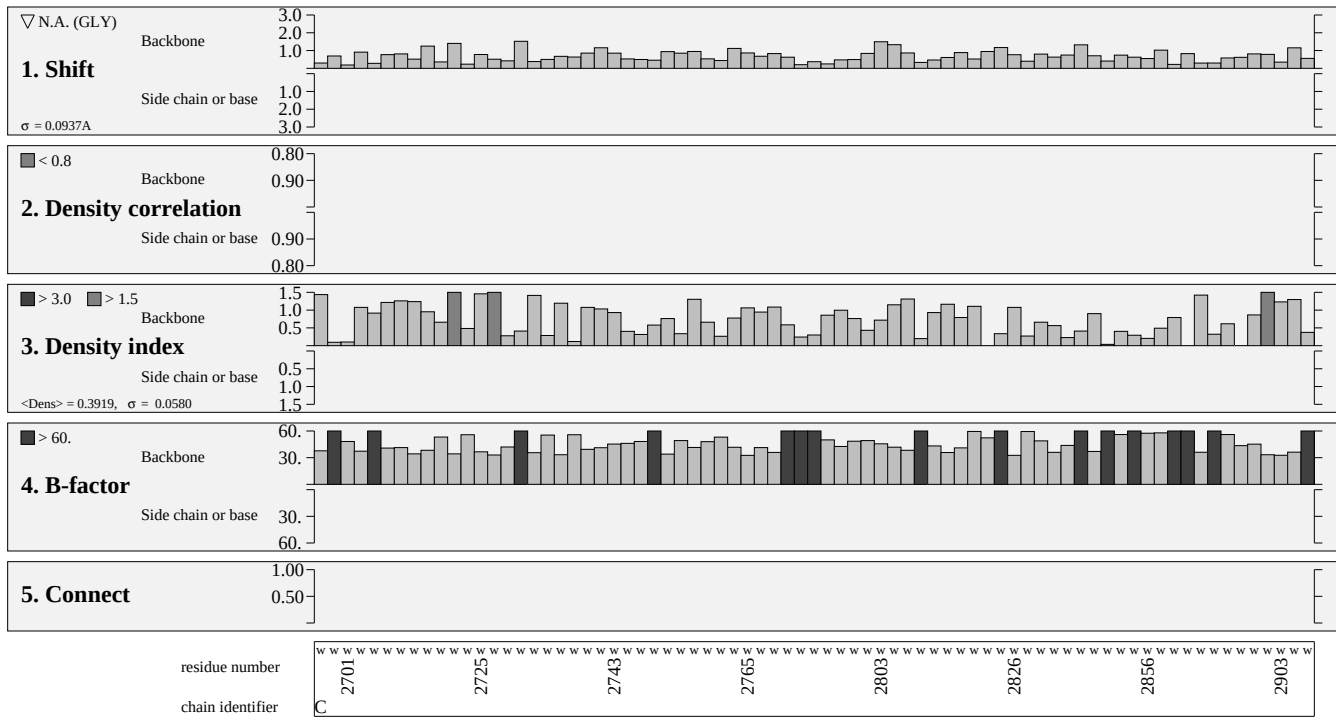
### Local estimation (19)



# Structure Factor Check

## 2FBW

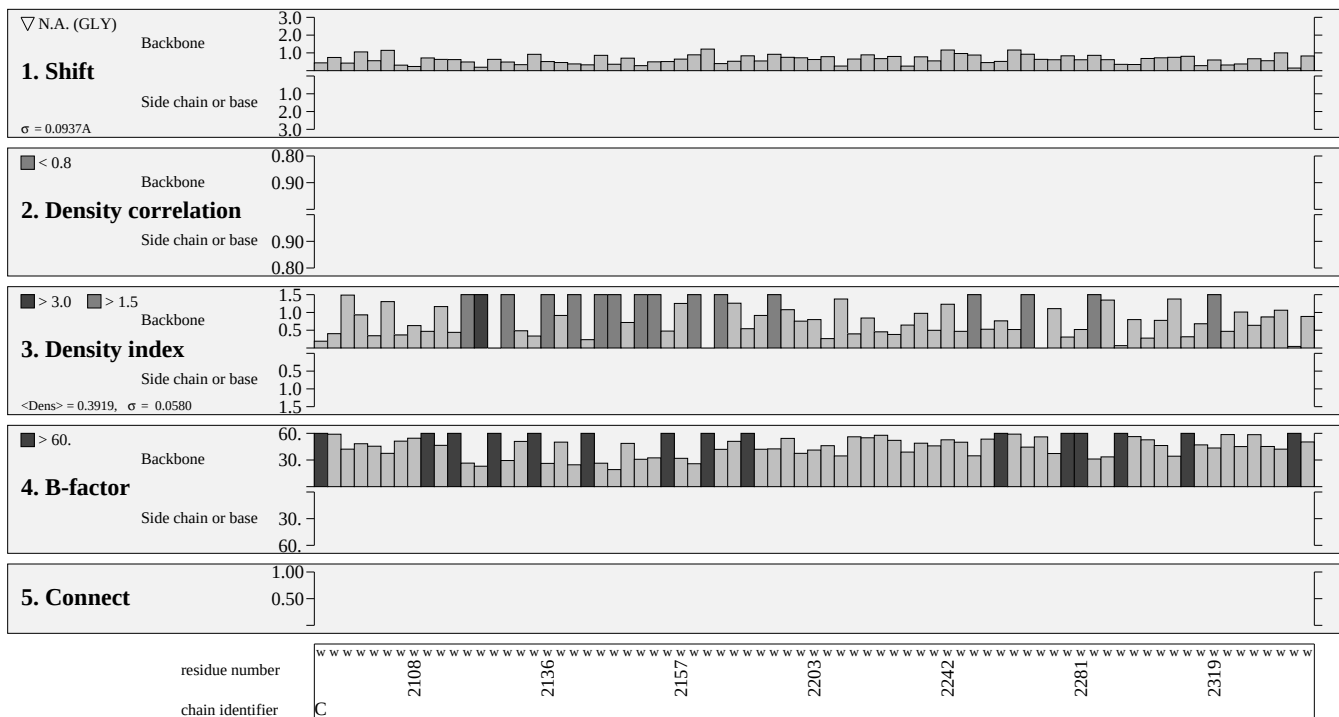
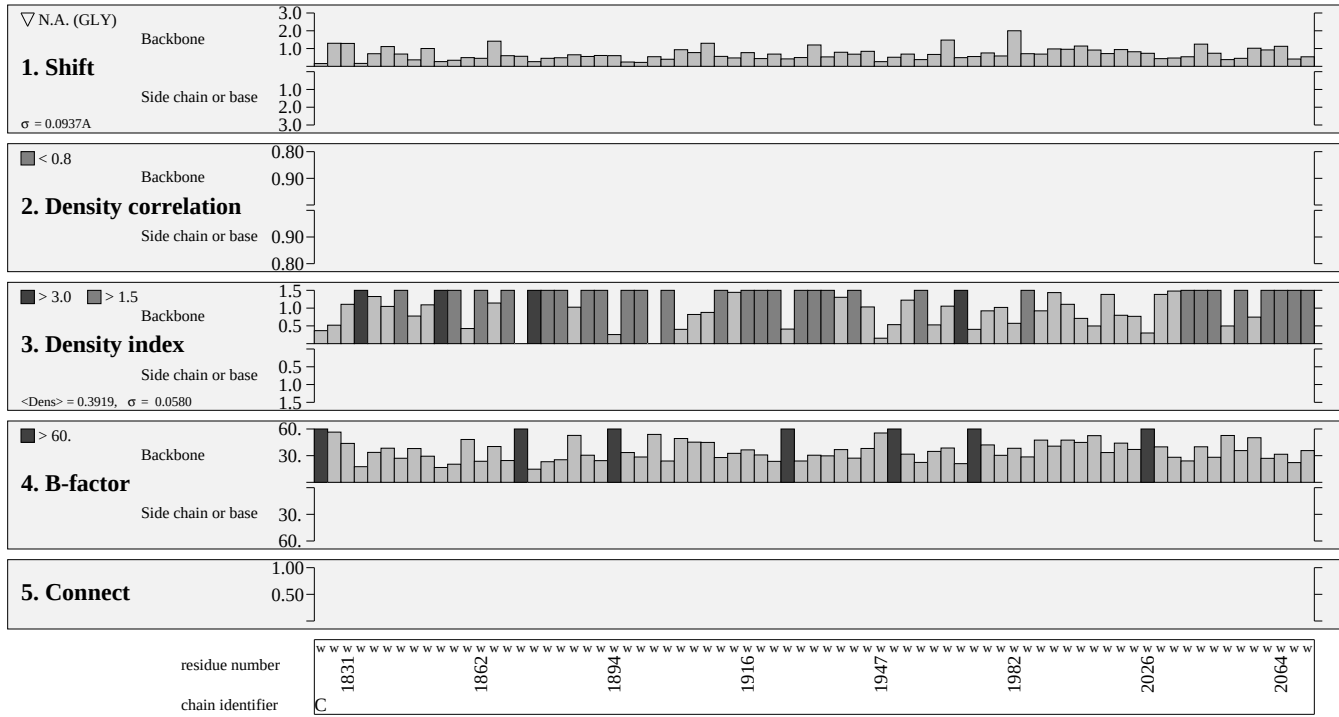
### Local estimation (20)



# Structure Factor Check

## 2FBW

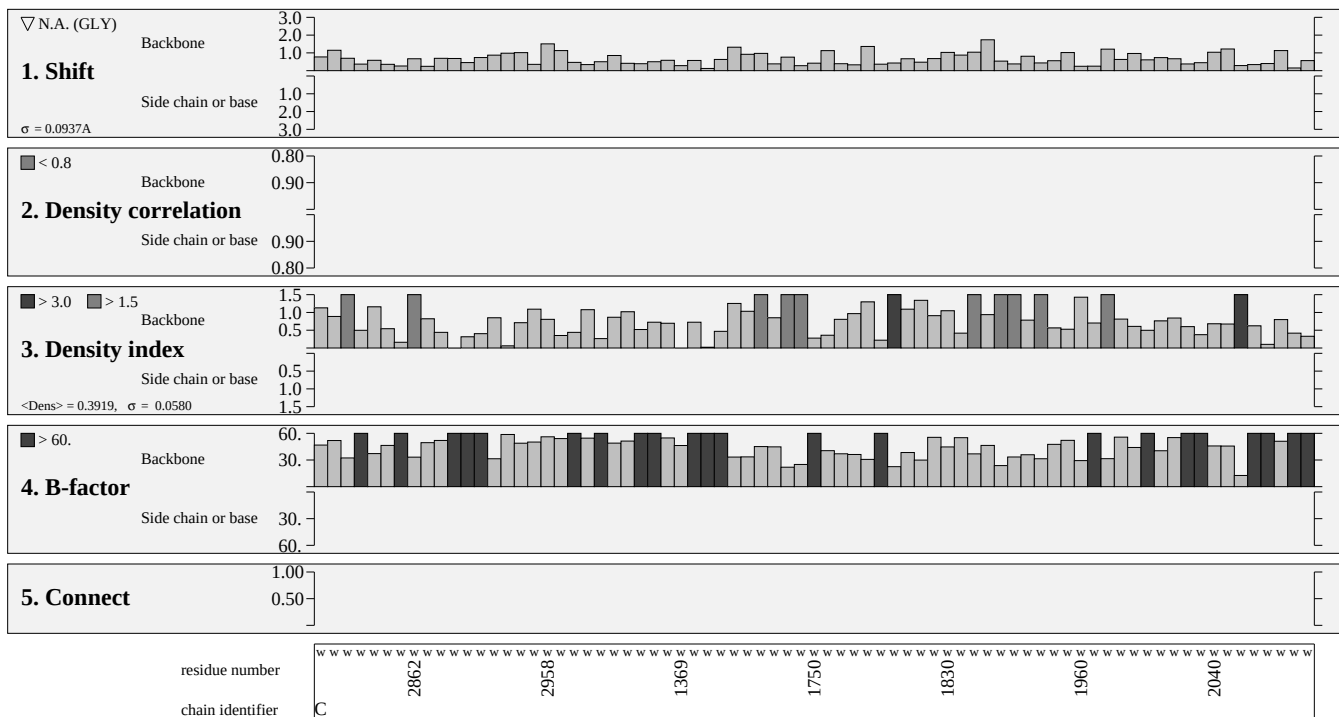
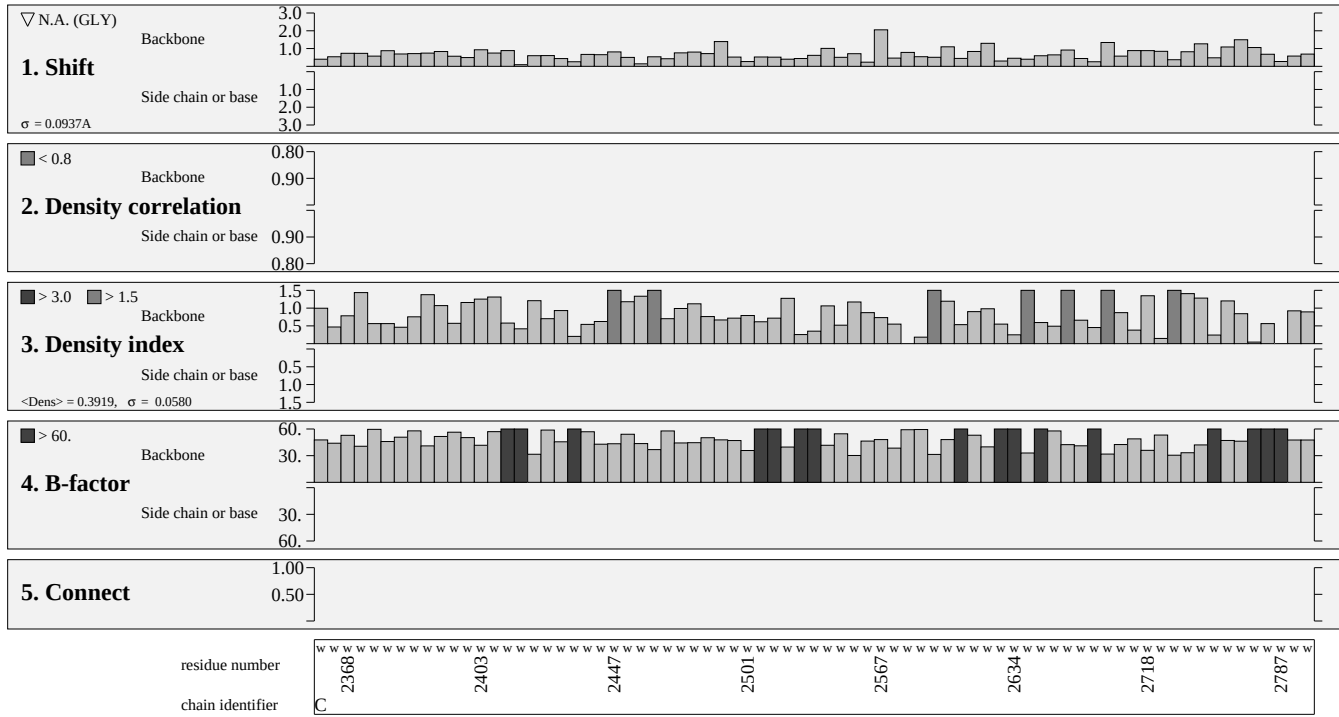
### Local estimation (21)



# Structure Factor Check

## 2FBW

### Local estimation (22)

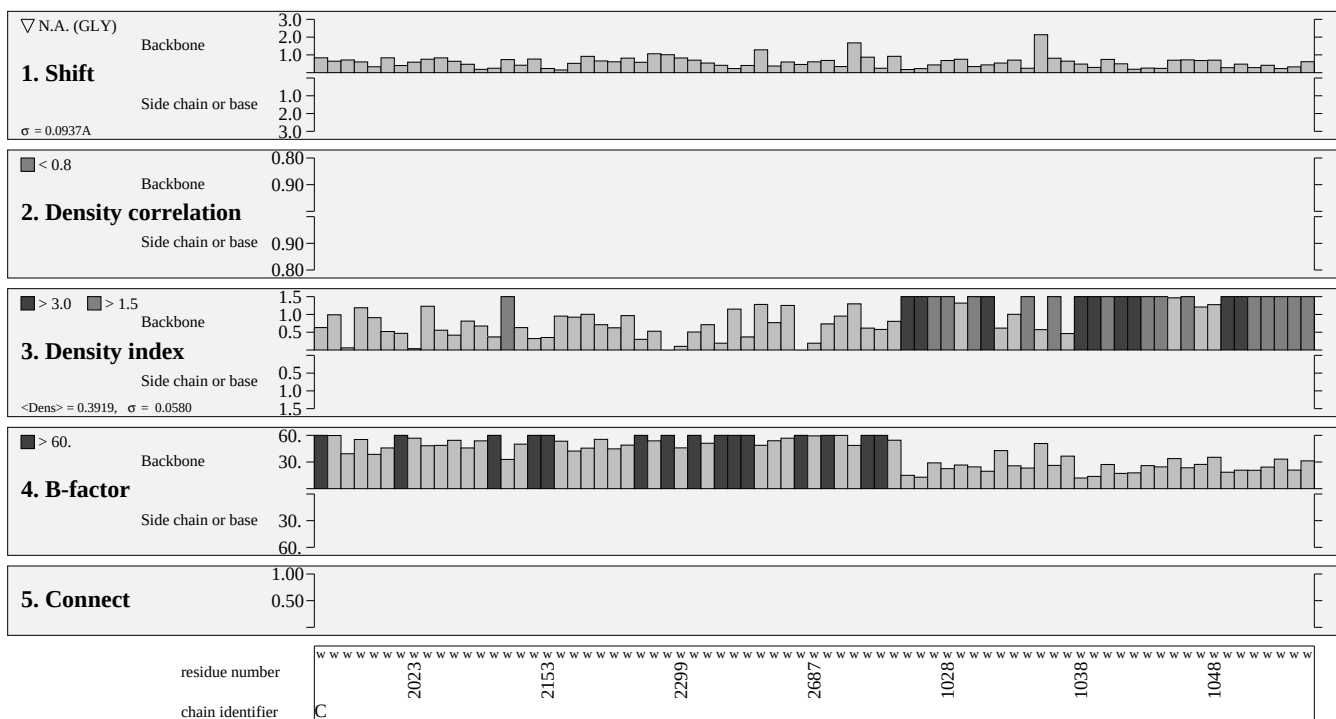
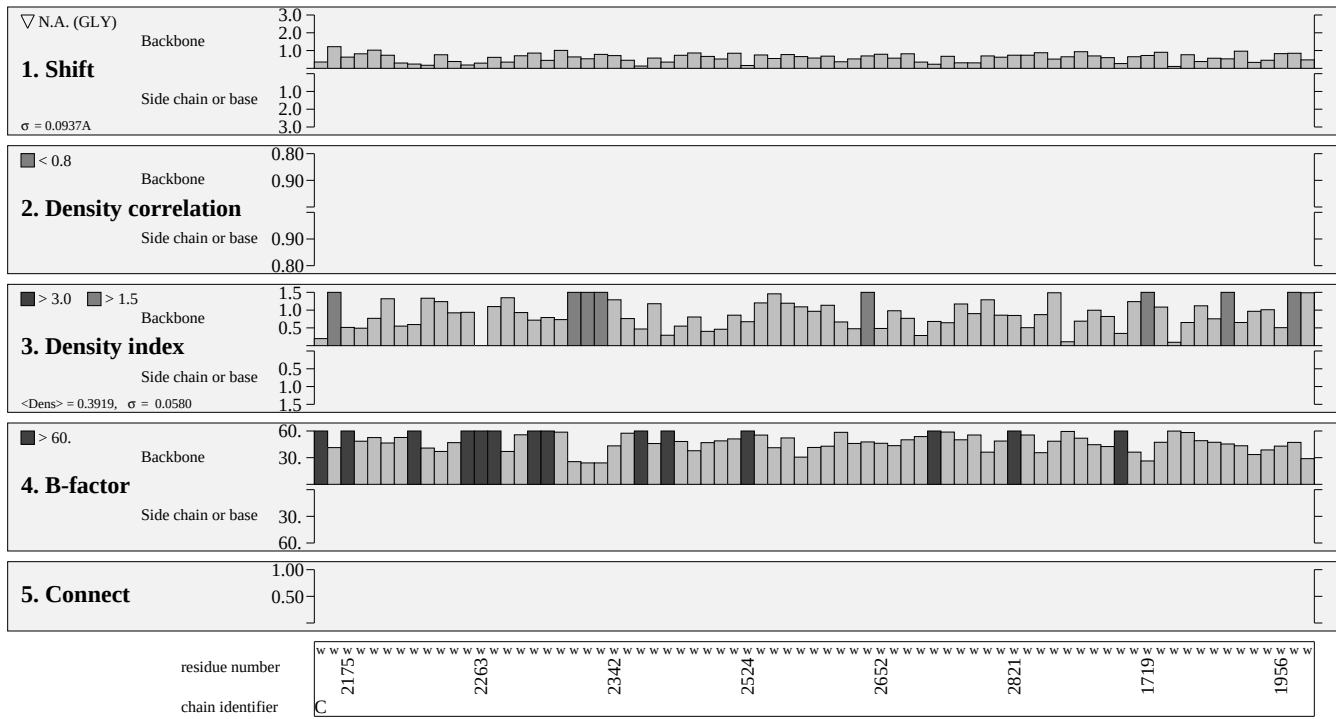




# Structure Factor Check

## 2FBW

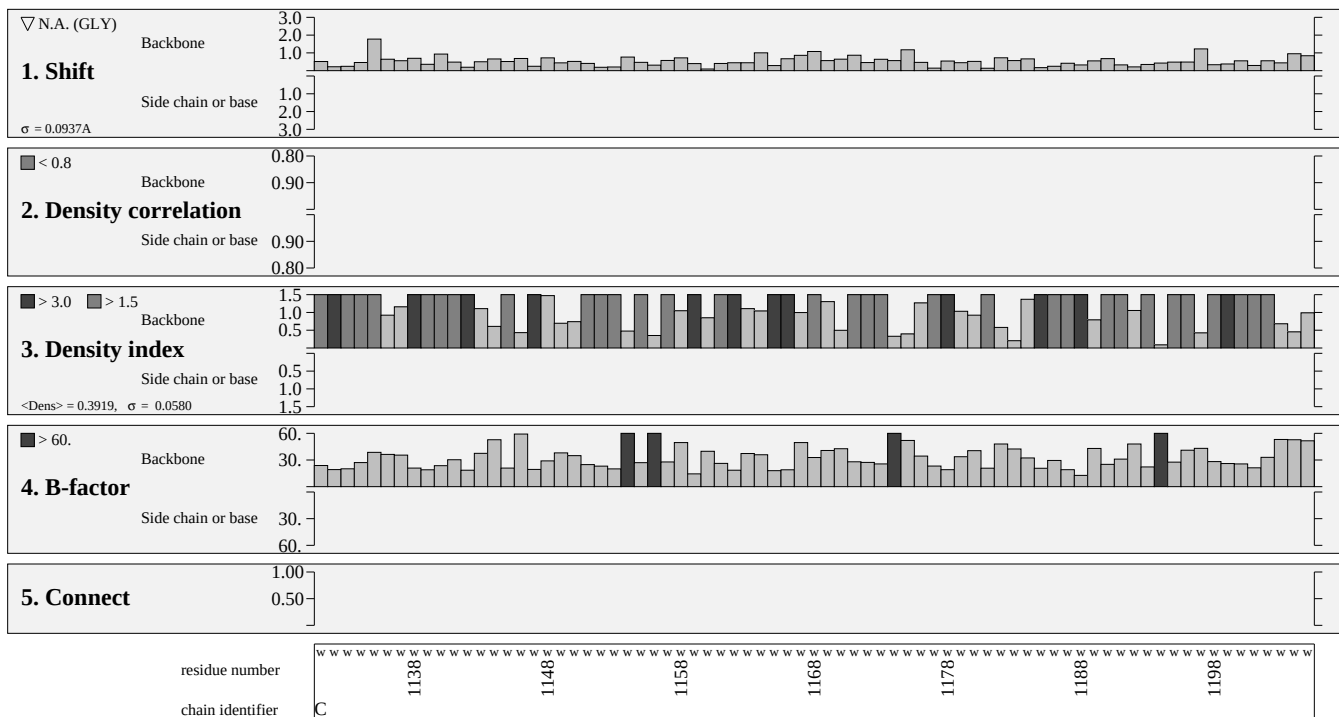
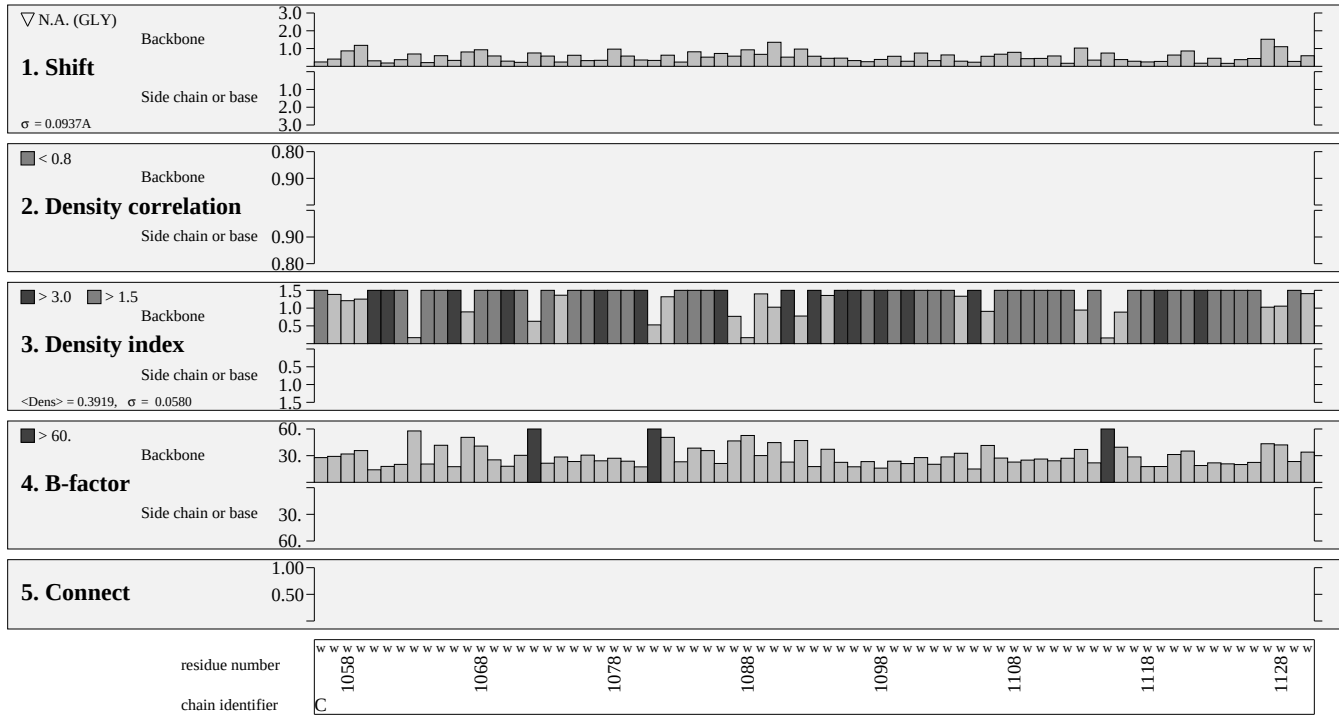
### Local estimation (23)



# Structure Factor Check

## 2FBW

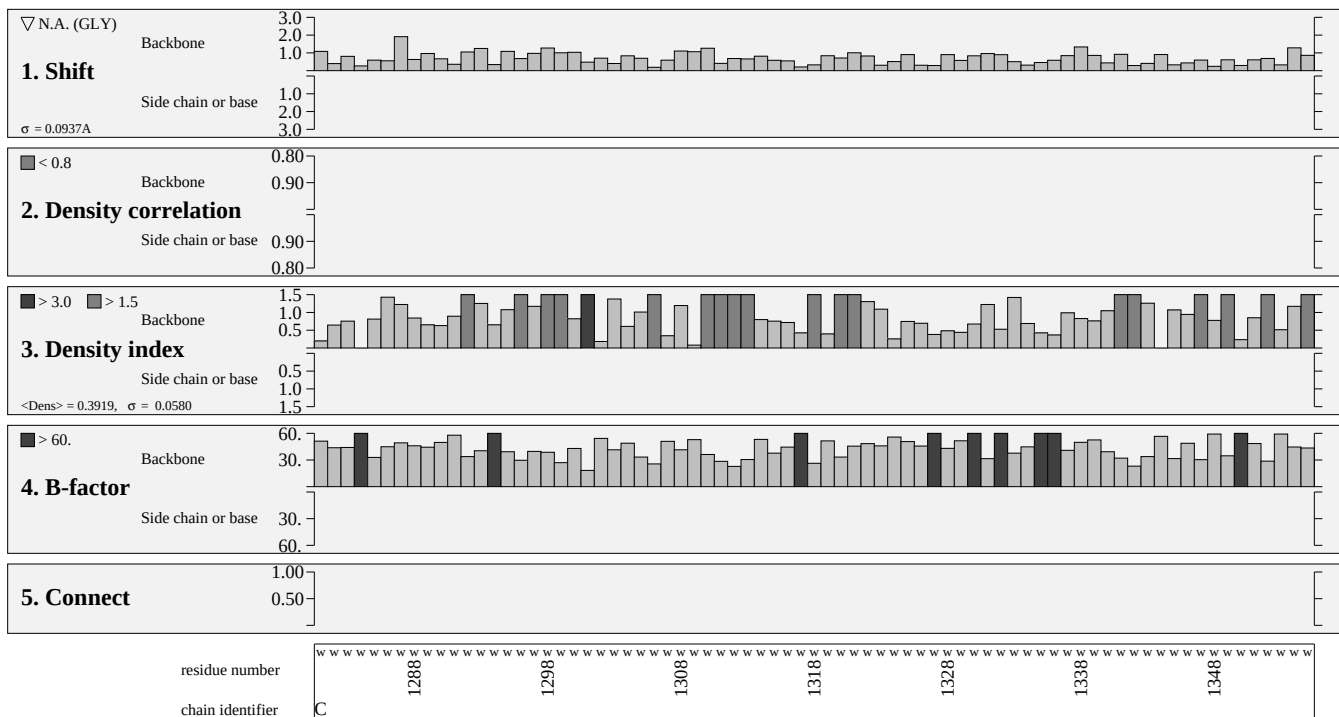
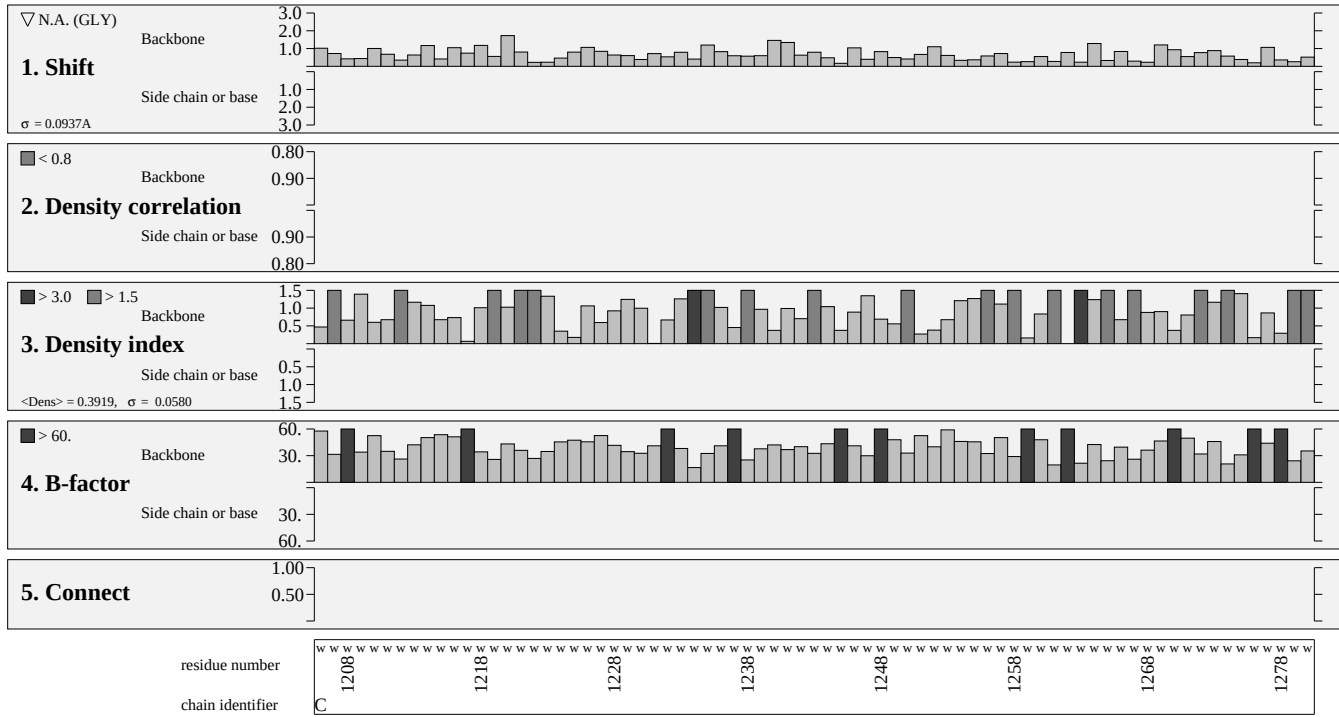
### Local estimation (24)



# Structure Factor Check

## 2FBW

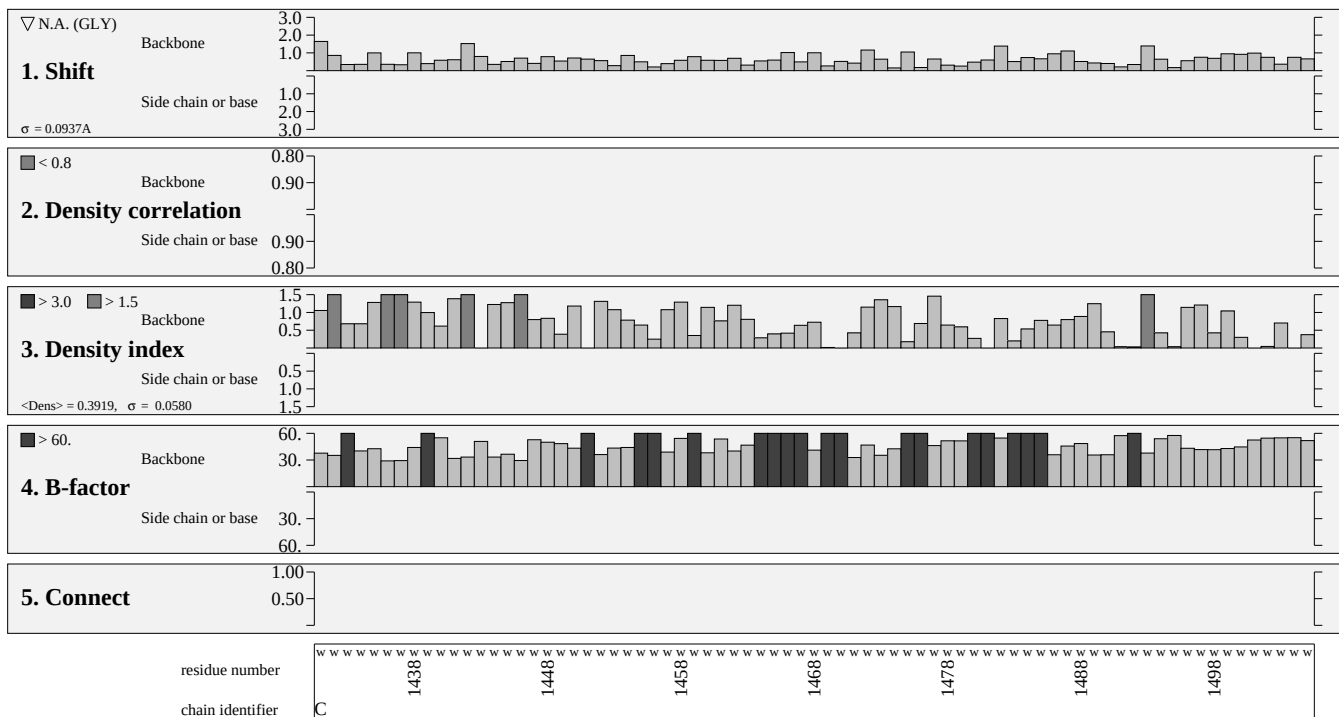
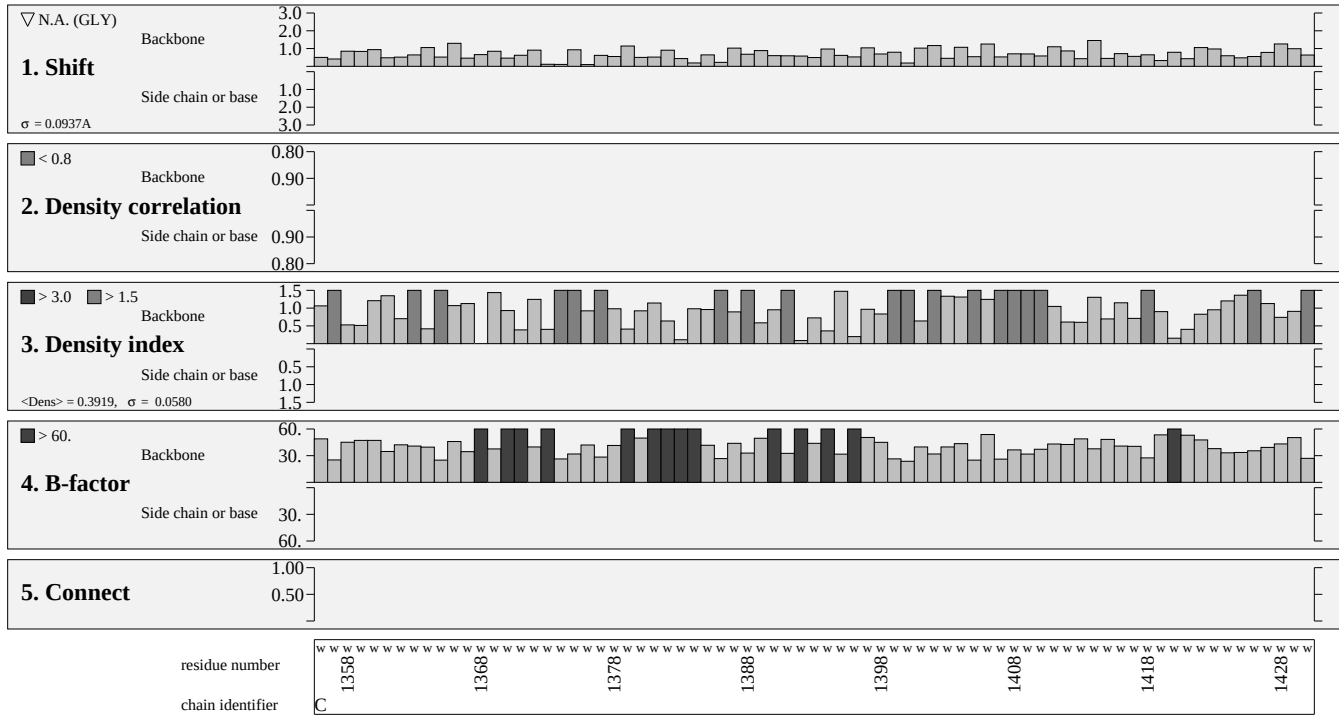
### Local estimation (25)



# Structure Factor Check

## 2FBW

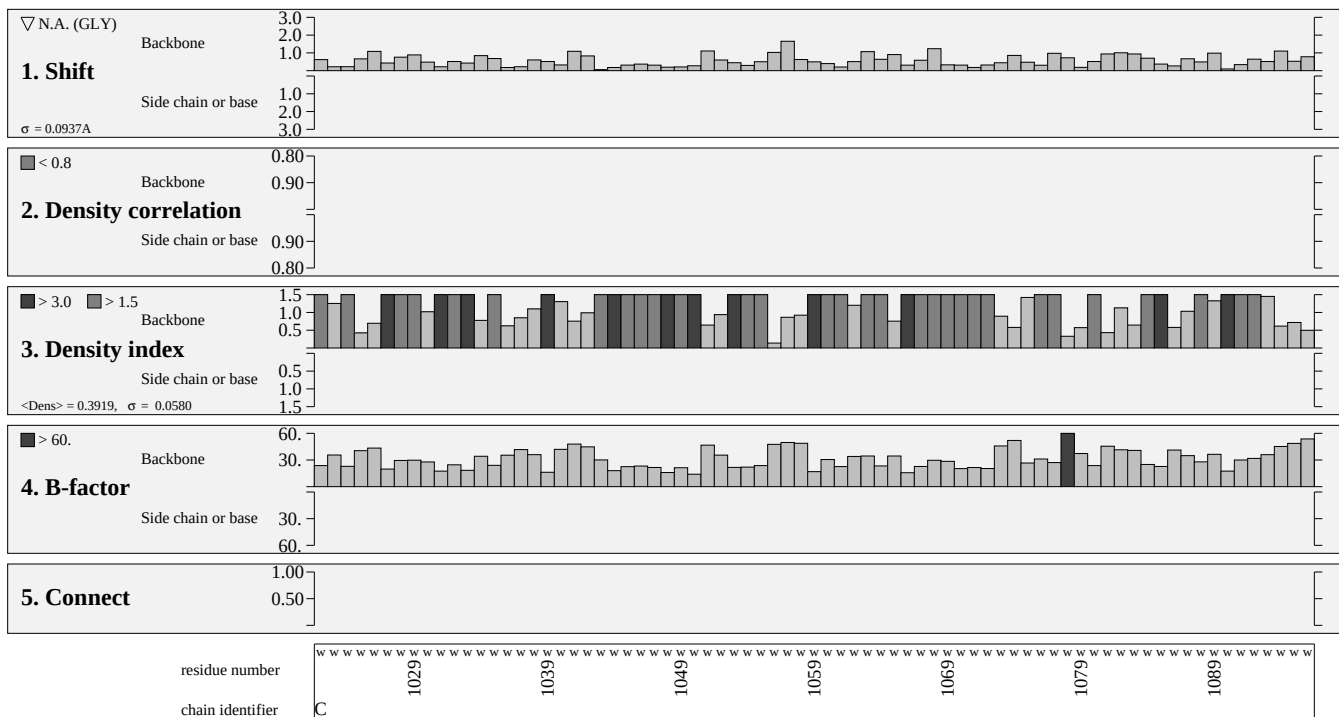
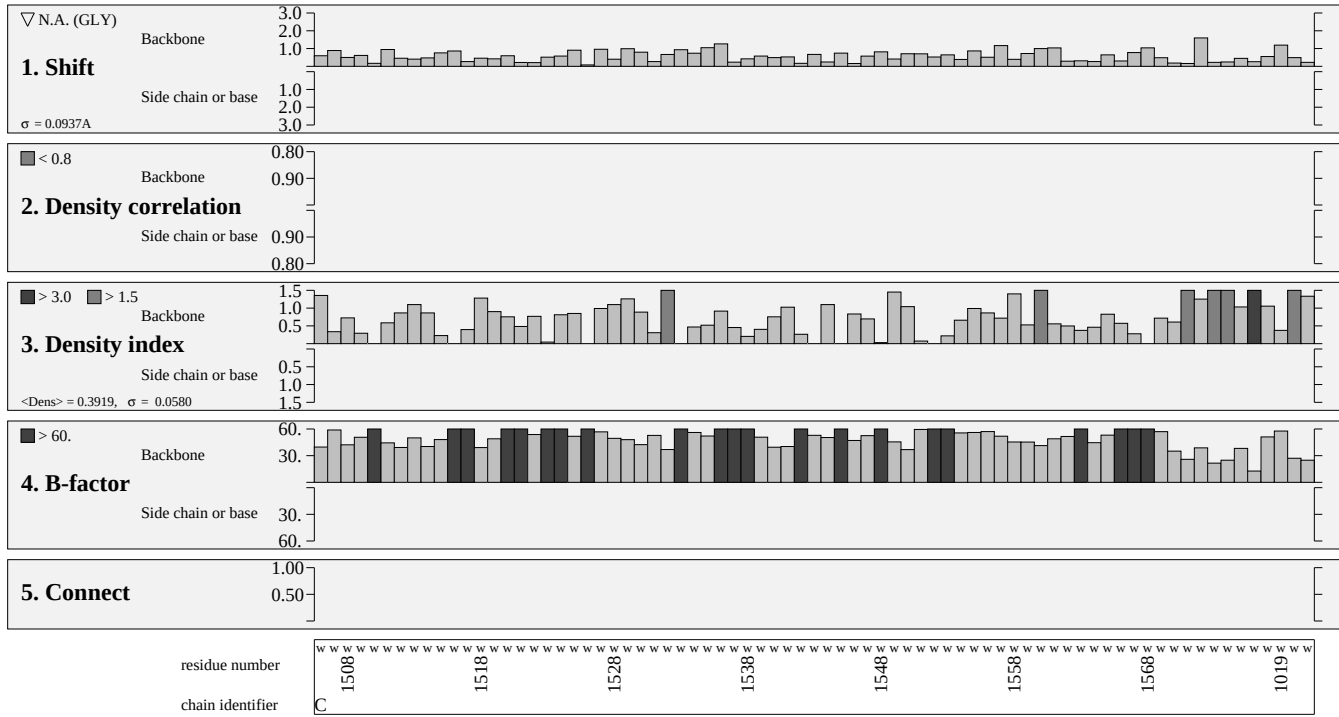
### Local estimation (26)



# Structure Factor Check

## 2FBW

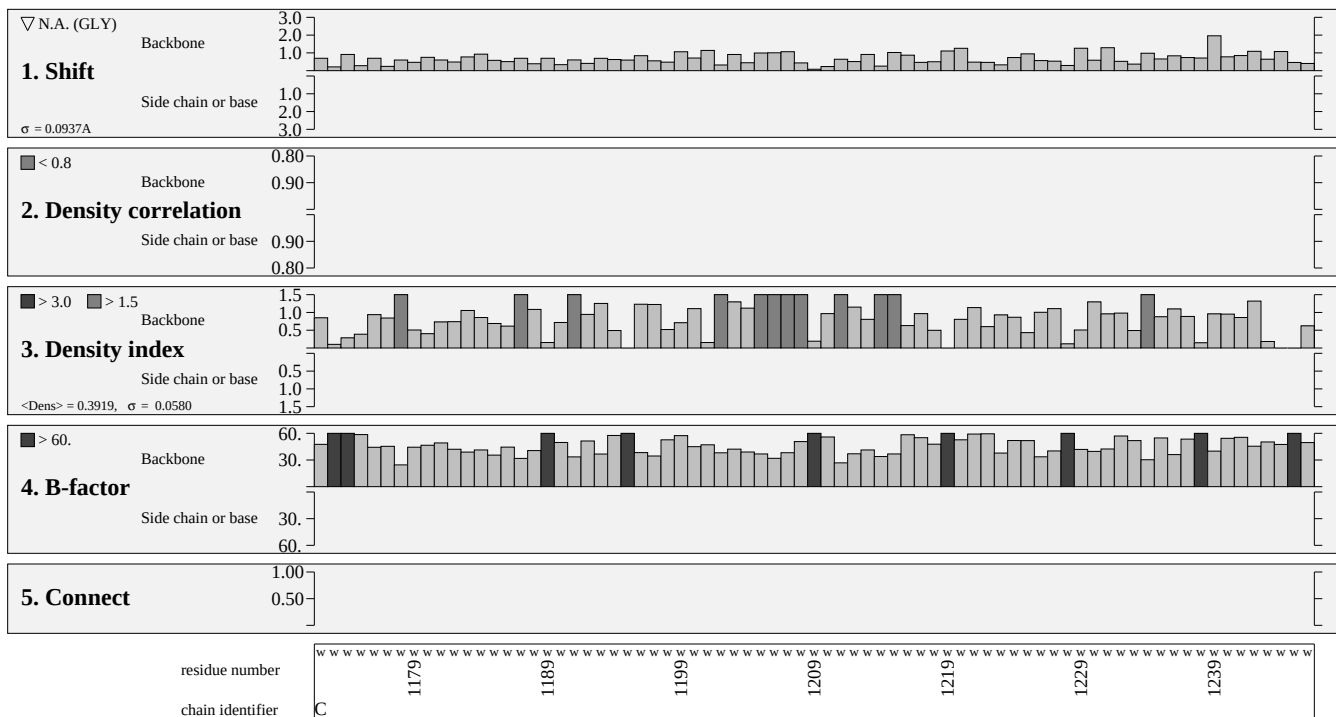
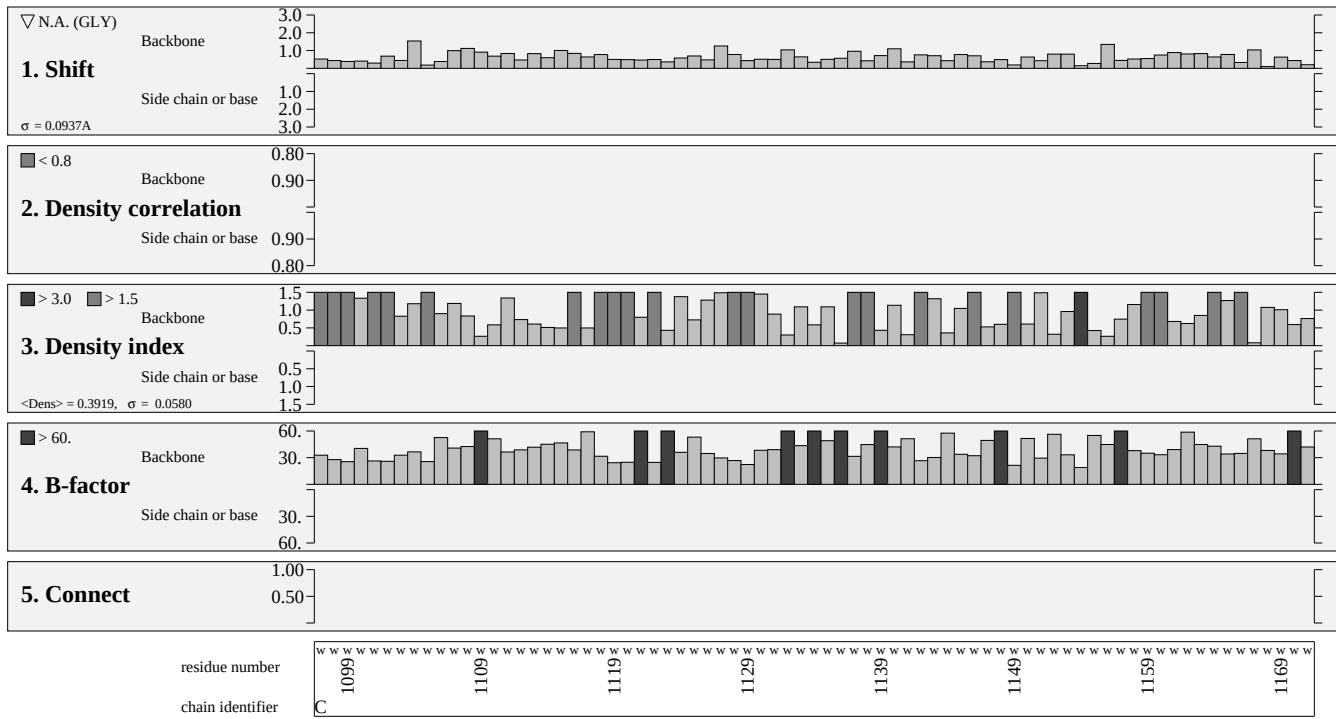
### Local estimation (27)



# Structure Factor Check

## 2FBW

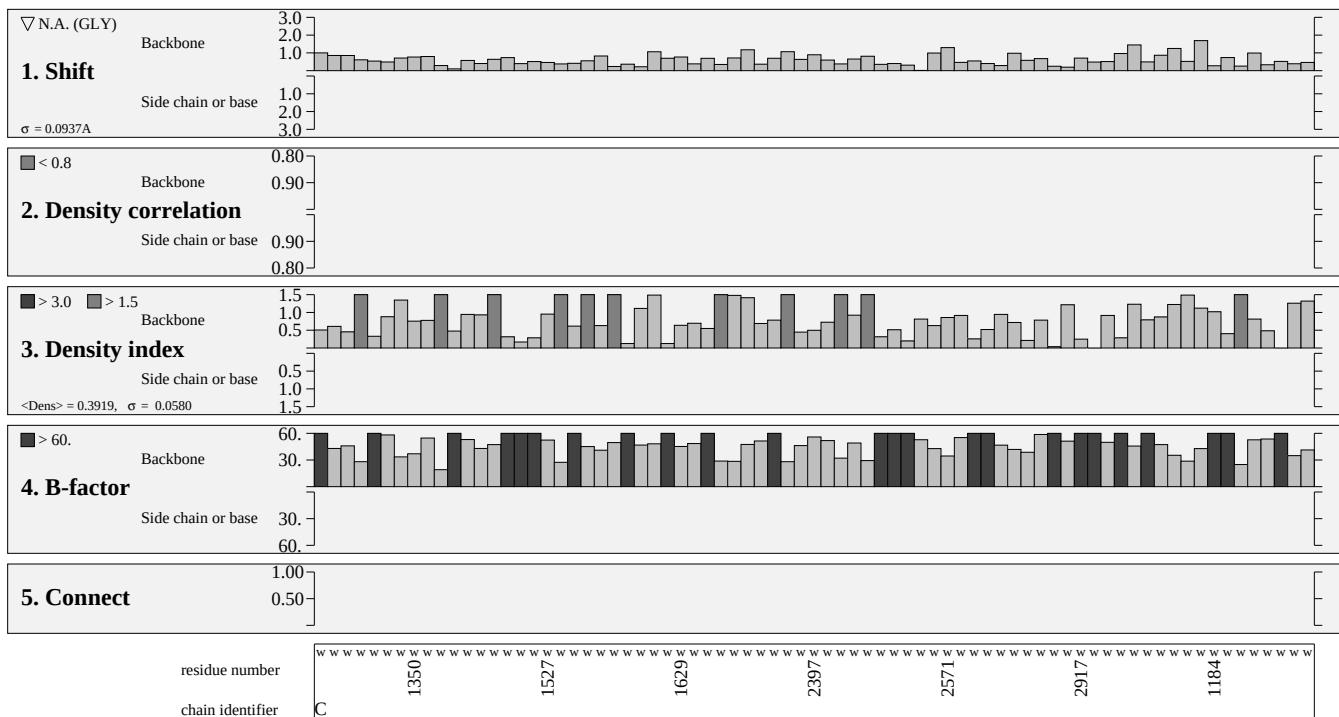
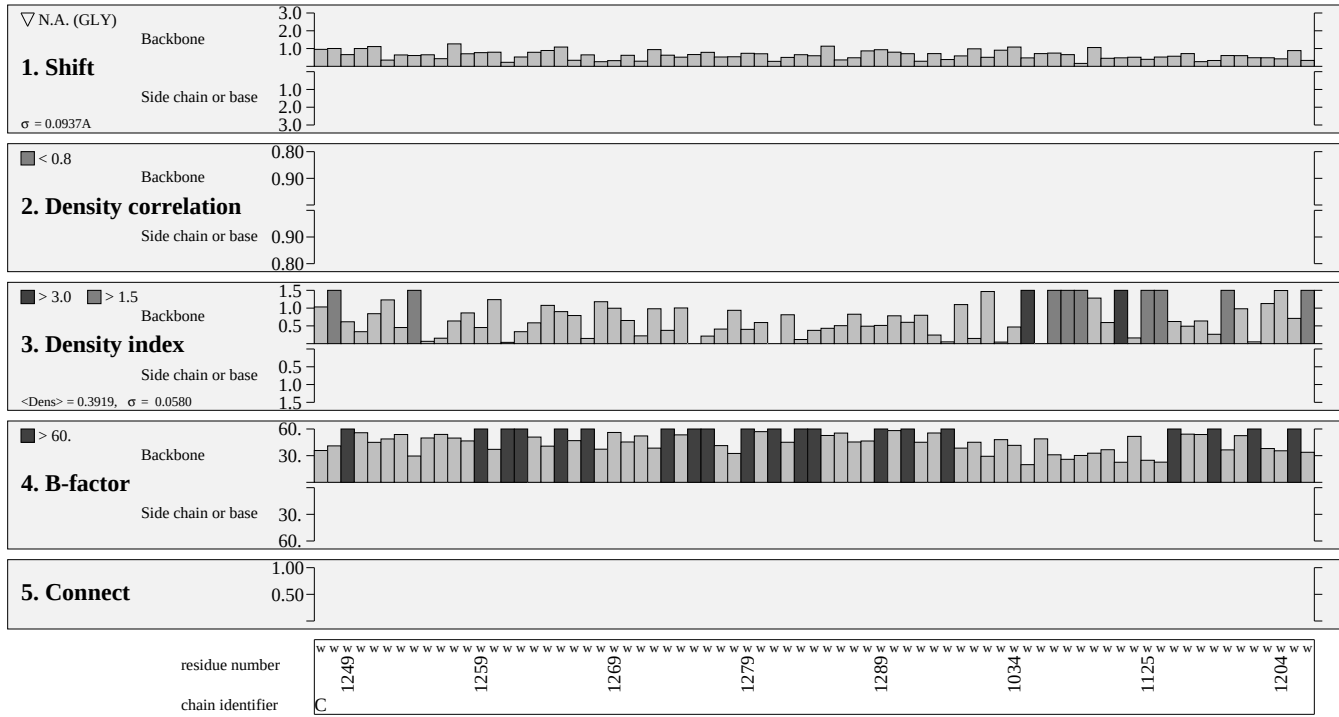
### Local estimation (28)



# Structure Factor Check

## 2FBW

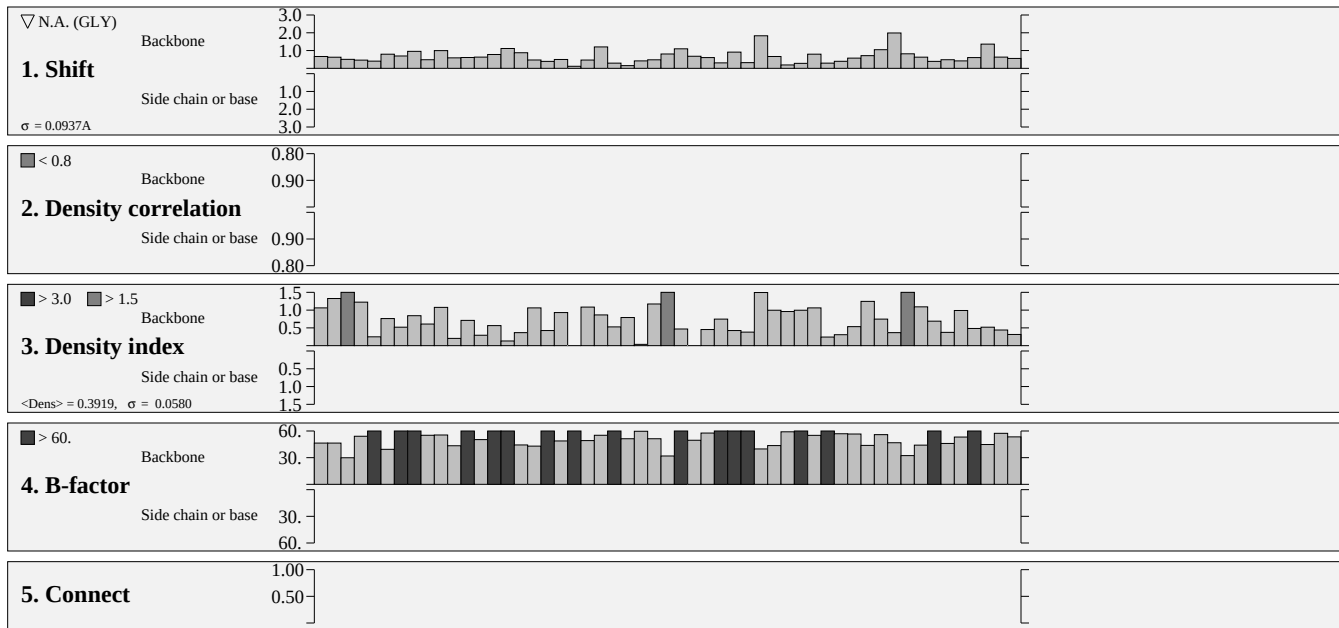
### Local estimation (29)



# Structure Factor Check

## 2FBW

### Local estimation (30)



|                  |      |      |      |      |      |      |
|------------------|------|------|------|------|------|------|
| residue number   | 1283 | 1348 | 1486 | 1598 | 2448 | 2976 |
| chain identifier | C    |      |      |      |      |      |