

# IUCrJ

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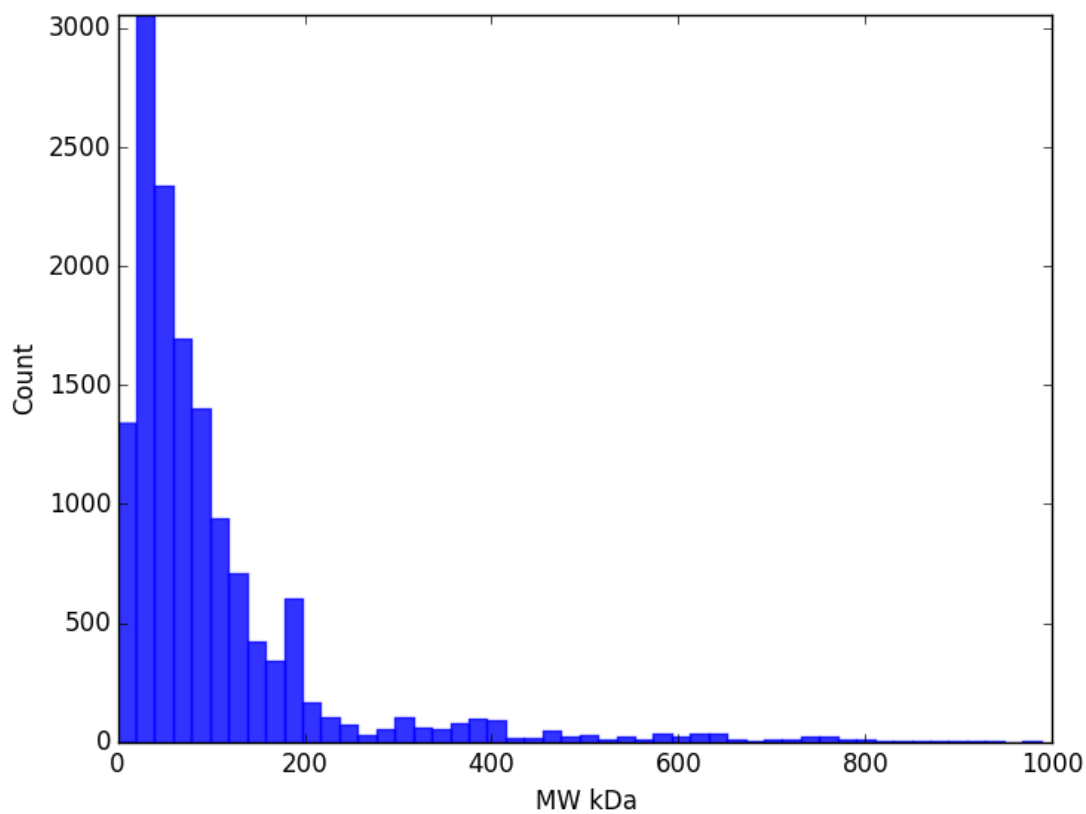
**Supporting information for article:**

**A comparative anatomy of protein crystals: lessons from the automatic processing of 56 000 samples**

**Olof Svensson, Maciej Gilski, Didier Nurizzo and Matthew W. Bowler**

**Table S1** Verification of calculated molecular weights compared to known samples.

Acronym	Actual MW (kDa)	Calculated MW (kDa)
Lyso	14.2	12
RICK	34.85	48
CRU	41.31	43
Erk2	42.24	41
DPP	162	200
TbrATP	922.6	800
GroEL	910	700

**Figure S1** Histogram of molecular weights calculated from samples processed on MASSIF-1.