

# Measures with error bars (here: $S_{CH}$ )

$$Q = \sum_i^N \frac{1}{N} \frac{|S_{CH,exp,i} - S_{CH,sim,i}|}{\sqrt{\Delta_{exp,i}^2 + \Delta_{sim,i}^2}}$$

“How many times the error is the deviation”

Suggestion: cap the scale at 3

**Option 1: Average over entire molecule**

- + one number
- tails will dominate

**Option 2: headgroup Q1 and tails (both) Q2**

- + more balanced
- two numbers
- doesn't separate tails

**Option 3: headgroup Q1**

**+ sn1 Q2 + sn2 Q3**

- + separates tails
- 3 numbers

# Measures without error bars (Form factor)

Can't use the SIMtoEXP (closest thing to a standard):

$$k_e = \frac{\sum_{i=1}^{N_q} \frac{|F_s(q_i)| |F_e(q_i)|}{(\Delta F_e(q_i))^2}}{\sum_{i=1}^{N_q} \frac{|F_e(q_i)|^2}{(\Delta F_e(q_i))^2}}$$

$$\chi^2 = \frac{\sqrt{\sum_{i=1}^{N_q} (|F_s(q_i)| - k_e |F_e(q_i)|)^2 / (\Delta F_e(q_i))^2}}{\sqrt{N_q - 1}}$$

- Experimental and simulation accuracy will differ based on curve location. Currently **both** unknown
- Ranking will be based on rmse
  - will depend on choice of q cutoff
  - Will depend on choice of how to scale
  - + will take heights and zeros into account in “same units”

# Suggestions

- Obtain the scaling from fit to first lobe
- Calculate rmse to  $q=0.5$ 
  - + fit based on most reliable data
  - fit and rmse not from same measure
- Obtain the scaling from minimizing the rmse up to  $q=0.5$ 
  - + fit and rmse consistent
  - all points treated as equally accurate in the fit