Co-solute	Effect on disordered proteins	Effect on denatured globular proteins	Effect on other systems
Dextrans	1. No effects on $p27^{KIP1}$ or c-FOS [3]. 2. Folding of TCAM RNAse [4].	1. Increased refolding (k_f) of lysosyme [1]. 2. Change to molten globular state of cyt C [2].	1. Decreased unfolding (k_u) of GAPDH [5].
Ficolls	1. No effects on $p27^{KIP1}$ or c-FOS [3]. 2. No effects on TCAM RNAse stability (K_{D-N}) [4].	1. Minor effect on FKB12 stability [6]. 2. Minor effect on refolding (k_f) of lysosymes [7].	
EG/PEGs	1. Accelerated fibrillation of α-synuclein [10].	1. Collapse of the coil state (decreased k_f) of CI2 at high PEG [8]. 2. Increased folding rate (k_f) of CI2 at low PEG [8]. 3. Decreased folding rate (k_f) of CspB (EG) [9].	1. Decreased unfolding (k_u) of GAPDH [5].
Glycerol	1. Accelerated fibrillation of α-synuclein [10].	1. Stabilised native state of RNAseA [11]. 2. Increased refolding(k_f) of GS with GroEL [12]. 3. Increased refolding (k_f) of MHD [13].	
Glucose	2. Induced collapse of α-synuclein [14].	1. Folding of Fe cyt C [14].	1. Stabilised dimeric chymotrypsin [15].
Sucrose	1. Contracted RCAM RNAse [17].	1. Increased refolding (k_f) of GS [12]. 2. Increased refolding (k_f) of CI2 [16]. 3. Increased refolding (k_f) of MHD [13].	2. Stabilised dimeric chymotrypsin [15].

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