

Searching for Silver Bullets

An Alternative Strategy for Crystallizing Macromolecules

Bob Cudney¹ and Alexander McPherson²

¹Hampton Research, Aliso Viejo, California ²University of California Irvine, Irvine, California

What we did

Based on a hypothesis that various small molecules might establish stabilizing, intermolecular, non covalent crosslinks in protein crystals and thereby promote lattice formation, we carried out three separate experiments. We assessed the impact of 200 chemicals on the propensity of 81 different proteins and viruses to crystallize. The experiments were comprised of 18,240 vapor diffusion trials. A salient feature of the experiments was that, aside from the inclusion of the reagent mixes, only two base crystallization conditions were used, 30% PEG 3350, and 50% Tacsimate™, both buffered at pH 7.

How we did it

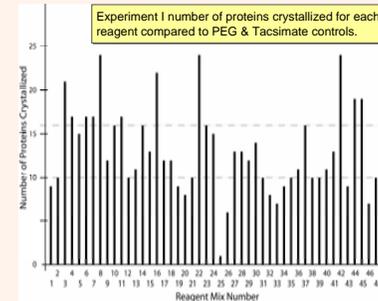
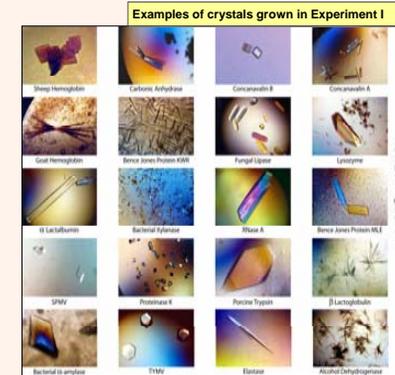
- All reagents were:
 - Manually dispensed sitting drop vapor diffusion
 - Incubated at 25° Celsius
 - Reviewed manually at 20 and 200x using polarized light at 4, 8 and 12 weeks
 - Expt I used 3 different reservoir concentrations, Expt II used 2, and Expt III used 1
- All experiments used two fundamental conditions:
 - 30% w/v PEG 3,350, 0.1 M Hepes pH 7.0
 - 50% Tacsimate™ pH 7.0
- A total of 200 chemicals compounds were tested.
- Screened a total of 81 total proteins in all 3 expts: Expt I=60; Expt II=67; Expt III=66

Expt I & II used 48 different reagent mixes, 2 of which were PEG and Tacsimate controls. The remaining 46 reagents were comprised of 3 to 20 different chemicals, on average 3 to 6 chemicals in each reagent.

Expt III used 24 different bioactive compound reagent mixes, 1 of which was a PEG control. Each reagent is comprised of 4 to 5 bioactive compounds such as coenzymes, prosthetic groups, inhibitors, drugs, effectors, nucleotides, amino acids/peptides, and sugars/oligosaccharides. Sample arrays set in duplicate.

A total of 18,240 experiment drops were set for three groups of experiments.

Experimental Results

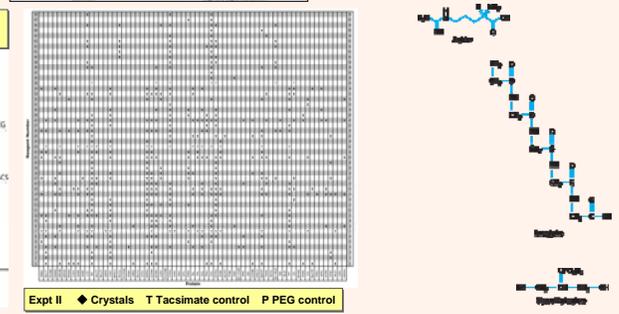
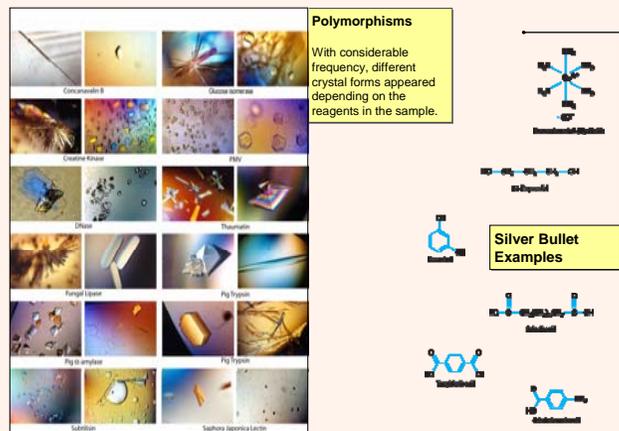


Experiment I

- spermidine, spermine, cadaverine, putrescine, 1,8 diamino octane
- 6-amino hexanoic acid, 3-aminopropionic acid, 4-aminobutyric acid
- oxamic acid, sulfanilic acid, 4-aminobenzoic acid
- Mix #2 and Mix #3
- glutamic acid, hexadecanoic acid, dodecanoic acid, fumaric acid
- oxalic acid, malic acid, oxaloacetic acid, terephthalic acid
- Mix #5 and Mix #6
- Pantothine, Inosine
- poly-L-lysine, poly-L-ornithine, poly-DL-alanine
- poly-L-glutamic acid, poly-DL-alanine
- Mix #9 and Mix #10
- Protamine
- heparin, dextran sulfate
- dipAH, dipAII
- ds-T4, dip12
- pyrophosphate, tetraphosphate
- ornithine, arginine, asparagine, glutamine
- Tacsimate, PEG 3350
- glycerol, sucrose, sorbitol
- TMAO, pectine
- sarcosine, glutamate, glycine, betaine
- hexanoic acid, glyoxylic acid, indolebutyric acid, D-hydroxyphenylacetic acid
- palmitic acid, octanoic acid, stearoylamine
- butyric acid, isosorbide, glycerol-2-phosphate
- potassium acetate, goldodium chloride, cobalt hexamine
- hexadecyl trimethylammonium bromide
- Tacsimate, Mix #8
- Tacsimate, poly-L-ornithine, lysine, glutamic acid, alanine
- Tacsimate, Mix #2
- Tacsimate, Mix #3
- Tacsimate, Mix #5
- Tacsimate, Mix #5
- Tacsimate, Mix #1
- Tacsimate, Mix #19
- Tacsimate, Mix #17
- Tacsimate, Mix #13
- PEG 3350
- Tacsimate, dipAH, dipAII
- Tacsimate
- Tacsimate, dip12, dipT4
- Mix #8, Mix #9
- Mix #8, Mix #9
- Mix #1, Mix #3
- Mix #1, Mix #3
- The 20 amino acids
- Tacsimate, the 20 amino acids
- Tacsimate, the 20 amino acids
- Tacsimate, proline

Experiment II

- Schardinger's dextrin
- NDSB-201
- NDSB-195, NDSB-201, NDSB-211, NDSB-221, NDSB-256
- lactic acid
- streptomycin
- lysozyme, phycoerythrin
- tetraphosphoric acid, cobalt hexamine
- malic acid, myo-inositol, phytic acid
- Mix #8 and Mix #9
- pipерidine ethane sulfonic acid, anthraquinone disulfonic acid
- inositol, tetrahydroborate
- pyromellitic acid, 2,2-thiodiglycolic acid, barbituric acid, terephthalic acid
- malonic acid, malic acid, glutaric acid, pimelic acid, succinic acid
- suberic acid, sebacic acid, hexadecanoic acid, dodecanoic acid
- malonic acid, phthalic acid, sebacic acid, hexadecanoic acid
- malic acid, glutaric acid, succinic acid, sebacic acid, dodecanoic acid, oxamic acid
- malonase, succinic, malicose, maltose, cellobiose
- arabinose, maltotriose, melitiose
- melitiose, starchyose
- cytidine, 2-hydroxypropyl-β-cyclodextrin, 2,6-di-O-methyl-β-cyclodextrin
- D, L, tetra, pentaglycine
- 3-phenylalanine, 1,2-hydroxyethyl-β-cyclodextrin, MPD, 1,6-hexanediol
- PEG 3350
- hexamethylenediamine, 1,4-diaminobutane, spermine, 1,8-diaminooctane, cadaverine
- pyrophosphate, phosphoric acid triethyl ester, phytic acid
- Mix #4 and Mix #10
- NDSB-201 and Mix #13
- Mix #15 and Mix #16
- phenyl urea, sodium L-pentanesulfonate, sulfanilic acid, salicin, pentamercaptoacetic acid
- oxamic acid, fumarate, fumaramide, putrescine, pentane dicarboxylic acid
- isoxanoic acid, PAA, sulfonic acid, 3-oxobutyric acid, pentane dicarboxylic acid
- phosphoglycinate, Anapoe-309, MEGA-9, nonyl-β-D-glucoside
- inosinic acid, malic acid, pyromelic acid, terephthalic acid
- phosphoglycolic, phytic acid, anthraquinone disulfonic acid, barbituric acid, tetraphosphoric acid
- Tacsimate and Mix #2
- Tacsimate and Mix #5
- Tacsimate and Mix #9
- Tacsimate and Mix #10
- Tacsimate and Mix #12
- Tacsimate and Mix #13
- Tacsimate and Mix #14
- Tacsimate and Mix #15
- Mix #1, Mix #3
- Mix #1, Mix #3
- The 20 amino acids
- Tacsimate, the 20 amino acids
- Tacsimate and Mix #24
- Tacsimate and Mix #21



Experiment III - Bioactive Compounds

- S-adenosylmethionine, β-glycerophosphate, starchyose
- bovine phosphotungstic pyruvic, melitiose
- FAD, phosphoglycolic acid, spermine
- Rhymine, pyrophosphate, glyoxylic acid, Schardinger's dextrin
- NAD, Inosine-1-β-diphosphate, spermidine
- γ-butyryl-β-phosphate, pyruvic acid, cyclodextrin
- ADP, AMP, LAMP
- GMP, GDP, nucleoside acid
- GTP, cTMP, chepironazine
- ATP, cCMP, estadiol, galactose
- Tacsimate
- spic acid, creatine, glutathione, parathetic acid
- acetaminophen, bay labate, glutate acid
- lavin mononucleotide, trolean, acetyl choline
- protonin A, GMP, lactic acid, glucose
- ADP, tetrahydrolic acid, succinic acid, glucose
- riboflavin, phosphoryl choline, raffinose
- GMP, cholesteryl, rhymine, oxamic acid
- phosphocreatine, benzylarginine ethyl ester, phenobarbital, tetraoctyl
- UTP, kamronin, melitiose, inosopin
- phosphorylthiose-1-phosphate, malic acid, n-acetyl-D-glucosamine
- UMP, ribose, phytic acid, pantoic acid
- The 20 amino acids
- PEG 3350

Reagent Mix Expt III

- Positive charge centers, Diols, Dicarboxylic acids, Mixed amino and carboxylic acids, Polar amino acids, Glycine oligomers, Organic acids, Charge symmetric, and Diamines

Condensation of Statistics

	Experiment			
	All	I	II	III
How many proteins were investigated?	81	60	67	66
How many proteins crystallized?	65	48	50	50
How many proteins crystallized in Polyethylene glycol controls?	20	11	14	13
in Tacsimate controls?	6	5	4	N/A
in both controls?	5	5	3	N/A
Total controls	31	21	21	13
How many proteins crystallized only in the presence of some reagent mix?	35	28	29	37
How many reagents exceeded the Polyethylene glycol controls?	11	7	4	4
Tacsimate controls?	7	1	N/A	N/A
How many proteins that did NOT crystallize in controls, crystallized in ONLY				
One reagent mix?	4	7	13	13
Two reagent mixes?	2	4	9	9
Three reagent mixes?	1	4	6	6

What happened

Overall, 65 proteins (85%) were crystallized. Most significant was that 35 of the 65 (54%) crystallized only in the presence of one or more reagent mixes, but not in control samples lacking any additives. Among the most promising types of reagent mixes were those composed of polyvalent, charged groups, such as di and tri carboxylic acids, diamino compounds, molecules bearing one or more sulfonyl or phosphate groups, and a broad range of common biochemicals, coenzymes, biological effectors, and ligands. Less promising reagent mixes were composed of osmolites, polyamines, detergents and sugars.

Summary

We propose that an alternate approach to crystallizing proteins might be developed, which employs a limited set of fundamental crystallization conditions combined with a broad screen of potentially useful small molecule additives.

What's next

The evaluation of more reagent mixes, the influence of pH and the use of common dehydrants.

Thank you Aaron Greenwood, Joe Luft & Peter Nguyen.