Corrigendum

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Corrigendum: Combining the Dihydrofolate Reductase Protein-Fragment Complementation Assay with Gene Deletions to Establish Genotype-to-Phenotype Maps of Protein Complexes and Interaction Networks

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When this protocol was first published, the concentrations for the reagents in the recipe entitled “Amino Acid Mix (10×)” were expressed as g/mL, but they should have been expressed as g/L. The authors apologize for this error. The HTML version of the recipe (doi: 10.1101/pdb.rec090308) has been corrected, as has the current PDF version of the protocol. A corrected version of the recipe is also included below.

Amino Acid Mix (10×)

Amino acid or nucleobase	Concentration (10×)
Adenine sulfate	0.4 g/L
Uracil	0.2 g/L
L-Tryptophan	0.4 g/L
L-Histidine hydrochloride	0.2 g/L
L-Arginine hydrochloride	0.2 g/L
L-Tyrosine	0.3 g/L
L-Leucine	0.6 g/L
L-Lysine hydrochloride	0.3 g/L
L-Phenylalanine	0.5 g/L
L-Glutamic acid	1.0 g/L
L-Asparagine	1.0 g/L
L-Valine	1.5 g/L
L-Threonine	2.0 g/L
L-Serine	3.75 g/L
L-Methionine	0.2 g/L

Dissolve the amino acids from the list above (except those to be excluded from any drop-out media) in distilled H₂O. Filter-sterilize, store at 4°C, and protect from light.

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