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## 1 Setup

This experiment tests the effectiveness of the Dijkstra reweighting scheme. It uses the same edge adding scheme (

Performance was evaluated using 5-fold CV with the GO label set from 2019 where all GO labels annotating less than 50 or greater than 1000 proteins are discarded (I use Lily’s code to do this).

## 2 Results

	Normalized DSD	Glide CW	Glide L3	Network Enhancement	No denoising
Dream 1	0.246	0.247	0.246	0.239	0.246
Dream 2	0.170	0.182	0.170	0.145	0.171
Dream 3	0.154	0.166	0.153	0.180	0.144
Dream 4	0.139	0.142	0.143	0.120	0.141
Dream 5	0.088	0.089	0.086	0.078	0.088
Dream 6	N/A	N/A	N/A	0.202	0.193
Average				0.161	