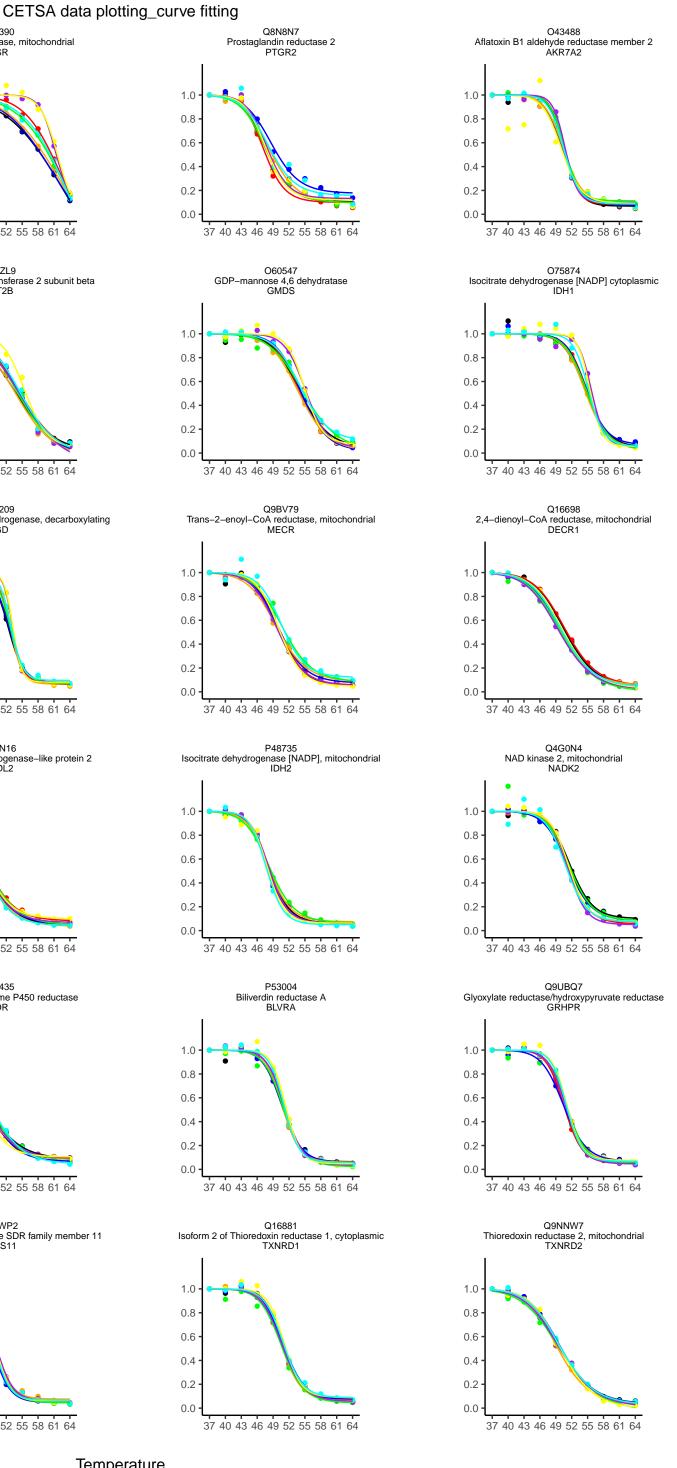
37 40 43 46 49 52 55 58 61 64

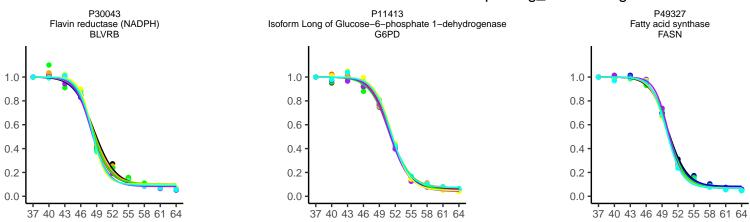


Temperature • Ctrl.1 • Ctrl.3 • Ctrl.5 • Ctrl.7

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37 40 43 46 49 52 55 58 61 64

CETSA data plotting_curve fitting



0.2

37 40 43 46 49 52 55 58 61 64

0.2

37 40 43 46 49 52 55 58 61 64

CETSA data plotting_curve fitting P36959 GMP reductase 1 GMPR Q8IW45 ATP-dependent (S)-NAD(P)H-hydrate dehydratase NAXD P04040 Catalase CAT O95544 Isoform 2 of NAD kinase NADK 1.0 8.0 8.0 8.0 0.8 0.6 0.6 0.6 0.6 0.4 0.4 0.4 0.4 0.2 0.2 0.2 0.2 0.0 0.0 0.0 0.0 37 40 43 46 49 52 55 58 61 64 37 40 43 46 49 52 55 58 61 64 37 40 43 46 49 52 55 58 61 64 37 40 43 46 49 52 55 58 61 64 O95825 Quinone oxidoreductase-like protein 1 CRYZL1 Q8NCW5 NAD(P)H-hydrate epimerase NAXE Q96A73 Putative monooxygenase p33MONOX KIAA1191 P48163 NADP-dependent malic enzyme ME1 1.0 1.0 1.0 1.0 8.0 8.0 0.8 8.0 0.6 0.6 0.6 0.6 0.4 0.4 0.4 0.4 0.2 0.2 0.2 0.2 0.0 0.0 0.0 0.0 37 40 43 46 49 52 55 58 61 64 37 40 43 46 49 52 55 58 61 64 37 40 43 46 49 52 55 58 61 64 37 40 43 46 49 52 55 58 61 64 Q7RTP6 Protein-methionine sulfoxide oxidase MICAL3 MICAL3 Q53H96 Pyrroline–5–carboxylate reductase 3 PYCR3 Q13630 GDP-L-fucose synthase TSTA3 1.0 1.0 1.0 8.0 8.0 8.0 0.6 0.6 0.6 0.4 0.4 0.4

0.2

37 40 43 46 49 52 55 58 61 64