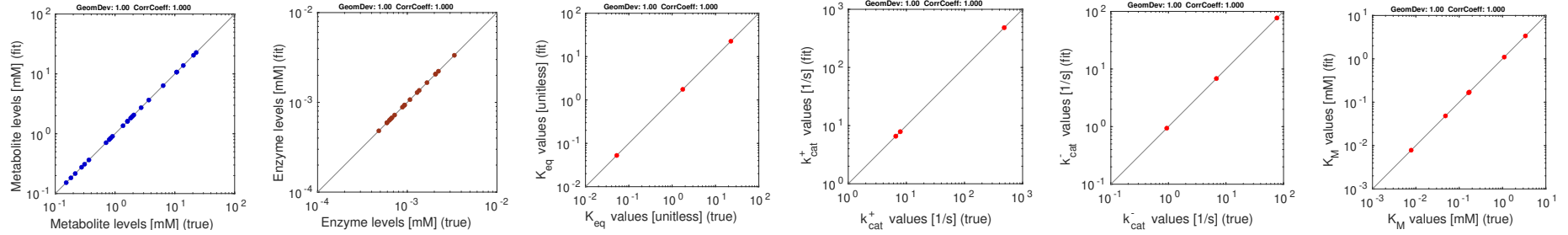


- 1 Three-chain model Simulations with $\alpha = 0.1$**
- 2 Three-chain point model Simulations with $\alpha = 0.1$, $\beta = 0$**

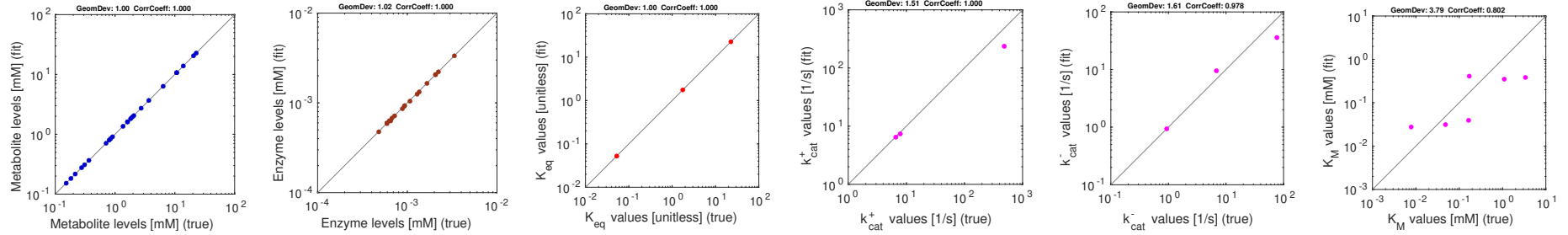
Three-chain model with artificial data (noise-free kinetic data, noise-free state data) - $\alpha = 0.1$, $\beta = 0$

(a) Metabolites (b) Enzymes (c) K_{eq} values (d) k_{cat}^+ values (e) k_{cat}^- values (f) K_M values

With kinetic data



With K_{eq} data only



Without kinetic data

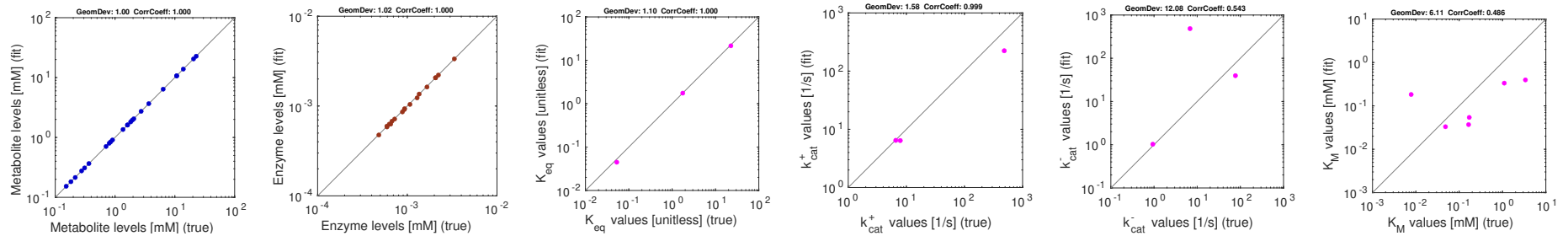


Figure 3: Model balancing results for Three-chain model with artificial data. The model structure is shown in Figure ?? . Each subfigure shows “true” values (x-axis) versus reconstructed values (y-axis). Similarities are quantified by geometric standard deviations (“GeomDev”) and Pearson correlation coefficients (“CorrCoeff”). (a) Metabolite levels. (b) Enzyme levels. (c)-(f) Different types of kinetic constants. Rows show different estimation scenarios (see Figure) Upper row: simple scenario S1 (noise-free artificial data, data for kinetic constants). Centre row: scenario S1K (noise-free artificial data, kinetic data given only for equilibrium constants). Lower row: scenario S2 (noise-free artificial data, no data for kinetic constants). Depending on the scenario, kinetic constants are either fitted (red dots) or predicted (magenta dots).

Three-chain model with artificial data (noisy kinetic data, noise-free state data) - $\alpha = 0.1$, $\beta = 0$

(a) Metabolites

(b) Enzymes

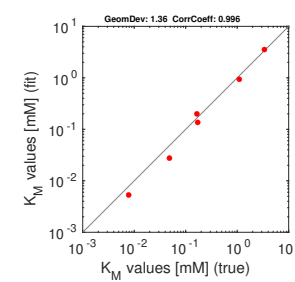
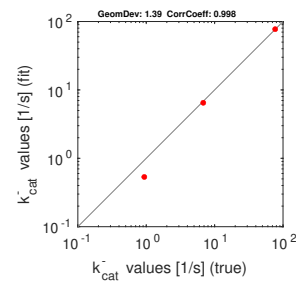
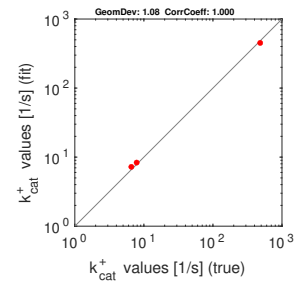
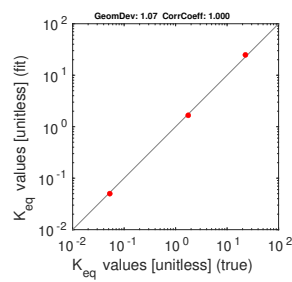
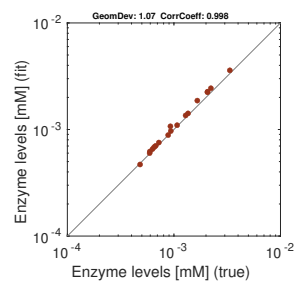
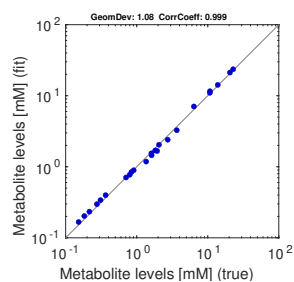
(c) K_{eq} values

(d) k_{cat}^+ values

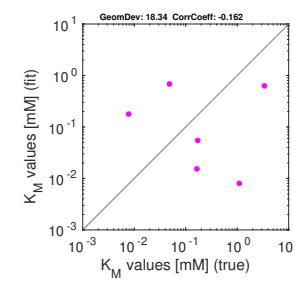
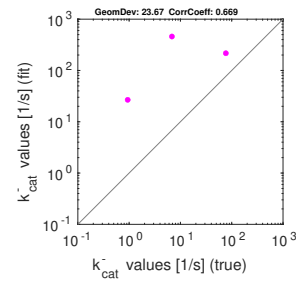
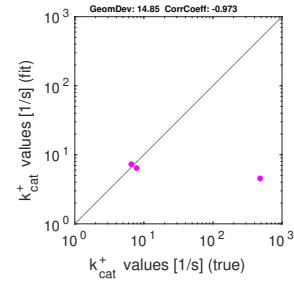
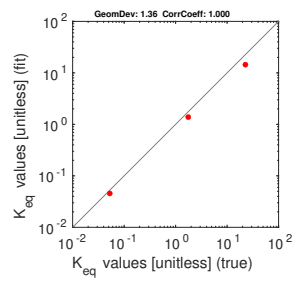
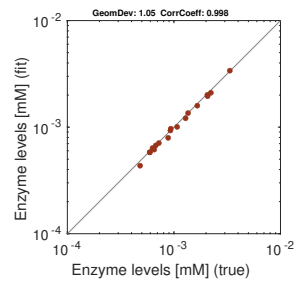
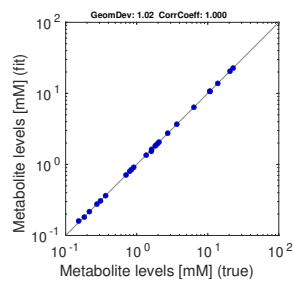
(e) k_{cat}^- values

(f) K_M values

With kinetic data



With K_{eq} data only



Without kinetic data

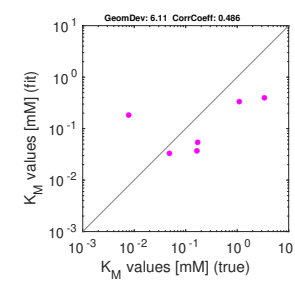
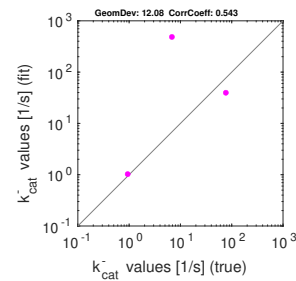
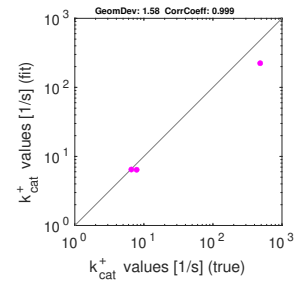
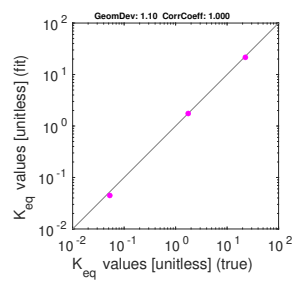
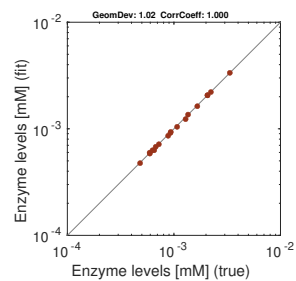
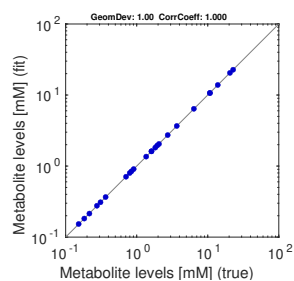


Figure 4: Same as Figure 8, with noisy kinetic data

Three-chain model with artificial data (noise-free kinetic data, noisy state data) - $\alpha = 0.1$, $\beta = 0$

(a) Metabolites

(b) Enzymes

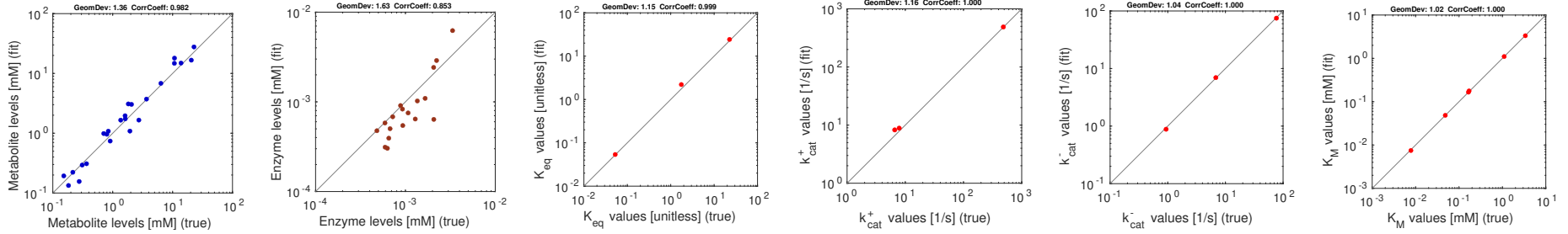
(c) K_{eq} values

(d) k_{cat}^+ values

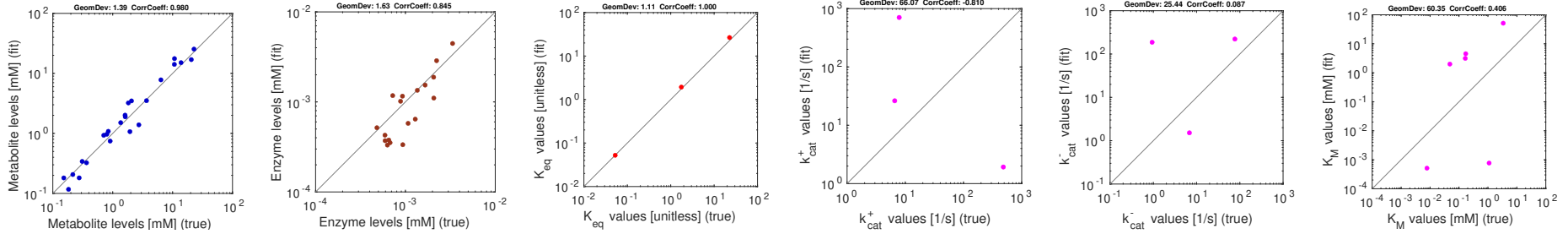
(e) k_{cat}^- values

(f) K_M values

With kinetic data



With K_{eq} data only



Without kinetic data

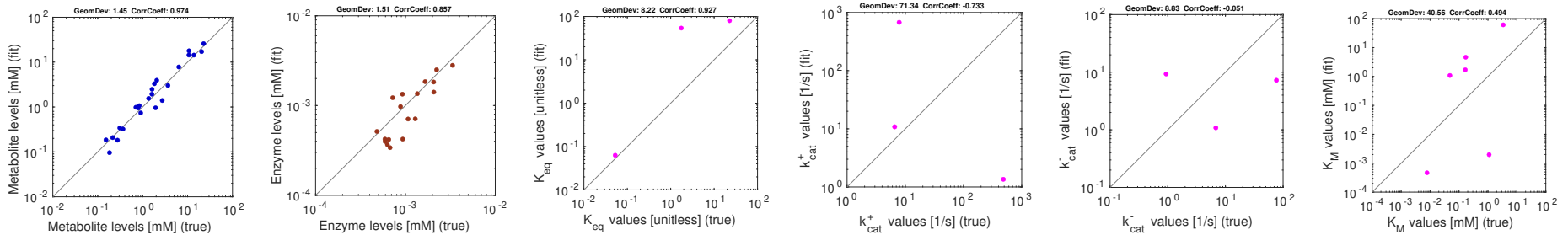


Figure 5: Results for Three-chain with noisy artificial data. Top row: estimation scenario S3 (noisy artificial data, data used for kinetic constants). Centre row: estimation scenario S3K (noisy artificial data, data for equilibrium constants only). Bottom row: estimation scenario S4 (noisy artificial data, no data for kinetic constants).

Three-chain model with artificial data (noisy kinetic data, noisy state data) - $\alpha = 0.1$, $\beta = 0$

(a) Metabolites

(b) Enzymes

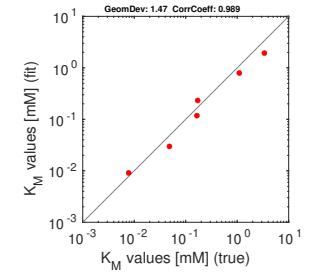
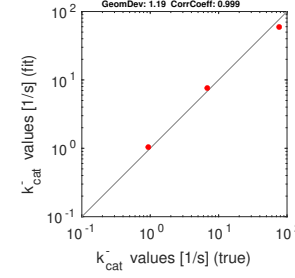
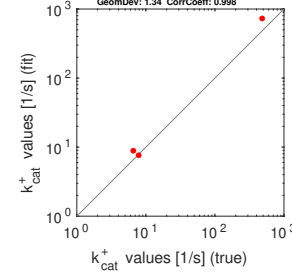
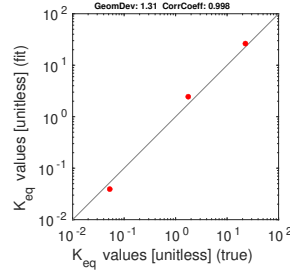
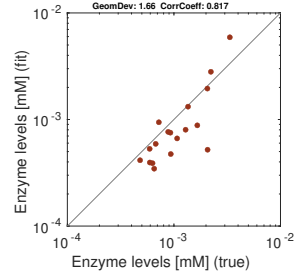
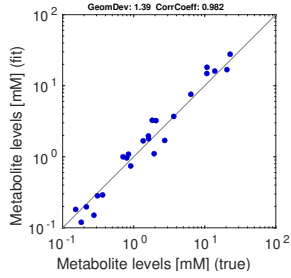
(c) K_{eq} values

(d) k_{cat}^+ values

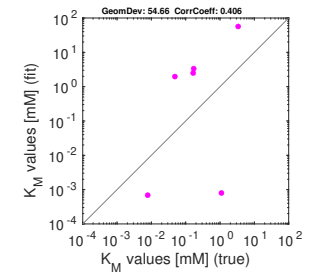
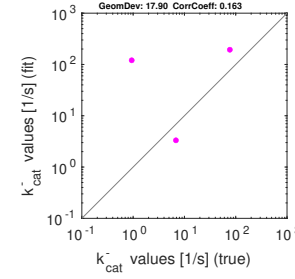
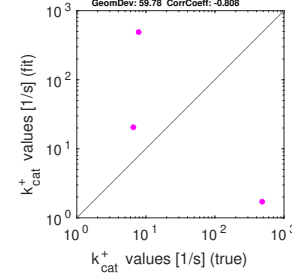
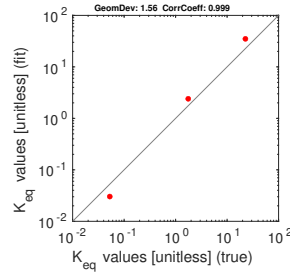
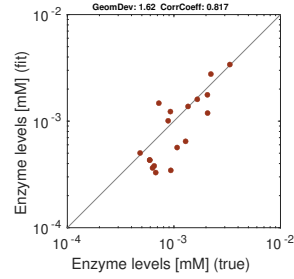
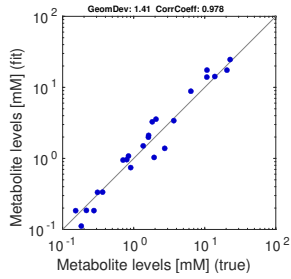
(e) k_{cat}^- values

(f) K_M values

With kinetic data



With K_{eq} data only



Without kinetic data

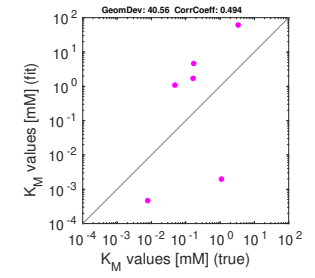
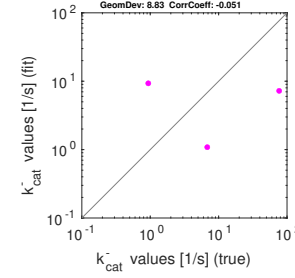
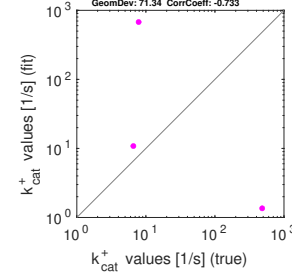
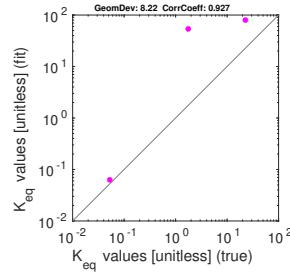
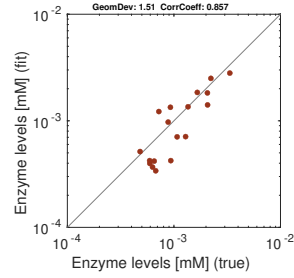
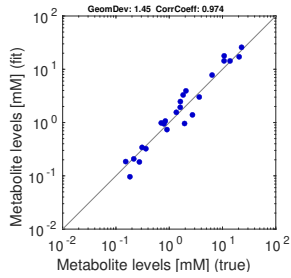
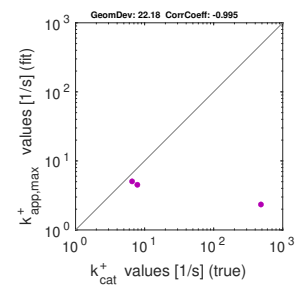


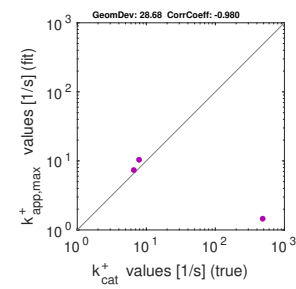
Figure 6: Same as Figure 10, with noisy kinetic data

k_{cat}^+ values

No metabolic noise

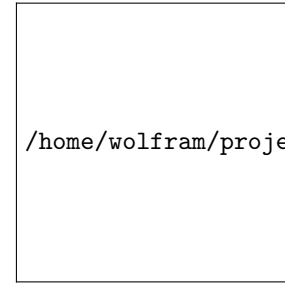


With metabolic noise



k_{cat}^- values

No metabolic noise



With metabolic noise

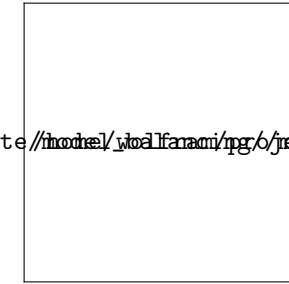


Figure 7: Catalytic constants in Three-chain (artificial data), estimated by kinetic profiling [?]. Note that k_{cat} values can only be estimated in the direction of fluxes (e.g. k_{cat}^+ for reactions with forward flux).

3 Three-chain model Simulations with $\alpha = 1$

4 Three-chain point model Simulations with $\alpha = 1$, $\beta = 0$

Three-chain model with artificial data (noise-free kinetic data, noise-free state data) - $\alpha = 1$, $\beta = 0$

(a) Metabolites

(b) Enzymes

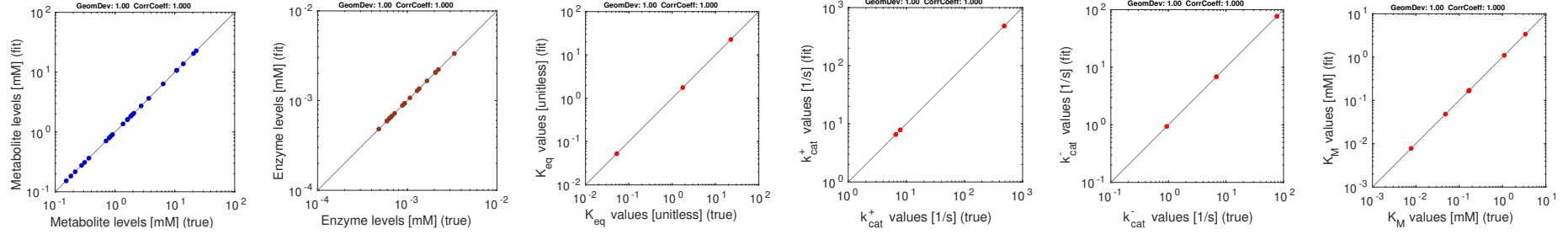
(c) K_{eq} values

(d) k_{cat}^+ values

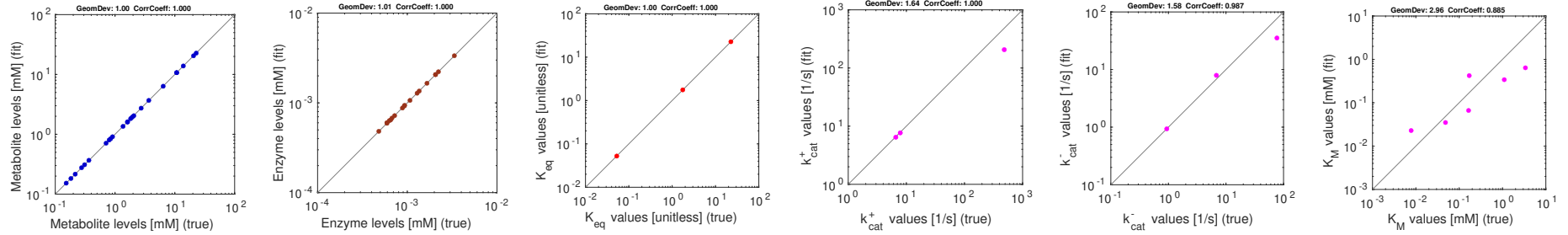
(e) k_{cat}^- values

(f) K_M values

With kinetic data



With K_{eq} data only



Without kinetic data

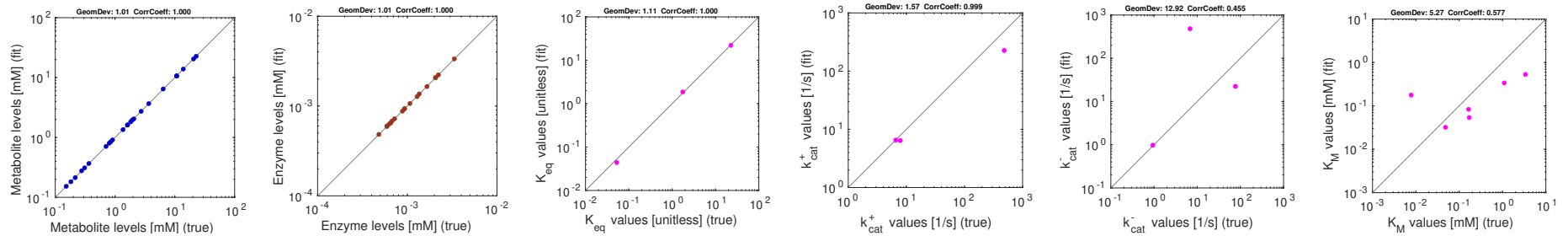


Figure 8: Model balancing results for Three-chain model with artificial data. The model structure is shown in Figure ?? . Each subfigure shows “true” values (x-axis) versus reconstructed values (y-axis). Similarities are quantified by geometric standard deviations (“GeomDev”) and Pearson correlation coefficients (“CorrCoeff”). (a) Metabolite levels. (b) Enzyme levels. (c)-(f) Different types of kinetic constants. Rows show different estimation scenarios (see Figure) Upper row: simple scenario S1 (noise-free artificial data, data for kinetic constants). Centre row: scenario S1K (noise-free artificial data, kinetic data given only for equilibrium constants). Lower row: scenario S2 (noise-free artificial data, no data for kinetic constants). Depending on the scenario, kinetic constants are either fitted (red dots) or predicted (magenta dots).

Three-chain model with artificial data (noisy kinetic data, noise-free state data) - $\alpha = 1$, $\beta = 0$

(a) Metabolites

(b) Enzymes

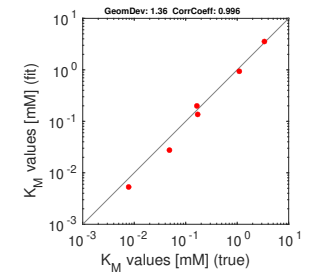
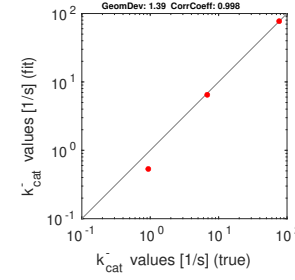
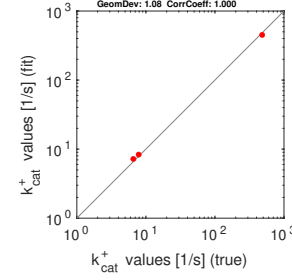
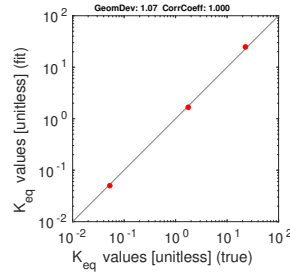
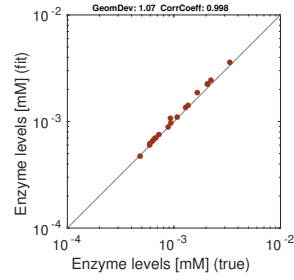
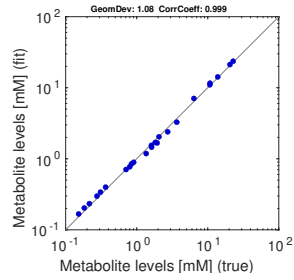
(c) K_{eq} values

(d) k_{cat}^+ values

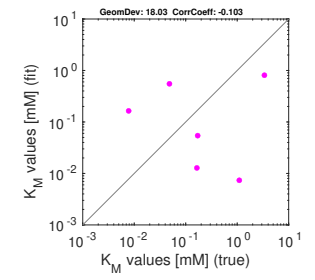
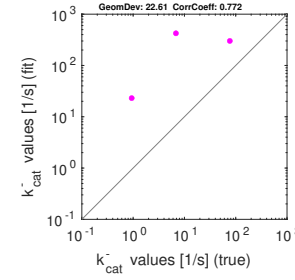
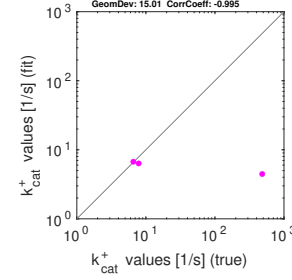
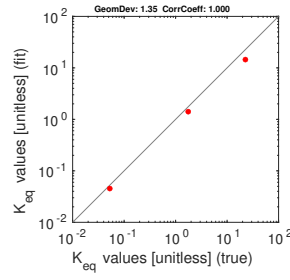
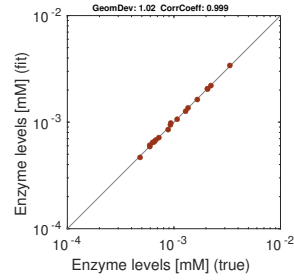
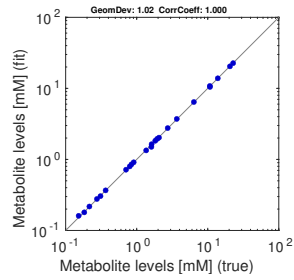
(e) k_{cat}^- values

(f) K_M values

With kinetic data



With K_{eq} data only



Without kinetic data

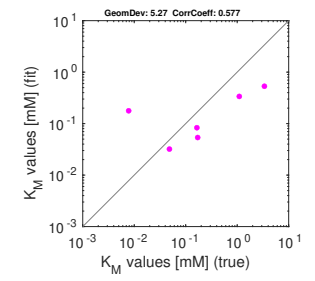
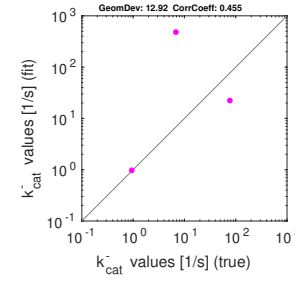
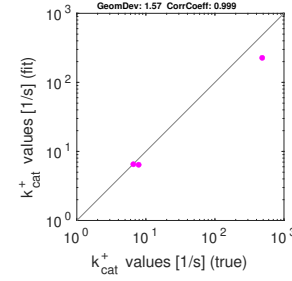
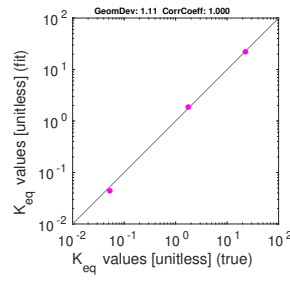
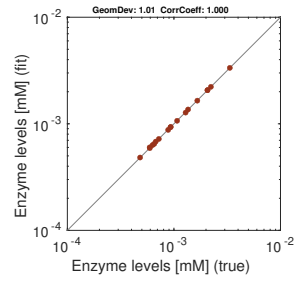
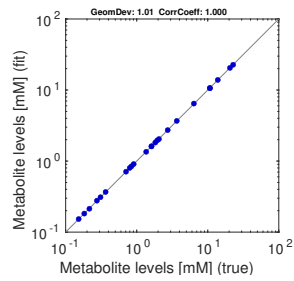


Figure 9: Same as Figure 8, with noisy kinetic data

Three-chain model with artificial data (noise-free kinetic data, noisy state data) - $\alpha = 1$, $\beta = 0$

(a) Metabolites

(b) Enzymes

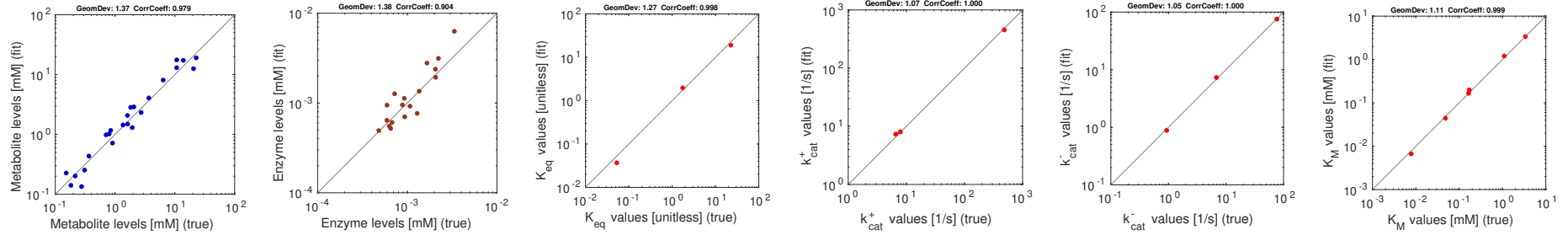
(c) K_{eq} values

(d) k_{cat}^+ values

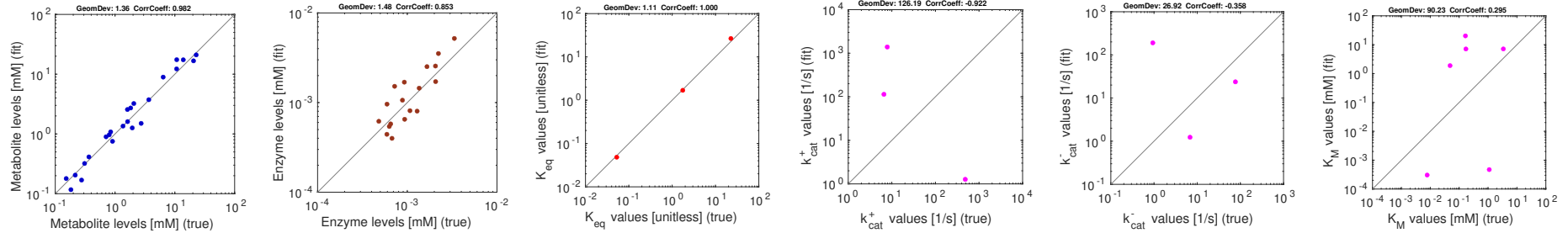
(e) k_{cat}^- values

(f) K_M values

With kinetic data



With K_{eq} data only



Without kinetic data

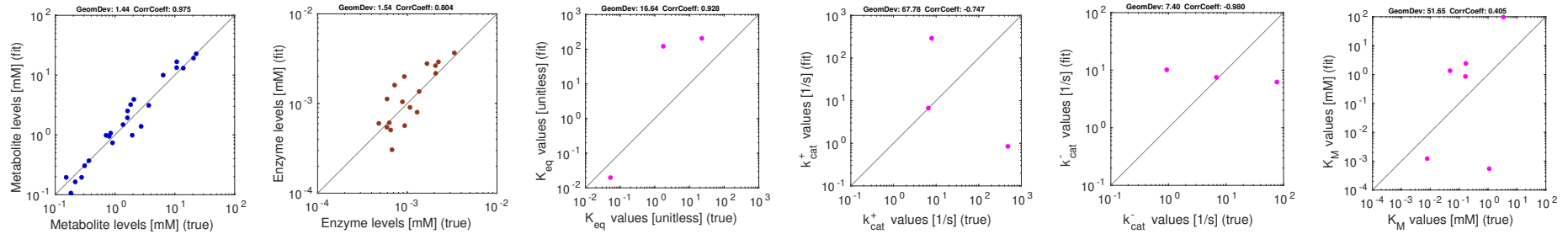


Figure 10: Results for Three-chain with noisy artificial data. Top row: estimation scenario S3 (noisy artificial data, data used for kinetic constants). Centre row: estimation scenario S3K (noisy artificial data, data for equilibrium constants only). Bottom row: estimation scenario S4 (noisy artificial data, no data for kinetic constants).

Three-chain model with artificial data (noisy kinetic data, noisy state data) - $\alpha = 1$, $\beta = 0$

(a) Metabolites

(b) Enzymes

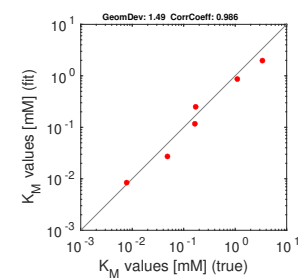
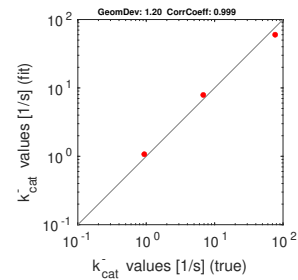
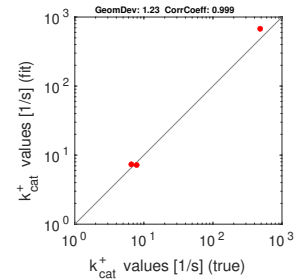
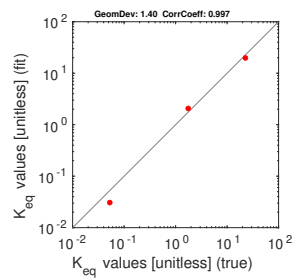
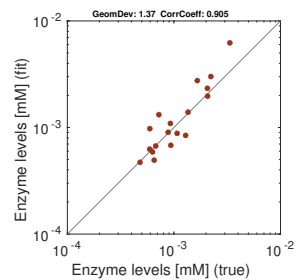
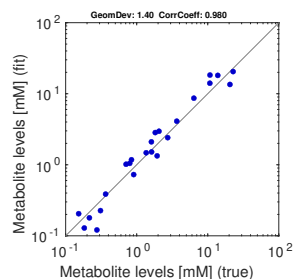
(c) K_{eq} values

(d) k_{cat}^+ values

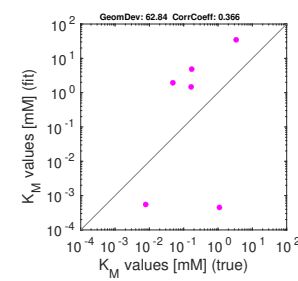
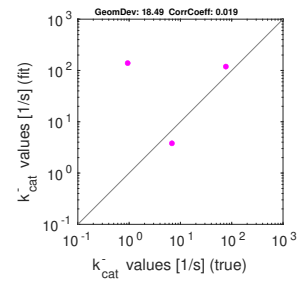
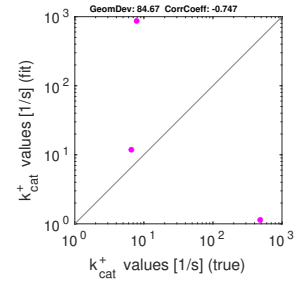
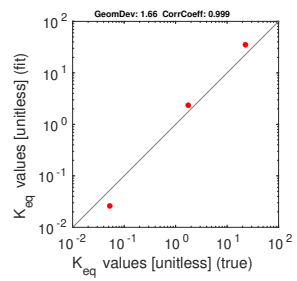
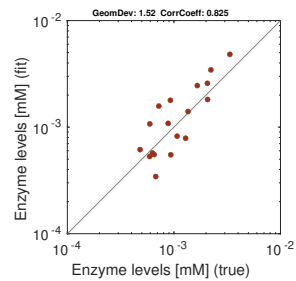
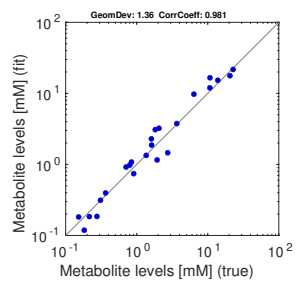
(e) k_{cat}^- values

(f) K_M values

With kinetic data



With K_{eq} data only



Without kinetic data

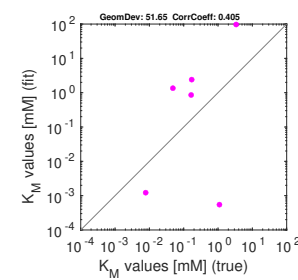
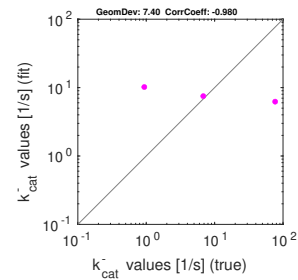
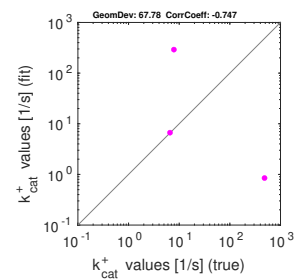
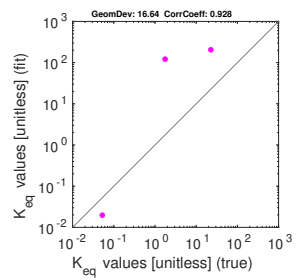
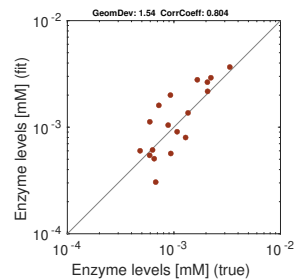
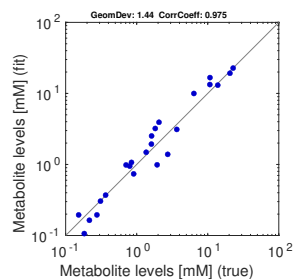


Figure 11: Same as Figure 10, with noisy kinetic data

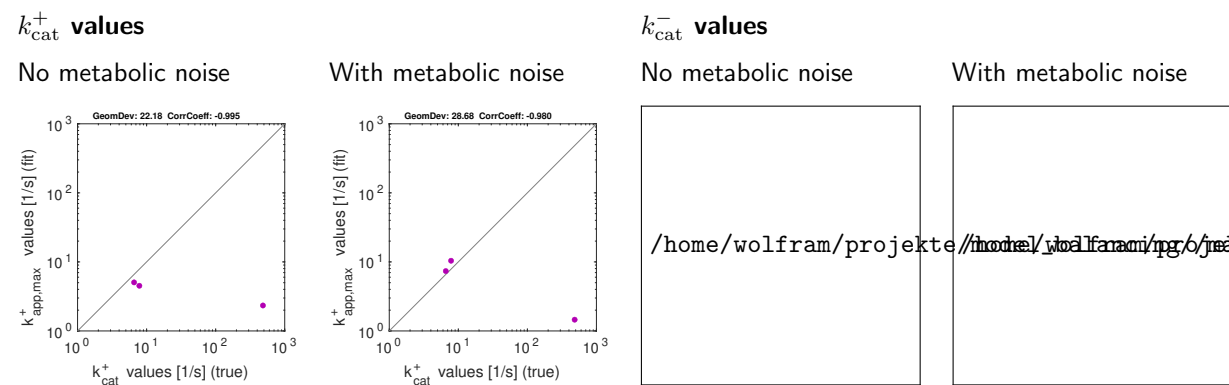


Figure 12: Catalytic constants in Three-chain (artificial data), estimated by kinetic profiling [?]. Note that k_{cat} values can only be estimated in the direction of fluxes (e.g. k_{cat}^+ for reactions with forward flux).