

Figure 2, predictions according to the reparameterized model (blue lines), model configuration i (magenta lines) and model configuration ii (red lines) compared to experimental data (grey diamonds) for the $\triangle G_p - J_p$ relation (**A**) and the ADP $-J_p$ relation (**B**). The simulations of the $\triangle G_p - J_p$ relation show that the reparameterized model did not follow the expected sigmoidal relation. In addition, panel B shows that the reparameterized model failed in reproducing the ADP $-J_p$ relation. In contrast, the $\triangle G_p - J_p$ transduction function predicted according to model configuration i and ii followed a sigmoidal relation. Furthermore, both model configurations reproduced the ADP $-J_p$ relation with a highly improved accuracy.