

# Supplemental Data and Plots

Linking Bioprocess Development to Quality-by-Design Milestones via Digital Twin Applications

## Model tables

Table S1: Overview of identified models based on DoE data. AIC was used as a primary identifier of selected model.  $R^2$ ,  $Q^2$ , RMSE, and  $p$ -values, as well as residual analysis, were used alongside process expertise to determine acceptance of the model within the IPM. All residual variation is accounted for within the prediction interval of the simulated values within the IPM. (+) indicates positive coefficient and (-) indicates negative coefficient.

Unit Op	Blinded	Response	Model	$R^2$	$Q^2$	RMSE	P	Parameters
CAP	CQA1 <sub>prod</sub>	Main SCX	Quadratic	0.45	0.34	1.119	0.019	(-) Load Density (+) Load Density <sup>2</sup>
	CQA1 <sub>imp</sub>	BPG	Linear	0.24	0.11	0.229	0.011	(-) Residence Time (+) Residence Time
	CQA2 <sub>imp</sub>	HMW SEC	Quadratic	0.67	0.57	0.309	0.0001	(-) Load Density (+) Load Density <sup>2</sup>
AT	CQA1 <sub>prod</sub>	Main SCX	Interaction/Quadratic	0.85	0.55	0.016	0.0001	(-) Concentration (+) pH AT (-) Hold Time AT (+) Concentration * pH AT (-) Concentration * Hold AT (+) pH AT * Hold AT (-) pH AT <sup>2</sup>
	CQA1 <sub>imp</sub>	BPG	Interaction/Quadratic	0.94	0.92	0.04	0.0001	(-) Concentration (+) pH AT (-) Hold Time AT (+) Neutr. pH (+) Concentration * pH AT (-) Concentration * Neutr. pH (+) pH AT * Hold AT (-) pH AT <sup>2</sup> (+) Hold AT <sup>2</sup>
	CQA2 <sub>imp</sub>	HMW SEC	Interaction/Quadratic	0.92	0.89	0.672	0.0001	(-) Concentration (+) pH AT (-) Hold Time AT (+) Neutr. Hold Time (-) pH AT * Neutr. Hold (+) Concentration <sup>2</sup> (+) Hold AT <sup>2</sup> (+) Neutr. Hold <sup>2</sup>
	CQA3 <sub>imp</sub>	LMW SEC	Interaction/Quadratic	0.43	0.12	0.031	0.0087	(-) Concentration (+) pH AT (-) Hold Time AT (+) Neutr. Hold Time (-) Neutr. pH (-) Concentration * pH AT (-) Hold AT <sup>2</sup> (+) Neutr. pH <sup>2</sup>
								(-) Concentration <sup>2</sup>

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Unit Op	Blinded	Response	Model	R <sup>2</sup>	Q <sup>2</sup>	RMSE	P	Parameters
AEX	CQA1 <sub>prod</sub>	Main SCX	Interaction	0.77	0.68	0.01	0.0001	(-) Load Density
								(-) Equil. pH
	CQA1 <sub>imp</sub>	BPG	Quadratic	0.53	0.43	0.103	0.002	(+) Load Density * Equil pH
								(-) Equil. pH <sup>2</sup>
								(+) Equil. pH
								(+) Equil. pH <sup>2</sup>
	CQA2 <sub>imp</sub>	HMW SEC	Quadratic	0.65	0.17	0.067	0.0001	(+) Equil. pH
								(-) Equil. Conductivity
								(+) Equil. pH <sup>2</sup>
								(+) Equil. pH
CQA3 <sub>imp</sub>	LMW SEC	Quadratic	0.54	0.41	0.037	0.0005	(-) Equil. Conductivity	
							(+) Equil. Conductivity <sup>2</sup>	
CEX	CQA1 <sub>prod</sub>	Main SCX	Interaction / Quadratic	0.76	0.64	0.001	0.0001	(+) Elu. Conductivity
								(+) Load Density
								(+) Load Conductivity
								(+) Load / Elu. pH
								(-) Elu. Cond. * Load / Elu. pH
								(+) Load Cond. <sup>2</sup>
								(-) Load / Elu. pH <sup>2</sup>
								(-) Elu. Conductivity
	CQA1 <sub>imp</sub>	BPG	Interaction / Quadratic	0.77	0.65	0.068	0.0001	(-) Load Density
								(-) Load / Elu. pH
								(+) Elu. Cond. * Load Density
								(+) Elu. Cond. * Load / Elu. pH
								(+) Elu Cond. <sup>2</sup>
								(-) Elu. Conductivity
								(-) Load / Elu. pH
								(-) Elu. Conductivity
CQA2 <sub>imp</sub>	HMW SEC	Linear	0.33	0.22	0.598	0.0005	(+) Load Conductivity	
							(-) Load / Elu. pH	
							(-) Elu. Conductivity	
							(+) Load / Elu. pH	
CQA3 <sub>imp</sub>	LMW SEC	Quadratic	0.4	0.31	0.434	0.0007	(+) Load Conductivity	
							(-) Load Cond. <sup>2</sup>	

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*Table S2: Overview of models showing a correlation between specific CQA clearances and CQA load (load models). (+) indicates a positive regression line between clearance and load, whereas (-) indicates a negative regression line between clearance and load*

Unit Operation	CQA	R <sup>2</sup>	Effect
AT	CQA1 <sub>imp</sub>	0.75	(+)
AEX	CQA2 <sub>imp</sub>	0.76	(+)
	CQA3 <sub>imp</sub>	0.70	(-)
DF	CQA3 <sub>imp</sub>	0.76	(+)
CEX	CQA1 <sub>imp</sub>	0.73	(-)
UFDF	CQA1 <sub>imp</sub>	0.95	(-)
	CQA2 <sub>imp</sub>	0.95	(+)

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## Linking Bioprocess Development to Quality-by-Design Milestones via Digital Twin Applications

### Simulation Trend Plots

For all plots below, 1000 runs were simulated over the full process(top) and plotted against the sum of existing real manufacturing data (bottom) as well as the Out-of-Acceptance limit (red). Additionally, given only this information, a PPK result is given for OOA, that is, how likely given the normal distribution around the real and simulated data would be OOA at drug substance, which can be used as a comparative diagnostic of IPM quality.

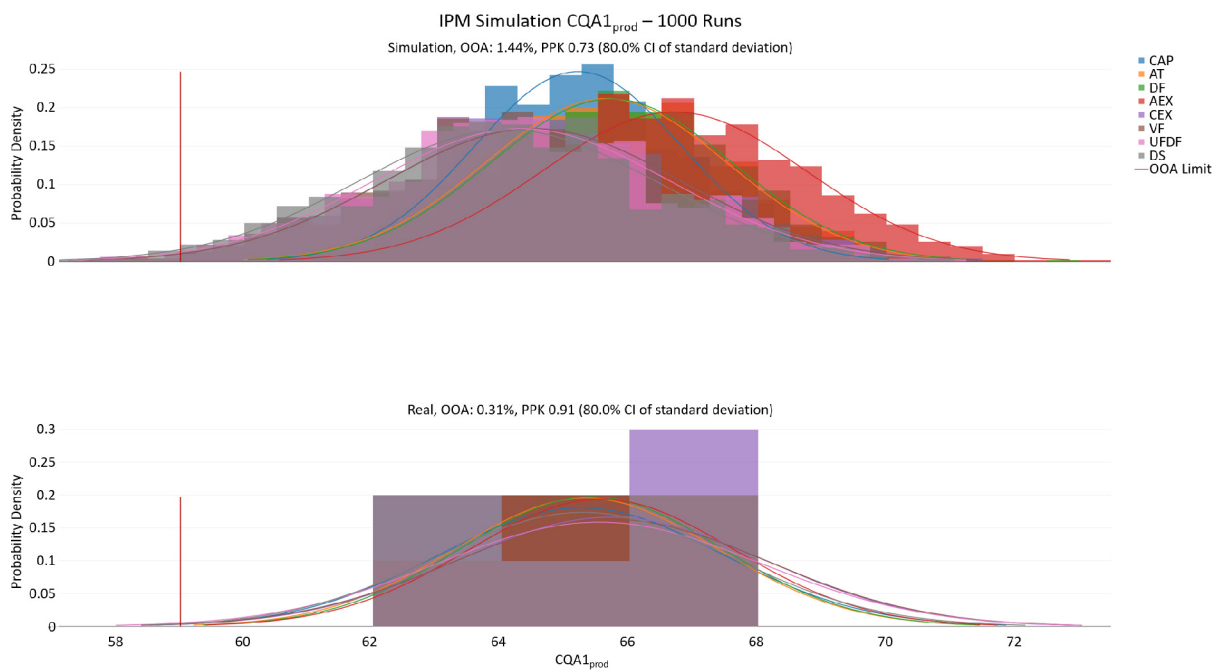


Figure S1: Simulation trend plot for  $CQA1_{prod}$

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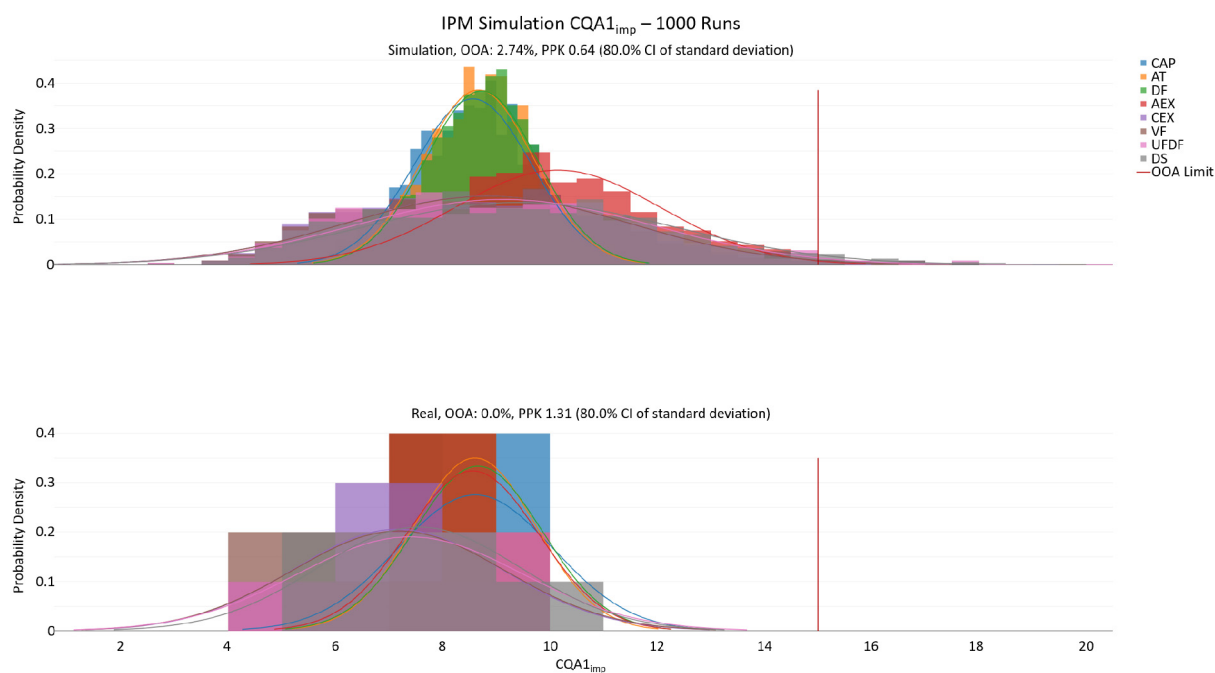


Figure S2: Simulation trend plot for  $CQA1_{imp}$

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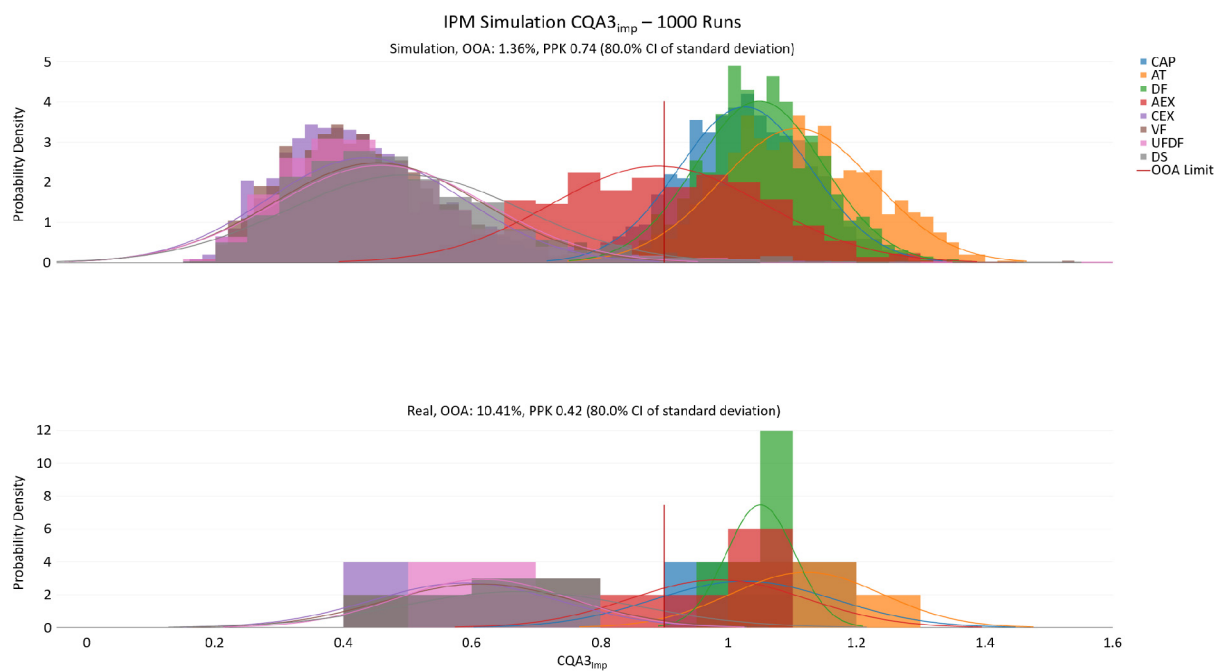


Figure S3: Simulation trend plot for  $CQA3_{imp}$

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## PSA CQA Plots

For all plots below, 1000 simulations per grid point were performed over the full process. The results in OOA are plotted across the relative screening range. The screening range is depicted such that 0 is always the set point condition and the remaining screening ranges are coded between -2, 2.

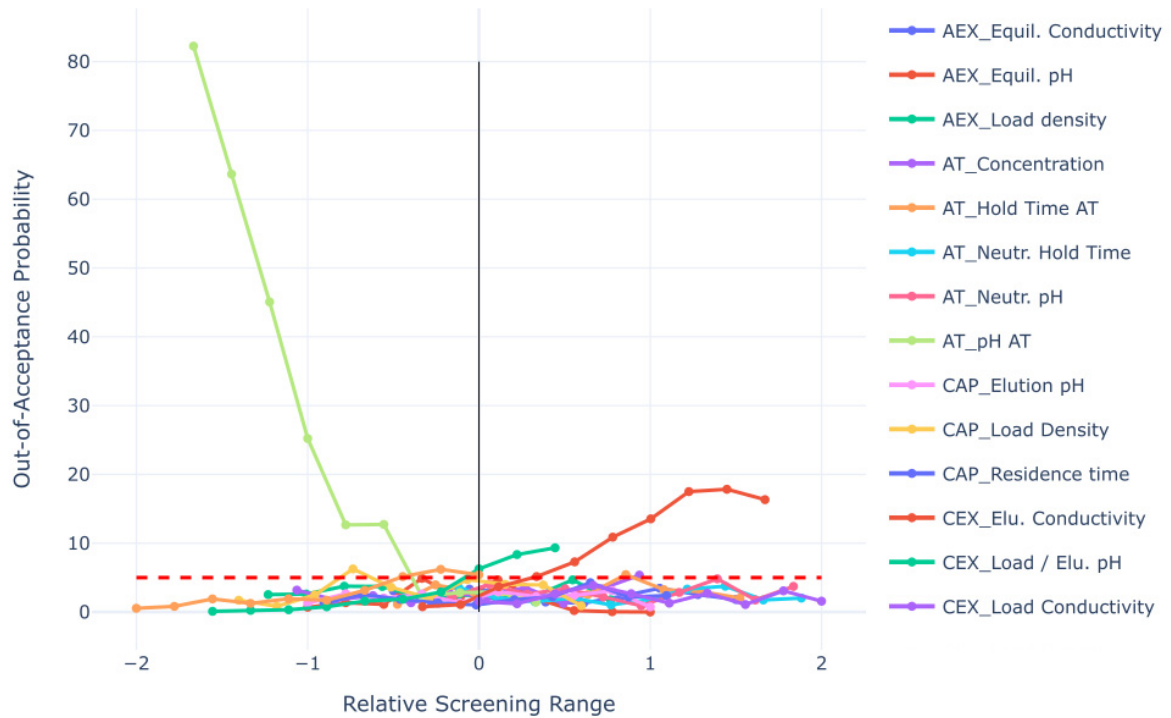


Figure S4: CQA1<sub>imp</sub> CQA PSA plot

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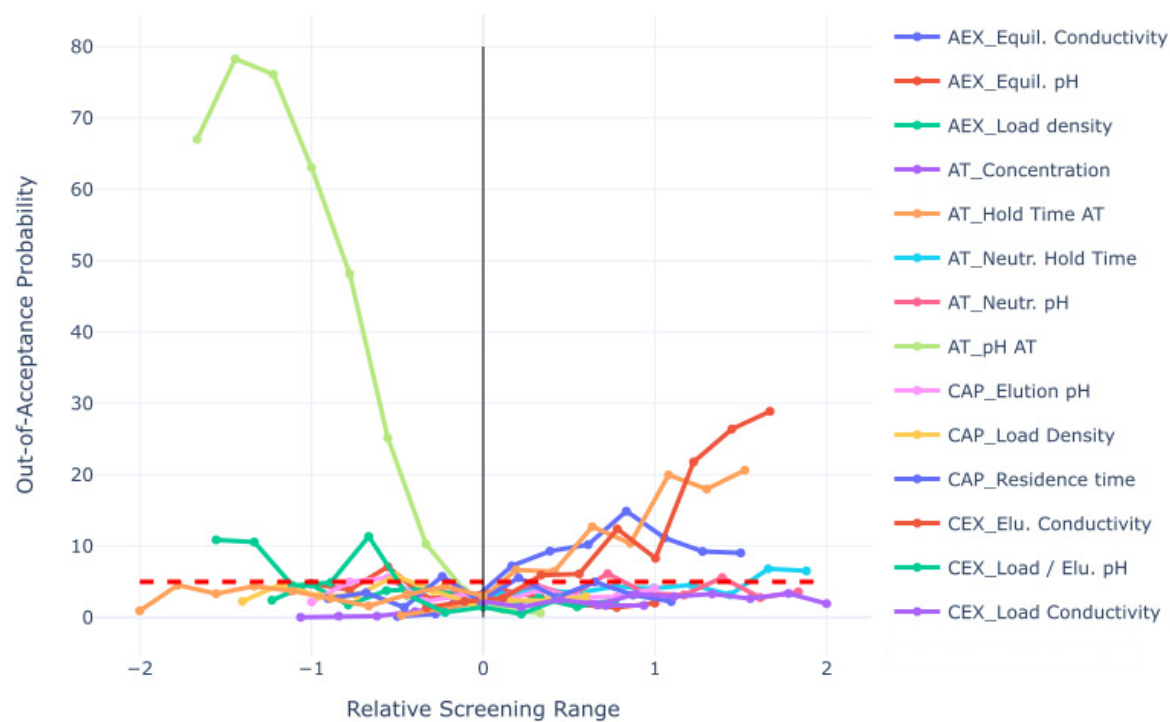


Figure S5: CQA2<sub>imp</sub> CQA PSA plot



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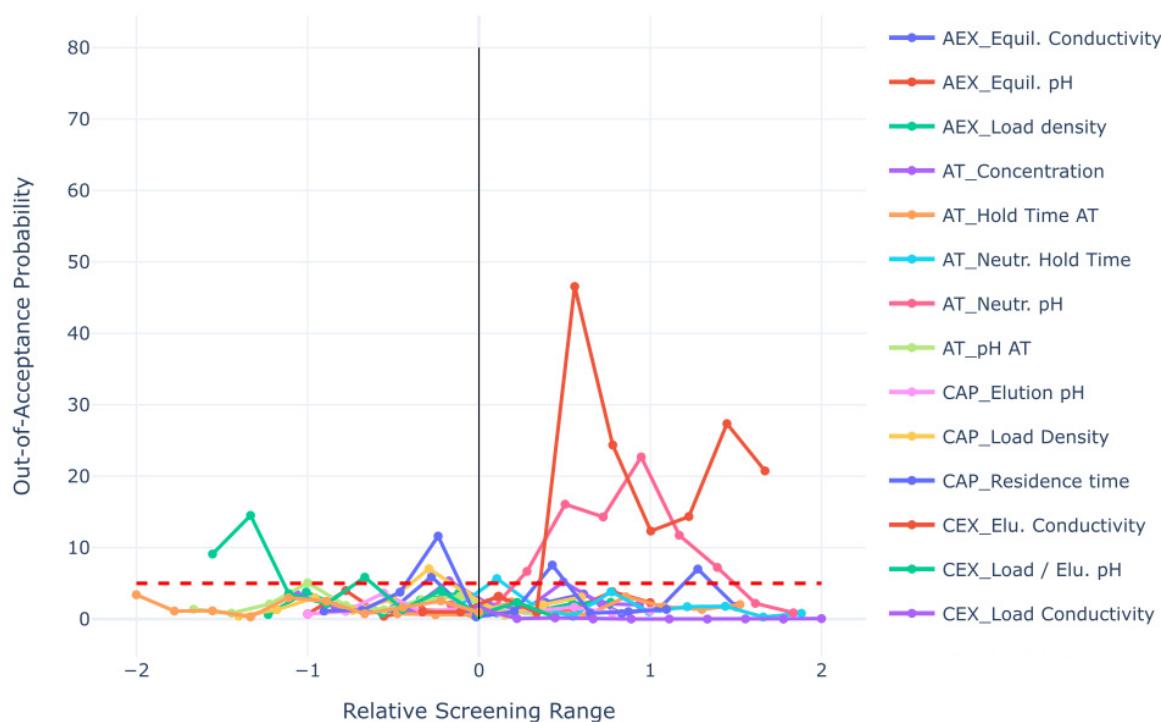


Figure S6: CQA3<sub>imp</sub> CQA PSA plot

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## PSA CPP Plots

For all plots below, 1000 simulations per grid point were performed over the full process. The results in OOA are plotted across the individual CPP range. The area is grey is excluded from the proposed manufacturing PAR range.

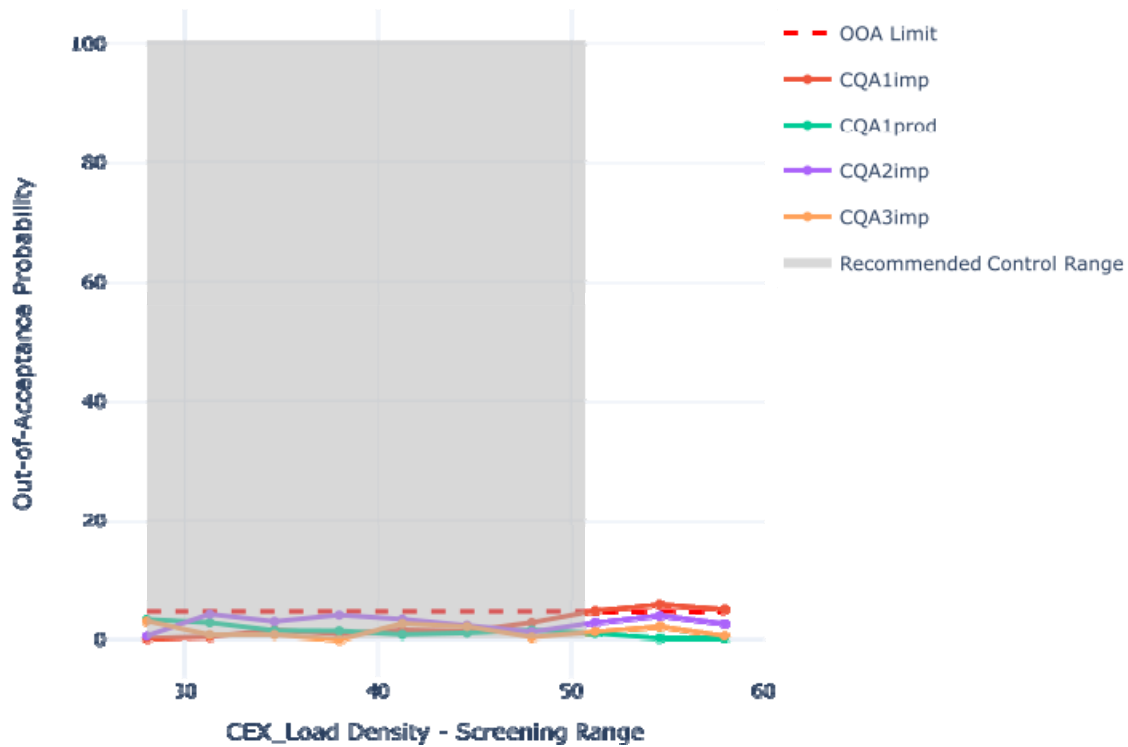


Figure S7: CEX\_Load Density CPP PSA plot

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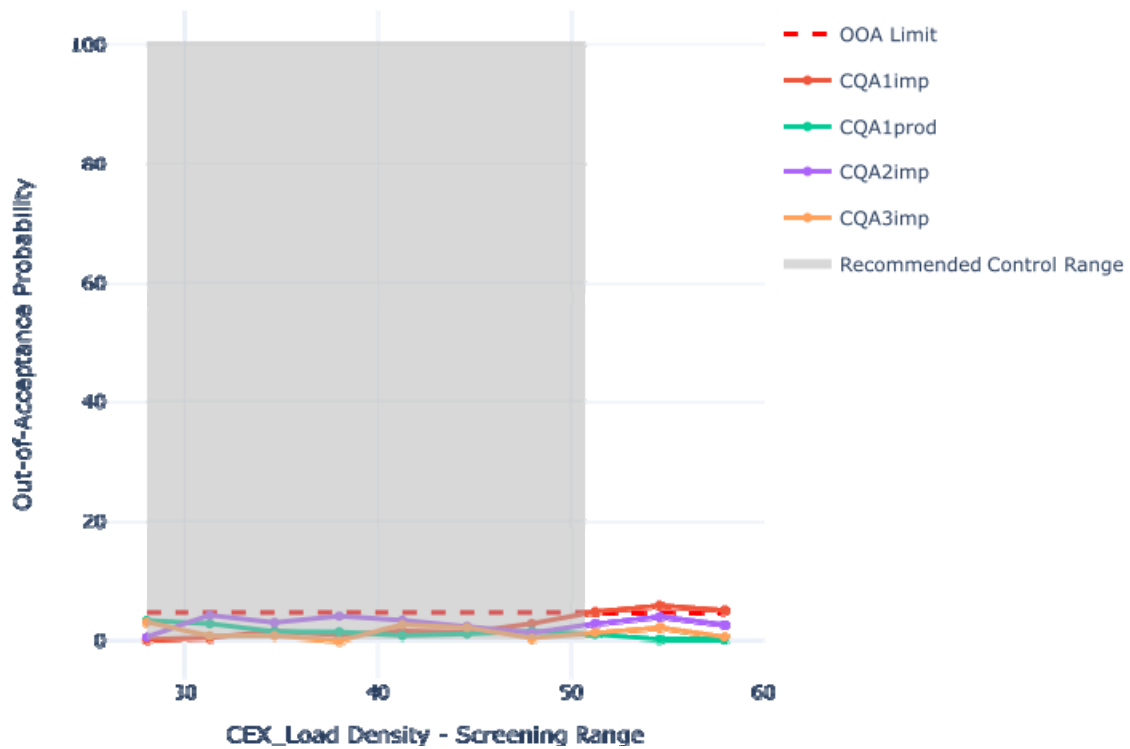


Figure S8: CEX\_Load Density CPP PSA Plot

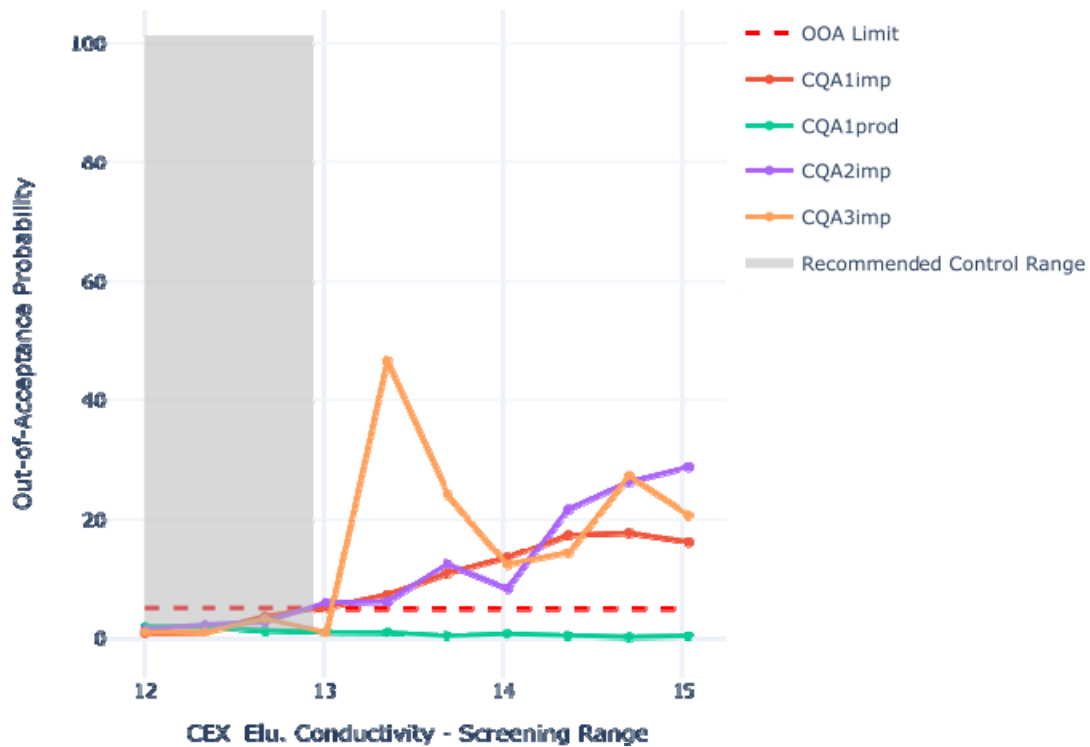


Figure S9: CEX\_Elu. Conductivity CPP PSA plot