Performance Overview
8 samples

	Contaminants	Peptide Intensity	Charge	Missed Cleavages	Missed Cleavages Var	ID rate over RT	MS2 OverSampling	Pep Missing Values
BSA1_F1	1.00	0.10	0.92	0.61	0.85	0.83	0.62	0.33
BSA1_F2	1.00	1.00	0.86	0.40	0.94	0.78	0.50	0.20
BSA2_F1	1.00	0.09	1.00	0.53	0.94	0.81	0.79	0.33
BSA2_F2	1.00	0.61	1.00	0.19	0.73	0.81	0.83	0.20
BSA3_F1	1.00	0.49	0.94	0.72	0.74	0.81	0.91	0.22
BSA3_F2	1.00	1.00	0.89	0.20	0.74	0.85	1.00	0.11

Created with MultiQC