

reporting_jurisdiction, county_names, other_jurisdiction, zipcode, population_served, sewage_travel_time, sample_location, sample_location_specify, institution_type, epa_id, wwtp_name, wwtp_jurisdiction, capacity_mgd, industrial_input, stormwater_input, influent_equilibrated, sample_type, composite_freq, sample_matrix, collection_storage_time, collection_storage_temp, pretreatment, pretreatment_specify, solids_separation, concentration_method, extraction_method, pre_conc_storage_time, pre_conc_storage_temp, pre_ext_storage_time, pre_ext_storage_temp, tot_conc_vol, ext_blank, rec_eff_target_name, rec_eff_spike_matrix, rec_eff_spike_conc, pasteurized, pcr_target, pcr_gene_target, pcr_gene_target_ref, pcr_type, lod_ref, hum_frac_target_mic, hum_frac_target_mic_ref, hum_frac_target_chem, hum_frac_target_chem_ref, other_norm_name, other_norm_ref, quant_stan_type, stan_ref, inhibition_method, num_no_target_control, sample_collect_date, sample_collect_time, time_zone, flow_rate, ph, conductivity, tss, collection_water_temp, equiv_sewage_amt, sample_id, lab_id, qc_ignore, dashboard_ignore, analysis_ignore, test_result_date, pcr_target_units, pcr_target_avg_conc, pcr_target_std_error, pcr_target_cl_95_lo, pcr_target_cl_95_up, pcr_target_below_lod, lod_sewage, ntc_amplify, rec_eff_percent, inhibition_detect, inhibition_adjust, hum_frac_mic_conc, hum_frac_mic_unit, hum_frac_chem_conc, hum_frac_chem_unit, other_norm_conc, other_norm_unit, quality_flag, major_lab_method, major_lab_method_desc