

NORS Data Dictionary for Flat Files - All Variables

Download variable order	Table/Tab Name	Variable Name
1	NORSMain	CDCID
2	NORSMain	eFORSID
3	NORSMain	StateID
4	NORSMain	NORSUserID
5	NORSMain	RecordStatus
6	NORSMain	CDCStatus
7	NORSMain	CDCWaterStatus
8	NORSMain	StartTimeStamp
9	NORSMain	FinalTimeStamp
10	NORSMain	PrimaryMode
11	NORSMain	InvestigationMethod1
12	NORSMain	InvestigationMethod2
13	NORSMain	InvestigationMethod3
14	NORSMain	InvestigationFlag
15	NORSMain	InvestigationComments
16	NORSMain	DateFirstIll
17	NORSMain	DateLastIll
18	NORSMain	InitialExposure
19	NORSMain	LastExposure
20	NORSMain	CDCReportDate

19	NORSMain	LocalReportDate
20	NORSMain	ExposureState
21	NORSMain	ExposureStateCount
22	NORSMain	MultiStateExposure
23	NORSMain	MultiStateResidence
24	NORSMain	State1
		State2
		State3
25	NORSMain	StateCount1
		StateCount2
		StateCount3
26	NORSMain	StateFlag
27	NORSMain	ExposureCounty
28	NORSMain	MultiCountyExposure
29	NORSMain	MultiCountyResidence
30	NORSMain	County1
		County2
		County3
31	NORSMain	CountyFlag
32	NORSMain	ExposureLocation
33	NORSMain	ConfirmedPrimary
34	NORSMain	ProbablePrimary
35	NORSMain	EstimatedPrimary
36	NORSMain	DeathsNum
37	NORSMain	DeathsInfo
38	NORSMain	HospitalNum

39	NORSMain	HospitalInfo
40	NORSMain	ERNum
41	NORSMain	ERInfo
42	NORSMain	HealthcareNum
43	NORSMain	HealthcareInfo
44	NORSMain	NumberMale
45	NORSMain	PercentMale
46	NORSMain	NumberFemale
47	NORSMain	PercentFemale
48	NORSMain	NumberSexUnknown
49	NORSMain	PercentSexUnknown
50	NORSMain	NumberAgeUnder1
51	NORSMain	PercentAgeUnder1
52	NORSMain	NumberAge1to4
53	NORSMain	PercentAge1to4
54	NORSMain	NumberAge5to9
55	NORSMain	PercentAge5to9
56	NORSMain	NumberAge10to19
57	NORSMain	PercentAge10to19
58	NORSMain	NumberAge20to49

59	NORSMain	PercentAge20to49
60	NORSMain	NumberAge50to74
61	NORSMain	PercentAge50to74
62	NORSMain	NumberAge75plus
63	NORSMain	PercentAge75plus
64	NORSMain	NumberAgeUnknown
65	NORSMain	PercentAgeUnknown
66	NORSMain	IncShort
67	NORSMain	IncShortUnit
68	NORSMain	IncMedian
69	NORSMain	IncMedianUnit
70	NORSMain	IncLong
71	NORSMain	IncLongUnit
72	NORSMain	IncubationNum
73	NORSMain	IncUnknown
74	NORSMain	DurShort
75	NORSMain	DurShortUnit
76	NORSMain	DurMedian
77	NORSMain	DurMedianUnit
78	NORSMain	DurLong

79	NORSMain	DurLongUnit
80	NORSMain	DurationNum
81	NORSMain	DurUnknown
82	NORSMain	Symptom1
83	NORSMain	Symptom2
84	NORSMain	Symptom3
85	NORSMain	SymptomCases1
86	NORSMain	SymptomCases2
87	NORSMain	SymptomCases3
88	NORSMain	SymptomInfo1
89	NORSMain	SymptomInfo2
90	NORSMain	SymptomInfo3
91	NORSMain	SymptomFlag
92	NORSMain	SecondaryFood
93	NORSMain	SecondaryWater
94	NORSMain	SecondaryAnimal
95	NORSMain	SecondaryPerson
96	NORSMain	SecondaryEnviron
97	NORSMain	SecondaryOther
98	NORSMain	ConfirmedSecondary
99	NORSMain	ProbableSecondary
100	NORSMain	TotalSecondary
101	NORSMain	TotalCases
102	NORSMain	NEARSID1
103	NORSMain	NEARSID2
104	NORSMain	NEARSID3

99	NORSMain	NEARSID4
100	NORSMain	OHHABSID1
101	NORSMain	OHHABSID2
102	NORSMain	TracebackConducted
103	NORSMain	TracebackSource1
		TracebackSource2
		TracebackSource3
104	NORSMain	TracebackType1
		TracebackType2
		TracebackType3
105	NORSMain	TracebackState1
		TracebackState2
		TracebackState3
106	NORSMain	TracebackCountry1
		TracebackCountry2
		TracebackCountry3
107	NORSMain	TracebackComments1
		TracebackComments2
		TracebackComments3
108	NORSMain	TracebackFlag
109	NORSMain	Recall
110	NORSMain	RecallItem
111	NORSMain	RecallComments
112	NORSMain	ReportingSite
113	NORSMain	AgencyName
114	NORSMain	AgencyContact
115	NORSMain	AgencyTitle
116	NORSMain	AgencyEmail
117	NORSMain	AgencyPhone
118	NORSMain	AgencyFax

119	NORSMain	GeneralComments
120	NORSMain	CDCRemarks
121	NORSMain	References
122	NORSMain	Attachment1 Attachment2
123	NORSMain	AttachmentName1 AttachmentName2
124	NORSMain	AttachmentUploadDate1 AttachmentUploadDate2
125	NORSMain	AttachmentFlag
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126	GenLab	CDCID
127	GenLab	StateID
128	GenLab	SpecimenCollected
129	GenLab	SampleTypeHuman
130	GenLab	SampleTypeHumanNum
131	GenLab	SampleTypeAnimal
132	GenLab	SampleTypeAnimalNum
133	GenLab	SampleTypeFood
134	GenLab	SampleTypeFoodNum
135	GenLab	SampleTypeWater
136	GenLab	SampleTypeWaterNum
137	GenLab	SampleTypeOther
138	GenLab	SampleTypeOtherNum

139	GenLab	TestedBacteria
140	GenLab	TestedViruses
141	GenLab	TestedParasites
142	GenLab	TestedChemicals
143	GenLab	TestedUnknown
144	GenLab	TestTypeChemical
145	GenLab	TestTypeCulture
146	GenLab	TestTypePCR
147	GenLab	TestTypeMicro
148	GenLab	TestTypeSero
149	GenLab	TestTypeTissue
150	GenLab	TestTypeOther
151	GenLab	TestTypeUnknown
152	GenLab	ASTPerformed
153	GenLab	ASTWhereClinical
154	GenLab	ASTWherePHLab
155	GenLab	ASTWhereNARMS
156	GenLab	ASTWhereOther
157	GenLab	ASTWhereUnknown
158	GenLab	ASTOutbreak

159	GenLab	EtiologyKnown
160	GenLab	GenusName1 GenusName2 GenusName3
161	GenLab	SpeciesName1 SpeciesName2 SpeciesName3
162	GenLab	SerotypeName1 SerotypeName2 SerotypeName3
163	GenLab	Polymerase1 Polymerase2 Polymerase3
164	GenLab	Capsid1 Capsid2 Capsid3
165	GenLab	OtherCharacteristics1 OtherCharacteristics2 OtherCharacteristics3
166	GenLab	Confirmed1 Confirmed2 Confirmed3
167	GenLab	NumberLabConfirmed1 NumberLabConfirmed2 NumberLabConfirmed3
168	GenLab	PatientSpecimen1 PatientSpecimen2 PatientSpecimen3
169	GenLab	FoodSpecimen1 FoodSpecimen2 FoodSpecimen3
170	GenLab	EnvironmentSpecimen1 EnvironmentSpecimen2 EnvironmentSpecimen3
171	GenLab	FoodWorkerSpecimen1 FoodWorkerSpecimen2 FoodWorkerSpecimen3
172	GenLab	WaterSample1 WaterSample2 WaterSample3
173	GenLab	AnimalSpecimen1 AnimalSpecimen2 AnimalSpecimen3
174	GenLab	EtiologyValue1 EtiologyValue2 EtiologyValue3
175	GenLab	EtiologyFlag
176	GenLab	CDCSystem1 CDCSystem2 CDCSystem3
177	GenLab	StateLabID1 StateLabID2 StateLabID3
178	GenLab	CDCLabOutbreakID1 CDCLabOutbreakID2 CDCLabOutbreakID3

179	GenLab	Enzyme1_1 Enzyme1_2 Enzyme1_3
180	GenLab	Enzyme2_1 Enzyme2_2 Enzyme2_3
181	GenLab	SequenceID1 SequenceID2 SequenceID3
182	GenLab	OtherMolecularDesignation1 OtherMolecularDesignation2 OtherMolecularDesignation3
183	GenLab	NARMSResistance1 NARMSResistance2 NARMSResistance3
184	GenLab	NARMSAST1 NARMSAST2 NARMSAST3
185	GenLab	SpecimenID1 SpecimenID2 SpecimenID3
186	GenLab	SourceType1 SourceType2 SourceType3
187	GenLab	ArchivedSpecimensTaken
188	GenLab	ArchivedNumberSpecimens
189	GenLab	IsolatesFlag
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190	Foodborne	CDCID
191	Foodborne	StateID
192	Foodborne	FoodVehicleUndetermined
193	Foodborne	ReasonFoodEpi
194	Foodborne	ReasonFoodLab
195	Foodborne	ReasonFoodTrace
196	Foodborne	ReasonFoodOther
197	Foodborne	FoodName1 FoodName2 FoodName3
198	Foodborne	FoodConfirmed1 FoodConfirmed2 FoodConfirmed3

199	Foodborne	ReasonSuspected1_1 ReasonSuspected2_1 ReasonSuspected3_1
200	Foodborne	IngredientName1_1 IngredientName2_1 IngredientName3_1
201	Foodborne	ContaminatedIngredient1_1 ContaminatedIngredient2_1 ContaminatedIngredient3_1
202	Foodborne	FoodExposed1 FoodExposed2 FoodExposed3
203	Foodborne	MethodOfProcessing1_1 MethodOfProcessing2_1 MethodOfProcessing3_1
204	Foodborne	MethodOfPreparation1_1 MethodOfPreparation2_1 MethodOfPreparation3_1
205	Foodborne	LevelofPreparation1_1 LevelofPreparation2_1 LevelofPreparation3_1
206	Foodborne	Imported1 Imported2 Imported3
207	Foodborne	CountryName1 CountryName2 CountryName3
208	Foodborne	ProduceAndSold1 ProduceAndSold2 ProduceAndSold3
209	Foodborne	FoodNameFlag
210	Foodborne	ArchivedTotalExpFood
211	Foodborne	WherePrepName1 WherePrepName2 WherePrepName3
212	Foodborne	WherePrepNameFlag
213	Foodborne	WherePrepComments
214	Foodborne	WhereEatenName1 WhereEatenName2 WhereEatenName3
215	Foodborne	WhereEatenNameFlag
216	Foodborne	WhereEatenComments
217	Foodborne	KitchenManager
218	Foodborne	ContributingFactorsUnknown

219	Foodborne	ContributingFactorName1 ContributingFactorName2
220	Foodborne	ContributingFactorName3 ContributingFactorType1 ContributingFactorType2
221	Foodborne	ContributingFactorType3 ContributingFactorTypeFlag
222	Foodborne	ContributingFactorNameFlag
223	Foodborne	ContaminationPoint
224	Foodborne	BeforePrepPreHarvest
225	Foodborne	BeforePrepProcessing
226	Foodborne	BeforePrepUnknown
227	Foodborne	EnvironmentalEvidence
228	Foodborne	EpidemiologyEvidence
229	Foodborne	LabEvidence
230	Foodborne	PriorExperience
231	Foodborne	FoodWorkerImplicated
232	Foodborne	FoodWorkerImplicatedName
233	Foodborne	MultipleSchools
234	Foodborne	NumofMultipleSchools
235	Foodborne	TotalEnrollment
236	Foodborne	UnknownEnrollmentNumber
237	Foodborne	GradeLevelK GradeLevel1 GradeLevel2
238	Foodborne	SchoolFundingName

239	Foodborne	SchoolFoodPrepName1 SchoolFoodPrepName2 SchoolFoodPrepName3
240	Foodborne	SchoolInspectedName
241	Foodborne	HACCP
242	Foodborne	NationalSchoolProgram
243	Foodborne	FoodItemPurchasedDonatedBy
244	Foodborne	PercentageGBConsumed
245	Foodborne	CaseReady
246	Foodborne	GBReground
247	Foodborne	GBRegroundComment
248	Foodborne	ArchivedPHAGEType1 ArchivedPHAGEType2 ArchivedPHAGEType3
249	Foodborne	ArchivedRDNCNumber1 ArchivedRDNCNumber2 ArchivedRDNCNumber3
250	Foodborne	EggName1 EggName2 EggName3
251	Foodborne	SEFoundOnFarm
252	Foodborne	EggComment

253		CDCID
	Setting	
254		StateID
	Setting	
255		MajorSetting
	Setting	
256		MajorSettingSpecify
	Setting	
257		GuestExposed
	Setting	
258		StaffExposed
	Setting	

259		GuestIll
	Setting	
260		StaffIll
	Setting	
261		GuestAR
	Setting	
262		StaffAR
	Setting	
263		OtherSetting1
	Setting	OtherSetting2
264		OtherSetting3
	Setting	OtherSettingFlag
	Setting	
265		OtherSettingDetails
	Setting	
266		ShigTravel
	Setting	
267		ShigTravelType
	Setting	
268		ShigImmun
	Setting	
269		ShigMSM
	Setting	
270	Animal	CDCID
271	Animal	StateID
272	Animal	AnimalUndetermined
273	Animal	AnimalReasonEpi
274	Animal	AnimalReasonLab
275	Animal	AnimalReasonTrace
276	Animal	AnimalReasonOther
277	Animal	AnimalTypeName1
		AnimalTypeName2
		AnimalTypeName3
278	Animal	AnimalTypeSpecify1
		AnimalTypeSpecify2
		AnimalTypeSpecify3

279	Animal	AnimalConfirmed1 AnimalConfirmed2
280	Animal	AnimalConfirmed3 AnimalVehicleEpi1 AnimalVehicleEpi2
281	Animal	AnimalVehicleEpi3 AnimalVehicleLab1 AnimalVehicleLab2
282	Animal	AnimalVehicleLab3 AnimalVehicleTrace1 AnimalVehicleTrace2
283	Animal	AnimalVehicleTrace3 AnimalVehicleOther1 AnimalVehicleOther2
284	Animal	AnimalVehicleOther3 AnimalSetting1 AnimalSetting2
285	Animal	AnimalSetting3 AnimalFeedImplicated
286	Animal	AnimalFeedType1 AnimalFeedType2
287	Animal	AnimalFeedType3 AnimalLivestockPet
288	Animal	AnimalCompendium
289	Animal	AnimalPrevention1 AnimalPrevention2
290	Animal	AnimalPrevention3 AnimalRemarks
291	Animal	AnimalFlag
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292	WaterGeneral_Etiology	CDCID
293	WaterGeneral_Etiology	StateID
294	WaterGeneral_Etiology	WaterExposure
295	WaterGeneral_Etiology	NumberExposed
296	WaterGeneral_Etiology	ComparisonGroup
297	WaterGeneral_Etiology	CommonSource
298	WaterGeneral_Etiology	EpiExposure1 EpiExposure2 EpiExposure3

299	WaterGeneral_Etiology	TotalExposed1 TotalExposed2 TotalExposed3
300	WaterGeneral_Etiology	IllExposed1 IllExposed2 IllExposed3
301	WaterGeneral_Etiology	NotExposed3 NotExposed4 NotExposed5
302	WaterGeneral_Etiology	IllNotExposed1 IllNotExposed2 IllNotExposed3
303	WaterGeneral_Etiology	EpiAttackRate1 EpiAttackRate2 EpiAttackRate3
304	WaterGeneral_Etiology	OddsRatio1 OddsRatio2 OddsRatio3
305	WaterGeneral_Etiology	RelativeRisk1 RelativeRisk2 RelativeRisk3
306	WaterGeneral_Etiology	pValue1 pValue2 pValue3
307	WaterGeneral_Etiology	ConfidenceInterval1 ConfidenceInterval2 ConfidenceInterval3
308	WaterGeneral_Etiology	AttackRateRes
309	WaterGeneral_Etiology	AttackRateNonRes
310	WaterGeneral_Etiology	InReportingState
311	WaterGeneral_Etiology	AssociatedEvent
312	WaterGeneral_Etiology	EventName
313	WaterGeneral_Etiology	EventStart
314	WaterGeneral_Etiology	EventEnd
315	WaterGeneral_Etiology	RouteIngestion
316	WaterGeneral_Etiology	RouteContact
317	WaterGeneral_Etiology	RouteInhalation
318	WaterGeneral_Etiology	RouteOther

319	WaterGeneral_Etiology	RouteUnknown
320	WaterGeneral_Etiology	EtiologyStatus1 EtiologyStatus2 EtiologyStatus3
321	WaterGeneral_Etiology	EtiologyGenus1 EtiologyGenus2 EtiologyGenus3
322	WaterGeneral_Etiology	EtiologySpecies1 EtiologySpecies2 EtiologySpecies3
323	WaterGeneral_Etiology	EtiologySero1 EtiologySero2 EtiologySero3
324	WaterGeneral_Etiology	EtiologyGeno1 EtiologyGeno2 EtiologyGeno3
325	WaterGeneral_Etiology	EtiologyPolymerase1 EtiologyPolymerase2 EtiologyPolymerase3
326	WaterGeneral_Etiology	EtiologyCapsid1 EtiologyCapsid2 EtiologyCapsid3
327	WaterGeneral_Etiology	EtiologyDetectedIn1 EtiologyDetectedIn2 EtiologyDetectedIn3
328	WaterGeneral_Etiology	TotalTested1 TotalTested2
329	WaterGeneral_Etiology	TotalPositive1 TotalPositive2 TotalPositive3
330	WaterGeneral_Etiology	EtiologyGroup1_1 EtiologyGroup2_1 EtiologyGroup1_2
331	WaterGeneral_Etiology	CDCLabSystem1 CDCLabSystem2 CDCLabSystem3
332	WaterGeneral_Etiology	CDCLabSystemID1 CDCLabSystemID2 CDCLabSystemID3
333	WaterGeneral_Etiology	StateLabSystemID1 StateLabSystemID2 StateLabSystemID3
334	WaterGeneral_Etiology	MolecularDesignation1_1 MolecularDesignation2_1 MolecularDesignation1_2
335	WaterGeneral_Etiology	NARMSResistance1 NARMSResistance2 NARMSResistance3
336	WaterGeneral_Etiology	NARMSAST1 NARMSAST2 NARMSAST3
337	WaterGeneral_Etiology	IsolateFlag
338	WaterGeneral_Etiology	ClinicalSpecimens

339	WaterGeneral_Etiology	SpecimenCount
340	WaterGeneral_Etiology	SpecimenType1 SpecimenType2 SpecimenType3
341	WaterGeneral_Etiology	SpecimenSubtype1 SpecimenSubtype2 SpecimenSubtype3
342	WaterGeneral_Etiology	SpecimenTestedBacteria1 SpecimenTestedBacteria2 SpecimenTestedBacteria3
343	WaterGeneral_Etiology	SpecimenTestedChemicals1 SpecimenTestedChemicals2 SpecimenTestedChemicals3
344	WaterGeneral_Etiology	SpecimenTestedFungi1 SpecimenTestedFungi2 SpecimenTestedFungi3
345	WaterGeneral_Etiology	SpecimenTestedParasites1 SpecimenTestedParasites2 SpecimenTestedParasites3
346	WaterGeneral_Etiology	SpecimenTestedViruses1 SpecimenTestedViruses2 SpecimenTestedViruses3
347	WaterGeneral_Etiology	SpecimenTestedOther1 SpecimenTestedOther2 SpecimenTestedOther3
348	WaterGeneral_Etiology	SpecimenTestedUnknown1 SpecimenTestedUnknown2 SpecimenTestedUnknown3
349	WaterGeneral_Etiology	EtiologyTestChem
350	WaterGeneral_Etiology	EtiologyTestCulture
351	WaterGeneral_Etiology	EtiologyTestPCR
352	WaterGeneral_Etiology	EtiologyTestMicro
353	WaterGeneral_Etiology	EtiologyTestSero
354	WaterGeneral_Etiology	EtiologyTestTissue
355	WaterGeneral_Etiology	EtiologyTestOther
356	WaterGeneral_Etiology	EtiologyTestUnknown
357	WaterGeneral_Etiology	AST
358	WaterGeneral_Etiology	ASTSetting1 ASTSetting2 ASTSetting3

359 WaterGeneral_Etiology ASTOutbreakStrains

360 WaterSamples CDCID

361 WaterSamples StateID

362 WaterSamples WaterTested

363 WaterSamples WaterSampleFlag

364 WaterSamples WaterSampleSourceFlag

365 WaterSamples WaterIndicatorFlag

366 WaterSamples WaterAnalysisFlag

367 WaterSamples SampleSource1_1
SampleSource2_1
SampleSource1_2

368 WaterSamples SampleDescription1
SampleDescription2
SampleDescription3

369 WaterSamples SampleDate1
SampleDate2
SampleDate3

370 WaterSamples Volume1
Volume2
Volume3

371 WaterSamples VolumeUnit1
VolumeUnit2
VolumeUnit3

372 WaterSamples Temp1
Temp2
Temp3

373 WaterSamples TempUnit1
TempUnit2
TempUnit3

374 WaterSamples Residual1
Residual2
Residual3

375 WaterSamples ResidualUnit1
ResidualUnit2
ResidualUnit3

376 WaterSamples Combined1
Combined2
Combined3

377 WaterSamples CombinedUnit1
CombinedUnit2
CombinedUnit3

378 WaterSamples pH1
pH2
pH3

379	WaterSamples	Turbidity1 Turbidity2 Turbidity3
380	WaterSamples	IndicatorType1_1 IndicatorType2_1 IndicatorType3_1
381	WaterSamples	IndicatorConc1_1 IndicatorConc2_1 IndicatorConc3_1
382	WaterSamples	IndicatorUnit1_1 IndicatorUnit2_1 IndicatorUnit3_1
383	WaterSamples	AnalysisGenus1_1 AnalysisGenus2_1 AnalysisGenus3_1
384	WaterSamples	AnalysisSpecies1_1 AnalysisSpecies2_1 AnalysisSpecies3_1
385	WaterSamples	AnalysisSero1_1 AnalysisSero2_1 AnalysisSero3_1
386	WaterSamples	AnalysisGeno1_1 AnalysisGeno2_1 AnalysisGeno3_1
387	WaterSamples	AnalysisPolymerase1_1 AnalysisPolymerase2_1 AnalysisPolymerase3_1
388	WaterSamples	AnalysisCapsid1_1 AnalysisCapsid2_1 AnalysisCapsid3_1
389	WaterSamples	AnalysisPFGE1_1 AnalysisPFGE2_1 AnalysisPFGE3_1
390	WaterSamples	AnalysisPositive1_1 AnalysisPositive2_1 AnalysisPositive3_1
391	WaterSamples	AnalysisConc1_1 AnalysisConc2_1 AnalysisConc3_1
392	WaterSamples	AnalysisUnit1_1 AnalysisUnit2_1 AnalysisUnit3_1
393	WaterSamples	AnalysisTestType1_1 AnalysisTestType2_1 AnalysisTestType3_1
394	WaterSamples	AnalysisTestMethod1_1 AnalysisTestMethod2_1 AnalysisTestMethod3_1
395	ImplicatedWater	CDCID
396	ImplicatedWater	StateID
397	ImplicatedWater	WaterTypeFlag
398	ImplicatedWater	RecWaterTreatmentFlag

399	ImplicatedWater	FillWaterFlag
400	ImplicatedWater	RecTreatedVenue1 RecTreatedVenue2 RecTreatedVenue3
401	ImplicatedWater	RecTreatedSubType1 RecTreatedSubType2 RecTreatedSubType3
402	ImplicatedWater	RecTreatedSetting1 RecTreatedSetting2 RecTreatedSetting3
403	ImplicatedWater	RecTreatment1_1 RecTreatment2_1 RecTreatment3_1
404	ImplicatedWater	RecTreatmentSub3_5 RecTreatmentSub4_5
405	ImplicatedWater	Chlorination2_5 Chlorination3_5 Chlorination4_5
406	ImplicatedWater	FillType2_5 FillType3_5 FillType4_5
407	ImplicatedWater	FillTreatment2_5 FillTreatment3_5 FillTreatment4_5
408	ImplicatedWater	FillTreatmentSub3_5 FillTreatmentSub4_5
409	ImplicatedWater	TreatedQuality
410	ImplicatedWater	TreatedQualText
411	ImplicatedWater	TreatedPoolOp
412	ImplicatedWater	RecUntreatedVenue1 RecUntreatedVenue2 RecUntreatedVenue3
413	ImplicatedWater	RecUntreatedSubType1 RecUntreatedSubType2 RecUntreatedSubType3
414	ImplicatedWater	RecUntreatedSetting1 RecUntreatedSetting2 RecUntreatedSetting3
415	ImplicatedWater	UntreatedQuality
416	ImplicatedWater	UntreatedQualText
417	ImplicatedWater	UntreatedEPA
418	ImplicatedWater	UntreatedEPAText

419	ImplicatedWater	DWSystem1 DWSystem2 DWSystem3
420	ImplicatedWater	EPAID1 EPAID2 EPAID3
421	ImplicatedWater	DWSource1 DWSource2 DWSource3
422	ImplicatedWater	DWDescription1 DWDescription2 DWDescription3
423	ImplicatedWater	DWSetting1 DWSetting2 DWSetting3
424	ImplicatedWater	DWTreatment1 DWTreatment2 DWTreatment3
425	ImplicatedWater	DWTreatmentSub1 DWTreatmentSub2 DWTreatmentSub3
426	ImplicatedWater	DrinkingMonth
427	ImplicatedWater	DrinkingMonthText
428	ImplicatedWater	DrinkingMCL
429	ImplicatedWater	DrinkingMCLText
430	ImplicatedWater	DrinkingYear
431	ImplicatedWater	DrinkingYearText
432	ImplicatedWater	IntendedUse1 IntendedUse2 IntendedUse3
433	ImplicatedWater	OtherType1 OtherType2 OtherType3
434	ImplicatedWater	SettingOther1 SettingOther2 SettingOther3
435	ImplicatedWater	TreatmentOther1 TreatmentOther2 TreatmentOther3
436	ImplicatedWater	TreatmentSubOther1 TreatmentSubOther2 TreatmentSubOther3
437	ImplicatedWater	WaterRemarks
438	ContributingFactors	CDCID

439	ContributingFactors	StateID
440	ContributingFactors	SourceFactor
441	ContributingFactors	TreatmentFactor
442	ContributingFactors	DistributionFactor
443	ContributingFactors	NonUtilityFactor
444	ContributingFactors	WaterFactor1 WaterFactor2 WaterFactor3
445	ContributingFactors	FactorStatus1 FactorStatus2 FactorStatus3
446	ContributingFactors	FactorType1 FactorType2 FactorType3
447	ContributingFactors	WaterFactorFlag

448	Admin	CDCID
449	Admin	StateID
450	Admin	Strength
451	Admin	MonthOnset
452	Admin	YearOnset
453	Admin	PublicationDate
454	Admin	CDCWaterCategory
455	Admin	ConfirmedIn
456	Admin	AgentCodeBacterial
457	Admin	AgentCodeFungal
458	Admin	AgentCodeViral

459	Admin	AgentCodeChemical
460	Admin	AgentCodeParasitic
461	Admin	AgentCodeOther
462	Admin	AgentCodeUnknown
463	Admin	IllnessAGI
464	Admin	IllnessARI
465	Admin	IllnessSkin
466	Admin	IllnessEye
467	Admin	IllnessEar
468	Admin	IllnesHep
469	Admin	IllnessLD
470	Admin	IllnessLepto
471	Admin	IllnessNeuro
472	Admin	IllnessPF
473	Admin	IllnessWound
474	Admin	IllnessLeg
475	Admin	IllnessOther
476	Admin	Deficiency1
477	Admin	Deficiency2
478	Admin	Deficiency3

479	Admin	Deficiency4
480	Admin	Deficiency5A
481	Admin	Deficiency5B
482	Admin	Deficiency5C
483	Admin	Deficiency6
484	Admin	Deficiency7
485	Admin	Deficiency8
486	Admin	Deficiency9
487	Admin	Deficiency10A
488	Admin	Deficiency10B
489	Admin	Deficiency11A
490	Admin	Deficiency11B
491	Admin	Deficiency11C
492	Admin	Deficiency11D
493	Admin	Deficiency11E
494	Admin	Deficiency12
495	Admin	Deficiency13A
496	Admin	Deficiency13B
497	Admin	Deficiency99A
498	Admin	Deficiency99B

499	Admin	Deficiency99C
500	Admin	Deficiency99D
501	Admin	OutbreakSummary
502	Admin	Coliform
503	Admin	ColiformCode
504	Admin	DisinfectionCode
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505	Efors	CDCID
506	Efors	StateID
507	Efors	AgeLessThan1
508	Efors	Age1to4
509	Efors	Age5to19
510	Efors	Age20to49
511	Efors	AgeGreaterThanEqual50
512	Efors	AgeUnknown
513	Efors	IllExposed
514	Efors	Total
515	Efors	ContributingFactorName1
		ContributingFactorName2
		ContributingFactorName3
516	Efors	ContributingFactorType1
		ContributingFactorType2
		ContributingFactorType3
517	Efors	FoodImplicatedID1
		FoodImplicatedID2
		FoodImplicatedID3
518	Efors	MethodOfPreparationName1_1
		MethodOfPreparationName2_1
		MethodOfPreparationName3_1
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519	Finalization_Issues	CDCID
520	Finalization_Issues	StateID
521	Finalization_Issues	DateFirstIll
522	Finalization_Issues	PrimaryMode
523	Finalization_Issues	RecordStatus
524	Finalization_Issues	ReportingSite
525	Finalization_Issues	AgencyName
526	Finalization_Issues	NORSUserID
527	Finalization_Issues	IssueStatus
528	Finalization_Issues	IssueName
529	Finalization_Issues	IssueResponse
530	Finalization_Issues	IssueDescription

Form Marker	Form-Page and Section	Form Name
G1	Page 1, CDC Use Only	52.12 and 52.13
N/A	Page 1, CDC Use Only	52.12 and 52.13
G2	Page 1	52.12 and 52.13
N/A	N/A	52.12 and 52.13
N/A	N/A	52.12 and 52.13
N/A	N/A	52.12 and 52.13
N/A	N/A	52.12
N/A	N/A	52.12 and 52.13
N/A	N/A	52.12 and 52.13
G3	Page 1, Primary mode of transmission	52.12 and 52.13
G4	52.13 Page 1, Investigation Methods	52.12 and 52.13
N/A	N/A	52.12 and 52.13
G5	Page 1, Investigation Methods	52.12 and 52.13
G6	Page 1, Dates	52.12 and 52.13
G7	Page 1, Dates	52.12 and 52.13
G8	Page 1, Dates	52.12 and 52.13
G9	Page 1, Dates	52.12 and 52.13
G10	Page 1, Dates	52.12 and 52.13

G11	Page 1, Dates	52.12 and 52.13
G12	Page 1, Geographic Location	52.12 and 52.13
N/A	Page 1, Geographic Location	52.12 and 52.13
G13	Page 1, Geographic Location	52.12 and 52.13
G14	Page 1, Geographic Location	52.12 and 52.13
G15	Page 1, Geographic Location	52.12 and 52.13
G16	Page 1, Geographic Location	52.12 and 52.13
N/A	N/A	52.12 and 52.13
G17	Page 1, Geographic Location	52.12 and 52.13
G18	Page 1, Geographic Location	52.12 and 52.13
	Page 1, Geographic Location	52.12 and 52.13
G20	Page 1, Geographic Location	52.12 and 52.13
N/A	N/A	52.12 and 52.13
G21	Page 1, Geographic Location	52.12 and 52.13
G22	Page 1, Primary Cases	52.12 and 52.13
G23	Page 1, Primary Cases	52.12 and 52.13
G24	Page 1, Primary Cases	52.12 and 52.13
G25	Page 1, Primary Cases	52.12 and 52.13
G26	Page 1, Primary Cases	52.12 and 52.13
G27	Page 1, Primary Cases	52.12 and 52.13

G28	Page 1, Primary Cases	52.12 and 52.13
G29	Page 1, Primary Cases	52.12 and 52.13
G30	Page 1, Primary Cases	52.12 and 52.13
G31	Page 1, Primary Cases	52.12 and 52.13
G32	Page 1, Primary Cases	52.12 and 52.13
G33	Page 1, Primary Cases	52.12 and 52.13
G34	Page 1, Primary Cases	52.12 and 52.13
G35	Page 1, Primary Cases	52.12 and 52.13
G36	Page 1, Primary Cases	52.12 and 52.13
G37	Page 1, Primary Cases	52.12 and 52.13
G38	Page 1, Primary Cases	52.12 and 52.13
G39	Page 1, Primary Cases	52.12 and 52.13
G40	Page 1, Primary Cases	52.12 and 52.13
G41	Page 1, Primary Cases	52.12 and 52.13
G42	Page 1, Primary Cases	52.12 and 52.13
G43	Page 1, Primary Cases	52.12 and 52.13
G44	Page 1, Primary Cases	52.12 and 52.13
G45	Page 1, Primary Cases	52.12 and 52.13
G46	Page 1, Primary Cases	52.12 and 52.13
G47	Page 1, Primary Cases	52.12 and 52.13

G48	Page 1, Primary Cases	52.12 and 52.13
G49	Page 1, Primary Cases	52.12 and 52.13
G50	Page 1, Primary Cases	52.12 and 52.13
G51	Page 1, Primary Cases	52.12 and 52.13
G52	Page 1, Primary Cases	52.12 and 52.13
G53	Page 1, Primary Cases	52.12 and 52.13
G54	Page 1, Primary Cases	52.12 and 52.13
G55	Page 2, Incubation Period	52.12 and 52.13
G56	Page 2, Incubation Period	52.12 and 52.13
G57	Page 2, Incubation Period	52.12 and 52.13
G58	Page 2, Incubation Period	52.12 and 52.13
G59	Page 2, Incubation Period	52.12 and 52.13
G60	Page 2, Incubation Period	52.12 and 52.13
G61	Page 2, Incubation Period	52.12 and 52.13
G62	Page 2, Incubation Period	52.12 and 52.13
G63	Page 2, Duration of Illness	52.12 and 52.13
G64	Page 2, Duration of Illness	52.12 and 52.13
G65	Page 2, Duration of Illness	52.12 and 52.13
G66	Page 2, Duration of Illness	52.12 and 52.13
G67	Page 2, Duration of Illness	52.12 and 52.13

G68	Page 2, Duration of Illness	52.12 and 52.13
G69	Page 2, Duration of Illness	52.12 and 52.13
G70	Page 2, Duration of Illness	52.12 and 52.13
G71	Page 2, Signs or Symptoms	52.12 and 52.13
G72	Page 2, Signs or Symptoms	52.12 and 52.13
G73	Page 2, Signs or Symptoms	52.12 and 52.13
N/A	N/A	N/A
G74	Page 2, Secondary Cases	52.12 and 52.13
G74	Page 2, Secondary Cases	52.12 and 52.13
G74	Page 2, Secondary Cases	52.12 and 52.13
G74	Page 2, Secondary Cases	52.12 and 52.13
G74	Page 2, Secondary Cases	52.12 and 52.13
G74	Page 2, Secondary Cases	52.12 and 52.13
G74	Page 2, Secondary Cases	52.12 and 52.13
G74	Page 2, Secondary Cases	52.12 and 52.13
G75	Page 2, Secondary Cases	52.12 and 52.13
G76	Page 2, Secondary Cases	52.12 and 52.13
G77	Page 2, Secondary Cases	52.12 and 52.13
G78	Page 2, Secondary Cases	52.12 and 52.13
G79	Page 2, Other CDC System IDs	52.12 and 52.13
G80	Page 2, Other CDC System IDs	52.12 and 52.13
G81	Page 2, Other CDC System IDs	52.12 and 52.13

G82	Page 2, Other CDC System IDs	52.12 and 52.13
G83	Page 2, Other CDC System IDs	52.12 and 52.13
G84	Page 2, Other CDC System IDs	52.12 and 52.13
G85	Page 2, Traceback	52.12 and 52.13
G86	Page 2, Traceback	52.12 and 52.13
G87	Page 2, Traceback	52.12 and 52.13
G88	Page 2, Traceback	52.12 and 52.13
G89	Page 2, Traceback	52.12 and 52.13
G90	Page 2, Traceback	52.12 and 52.13
N/A	N/A	N/A
G91	Page 2, Recall	52.12 and 52.13
G92	Page 2, Recall	52.12 and 52.13
G93	Page 2, Recall	52.12 and 52.13
G94	Page 2, Reporting Agency	52.12 and 52.13
G95	Page 2, Reporting Agency	52.12 and 52.13
G96	Page 2, Reporting Agency	52.12 and 52.13
G97	Page 2, Reporting Agency	52.12 and 52.13
G98	Page 2, Reporting Agency	52.12 and 52.13
G99	Page 2, Reporting Agency	52.12 and 52.13
G100	Page 2, Reporting Agency	52.12 and 52.13

G101	Page 2, General Remarks	52.12 and 52.13
N/A	N/A	N/A
N/A	N/A	N/A
N/A	N/A	N/A
N/A	N/A	N/A
N/A	N/A	N/A
N/A	N/A	N/A

G1	Page 1, CDC Use Only	N/A
G2	Page 1	52.12 and 52.13
GL1	Page 3, Etiology Section	52.13
GL2	Page 3, Etiology Section	52.13
GL3	Page 3, Etiology Section	52.13
GL4	Page 3, Etiology Section	52.13
GL5	Page 3, Etiology Section	52.13
GL6	Page 3, Etiology Section	52.13
GL7	Page 3, Etiology Section	52.13
GL8	Page 3, Etiology Section	52.13
GL9	Page 3, Etiology Section	52.13
GL10	Page 3, Etiology Section	52.13
GL11	Page 3, Etiology Section	52.13

GL12	Page 3, Etiology Section	52.13
GL13	Page 3, Etiology Section	52.13
GL14	Page 3, Etiology Section	52.13
GL15	Page 3, Etiology Section	52.13
GL16	Page 3, Etiology Section	52.13
GL17	Page 3, Etiology Section	52.13
GL18	Page 3, Etiology Section	52.13
GL19	Page 3, Etiology Section	52.13
GL20	Page 3, Etiology Section	52.13
GL21	Page 3, Etiology Section	52.13
GL22	Page 3, Etiology Section	52.13
GL23	Page 3, Etiology Section	52.13
GL24	Page 3, Etiology Section	52.13
GL25	Page 3, Etiology Section	52.13
GL26	Page 3, Etiology Section	52.13
GL27	Page 3, Etiology Section	52.13
GL28	Page 3, Etiology Section	52.13
GL29	Page 3, Etiology Section	52.13
GL30	Page 3, Etiology	52.13
GL31	Page 3, Etiology Section	52.13

GL32	Page 3, Etiology Section	52.13
GL33	Page 3, Etiology Section	52.13
GL34	Page 3, Etiology Section	52.13
GL35	Page 3, Etiology Section	52.13
GL35	Page 3, Etiology Section	52.13
GL35	Page 3, Etiology Section	52.13
GL36	Page 3, Etiology Section	52.13
GL37	Page 3, Etiology Section	52.13
GL38	Page 3, Etiology Section	52.13
GL39	Page 3, Etiology Section	52.13
GL40	Page 3, Etiology Section	52.13
GL41	Page 3, Etiology Section	52.13
GL42	Page 3, Etiology Section	52.13
GL43	Page 3, Etiology Section	52.13
GL44	Page 3, Etiology Section	52.13
N/A	N/A	N/A
N/A	N/A	N/A
GL45	Page 3, Isolates/Strains	52.13
GL46	Page 3, Isolates/Strains	52.13
GL47	Page 3, Isolates/Strains	52.13

GL48	Page 3, Isolates/Strains	52.13
GL49	Page 3, Isolates/Strains	52.13
GL50	Page 3, Isolates/Strains	52.13
GL51	Page 3, Isolates/Strains	52.13

N/A	N/A	N/A
N/A	N/A	N/A
N/A	N/A	N/A

G1	Page 1, CDC Use Only	N/A
G2	Page 1	52.12 and 52.13
F1	Page 4, Food	52.13
F2	Page 4, Food	52.13
F3	Page 4, Food	52.13
F4	Page 4, Food	52.13
F5	Page 4, Food	52.13
F6	Page 4, Food	52.13
F7	Page 4, Food	52.13

F8	Page 4, Food	52.13
F9	Page 4, Food	52.13
F10	Page 4, Food	52.13
F11	Page 4, Food	52.13
F12	Page 4, Food	52.13
F13	Page 4, Food	52.13
F14	Page 4, Food	52.13
F15	Page 4, Food	52.13
F16	Page 4, Food	52.13
F17	Page 4, Food	52.13
N/A	N/A	52.13
N/A	N/A	52.13
F18	Page 5, Location where food was prepared	52.13
N/A	N/A	52.13
F19	Page 5, Location where food was prepared	52.13
F20	Page 5, Location of exposure	52.13
N/A	N/A	52.13
F21	Page 5, Location of exposure	52.13
F22	Page 5, Location of exposure	52.13
F23	Page 5, Contributing Factors	52.13

F24	Page 5, Contributing factors	52.13
F25	Page 5, Contributing factors	52.13
N/A	N/A	52.13
N/A	N/A	52.13
F26	Page 5, Point of Contamination	52.13
F27	Page 5, Point of Contamination	52.13
F28	Page 5, Point of Contamination	52.13
F29	Page 5, Point of Contamination	52.13
F30	Page 5, Point of Contamination	52.13
F31	Page 5, Point of Contamination	52.13
F32	Page 5, Point of Contamination	52.13
F33	Page 5, Point of Contamination	52.13
F34	Page 5, Contributing Factors	52.13
F35	Page 5, Contributing Factors	52.13
F36	Page 6, School questions	52.13
F37	Page 6, School questions	52.13
F38	Page 6, School questions	52.13
F39	Page 6, School questions	52.13
F40	Page 6, School questions	52.13
F41	Page 6, School questions	52.13

F42	Page 6, School questions	52.13
F43	Page 6, School questions	52.13
F44	Page 6, School questions	52.13
F45	Page 6, School questions	52.13
F46	Page 6, School questions	52.13
F47	Page 6, Ground Beef	52.13
F48	Page 6, Ground Beef	52.13
F49	Page 6, Ground Beef	52.13
F50	Page 6, Ground Beef	52.13
N/A	Page 6, Additional Salmonella questions	52.13
N/A	Page 6, Additional Salmonella questions	52.13
F51	Page 6, Eggs	52.13
F52	Page 6, Eggs	52.13
F53	Page 6, Eggs	52.13
<hr/>		
G1	Page 1, CDC Use Only	N/A
G2	Page 1	52.12 and 52.13
S1	Page 4, Major Setting of Exposure	N/A
S2	Page 4, Major Setting of Exposure	N/A
S3	Page 4, Attack Rates for Major Setting of Exposure	N/A
S4	Page 4, Attack Rates for Major Setting of Exposure	N/A

S5	Page 4, Attack Rates for Major Setting of Exposure	N/A
S6	Page 4, Attack Rates for Major Setting of Exposure	N/A
S7	Page 4, Attack Rates for Major Setting of Exposure	N/A
S8	Page 4, Attack Rates for Major Setting of Exposure	N/A
S9	Page 4, Other Settings of Exposure	N/A
N/A	N/A	N/A
S10	Page 4, Attack Rates for Major Setting of Exposure	N/A
SH1	Page 3, Additional Shigella Questions	52.13
SH2	Page 3, Additional Shigella Questions	52.13
SH3	Page 3, Additional Shigella Questions	52.13
SH4	Page 3, Additional Shigella Questions	52.13
<hr/>		
G1	Page 1, CDC Use Only	N/A
G2	Page 1	52.12 and 52.13
A1		52.13
A2	Page 3, Animal Contact	52.13
A3	Page 3, Animal Contact	52.13
A4	Page 3, Animal Contact	52.13
A5	Page 3, Animal Contact	52.13
A6	Page 3, Animal Contact	52.13
A7	Page 3, Animal Contact	52.13

A8	Page 3, Animal Contact	52.13
A9	Page 3, Animal Contact	52.13
A10	Page 3, Animal Contact	52.13
A11	Page 3, Animal Contact	52.13
A12	Page 3, Animal Contact	52.13
A13	Page 3, Animal Contact	52.13
A14	Page 3, Animal Contact	52.13
A15	Page 3, Animal Contact	52.13
A16	Page 3, Animal Contact	52.13
A17	Page 3, Animal Contact	52.13
A18	Page 3, Animal Contact	52.13
A19	Page 3, Animal Contact	52.13
N/A	N/A	N/A

G1	Page 1, CDC Use Only	52.12 and 52.13
G2	Page 1	52.12 and 52.13
W1	Page 3, Type of Water Exposure	52.12
W2	Page 3, Epidemiologic Data	52.12
W3	Page 3, Epidemiologic Data	52.12
W4	Page 3, Epidemiologic Data	52.12
W5	Page 3, Epidemiologic Data	52.12

W6	Page 3, Epidemiologic Data	52.12
W7	Page 3, Epidemiologic Data	52.12
W8	Page 3, Epidemiologic Data	52.12
W9	Page 3, Epidemiologic Data	52.12
W10	Page 3, Epidemiologic Data	52.12
W11	Page 3, Epidemiologic Data	52.12
W12	Page 3, Epidemiologic Data	52.12
W13	Page 3, Epidemiologic Data	52.12
W14	Page 3, Epidemiologic Data	52.12
W15	Page 3, Epidemiologic Data	52.12
W16	Page 3, Epidemiologic Data	52.12
W17	Page 3, Geographic Location	52.12
W18	Page 3, Associated Events	52.12
W19	Page 3, Associated Events	52.12
W20	Page 3, Associated Events	52.12
W21	Page 3, Associated Events	52.12
W22	Page 3, Route of Entry	52.12
W22	Page 3, Route of Entry	52.12
W22	Page 3, Route of Entry	52.12
W22	Page 3, Route of Entry	52.12

W22	Page 3, Route of Entry	52.12
W23	Page 4, Outbreak Etiology	52.12
W24	Page 4, Outbreak Etiology	52.12
W25	Page 4, Outbreak Etiology	52.12
W26	Page 4, Outbreak Etiology	52.12
W27	Page 4, Outbreak Etiology	52.12
W26	Page 4, Outbreak Etiology	52.12
W26	Page 4, Outbreak Etiology	52.12
W28	Page 4, Outbreak Etiology	52.12
W29	Page 4, Outbreak Etiology	52.12
W30	Page 4, Outbreak Etiology	52.12
N/A	N/A	N/A
W31	Page 4, Outbreak Isolates	52.12
W32	Page 4, Outbreak Isolates	52.12
W33	Page 4, Outbreak Isolates	52.12
W34	Page 4, Outbreak Isolates	52.12
N/A	N/A	N/A
W36	Page 4, Clinical Specimens	52.12

W37	Page 4, Clinical Specimens	52.12
W38	Page 4, Clinical Specimens	52.12
W39	Page 4, Clinical Specimens	52.12
W40	Page 4, Clinical Specimens	52.12
W40	Page 4, Clinical Specimens	52.12
W40	Page 4, Clinical Specimens	52.12
W40	Page 4, Clinical Specimens	52.12
W40	Page 4, Clinical Specimens	52.12
W40	Page 4, Clinical Specimens	52.12
W40	Page 4, Clinical Specimens	52.12
W40	Page 4, Clinical Specimens	52.12
W41	Page 4, Testing Information	52.12
W41	Page 4, Testing Information	52.12
W41	Page 4, Testing Information	52.12
W41	Page 4, Testing Information	52.12
W41	Page 4, Testing Information	52.12
W41	Page 4, Testing Information	52.12
W41	Page 4, Testing Information	52.12
W41	Page 4, Testing Information	52.12
W41	Page 4, Testing Information	52.12
W42	Page 4, Testing Information	52.12
W43	Page 4, Testing Information	52.12

W44	Page 4, Testing Information	52.12
G1	Page 1, CDC Use Only	52.12 and 52.13
G2	Page 1	52.12 and 52.13
W45	Page 5, Water Samples	52.12
N/A	N/A	N/A
N/A	N/A	N/A
N/A	N/A	N/A
N/A	N/A	N/A
W46	Page 5, Water Samples	52.12
W47	Page 5, Water Samples	52.12
W48	Page 5, Water Samples	52.12
W49	Page 5, Water Samples	52.12
W50	Page 5, Water Samples	52.12
W51	Page 5, Water Samples	52.12
W52	Page 5, Water Samples	52.12
W53	Page 5, Water Samples	52.12
W54	Page 5, Water Samples	52.12
W55	Page 5, Water Samples	52.12
W56	Page 5, Water Samples	52.12
W57	Page 5, Water Samples	52.12

W58	Page 5, Water Samples	52.12
W59	Page 5, Water Samples - Water Quality Indicators	52.12
W60	Page 5, Water Samples - Water Quality Indicators	52.12
W61	Page 5, Water Samples - Water Quality Indicators	52.12
W62	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
W63	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
W64	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
W65	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
W64	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
W64	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
W66	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
W67	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
W68	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
W69	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
W70	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
W71	Page 5, Water Samples - Microbiology or Chemical/Toxin Analysis	52.12
G1	Page 1, CDC Use Only	N/A
G2	Page 1	52.12 and 52.13
N/A	N/A	N/A
N/A	N/A	N/A

N/A	N/A	N/A
W72	Page 6, Recreational Water Venue Description	52.12
W73	Page 5, Vehicle Description (Treated Rec Water)	52.12
W74	Page 5, Vehicle Description (Treated Rec Water)	52.12
W75	Page 5, Vehicle Description (Treated Rec Water)	52.12
W76	Page 5, Vehicle Description (Treated Rec Water)	52.12
W77	Page 5, Vehicle Description (Treated Rec Water)	52.12
W78	Page 5, Vehicle Description (Treated Rec Water)	52.12
W79	Page 5, Vehicle Description (Treated Rec Water)	52.12
W80	Page 5, Vehicle Description (Treated Rec Water)	52.12
W81	Page 5, Recreational Water Quality (Treated)	52.12
W82	Page 5, Recreational Water Quality (Treated)	52.12
W83	Page 5, Recreational Water Quality (Treated)	52.12
W87	Page 7, Vehicle Description (Untreated Rec Water)	52.12
W88	Page 7, Vehicle Description (Untreated Rec Water)	52.12
W89	Page 7, Vehicle Description (Untreated Rec Water)	52.12
W90	Page 8, Recreational Water Quality (Untreated)	52.12
W91	Page 8, Recreational Water Quality (Untreated)	52.12
W92	Page 8, Recreational Water Quality (Untreated)	52.12
W93	Page 8, Recreational Water Quality (Untreated)	52.12

W94	Page 9, Vehicle Description (Drinking Water)	52.12
W95	Page 9, Vehicle Description (Drinking Water)	52.12
W96	Page 9, Vehicle Description (Drinking Water)	52.12
W97	Page 9, Vehicle Description (Drinking Water)	52.12
W98	Page 9, Vehicle Description (Drinking Water)	52.12
W99	Page 9, Vehicle Description (Drinking Water)	52.12
W100	Page 9, Vehicle Description (Drinking Water)	52.12
W101	Page 9, Drinking Water Quality	52.12
W102	Page 9, Drinking Water Quality	52.12
W103	Page 9, Drinking Water Quality	52.12
W104	Page 9, Drinking Water Quality	52.12
W105	Page 9, Drinking Water Quality	52.12
W106	Page 9, Drinking Water Quality	52.12
W111	Page 13, Intent for Use	52.12
W112	Page 13, Vehicle Description (WNID/WUI)	52.12
W113	Page 13, Vehicle Description (WNID/WUI)	52.12
W114	Page 13, Vehicle Description (WNID/WUI)	52.12
W115	Page 13, Vehicle Description (WNID/WUI)	52.12
W86	Page 7, Remarks (Treated Rec Water); Page 8, Remarks (Untreated Rec	52.12
G1	Page 1, CDC Use Only	52.12 and 52.13

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G1	Page 1, CDC Use Only	N/A
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G2	Page 1	52.12 and 52.13
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G1	Page 1, CDC Use Only	52.12 and 52.13
G2	Page 1	52.12 and 52.13
G6	Page 1, Dates	52.12 and 52.13
G3	Page 1, Primary mode of transmission	52.12 and 52.13
N/A	N/A	52.12 and 52.13
G94	Page 2, Reporting Agency	52.12 and 52.13
G95	Page 2, Reporting Agency	52.12 and 52.13
N/A	N/A	52.12 and 52.13
N/A	N/A	N/A
N/A	N/A	N/A
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NORS Interface- Tab and Section	Format
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General, Investigation Methods & Dates	Text
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General, Investigation Methods & Dates	MM/DD/YYYY
General, Investigation Methods & Dates	MM/DD/YYYY

General, Investigation Methods & Dates MM/DD/YYYY

General, Geographic Location Text

General, Geographic Location Number

General, Geographic Location True/False, 1/0, or -1/0 ,

General, Geographic Location True/False, 1/0, or -1/0

General, Geographic Location Text

General, Geographic Location Number

N/A True/False, 1/0, or -1/0

General, Geographic Location Text

General, Geographic Location True/False, 1/0, or -1/0

General, Geographic Location True/False, 1/0, or -1/0

General, Geographic Location Text

N/A True/False, 1/0, or -1/0

General, Geographic Location Text

General, Primary Cases Number

General, Primary Cases Number

General, Primary Cases Number

General, Primary Cases Number

General, Primary Cases Number

General, Primary Cases Number

General, Primary Cases	Number
General, Primary Cases	Number
General, Primary Cases	Number
General, Primary Cases	Number
General, Primary Cases	Number
General, Primary Cases	Number
General, Primary Cases	Number
General, Incubation & Duration	Number
General, Incubation & Duration	Text
General, Incubation & Duration	Number
General, Incubation & Duration	Text
General, Incubation & Duration	Number
General, Incubation & Duration	Text
General, Incubation & Duration	Number
General, Incubation & Duration	True/False, 1/0, or -1/0
Gen-Pt2, Duration of Illness	Number
Gen-Pt2, Duration of Illness	Text
Gen-Pt2, Duration of Illness	Number
Gen-Pt2, Duration of Illness	Text
Gen-Pt2, Duration of Illness	Number

Gen-Pt2, Duration of Illness	Text
Gen-Pt2, Duration of Illness	Number
Gen-Pt2, Duration of Illness	True/False, 1/0, or -1/0
General, Signs or Symptoms	Text
General, Signs or Symptoms	Number
General, Signs or Symptoms	Number
N/A	True/False, 1/0, or -1/0
General, Secondary Cases	True/False, 1/0, or -1/0
General, Secondary Cases	True/False, 1/0, or -1/0
General, Secondary Cases	True/False, 1/0, or -1/0
General, Secondary Cases	True/False, 1/0, or -1/0
General, Secondary Cases	True/False, 1/0, or -1/0
General, Secondary Cases	True/False, 1/0, or -1/0
General, Secondary Cases	Number
General, Secondary Cases	Number
General, Secondary Cases	Number
General, Secondary Cases	Number
General, Other CDC Systems, Traceback, & Recall	Text
General, Other CDC Systems, Traceback, & Recall	Text
General, Other CDC Systems, Traceback, & Recall	Text

General, Agency & Remarks	Text
General, Agency & Remarks	Text
General, Agency & Remarks	Text
Attachments	True/False, 1/0, or -1/0
N/A	Text
N/A	MM/DD/YYYY
N/A	True/False, 1/0, or -1/0
<hr/>	
N/A	Number
N/A	Text
Etiology, General Etiology	0/1/7/Null (blank)
Etiology, General Etiology	0/1/7/Null (blank)
Etiology, General Etiology	Number
Etiology, General Etiology	0/1/7/Null (blank)
Etiology, General Etiology	Number
Etiology, General Etiology	0/1/7/Null (blank)
Etiology, General Etiology	Number
Etiology, General Etiology	0/1/7/Null (blank)
Etiology, General Etiology	Number
Etiology, General Etiology	0/1/7/Null (blank)
Etiology, General Etiology	Number

Etiology, General Etiology	True/False, 1/0, or -1/0
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Number
Etiology, General Etiology	True/False, 1/0, or -1/0
Etiology, General Etiology	True/False, 1/0, or -1/0
Etiology, General Etiology	True/False, 1/0, or -1/0
Etiology, General Etiology	True/False, 1/0, or -1/0
Etiology, General Etiology	True/False, 1/0, or -1/0
Etiology, General Etiology	True/False, 1/0, or -1/0
Etiology, General Etiology	Text
Etiology, General Etiology	True/False, 1/0, or -1/0
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text

Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	Text
Etiology, General Etiology	0/1/7/Null (blank)
Etiology, General Etiology	Number
Etiology, General Etiology	True/False, 1/0, or -1/0

N/A	Number
N/A	Text
Food	True/False, 1/0, -1/0
Food	True/False, 1/0, -1/0
Food	True/False, 1/0, -1/1
Food	True/False, 1/0, -1/2
Food	True/False, 1/0, -1/3
Food	Text
Food	Text

Food	Text
Food	Text
Food	True/False, 1/0, or -1/0
Food	Number
Food	Text
Food	Text
Food	Text
Food	0/1/9/Null (blank)
Food	Text
Food	0/1/7/Null (blank)
N/A	True/False, 1/0, or -1/0
N/A	Number
Factors	Text
N/A	True/False, 1/0, or -1/0
Factors	Text
Factors	Text
N/A	True/False, 1/0, or -1/0
Factors	Text
Food	0/1/7/Null (blank)
Food	True/False, 1/0, -1/0

Factors	Text
Factors	Text
N/A	True/False, 1/0, or -1/0
N/A	True/False, 1/0, or -1/0
Factors	Text
Factors	True/False, 1/0, -1/0
Factors	True/False, 1/0, -1/0
Factors	True/False, 1/0, -1/0
Factors	True/False, 1/0, -1/0
Factors	True/False, 1/0, -1/0
Factors	True/False, 1/0, -1/0
Factors	True/False, 1/0, -1/0
Food	0/1/7/Null (blank)
Food	Text
School	True/False, 1/0, -1/0
School	Number
School	Number
School	True/False, 1/0, -1/0
School	True/False, 1/0, -1/0
School	Text

School	Text
School	Text
School	0/1/7/Null (blank)
School	0/1/7/Null (blank)
School	Text
Ground Beef	Number
Ground Beef	0/1/7/Null (blank)
Ground Beef	0/1/7/Null (blank)
Ground Beef	Text
Salmonella	Text
Salmonella	Text
Eggs	Text
Eggs	0/1/7/Null (blank)
Eggs	Text
N/A	Number
N/A	Text
P2P, Major setting of exposure	Text
P2P, Attack rates for major setting of exposure	Text
P2P, Attack rates for major setting of exposure	Number
P2P, Attack rates for major setting of exposure	Number

P2P, Attack rates for major setting of exposure	Number
P2P, Attack rates for major setting of exposure	Number
P2P, Attack rates for major setting of exposure	Number
P2P, Attack rates for major setting of exposure	Number
P2P, Attack rates for major setting of exposure	Text
N/A	True/False, 1/0, or -1/0
P2P, Attack rates for major setting of exposure	Text
Settings, Shigella	0/1/7/Null (blank)
Settings, Shigella	Text
Settings, Shigella	0/1/7/Null (blank)
Settings, Shigella	0/1/7/Null (blank)
<hr/>	
N/A	Number
N/A	Text
Animal, Animal Contact	True/False, 1/0, or -1/0
Animal, Animal Contact	True/False, 1/0, or -1/0
Animal, Animal Contact	True/False, 1/0, or -1/0
Animal, Animal Contact	True/False, 1/0, or -1/0
Animal, Animal Contact	True/False, 1/0, or -1/0
Animal, Animal Contact	Text
Animal, Animal Contact	Text

Animal, Animal Contact

Animal, Animal Contact True/False, 1/0, or -1/0

Animal, Animal Contact True/False, 1/0, or -1/0

Animal, Animal Contact True/False, 1/0, or -1/0

Animal, Animal Contact True/False, 1/0, or -1/0

Animal, Animal Contact Text

Animal, Animal Contact 0/1/7/Null (blank)

Animal, Animal Contact Text

Animal, Animal Contact 0/1/7/Null (blank)

Animal, Animal Contact 0/1/7/Null (blank)

Animal, Animal Contact Text

Animal, Animal Contact Text

N/A Text

N/A Number

N/A Text

Water, Type of Water Exposure Text

Water, Epidemiologic Data Number

Water, Epidemiologic Data 0/1/7/Null (blank)

Water, Epidemiologic Data 0/1/7/Null (blank)

Water, Epidemiologic Data Text

Water, Epidemiologic Data	Number
Water, Epidemiologic Data	Number
Water, Epidemiologic Data	Number
Water, Epidemiologic Data	Number
Water, Epidemiologic Data	Percentage, 0.00-100.00
Water, Epidemiologic Data	Number
Water, Epidemiologic Data	Number
Water, Epidemiologic Data	Number
Water, Epidemiologic Data	Text
Water, Epidemiologic Data	Percentage, 0.00-100.00
Water, Epidemiologic Data	Percentage, 0.00-100.00
Water, Epidemiologic Data	Percentage, 0.00-100.00
Water, Epidemiologic Data	0/1/7/Null (blank)
Water, Epidemiologic Data	Text
Water, Epidemiologic Data	MM/DD/YYYY Time
Water, Epidemiologic Data	MM/DD/YYYY Time
Water, Epidemiologic Data	True/False, 1/0, or -1/0
Water, Epidemiologic Data	True/False, 1/0, or -1/0
Water, Epidemiologic Data	True/False, 1/0, or -1/0
Water, Epidemiologic Data	True/False, 1/0, or -1/0

Water, Epidemiologic Data	True/False, 1/0, or -1/0
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Number
Etiology & Lab, General Etiology	Number
N/A	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
Etiology & Lab, General Etiology	Text
N/A	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	0/1/7/Null (blank)

Etiology & Lab, Clinical Specimens	Number
Etiology & Lab, Clinical Specimens	Text
Etiology & Lab, Clinical Specimens	Text
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	True/False, 1/0, or -1/0
Etiology & Lab, Clinical Specimens	0/1/7/Null (blank)
Etiology & Lab, Clinical Specimens	Text

Etiology & Lab, Clinical Specimens	0/1/7/Null (blank)
N/A	Number
N/A	Text
Etiology & Lab, Water Samples	0/1/7/Null (blank)
N/A	True/False, 1/0, or -1/0
N/A	True/False, 1/0, or -1/0
N/A	True/False, 1/0, or -1/0
N/A	True/False, 1/0, or -1/0
Etiology & Lab, Water Samples	Text
Etiology & Lab, Water Samples	Text
Etiology & Lab, Water Samples	MM/DD/YYYY Time
Etiology & Lab, Water Samples	Number
Etiology & Lab, Water Samples	Text
Etiology & Lab, Water Samples	Number
Etiology & Lab, Water Samples	Text
Etiology & Lab, Water Samples	Number
Etiology & Lab, Water Samples	Text
Etiology & Lab, Water Samples	Number
Etiology & Lab, Water Samples	Text
Etiology & Lab, Water Samples	Number

Etiology & Lab, Water Samples Number

Etiology & Lab, Water Samples Text

Etiology & Lab, Water Samples Number

Etiology & Lab, Water Samples Text

Etiology & Lab, Water Samples Text

Etiology & Lab, Water Samples Text

Etiology & Lab, Water Samples Text

Etiology & Lab, Water Samples Text

Etiology & Lab, Water Samples Text

Etiology & Lab, Water Samples Text

Etiology & Lab, Water Samples Text

Etiology & Lab, Water Samples True/False, 1/0, or -1/0

Etiology & Lab, Water Samples Number

Etiology & Lab, Water Samples Text

Etiology & Lab, Water Samples Text

Etiology & Lab, Water Samples Text

N/A Number

N/A Text

N/A True/False, 1/0, or -1/0

N/A True/False, 1/0, or -1/0

N/A	True/False, 1/0, or -1/0
Rec. Treated,Water Venue	Text
Rec. Treated,Water Venue	Text
Rec. Treated,Water Venue	Text
Rec. Treated,Water Venue	Text
Rec. Treated,Water Venue	Text
Rec. Treated,Water Venue	Text
Rec. Treated,Water Venue	Text
Rec. Treated,Water Venue	Text
Rec. Treated,Water Venue	Text
Rec. Treated, Water Quality	0/1/7/9/Null (blank)
Rec. Treated, Water Quality	Text
Rec. Treated, Water Quality	0/1/7/Null (blank)
Water-Untreated, Vehicle Description	Text
Water-Untreated, Vehicle Description	Text
Water-Untreated, Vehicle Description	Text
Rec. Untreated, Water Quality	0/1/7/9/Null (blank)
Rec. Untreated, Water Quality	Text
Rec. Untreated, Water Quality	0/1/7/9/Null (blank)
Rec. Untreated, Water Quality	Text

Drinking, Water System	Text
Drinking, Water System	Text
Drinking, Water System	Text
Drinking, Water System	Text
Drinking, Water System	Text
Drinking, Water System	Text
Drinking, Water System	Text
Drinking, Water Quality	0/1/7/9/Null (blank)
Drinking, Water Quality	Text
Drinking, Water Quality	0/1/7/9/Null (blank)
Drinking, Water Quality	Text
Drinking, Water Quality	0/1/7/9/Null (blank)
Drinking, Water Quality	Text
Other/Unknown, Intent for Use	Text
Other/Unknown, Water Description	Text
Other/Unknown, Water Description	Text
Other/Unknown, Water Description	Text
Other/Unknown, Water Description	Text
(Rec. Treated, Rec. Untreated, Drinking, Other/Unknown), Remarks	Text
N/A	Number

N/A	Text
Drinking, Contributing Factors	0/1/7/Null (blank)
Drinking, Contributing Factors	0/1/7/Null (blank)
Drinking, Contributing Factors	0/1/7/Null (blank)
Drinking, Contributing Factors	0/1/7/Null (blank)
Rec. Treated, Contributing Factors; Rec. Untreated, Contributing Factors; Drinking, Rec. Treated, Contributing Factors; Rec. Untreated, Contributing Factors; Drinking, Rec. Treated, Contributing Factors; Rec. Untreated, Contributing Factors; Drinking, N/A	Text True/False, 1/0, or -1/0 Text True/False, 1/0, or -1/0

N/A	Number
N/A	Text
Water, Water Admin I	Text
Water, Water Admin I	Number
Water, Water Admin I	Number
Water, Water Admin I	Text
Water, Water Admin I	Text
Water, Water Admin I	Text
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0

Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin I	True/False, 1/0, or -1/0
Water, Water Admin II	True/False, 1/0, or -1/0
Water, Water Admin II	True/False, 1/0, or -1/0
Water, Water Admin II	True/False, 1/0, or -1/0

Water, Water Admin II True/False, 1/0, or -1/0

Water, Water Admin II True/False, 1/0, or -1/0

Water, Water Admin II Text

Water, Water Admin II True/False, 1/0, or -1/0

Water, Water Admin II Text

Water, Water Admin II Text

N/A Number

N/A Text

eFORS, Additional eFORS Data Number

eFORS, Additional eFORS Data Number

eFORS, Additional eFORS Data Number

eFORS, Additional eFORS Data Number

eFORS, Additional eFORS Data Number

eFORS, Additional eFORS Data Number

eFORS, Additional eFORS Data Number

eFORS, Additional eFORS Data Number

eFORS, Additional eFORS Data Text

eFORS, Additional eFORS Data Text

N/A Number

eFORS, Additional eFORS Data Text

N/A	Number
N/A	Text
General, Investigation Methods & Dates	MM/DD/YYYY
N/A	Text
N/A	Text
General, Agency & Remarks	Text
General, Agency & Remarks	Text
N/A	Text
Finalize	Text
Finalize	Text
Finalize	Text
Finalize	Text

Comments

The CDCID is an autogenerated unique ID that represents a single NORS report. Links record from a single NORS report across all tables of the flattened database.

This is the unique ID previously assigned to a report originally entered in eFORS. All previous eFORS reports have been migrated into NORS.

The report ID entered by the reporting site. The StateID is unique to each report created by a NORS reporting agency.

User ID of person who created the report

Indicates whether the report is active, finalized, or deleted

NORS value for internal data management processes. Indicates whether a report has been excluded (e.g., due to multistate consolidation), cleaned, or closed out (data cleaning status applied for NORS reports for some primary modes)

NORS value that indicates whether CDC has classified a waterborne disease outbreak report as excluded, cleaned, or closed out as a result of routine data cleaning and review activities.

Date NORS report was created.

Date NORS report was most recently finalized, if any.

Primary mode of transmission

Name of method used to investigate outbreak

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

Reporting site comments relating to the selected Investigation Method(s). See Investigation Method table/tab for selected investigation method(s)

Earliest date of reported illness onset

Latest date of reported illness onset

Earliest date of reported exposure

Latest date of reported exposure

Date of report to CDC other than the NORS report

Date of notification to State/Territory/Local Health authorities

The state where the exposure occurred. For a single state of exposure, the state will be listed. For multiple states of exposure, "multistate exposure" will be listed.

The case count for the exposure state.

Exposure occurred in multiple states. 0="False", 1 or -1="True" Format depends on export and import program

Exposure occurred in a single state, but cases resided in another state or multiple states. 0="False", 1 or -1="True" Format depends on export and import program

Name of each state involved in the outbreak. If multiple states were involved but there was only one exposure state, it will be included in the list of states.

The case count for each state for multistate exposure or multistate residency outbreaks. This information can be entered on the reporting form when additional states are reported as part of a multistate exposure or multistate residency outbreak.

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

The county where the exposure occurred for a single state exposure outbreak. For a single county of exposure, the county will be listed. For multiple counties of exposure, "multicounty exposure" will be listed.'

Exposure occurred in multiple counties. 0="False", 1 or -1="True" Format depends on export and import program

Exposure occurred in a single county, but cases resided in another county or multiple counties. 0="False", 1 or -1="True" Format depends on export and import program

Name of each county involved in the outbreak. If multiple counties were involved but there was only one exposure county, it will be included in the list of counties.

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

Name of city, town, or place of exposure

Number of laboratory confirmed primary cases

Number of probable primary cases

Estimated total number of primary cases, including lab-confirmed and probable, based on the outbreak-specific definition.

Number of primary cases who died

Total number of primary cases for whom information on survival is available

Number of primary cases who were hospitalized

Total number of primary cases for whom information on hospitalization is available

Number of primary cases who visited an emergency room

Total number of primary cases for whom information on ER visit is available

Number of primary cases who visited a health care provider

Total number of primary cases for whom information on health care visit is available

Number of primary cases who are male

Percentage of primary cases who are male

Number of primary cases who are female

Percentage of primary cases who are female

Number of primary cases of unknown sex

Percentage of primary cases of unknown sex

Number of primary cases age less than one year

Percentage of primary cases aged less than one year

Number of primary cases age one to four years

Percentage of primary cases age one to four years

Number of primary cases age five to nine years

Percentage of primary cases age five to nine years

Number of primary cases age ten to nineteen years

Percentage of primary cases age 10 to 19 years

Number of primary cases age 20 to 49 years

Percentage of primary cases age 20 to 49 years

Number of primary cases age 50 to 74 years

Percentage of primary cases age 50 to 74 years

Number of primary cases over 75 years of age

Percentage of primary cases over 75 years of age

Number of primary cases of unknown age

Percentage of primary cases of unknown age

Shortest incubation period, in selected units

Units of shortest incubation time

Median incubation period, in selected units

Units of median incubation time

Longest incubation period, in selected units

Units of longest incubation time

Number of primary cases for whom information on incubation period is available.

Information on incubation period is unknown. 0="False", 1 or -1 ="True" Format depends on export and import program

Shortest duration time, in selected units

Units of shortest duration time

Median duration time, in selected units

Units of median duration time

Longest duration time, in selected units

Units of longest duration time

Number of primary cases for whom information on duration of illness is available.

Information on duration of illness is unknown. 0="False", 1 or -1 ="True" Format depends on export and import program

Name of sign or symptom

Number of primary cases with sign or symptom

Total number of primary cases for whom information on specified sign or symptom is available

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

Food reported as a confirmed or suspected mode of secondary transmission. 1='confirmed or suspected secondary mode' 0='not a confirmed or suspected secondary mode'

Water reported as a confirmed or suspected mode of secondary transmission. 1='confirmed or suspected secondary mode' 0='not a confirmed or suspected secondary mode'

Animal contact reported as a confirmed or suspected mode of secondary transmission. 1='confirmed or suspected secondary mode' 0='not a confirmed or suspected secondary mode'

Person-to-person transmission reported as a confirmed or suspected mode of secondary transmission. 1='confirmed or suspected secondary mode' 0='not a confirmed or suspected secondary mode'

Environmental contact reported as a confirmed or suspected mode of secondary transmission. 1='confirmed or suspected secondary mode' 0='not a confirmed or suspected secondary mode'

Another or an undetermined exposure reported as a confirmed or suspected mode of secondary transmission. 1='confirmed or suspected secondary mode' 0='not a confirmed or suspected secondary mode'

Number of laboratory confirmed secondary cases

Number of probable secondary cases

Total number of laboratory confirmed and probable secondary cases

Estimated total number of secondary cases, including lab-confirmed and probable, based on the outbreak-specific definition

This field is used to link NORS reports to environmental health investigation reports in the National Environmental Assessment Reporting System (NEARS) (formerly Environmental Health Specialists Network)

This field is used to link NORS reports to environmental health investigation reports in the National Environmental Assessment Reporting System (NEARS) (formerly Environmental Health Specialists Network)

This field is used to link NORS reports to environmental health investigation reports in the National Environmental Assessment Reporting System (NEARS) (formerly Environmental Health Specialists Network)

This field is used to link NORS reports to environmental health investigation reports in the National Environmental Assessment Reporting System (NEARS) (formerly Environmental Health Specialists Network)

This field is used to link NORS reports to reports in the One Health Harmful Algal Bloom System (OHHABS)

This field is used to link NORS reports to reports in the One Health Harmful Algal Bloom System (OHHABS)

Traceback was conducted. 0="False", 1 or -1 ="True". Format depends on export and import program

Name of source

Source type

State of source

Country of source

Additional remarks concerning traceback

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

A food or bottled water product was recalled. 0="False", 1="True". Format depends on export and import program

Type of item recalled

Comments about the recall.

Reporting site of the report author

Name of the NORS agency reporting the outbreak

Name of primary agency contact

Title of agency contact

Email address of agency contact

Phone number of agency contact

Fax number of agency contact

General comments about the outbreak

General comments added by CDC NORS managing administrators

Publications or other references related to outbreak added by CDC

Indicates whether or not one or more attachments have been included in the NORS report. 0="False", 1 or -1="True" Format depends on export and import program

The filename of the attachment(s) to the NORS report, if any

Upload date of the attachment(s)

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

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Indicates whether any specimens or environmental, food, or water samples were collected and tested. 0="False", 1 or -1="True" Format depends on export and import program

Indicates whether human samples were tested. 0="False", 1 or -1="True" Format depends on export and import program

Number of human specimens tested

Indicates whether animal samples were tested. 0="False", 1 or -1="True" Format depends on export and import program

Number of animal specimens tested

Indicates whether food samples were tested. 0="False", 1 or -1="True" Format depends on export and import program

Number of food specimens tested

Indicates whether water samples were tested. 0="False", 1 or -1="True" Format depends on export and import program

Number of water specimens tested

Indicates whether other samples were tested, specify in general remarks section. 0="False", 1 or -1="True" Format depends on export and import program

Number of other samples that were tested.

This field indicates whether collected specimens, if any, were tested for bacterial etiologies. 0="False", 1 or -1="True" Format depends on export and import program

This field indicates whether collected specimens, if any, were tested for viral etiologies. 0="False", 1 or -1="True" Format depends on export and import program

This field indicates whether collected specimens, if any, were tested for parasitic etiologies. 0="False", 1 or -1="True" Format depends on export and import program

This field indicates whether collected specimens, if any, were tested for chemical etiologies. 0="False", 1 or -1="True" Format depends on export and import program

This field indicates that collected specimens/samples were tested, but for what they were tested is unknown. 0="False", 1 or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using chemical testing methods. 0="False", 1 or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using culture methods. 0="False", 1 or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using DNA or RNA amplification methods. 0="False", 1 or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using microscopy methods. 0="False", 1 or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using serological or immunological methods. 0="False", 1 or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using tissue culture infectivity assay. 0="False", 1 or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using another method. 0="False", 1 or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using an unknown methods. 0="False", 1 or -1="True" Format depends on export and import program

Indicates whether any samples were tested for antimicrobial resistance. 0="No", 1="Yes", 7="Unknown", null (blank) = no response

Indicates if AST was done by a clinical lab (e.g., hospital). 0="False", 1 or -1="True" Format depends on export and import program

Indicates if AST was done by a public health lab. 0="False", 1 or -1="True" Format depends on export and import program

Indicates if AST was done by the National Antimicrobial Resistance Monitoring System (NARMS). 0="False", 1 or -1="True" Format depends on export and import program

Indicates if AST was done by another lab. 0="False", 1 or -1="True" Format depends on export and import program

Indicates if AST testing is unknown. 0="False", 1 or -1="True" Format depends on export and import program

Indicates whether any antimicrobial resistant strains were associated with the outbreak. 0="False", 1="True", 7="Unknown", null (blank) = no response

Indicates that there is at least one confirmed or suspected outbreak etiology (based on either laboratory confirmation or epidemiological evidence). 0="False", 1 or -1 ="True", Format depends on export and import program.

Genus of identified etiology

Species of identified etiology

Serotype of identified etiology

Polymerase type of a norovirus strain identified as an outbreak etiology

Capsid type of a norovirus strain identified as an outbreak etiology

Other characteristics of identified etiology

Indicates whether or not the listed etiology is laboratory 'Confirmed' or 'Suspected'.

Number of lab confirmed cases

Etiology was identified/detected in a patient specimen. 0="False", 1 or -1 ="True". Format depends on export and import program

Etiology was identified/detected in a food specimen. 0="False", 1 or -1 ="True". Format depends on export and import program

Etiology was identified/detected in an environmental specimen. 0="False", 1 or -1 ="True". Format depends on export and import program

Etiology was identified/detected in a food worker specimen. 0="False", 1 or -1 ="True". Format depends on export and import program

Etiology was identified/detected in a water sample specimen. 0="False", 1 or -1 ="True". Format depends on export and import program

Etiology was identified/detected in an animal specimen. 0="False", 1 or -1 ="True". Format depends on export and import program

Indicates whether listed etiology is bacterial, chemical, viral, parasitic, or hepatitis. Value is automatically assigned for analysis purposes.

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

The CDC laboratory system with associated isolate or specimen data.

State PulseNet laboratory ID, CaliciNet key, Accession ID, or PulseNet key

PulseNet cluster code or CaliciNet outbreak number

CDC PulseNet pattern designation for enzyme 1. The numeral before the underscore identifies the pattern designation and the numeral after the underscore identifies the enzyme with which the pattern is associated. (e.g., FoodName1, Ingredient1_1, Ingredient2_1, Ingredient3_1, FoodName2, Ingredient1_2, Ingredient2_2, CDC PulseNet pattern designation for enzyme 2. The numeral before the underscore identifies the pattern designation and the numeral after the underscore identifies the enzyme with which the pattern is associated. (e.g., FoodName1, Ingredient1_1, Ingredient2_1, Ingredient3_1, FoodName2, Ingredient1_2, Ingredient2_2, Whole genome sequencing ID/CaliciNet sequenced region

Other molecular designation/ CaliciNet genotype/ PulseNet species or serotype

CDC NARMS predicted resistance pattern based on resistance determinants.

CDC NARMS antimicrobial susceptibility testing (AST) results.

State Lab Specimen ID from the CaliciNet database

Source Type for the specimen from the CaliciNet database

NORS question collected between Jan 2009 and Sept 2015. This question is no longer being collected, but the data entered during this period has been archived for reference.. If etiology is unknown, were patient specimens collected? 0="False", 1="True", 7="Unknown", null (blank) = no response

NORS question collected between Jan 2009 and Sept 2015. This question is no longer being collected, but the data entered during this period has been archived for reference. Number of specimens collected for outbreak of unknown etiology.

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

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Indicates if a food vehicle was not identified for the outbreak. Format depends on export and import program, 0="False", 1 or -1 ="True".

Indicates whether epidemiologic evidence determined that the outbreak was foodborne, with an undetermined vehicle.

Indicates whether laboratory evidence determined that the outbreak was foodborne, with an undetermined vehicle.

Indicates whether traceback or environmental evidence determined that the outbreak was foodborne, with an undetermined vehicle.

Indicates whether other evidence determined that the outbreak was foodborne, with an undetermined vehicle.

Implicated food(s)

Indicates whether the vehicle was confirmed or suspected.

For the implicated food, indicates reason(s) suspected or confirmed.

For the implicated food suspected or investigated, indicates a single ingredient

Among the ingredients previously listed, indicates a contaminated ingredient(s), 0="False", 1 or -1 ="True"
Format depends on export and import program

Number of cases exposed to the implicated food

For the implicated food, indicates method of processing

For the implicated food, indicates method of preparation

For the implicated food, indicates level of preparation

Indicates if the implicated food was imported into the US, 0="No", 1 or -1="Yes" 9="Unknown", Null (blank)=No Response

If the contaminated food was imported, indicate the name of the country if known.

Indicates if the food product was both produced under domestic regulatory oversight (commercial product produced within US that is regulated by the FDA) and sold, 0="No", 1="Yes", 7="Unknown", Null (blank)=No Response

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

NORS question collected between Jan 2009 and May 2011. This question is no longer being collected, but the data entered during this period has been archived for reference. This variable indicates the number of cases exposed to at least one of the implicated foods reported.

Location of exposure of where food was prepared

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

Comments regarding the location of preparation

Location of exposure of where food was eaten

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

Comments regarding the location of exposure

Indicates whether a kitchen manager at the location of preparation was certified in food safety (e.g., ServSafe). 0="No", 1="True", 7="Unknown", null (blank) = no response

Indicates if contributing factor is unknown, 0="False", 1 or -1="True". Format depends on export and import program

Name of contributing factor

Type of contributing factor

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

Indicates if point of contamination occurred "Before preparation," "Preparation," or "Unknown."

If point of contamination occurred 'before preparation', variable indicates during preharvest, 0="False", 1 or -1="True". Format depends on export and import program

If point of contamination occurred 'before preparation', variable indicates during processing, 0="False", 1 or -1="True". Format depends on export and import program

If point of contamination occurred 'before preparation', variable indicates when is unknown, 0="False", 1 or -1="True". Format depends on export and import program

Environmental evidence is the reason why the confirmed or suspected point of contamination was assumed, 0="False", 1 or -1="True". Format depends on export and import program

Indicates if epidemiologic evidence is the reason why the confirmed or suspected point of contamination was assumed, 0="False", 1 or -1="True". Format depends on export and import program

Lab evidence is the reason why the confirmed or suspected point of contamination was assumed, 0="False", 1 or -1="True". Format depends on export and import program

Indicates if prior experience is the reason why the confirmed or suspected point of contamination was assumed, 0="False", 1 or -1="True". Format depends on export and import program

Indicates if food worker was implicated, 0="No", 1="True", 7="Unknown", null (blank) = no response

If food worker was implicated, indicates type of evidence that implicated the food worker

Did the outbreak involve a single or multiple schools? 0="False", 1 or -1="True", Format depends on export and import program

Number of schools if multiple schools were involved in the outbreak

Approximate number of students enrolled in the school

Indicates if the number of students enrolled is unknown. Format depends on import and export program, 0="False", 1 or -1="True".

Indicates grade level of the students in the outbreak

Primary funding of involved schools

Indicates the preparation of the implicated item

Indicates how many times the school cafeteria or kitchen has been inspected in the last 12 months by state, county, or local health departments.

Indicate whether the school involved in the outbreak has a HACCP plan in place for the school feeding program, 0="No", 1="Yes", 7="Unknown or Undetermined", Null (blank)= No Response

Was implicated food item provided to the school through the National School Lunch/Breakfast Program? 0="No", 1="Yes", 7="Unknown or Undetermined", Null (blank)= No Response

If school participated in the National School Lunch/Breakfast program, indicates the source of the implicated food items

Percentage of ill persons that ate ground beef raw or undercooked

Indicates whether the ground beef was case ready, 0="No", 1="Yes", 7="Unknown", Null (blank)= No Response

Indicates if the beef was ground or reground by the retailer, 0="No", 1="Yes", 7="Unknown", Null (blank)= No Response

Additional comments if beef was ground or reground

Phage types of patient isolates. NORS question collected between Jan 2009 and March 2017. This question is no longer being collected, but the data entered during this period has been archived for reference.

If is RDNC ('Reacts, Does Not Conform), then what is the number? NORS question collected between Jan 2009 and March 2017. This question is no longer being collected, but the data entered during this period has been archived for reference.

Indicate if the eggs were 'in shell, unpasteurized,' 'in shell, pasteurized,' 'packaged liquid or dry,' 'stored with inadequate refrigeration during or after sale,' 'consumed raw,' 'consumed undercooked,' or 'pooled'

Indicates if Salmonella enteritis was identified at the farm where the eggs originated, 0="No", 1="Yes", 7="Unknown", Null (blank)= No Response

Egg comments/remarks

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Name of major setting where exposure occurred.

Additional details regarding major setting of exposure

Number of residents, guests, passengers, patients, etc. who were exposed in the major setting.

Number of staff, crew, etc. who were exposed in the major setting.

Number of residents, guests, passengers, patients, etc. who were exposed in the major setting and became ill during the outbreak.

Number of staff, crew, etc. who were exposed in the major setting and became ill during the outbreak.

The attack rate among residents and guests, calculated by dividing the number of residents and guests ill by the number exposed.

The attack rate among staff and crew, calculated by dividing the number of staff and crew ill by the number exposed.

The name(s) of any additional settings of exposure for the outbreak

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

Additional details regarding other settings of exposure

Indicates whether any case-patients report travel prior to illness onset.

Indicates whether travel was international, domestic, both, or unknown.

Indicates whether any confirmed, suspected, or probable case-patients were immunocompromised.

Indicates whether any confirmed, suspected, or probable cases were among men who have sex with men.

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Indicates if an animal vehicle was not identified for the outbreak. Format depends on import and export program, 0="False", 1 or -1="True".

Indicates whether epidemiologic evidence determined that the outbreak was animal contact, with an undetermined vehicle.

Indicates whether laboratory evidence determined that the outbreak was animal contact, with an undetermined vehicle.

Indicates whether traceback or environmental evidence determined that the outbreak was animal contact, with an undetermined vehicle.

Indicates whether other evidence determined that the outbreak was animal contact, with an undetermined vehicle.

Type(s) of animals involved

Specific animal type

Indicate whether the animal vehicle was confirmed or suspected.

Indicates whether epidemiologic evidence helped identify the confirmed or suspected animal vehicle.

Indicates whether laboratory evidence helped identify the confirmed or suspected animal vehicle.

Indicates whether environmental or traceback evidence helped identify the confirmed or suspected animal vehicle.

Indicates whether other evidence helped identify the confirmed or suspected animal vehicle.

Name(s) of setting(s) of exposure.

Indicates whether pet food or animal feed was implicated as a potential source of the outbreak. 0= "No", 1= "Yes", 7= "Unknown", Null (blank)= No Response

Indicates the type of pet food implicated.

Indicates whether any cases had exposure to livestock or household pets that were experiencing diarrhea. 0= "No", 1= "Yes", 7= "Unknown", Null (blank)= No Response

Indicates whether the Compendium of Measures to Prevent Disease Associated with Animals in Public Settings was used in the investigation. 0= "No", 1= "Yes", 7= "Unknown", Null (blank)= No Response

Indicates what preventions measures or recommendations were used to stop the outbreak.

Animal remarks/comments

Indicates whether or not additional rows of data are available in NORS that could not be downloaded in the flat file structure due to size limits. 0="False", 1="True" Format depends on export and import program

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The implicated type of water exposure for a waterborne disease outbreak.

Estimated total number of persons with the primary water exposure, as determined by the epidemiologic investigation.

Question: Were data collected from comparison groups to estimate risk? 0= "No", 1= "Yes", 7= "Unknown", Null (blank)= No Response

Question: If response to EstimatedRisk=0 (No) or Unknown (7), was water the only common source shared by persons who were ill? 0= "No", 1= "Yes", 7= "Unknown", Null (blank)= No Response

A vehicle of pathogen transmission, such as an exposure (e.g., swimming pool use) or setting (e.g., hotel stay), measured in an epidemiologic study.

The total number of people that reported a specific exposure in an epidemiologic study

The total number of people that reported a specific exposure who also met the case definition in an epidemiologic study.

The total number of people that did not report a specific exposure in an epidemiologic study

The total number of people that did not report a specific exposure and did meet the case definition in an epidemiologic study.

The percentage of people that reported a specific exposure and met the case definition in an epidemiologic study. This attack rate is automatically calculated by NORS using the following equation:

$AttackRate = \frac{IllExposed}{TotalExposed}$

The odds ratio, with reference to a vehicle of transmission investigated in a waterborne disease outbreak. Displays value up to two decimal places.

The relative risk, with reference to a vehicle of transmission investigated in a waterborne disease outbreak. Displays value up to two decimal places.

The p-value, with reference to a vehicle of transmission investigated in a waterborne disease outbreak. Displays value up to four decimal places.

The confidence interval, with reference to a vehicle of transmission investigated in a waterborne disease outbreak. The format for this variable is ##.## - ##.##.

Provides information from the epidemiologic investigation about the attack rate for cases who were residents of the reporting state.

Provides information from the epidemiologic investigation about the attack rate for cases who were not residents of the reporting state.

The percentage of primary cases in a waterborne disease outbreak that live in the reporting state.

Question: Was exposure associated with a specific event or gathering? 0= "No", 1= "Yes", 7= "Unknown", Null (blank)= No Response

The type of event or gathering (If the water exposure implicated in the outbreak investigation was associated with a specific event or gathering).

Date that the event ended.

Date that the event started.

Indicates ingestion is a route of entry for the waterborne disease outbreak. An outbreak may have more than one route of entry.

Indicates contact is a route of entry for the waterborne disease outbreak. An outbreak may have more than one route of entry.

Indicates inhalation is a route of entry for the waterborne disease outbreak. An outbreak may have more than one route of entry.

Indicates 'other' as a route of entry for the waterborne disease outbreak. An outbreak may have more than one route of entry.

Indicates route of entry for the waterborne disease outbreak is unknown.

Designates whether or not a reported etiologic agent for a waterborne disease outbreak is a confirmed or a suspected etiology.

Genus reported in the Etiology table for waterborne disease outbreaks. This variable is also used to report chemicals and toxins. The Etiology table may be used to report confirmed and suspected etiologies, including when an outbreak etiology is unknown.

Species reported in the Etiology table for waterborne disease outbreaks.

Serotype/Serogroup/Serovar reported in the Etiology table for waterborne disease outbreaks.

Genotype/Subtype reported in the Etiology table for waterborne disease outbreaks.

Polymerase type of a norovirus strain identified as an outbreak etiology

Capsid type of a norovirus strain identified as an outbreak etiology

Information on what types of specimens or samples this microbial pathogen or chemical/toxin was detected in (e.g., clinical specimens)

The number of people tested for an etiologic agent in a waterborne disease outbreak.

The number of people tested for an etiologic agent in a waterborne disease outbreak who tested positive. The number should be less than or equal to the number of people reported in TotalTested

Indicates whether listed etiology is bacterial, chemical, viral, parasitic, etc. Value is automatically assigned for analysis purposes. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more variables ending in The CDC system that contains data about this isolate (e.g., PulseNet, CaliciNet)

A unique identifier by the CDC laboratory to assist with data management (e.g., PulseNet Tracking Number)

A unique identifier assigned by the state laboratory to assist with tracking or other aspects of data management.

The specimen profile for a clinical isolate. For example, the CDC PulseNet pattern designation for enzyme 1. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more variables ending in that same number CDC NARMS predicted resistance pattern based on resistance determinants.

CDC NARMS antimicrobial susceptibility testing (AST) results.

Indicates additional isolate information is available in NORS and was not downloaded due to file size restrictions

Question: Were clinical diagnostic specimens taken from persons? 0= "No", 1= "Yes", 7= "Unknown", Null (blank)= No Response

The number of persons from whom clinical specimens were collected, if clinical diagnostic specimens were collected (waterborne disease outbreak).

The type(s) of clinical specimens collected for a waterborne disease outbreak. Autopsy and biopsy specimens may have additional descriptions in the SpecimenSubtype field.

A variable that provides more information about autopsy and biopsy specimens collected in a waterborne disease outbreak.

Indicates that a clinical specimen was tested for bacteria in a waterborne disease outbreak.

Indicates that a clinical specimen was tested for chemicals in a waterborne disease outbreak.

Indicates that a clinical specimen was tested for fungi in a waterborne disease outbreak.

Indicates that a clinical specimen was tested for parasites in a waterborne disease outbreak.

Indicates that a clinical specimen was tested for viruses in a waterborne disease outbreak.

Indicates that a clinical specimen was tested for an etiologic agent category not listed in a waterborne disease outbreak.

Indicates that the etiologic agent category that a clinical specimen was tested for in a waterborne disease outbreak is unknown.

Indicates that specimens/samples were tested using chemical testing methods. 0="False", 1or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using culture methods. 0="False", 1or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using DNA or RNA amplification methods. 0="False", 1or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using microscopy methods. 0="False", 1or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using serological or immunological methods. 0="False", 1or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using tissue culture infectivity assay. 0="False", 1or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using another method. 0="False", 1or -1="True" Format depends on export and import program

Indicates that specimens/samples were tested using an unknown methods. 0="False", 1or -1="True" Format depends on export and import program

Indicates whether any samples were tested for antimicrobial resistance. 0="No", 1="Yes", 7="Unknown", null (blank) = no response

Indicates where AST was performed ('Clinical Lab', 'Public Health Lab', 'CDC-NARMS', 'Other', 'Unknown')

Question: "If yes, were any antimicrobial resistant strains associated with the outbreak?" 0= "No", 1= "Yes", 7= "Unknown", Null (blank)= No

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Question: Was water tested? 0= "No", 1= "Yes", 7= "Unknown", Null (blank)= No Response

Indicates additional water sample information is available in NORS and was not downloaded due to file size restrictions

Indicates additional sample source information is available in NORS and was not downloaded due to file size restrictions

Indicates additional indicator information is available in NORS and was not downloaded due to file size restrictions

Indicates additional isolate information is available in NORS and was not downloaded due to file size restrictions

A description of the source or location of a water sample (e.g., pool, lake, hot water tank, cooling tower).

Written description of the sample (e.g., more detailed information about where it was collected)

Date that the water sample was collected and onsite measurements such as temperature, volume and pH were taken.

The volume of water collected for a water sample.

The unit of measure for the Volume variable.

The temperature of the water sample.

The unit of measure for the Temperature variable.

The residual/free disinfectant level measured for a water sample.

The unit of measure associated with the residual/free disinfectant level measured for a water sample.

The combined disinfectant level measured for a water sample.

The unit of measure associated with the combined disinfectant level measured for a water sample.

The pH measured for a water sample. Displays value up to two decimal places.

The turbidity (in NTU) measured for a water sample.

The type of water quality indicator (e.g., fecal coliforms) measured for a water sample. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more variables ending in that same number (e.g., Concentration reported for IndicatorType based on results from laboratory testing. See IndicatorUnit for the measurement unit associated with this value. Displays as a whole number. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it Unit of measurement (e.g., CFU/100mL) for IndicatorConc based on results from laboratory testing. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more variables ending in that same number Genus reported for water samples collected during waterborne disease outbreak investigations. This variable is also used to report chemicals and toxins. Positive and negative test results may be reported. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies Species reported for water samples collected during waterborne disease outbreak investigations. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more variables ending in that same number (e.g., Serotype/Serogroup/Serovar reported for water samples collected during waterborne disease outbreak investigations. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more variables ending in that Genotype/Subtype reported for water samples collected during waterborne disease outbreak investigations. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more variables ending in that same number Polymerase type of a norovirus strain identified in a water sample

Capsid type of a norovirus strain identified in a water sample

The PFGE pattern for an isolate obtained from a water sample. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more variables ending in that same number (e.g., MolecularDesignation1_1 , Indicates whether or not an water sample tested positive for a microorganism, chemical or toxin. Value is automatically assigned for analysis purposes. 0="False, or Test Results Negative", 1="Yes, or Test Results Positive" Format depends on export and import program. For variables ending in the format #_#, the first Concentration reported for a water sample that was analyzed for microorganisms, chemicals or toxins. See AnalysisUnit for the measurement unit associated with this value. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and Unit of measurement (e.g., CFU/100mL) for AnalysisConc based on results from laboratory testing for microorganisms, chemicals or toxins. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more A description of the laboratory methods (e.g., microscopy, RNA/DNA tests) used to test for a microorganism, chemical or toxin in a water sample. For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more The standardized laboratory method used to test a water sample for a microorganism, chemical or toxin. Abbreviated format used in NORS has the information ordered as follows: Method source, Method number and Method summary. E.g., ASTM D1293-pH (routine/continuous measurement). For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more The CDCID is an autogenerated unique ID that represents a single NORS report. Links record from a single NORS report across all tables of the flattened database.

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Indicates whether additional water type information is available in NORS and was not downloaded due to file size restrictions

Indicates whether additional treatment information is available in NORS and was not downloaded due to size restrictions

Indicates whether additional fill water information is available in NORS and was not downloaded due to size restrictions

Water venue of exposure (e.g. spa/whirlpool/hot tub; pool-swimming pool; pool-waterpark)

Water venue subtype (i.e. indoor, outdoor, or unknown)

Setting of exposure (e.g. club, requiring membership; hotel/motel/lodge/inn; waterpark)

The usual type of water treatment provided at venue (e.g. no treatment, coagulation, disinfection, flocculation, filtration [pool], unknown). For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more variables

Venue treatment subtype (disinfection or pool filtration: e.g., UV; chlorine dioxide; bag filter; cartridge filter; unknown). For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back

Chlorination subtype (chlorine disinfection only: e.g., gaseous; sodium hypochlorite; cyanurates/stabilized chlorine). For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back

Fill water type (e.g., public water supply; sea water; untreated ground or surface water; unknown). For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more variables ending in that same number (e.g.,

If public water supply, usual water treatment provided before coming to the venue (e.g., no treatment; disinfection; filtration [treatment plant]; unknown). For variables ending in the format #_#, the first number uniquely identifies its

If public water supply, fill water treatment subtype (disinfection or filtration: e.g., UV; chlorine dioxide; bag filter; cartridge filter; unknown). For variables ending in the format #_#, the first number uniquely identifies its order in a set. The second number identifies the set to which it belongs and relates back to one or more

Question: Did the venue meet state or local recreational water quality regulations? 0= "No", 1= "Yes", 7= "Unknown", 9= "Not Applicable", Null (blank)= No Response

Additional description, if TreatedQuality=0 (No)

Question: Was there a pool operator on the payroll with state-approved training or certification? 0= "No", 1= "Yes", 7= "Unknown", Null (blank)= No Response

Water venue implicated (e.g., canal; lake; river/stream; ocean)

IF SPRING OR HOT SPRING, water venue subtype description (i.e. 'indoor', 'outdoor' or 'unknown')

Setting of exposure (e.g., beach-public; camp/cabin/recreational area)

Question: Did the venue meet state or local recreational water quality regulations? 0= "No", 1= "Yes", 7= "Unknown", 9= "Not Applicable", Null (blank)= No Response

Additional description, if UntreatedQuality=0 (No)

Question: Did the venue meet Environmental Protection Agency (EPA) recreational water quality standards? 0= "No", 1= "Yes", 7= "Unknown", 9= "Not Applicable", Null (blank)= No Response

Additional description, if UntreatedEPA=0 (No)

Water system (e.g., commercially-bottled water, community water system, individual water system)

Public water system EPA ID number

Water source (select ground water, surface water or unknown)

Water source description (e.g. spring; well; lake)

Setting of exposure (e.g., airport, mobile home park)

USUAL water treatment provided (e.g.,no treatment, disinfection, home filtration)

Water treatment subtype (disinfection or filtration: e.g., boiling; chlorine; rapid sand filter; reserve osmosis)

Question: Did the drinking water system have any monitoring violations in the 1 month prior to the outbreak? 0= "No", 1= "Yes", 7= "Unknown", 9= "Not Applicable", Null (blank)= No Response

Additional description, if DrinkingMonth = 1 (Yes)

Question: Did the drinking water system have any maximum contaminant level (MCL) violations in the 1 month prior to the outbreak? 0= "No", 1= "Yes", 7= "Unknown", 9= "Not Applicable", Null (blank)= No Response

Additional description, if DrinkingMCL = 1 (Yes)

Question: Did the drinking water system have any violations in the 12 months prior to the outbreak? 0= "No", 1= "Yes", 7= "Unknown", 9= "Not Applicable", Null (blank)= No Response

Additional description, if PreviousYearResponse = 1 (Yes)

What was the intended use for the implicated water? (i.e. cooling/air conditioning, mister, ornamental, industrial/occupational, agricultural irrigation, wast water, other (specify), unknown)

Water type (e.g., cooling tower; drainage ditch; fountain - ornamental)

Setting of exposure (e.g, airport; hospital/health care facility; nursing home; park-state park)

Usual water treatment provided (e.g., no treatment; disinfection; settling/sedimentation)

Water treatment subtype (disinfection or filtration: e.g., boiling; chlorine; rapid sand filter; reverse osmosis)

Remarks field at the end of the NORS report for additional information about the waterborne disease outbreak investigation.

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Indicates a problem with the source water contributed to the outbreak 0='No', 1='Yes' 7='Unknown'

Indicates a problem with the water treatment contributed to the outbreak 0='No', 1='Yes' 7='Unknown'

Indicates a problem with the water distribution system contributed to the outbreak 0='No', 1='Yes' 7='Unknown'

Indicates a problem with the water outside of a water utility jurisdiction contributed to the outbreak 0='No', 1='Yes' 7='Unknown'

Contributing factor

Is the contributing factor confirmed or suspected? 1='Confirmed', 0='Suspected'

What type of water exposure is associated with the selected contributing factor? ('Treated', 'Untreated', 'Other/Unknown', 'Drinking-source water', 'Drinking-treatment', 'Drinking-distribution and storage', 'Drinking-point-of-use')

Indicates whether additional information is available in NORS and was not downloaded due to size restrictions

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Strength of evidence classification assigned to the outbreak report

Numeric variable for month of onset.

4 digit year of outbreak onset, based on date first ill.

Surveillance period for the CDC surveillance summary publication that included this report.

Categorical variable for water exposure.

Description of which samples had confirmed coliforms (Clinical Specimens, Water Samples, Both)

Indicates Bacterial etiologic agent

Indicates Fungal etiologic agent

Indicates Viral etiologic agent

Indicates Chemical etiologic agent

Indicates Parasitic etiologic agent

Indicates other etiologic agent not listed by available categories.

Indicates etiologic agent is unknown

Indicates AGI as predominant illness according to surveillance system definitions. (e.g. SKIN='Skin infection, burn, or other condition', AGI='Acute Gastrointestinal Illness', NEURO='Neurologic Symptoms/Conditions', EYE='Keratitis, Infection, Irritation', ARI='Acute Respiratory Illness, HEP='Hepatitis', EAR='Ear Infection, Irritation', Indicates ARI as predominant illness according to surveillance system definitions. (e.g. SKIN='Skin infection, burn, or other condition', AGI='Acute Gastrointestinal Illness', NEURO='Neurologic Symptoms/Conditions', EYE='Keratitis, Infection, Irritation', ARI='Acute Respiratory Illness, HEP='Hepatitis', EAR='Ear Infection, Irritation', Indicates Skin as predominant illness according to surveillance system definitions. (e.g. SKIN='Skin infection, burn, or other condition', AGI='Acute Gastrointestinal Illness', NEURO='Neurologic Symptoms/Conditions', EYE='Keratitis, Infection, Irritation', ARI='Acute Respiratory Illness, HEP='Hepatitis', EAR='Ear Infection, Irritation', Indicates Eye as predominant illness according to surveillance system definitions. (e.g. SKIN='Skin infection, burn, or other condition', AGI='Acute Gastrointestinal Illness', NEURO='Neurologic Symptoms/Conditions', EYE='Keratitis, Infection, Irritation', ARI='Acute Respiratory Illness, HEP='Hepatitis', EAR='Ear Infection, Irritation', Indicates Ear as predominant illness according to surveillance system definitions. (e.g. SKIN='Skin infection, burn, or other condition', AGI='Acute Gastrointestinal Illness', NEURO='Neurologic Symptoms/Conditions', EYE='Keratitis, Infection, Irritation', ARI='Acute Respiratory Illness, HEP='Hepatitis', EAR='Ear Infection, Irritation', Indicates Hep as predominant illness according to surveillance system definitions. (e.g. SKIN='Skin infection, burn, or other condition', AGI='Acute Gastrointestinal Illness', NEURO='Neurologic Symptoms/Conditions', EYE='Keratitis, Infection, Irritation', ARI='Acute Respiratory Illness, HEP='Hepatitis', EAR='Ear Infection, Irritation', Indicates LD as predominant illness according to surveillance system definitions. (e.g. SKIN='Skin infection, burn, or other condition', AGI='Acute Gastrointestinal Illness', NEURO='Neurologic Symptoms/Conditions', EYE='Keratitis, Infection, Irritation', ARI='Acute Respiratory Illness, HEP='Hepatitis', EAR='Ear Infection, Irritation', Indicates Lepto as predominant illness according to surveillance system definitions. (e.g. SKIN='Skin infection, burn, or other condition', AGI='Acute Gastrointestinal Illness', NEURO='Neurologic Symptoms/Conditions', EYE='Keratitis, Infection, Irritation', ARI='Acute Respiratory Illness, HEP='Hepatitis', EAR='Ear Infection, Irritation', Indicates Neuro as predominant illness according to surveillance system definitions. (e.g. SKIN='Skin infection, burn, or other condition', AGI='Acute Gastrointestinal Illness', NEURO='Neurologic Symptoms/Conditions', EYE='Keratitis, Infection, Irritation', ARI='Acute Respiratory Illness, HEP='Hepatitis', EAR='Ear Infection, Irritation', Indicates PF as predominant illness according to surveillance system definitions. (e.g. SKIN='Skin infection, burn, or other condition', AGI='Acute Gastrointestinal Illness', NEURO='Neurologic Symptoms/Conditions', EYE='Keratitis, Infection, Irritation', ARI='Acute Respiratory Illness, HEP='Hepatitis', EAR='Ear Infection, Irritation', Indicates LD as predominant illness according to surveillance system definitions. (e.g. SKIN='Skin infection, burn, or other condition', AGI='Acute Gastrointestinal Illness', NEURO='Neurologic Symptoms/Conditions', EYE='Keratitis, Infection, Irritation', ARI='Acute Respiratory Illness, HEP='Hepatitis', EAR='Ear Infection, Irritation', Indicates Leg (Legionella) as predominant illness according to surveillance system definitions.

Indicates an additional predominant illness, not listed by provided categories according to surveillance system definitions.

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data:
1='Untreated surface water'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
2='Untreated ground water'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
3='Treatment deficiency (e.g., temporary interruption of disinfection, chronically inadequate disinfection, or inadequate or no filtration)'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
4='Distribution system deficiency, including storage (e.g., cross-connection, backflow, contamination of water mains during construction or repair)'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
5a='Legionella spp. in drinking water system'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
5b='Legionella spp. in other non-recreational water system'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
5c='Legionella spp. in unknown water use'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
6='Plumbing system deficiency after the water meter or property line (e.g., cross-connection, backflow, or corrosion products)'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
7='Deficiency in building/home-specific water treatment after the water meter or property line'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
8='Deficiency or contamination of equipment using or distributing water (e.g., drink-mix machines)'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
9='Contamination or treatment deficiency during commercial bottling'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
10a='Contamination during shipping, hauling, or storage: Drinking-tap water '

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
10b='Contamination during shipping, hauling, or storage: Drinking-commercially-bottled water'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
11a='Contamination at point of use: Tap'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
11b='Contamination at point of use: Hose'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
11c='Contamination at point of use: Commercially-bottled water'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
11d='Contamination at point of use: Container, bottle, or pitcher'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
11e='Contamination at point of use: Unknown'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
12='Drinking or contact with other non-recreational water'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
13a='Current treatment processes not expected to remove a chemical contaminant (e.g., pesticide contamination of ground water treated with disinfection only): Surface water'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
13b='Current treatment processes not expected to remove a chemical contaminant (e.g., pesticide contamination of ground water treated with disinfection only): Ground water'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
99a='Unknown/Insufficient information: Drinking water-tap water'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
99b='Unknown/Insufficient information: Drinking water- commercially-bottled water'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
99c='Unknown/Insufficient information: Other non-recreational water'

Deficiency classification(s) assigned to an outbreak based on a formal review of the outbreak report data.
99d='Unknown/Insufficient information: Unknown water use'

Summary description of the outbreak

Presence/Absence of coliforms. 0="False", 1="True" Format depends on export and import program

Description of when coliforms were found in relation to the outbreak, if applicable ('Before Outbreak', 'During Outbreak', 'After Outbreak')

Description of disinfection time in relation to outbreak, if applicable ('Before Outbreak', 'During Outbreak', 'After Outbreak')

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Approximate percentage of cases in each age group among the total number of primary cases for whom information is available

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Number of persons who were exposed and became ill

Total number of persons exposed to the implicated vehicle. The attack rate is applied to persons in a cohort who were exposed to the implicated vehicle. If the vehicle is unknown, then the attack rate should not be calculated.

Name of contributing factor

Type of contributing factor

NORS value for internal data management processes. Identifies the individual implicated food during the outbreak; each outbreak may have multiple implicated foods. Variable is a primary key.

For the implicated food, indicates method of preparation.

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Earliest date of reported illness onset

Primary mode of transmission

Indicates whether the report is active, finalized, or deleted

Reporting site of the report author

Name of the NORS agency reporting the outbreak

User ID of person who created the report

Indicates whether the finalization issue for the report is open or closed

Name of the finalization issue flagged for the report

Response to finalization issue, if addressed (e.g., "Verified" for a checkbox, or free text for explanations)

Description of the finalization issue flagged for the report
