

2014-0099 CDAS File Data Dictionary

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Document Summary

Property	Value
Document Title	2014-0099 CDAS File: Data Dictionary
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Sections	5
Entries	78
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2014-0099 CDAS File: Data Dictionary

Section 1: Identifiers

Class	Variable	Label	Description	Format Text
01. Principal	plco_id	PLCO ID		Char
01. Principal	selection_build	Masterfile Build Used for Population Selection		Char

Section 2: Study

Class	Variable	Label	Description	Format Text
01. Principal	is_case	Case Status		0="Non-Case" 1="Case"

Section 3: Match

Class	Variable	Label	Description	Format Text
01. Principal	match_agelevel	Matched Age Level at Randomization		0=" <= 59" 1="60-64" 2="65-69" 3="70-74"
01. Principal	match_drawmonth	Matched Month of Earliest Draw		.M="Missing"
01. Principal	match_gender	Matched Gender		"F"="Female" "M"="Male"
01. Principal	match_race7	Matched Race		1="White, Non-Hispanic" 2="Black, Non-Hispanic" 3="Hispanic" 4="Asian"
01. Principal	match_sample_startyr	Matched Earliest Study Year of Draw		Numeric

Section 4: Specimen

Class	Variable	Label	Description	Format Text
01. Principal	sample_drawdays0	T0 Days from Randomization to Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_drawdays1	T1 Days from Randomization to Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_drawdays2	T2 Days from Randomization to Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_drawdays4	T4 Days from Randomization to Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_drawdays5	T5 Days from Randomization to Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_studyyr0	T0 Study Year of Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_studyyr1	T1 Study Year of Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_studyyr2	T2 Study Year of Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_studyyr4	T4 Study Year of Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_studyyr5	T5 Study Year of Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_time0	T0 Time of Day of Blood Collection	Time of day is the hour within which the blood was collected, based on the 24 hour clock.	numeric .N="Not Applicable"
01. Principal	sample_time1	T1 Time of Day of Blood Collection	Time of day is the hour within which the blood was collected, based on the 24 hour clock.	numeric .N="Not Applicable"
01. Principal	sample_time2	T2 Time of Day of Blood Collection	Time of day is the hour within which the blood was collected, based on the 24 hour clock.	numeric .N="Not Applicable"
01. Principal	sample_time4	T4 Time of Day of Blood Collection	Time of day is the hour within which the blood was collected, based on the 24 hour clock.	numeric .N="Not Applicable"
01. Principal	sample_time5	T5 Time of Day of Blood Collection	Time of day is the hour within which the blood was collected, based on the 24 hour clock.	numeric .N="Not Applicable"
01. Principal	sample_yr0	T0 Calendar Year of Blood Collection		Numeric .N="Not Applicable"

Class	Variable	Label	Description	Format Text
01. Principal	sample_yr1	T1 Calendar Year of Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_yr2	T2 Calendar Year of Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_yr4	T4 Calendar Year of Blood Collection		Numeric .N="Not Applicable"
01. Principal	sample_yr5	T5 Calendar Year of Blood Collection		Numeric .N="Not Applicable"
09. Secondary	sample_flaw_stat us0	Worst Flaw Status for Vial 0		.N="Not Applicable" 0="Empty/Lost/Broken or otherwise Unusable" 1="Icteric, Lipemic, or Uncontrolled Thaw" 3="Vial is both Hemolyzed and Controlled Thaw" 4="Controlled Thaw - not Hemolyzed"
09. Secondary	sample_flaw_stat us1	Worst Flaw Status for Vial 1		.N="Not Applicable" 0="Empty/Lost/Broken or otherwise Unusable" 4="Controlled Thaw - not Hemolyzed" 5="Hemolyzed - no Thaws"
09. Secondary	sample_flaw_stat us2	Worst Flaw Status for Vial 2		.N="Not Applicable" 0="Empty/Lost/Broken or otherwise Unusable" 4="Controlled Thaw - not Hemolyzed"
09. Secondary	sample_flaw_stat us4	Worst Flaw Status for Vial 4		.N="Not Applicable" 0="Empty/Lost/Broken or otherwise Unusable" 4="Controlled Thaw - not Hemolyzed"
09. Secondary	sample_flaw_stat us5	Worst Flaw Status for Vial 5		.N="Not Applicable" 0="Empty/Lost/Broken or otherwise Unusable" 4="Controlled Thaw - not Hemolyzed"

Section 5: Results

Class	Variable	Label	Description	Format Text
01. Principal	il10_pg_ml_1	Interleukin 10 (pg/mL) T1		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il10_pg_ml_2	Interleukin 10 (pg/mL) T2		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il10_pg_ml_4	Interleukin 10 (pg/mL) T4		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il10_pg_ml_5	Interleukin 10 (pg/mL) T5		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il12p70_pg_ml_1	Interleukin 12 p70 (pg/mL) T1		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il12p70_pg_ml_2	Interleukin 12 p70 (pg/mL) T2		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il12p70_pg_ml_4	Interleukin 12 p70 (pg/mL) T4		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il12p70_pg_ml_5	Interleukin 12 p70 (pg/mL) T5		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il13_pg_ml_1	Interleukin 13 (pg/mL) T1		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il13_pg_ml_2	Interleukin 13 (pg/mL) T2		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il13_pg_ml_4	Interleukin 13 (pg/mL) T4		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il13_pg_ml_5	Interleukin 13 (pg/mL) T5		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"

Class	Variable	Label	Description	Format Text
01. Principal	il4_pg_ml_1	Interleukin 4 (pg/mL) T1		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il4_pg_ml_2	Interleukin 4 (pg/mL) T2		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il4_pg_ml_4	Interleukin 4 (pg/mL) T4		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il4_pg_ml_5	Interleukin 4 (pg/mL) T5		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il6_pg_ml_1	Interleukin 6 (pg/mL) T1		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il6_pg_ml_2	Interleukin 6 (pg/mL) T2		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il6_pg_ml_4	Interleukin 6 (pg/mL) T4		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	il6_pg_ml_5	Interleukin 6 (pg/mL) T5		Numeric .= "Missing" .M= "Missing" .O= "Out of Range"
01. Principal	phadiatop_cat1	Phadiatop Category T1		Char " " = "Missing"
01. Principal	phadiatop_cat2	Phadiatop Category T2		Char " " = "Missing"
01. Principal	phadiatop_cat4	Phadiatop Category T4		Char " " = "Missing"
01. Principal	phadiatop_cat5	Phadiatop Category T5		Char " " = "Missing"
01. Principal	phadiatop_pau_l_1	Phadiatop (pau/L) T1		Numeric .= "Missing" .M= "Missing"
01. Principal	phadiatop_pau_l_2	Phadiatop (pau/L) T2		Numeric .= "Missing" .M= "Missing"
01. Principal	phadiatop_pau_l_4	Phadiatop (pau/L) T4		Numeric .= "Missing" .M= "Missing"
01. Principal	phadiatop_pau_l_5	Phadiatop (pau/L) T5		Numeric .= "Missing" .M= "Missing"

Class	Variable	Label	Description	Format Text
01. Principal	tgf_beta1_pg_ml_1	Transforming Growth Factor Beta 1 (pg/mL) T1		Numeric .= "Missing" .M= "Missing"
01. Principal	tgf_beta1_pg_ml_2	Transforming Growth Factor Beta 1 (pg/mL) T2		Numeric .= "Missing" .M= "Missing"
01. Principal	tgf_beta1_pg_ml_4	Transforming Growth Factor Beta 1 (pg/mL) T4		Numeric .= "Missing" .M= "Missing"
01. Principal	tgf_beta1_pg_ml_5	Transforming Growth Factor Beta 1 (pg/mL) T5		Numeric .= "Missing" .M= "Missing"
01. Principal	tnfa_pg_ml_1	Tumor Necrosis Factor Alpha (pg/mL) T1		Numeric .= "Missing" .M= "Missing"
01. Principal	tnfa_pg_ml_2	Tumor Necrosis Factor Alpha (pg/mL) T2		Numeric .= "Missing" .M= "Missing"
01. Principal	tnfa_pg_ml_4	Tumor Necrosis Factor Alpha (pg/mL) T4		Numeric .= "Missing" .M= "Missing"
01. Principal	tnfa_pg_ml_5	Tumor Necrosis Factor Alpha (pg/mL) T5		Numeric .= "Missing" .M= "Missing"
01. Principal	total_ige_cat1	Total Immunoglobulin E Category T1		Char " " = "Missing"
01. Principal	total_ige_cat2	Total Immunoglobulin E Category T2		Char " " = "Missing"
01. Principal	total_ige_cat4	Total Immunoglobulin E Category T4		Char " " = "Missing"
01. Principal	total_ige_cat5	Total Immunoglobulin E Category T5		Char " " = "Missing"
01. Principal	total_ige_ku_I0	Total Immunoglobulin E (ku/L) T0		Numeric .N= "Not Applicable"
01. Principal	total_ige_ku_I1	Total Immunoglobulin E (ku/L) T1		Numeric .N= "Not Applicable"
01. Principal	total_ige_ku_I2	Total Immunoglobulin E (ku/L) T2		Numeric .N= "Not Applicable"
01. Principal	total_ige_ku_I4	Total Immunoglobulin E (ku/L) T4		Numeric .N= "Not Applicable"
01. Principal	total_ige_ku_I5	Total Immunoglobulin E (ku/L) T5		Numeric .N= "Not Applicable"