

T1DGC.2011.03_TAQMAN Duplicates List Data Dictionary

VARIABLE	ALIAS	TYPE	RANGE
family1	family indicator variable for first pair-wise sample	numeric integer	0=case-control data 1=family data
cohort1	cohort label for first pair wise sample	character categorical	AP=Asia Pacific BDA=BDA/Warren I DAN-SDC=Danish cases-controls GRID=UK GRID HBDI=HBDI UK=United Kingdom
famid1	pedigree: family ID for pair-wise sample. For case-control samples, value is the same as analytic_id1.	numeric continuous	100000+ 200000+ 400000+ 500000+ 20000000+ 40000000+ 50000000+
analytic_id1	analytic id for first pair-wise sample	numeric continuous	10000000+ 20000000+ 40000000+ 50000000+
match1	indicator variable for individuals matched to more than one sample	numeric integer	1-22
family2	family indicator variable for second pair-wise sample	numeric integer	0=case/control 1=family data
cohort2	cohort label for second pair-wise sample	character categorical	AP=Asia Pacific BDA=BDA/Warren I DAN-SDC=Danish cases-controls GRID=UK GRID HBDI=HBDI UK=United Kingdom
famid2	pedigree: family ID for pair wise sample. For case-control samples, value is the same as analytic_id2.	numeric continuous	100000+ 200000+ 400000+ 500000+ 20000000+ 40000000+ 50000000+
analytic_id2	analytic id for second pair-wise sample.	numeric continuous	10000000+ 20000000+ 40000000+ 50000000+
match2	Indicator variable for individuals matched to more than one sample	numeric integer	1-22

VARIABLE	ALIAS	TYPE	RANGE
miss1	count of missing markers for first pair-wise sample only	numeric integer	0-16
miss2	count of missing markers for second pair-wise sample only	numeric integer	0-15
bothm	count of missing markers for both pair wise samples	numeric integer	0-16
matchcnt	count of concordant non-missing markers between pair-wise samples	numeric integer	25-48
matchrate	concordance rate between pair-wise samples using non-missing markers	numeric continuous	0.94-1.0