

HFCeS

HaplotypeFrequency
Creation Service

Input data

ASCII File, XML

Structure

Mandatory	XML Tag	Values / Kind of Val.	Description
Yes	POP_ID	URI from POP DB	Population Identifier
Yes	GT_List	List of GTL	List of Genotypes
Yes	GTL	Record of GTL_Name, GTL_M_List	GTL_M_List is optional
Yes	GTL_Name	GL String	Raw Data
No	GTL_M_List	List of GTLP_Meth	Additional Information on GT, Typing
No	GTLP_Meth	Pair of GTLP_M_Data, GTLP_M_Value	
No	GTLP_M_Data	Free Text	
No	GTLP_M_Value	Free Text	
Yes	GT_Lic	License ID	License under which GT data is available
Yes	METHOD_LIST	List of METHOD	
Yes	METHOD	Record of METH_Type, METH_Value, METH_Comment, METH_Ref	
Yes	METH_Type	Free Text	
Yes	METH_Value	Free Text	
No	METH_Comment	Free Text	
No	METH_REF	Free Text	To be used to refer to an external source
No	HFCeS_COHORT_ID		As an alternative to GT List
No	HFCeS_METHOD_ID		As an alternative to Method List

Output

- An Inputfile to HFCuS
- HFCeS_METHOD_ID
- HFCeS_COHORT_ID

HT Creation (ASCII, XML)

Input Creation Service

Mandatory

TAG	Description	Kind of Value
-----	-------------	---------------

<p>GT Meta Data List GT Meta Data Type GT Meta Data Value</p>	<p>Eg. Typing Method IMGT version...</p>	<p>Text fields</p>
<p>Pop_ID</p>	<p>Population ID</p>	<p>URI</p>
<p>Pop-Desc</p>	<p>Description of Pop</p>	<p>Free Text</p>
<p>GT List</p>	<p>Genotypes (Raw Data)</p>	<p>List of GT</p>
<p>GT</p>	<p>One Genotype</p>	<p>GL - String</p>
<p>Cohort ID</p>	<p>Sample ID</p>	<p>INTEGER</p>
<p>License HT</p>	<p>Useability of Data of HT</p>	<p>List of Licenses</p>
<p>Resolution</p>		<p>GP, guma, gums, u-Field, Serology</p>
<p>Method-ID</p>		
<p>Method</p>	<p>Description of Method</p>	<p>INTEGER</p>
<p>Method-Parameter</p>		<p>Text field</p>
<p>Method-Parameter Value</p>	<p>Part of ↑</p>	<p>Free text Free text, likely TR</p>

rice