

Clinical Interpretation Data Model Schema

id <i>String</i>	The interpretation ID.																				
uuid <i>String</i>	A universally unique identifier for the interpretation.																				
description <i>String</i>	The interpretation description.																				
clinicalAnalysisId <i>String</i>	The clinical analysis ID.																				
methods <i>List<Interpretation Method></i>	List of interpretation methods.<table><tr><td>name <i>String</i></td><td>Interpretation method name, .e.g.: tiering, zetta, team,...</td></tr><tr><td>filters <i>Map<String, Object></i></td><td>Filters.</td></tr><tr><td>panels <i>List<DiseasePanel></i></td><td>Panels.</td></tr><tr><td>dependencies <i>List<Software></i></td><td> List of software dependencies.<table><tr><td>name <i>String</i></td><td>Name.</td></tr><tr><td>version <i>String</i></td><td>Version.</td></tr><tr><td>repository <i>String</i></td><td>Repository.</td></tr><tr><td>commit <i>String</i></td><td>Commit.</td></tr><tr><td>website <i>String</i></td><td>Website.</td></tr><tr><td>params <i>Map<String, String></i></td><td>Parameters.</td></tr></table> </td></tr></table>	name <i>String</i>	Interpretation method name, .e.g.: tiering, zetta, team,...	filters <i>Map<String, Object></i>	Filters.	panels <i>List<DiseasePanel></i>	Panels.	dependencies <i>List<Software></i>	List of software dependencies.<table><tr><td>name <i>String</i></td><td>Name.</td></tr><tr><td>version <i>String</i></td><td>Version.</td></tr><tr><td>repository <i>String</i></td><td>Repository.</td></tr><tr><td>commit <i>String</i></td><td>Commit.</td></tr><tr><td>website <i>String</i></td><td>Website.</td></tr><tr><td>params <i>Map<String, String></i></td><td>Parameters.</td></tr></table>	name <i>String</i>	Name.	version <i>String</i>	Version.	repository <i>String</i>	Repository.	commit <i>String</i>	Commit.	website <i>String</i>	Website.	params <i>Map<String, String></i>	Parameters.
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params <i>Map<String, String></i>	Parameters.																				
primaryFindings <i>List<ClinicalVariant></i>	List of primary findings. See clinical variant data model schema .																				
secondaryFindings <i>List<ClinicalVariant></i>	List of secondary findings. See clinical variant data model schema .																				

analyst <i>Analyst</i>	Analyst carrying out the interpretation. <table> <tr> <td>name <i>String</i></td><td>Name.</td></tr> <tr> <td>email <i>String</i></td><td>E-mail.</td></tr> <tr> <td>company <i>String</i></td><td>Company.</td></tr> </table>	name <i>String</i>	Name.	email <i>String</i>	E-mail.	company <i>String</i>	Company.		
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email <i>String</i>	E-mail.								
company <i>String</i>	Company.								
comments <i>List<Comment></i>	List of comments. <table> <tr> <td>author <i>String</i></td><td>Author.</td></tr> <tr> <td>type <i>String</i></td><td>Type.</td></tr> <tr> <td>message <i>String</i></td><td>Message.</td></tr> <tr> <td>date <i>String</i></td><td>Date.</td></tr> </table>	author <i>String</i>	Author.	type <i>String</i>	Type.	message <i>String</i>	Message.	date <i>String</i>	Date.
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description <i>String</i>	Description.								
date <i>String</i>	Date.								
creationDate <i>String</i>	Creation date.								
modificationDate <i>String</i>	Modification date.								
version <i>int</i>	Version.								
attributes Map<String, Object>	Map of attributes. Users can add custom information in this field. OpenCGA uses this field to store the ClinicalAnalysis object under the key OPENCGA_CLINICAL_ANALYSIS.								