

# Genomic Vocabularies

Ontologies incorporated:

- **HGNC (HUGO Gene Nomenclature Committee)**
- **ClinVar**
- **CIcIC (Clinical Interpretation of Variants in Cancer)**
- **NCIt (National Cancer Institute Thesaurus)**
- **CGI (Cancer Genome Interpreter)**
- **CAP (College of American Pathologists Cancer Checklists)**
- **JAX CKB (The Clinical Knowledgebase by The Jackson Laboratory)**

Ontologies provided HGVS expressions which were split on refseq, version of refseq, sequence type and locations.

concept_name	vocabulary_id	concept_code	gene	refseq	version	seqtype	variant
ABL1:V299L	CGI	ABL1:V299L	ABL1	ABL1		c	895G>C
BCR-ABL V299L	CIViC	1231	ABL1	ENST00000318560	.5	c	895G>C
BCR-ABL V299L	CIViC	1231	ABL1	NM_005157	.5	c	895G>C
NM_005157.6(ABL1):c.895G>C (p. Val299Leu)	ClinVar	362996	ABL1	NM_005157	.6	c	895G>C
V299L	JAX	3783	ABL1	ABL1		c	895G>C

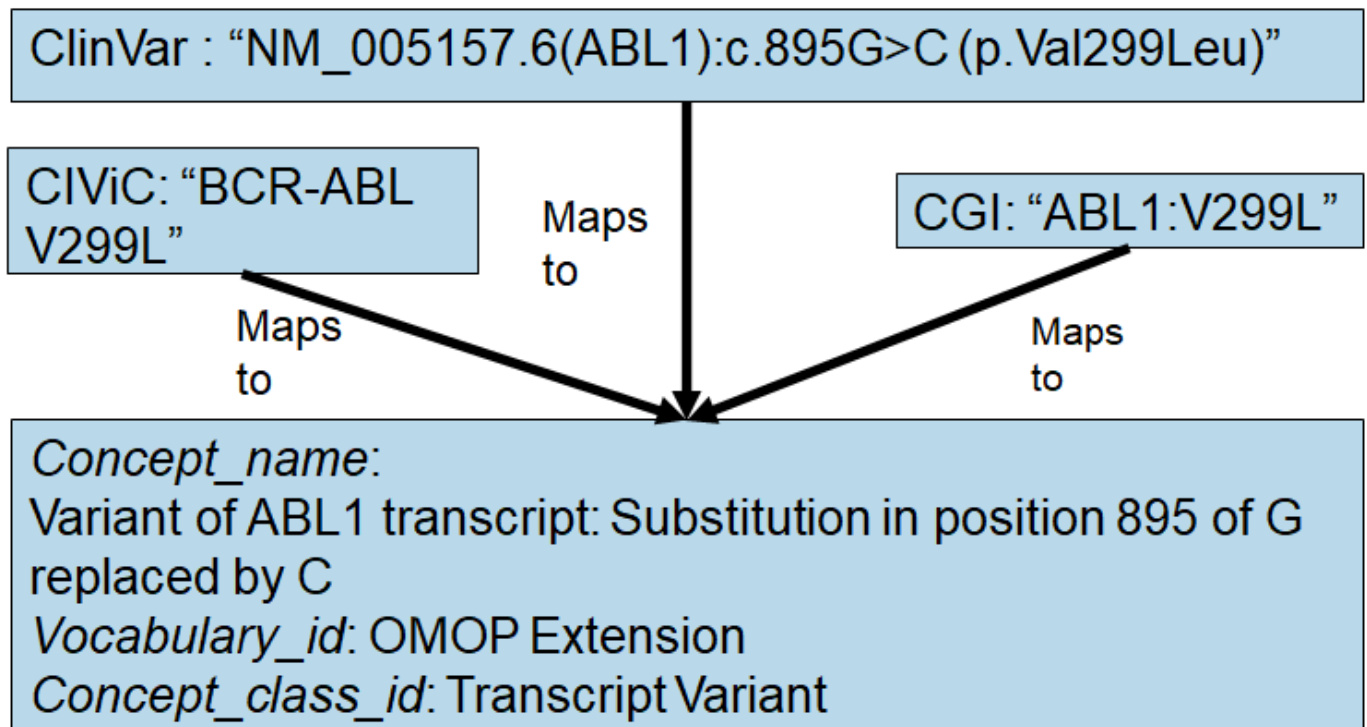
We developed a simple consolidation approach of single nucleotide polymorphism, insertions, deletions and duplications based on gene symbol, sequence type (g, c, p) and locations.

concept_name	vocabulary_id	concept_code	gene	seqtype	variant
ABL1:V299L	CGI	ABL1:V299L	ABL1	c	895G>C
BCR-ABL V299L	CIViC	1231	ABL1	c	895G>C
BCR-ABL V299L	CIViC	1231	ABL1	c	895G>C
NM_005157.6(ABL1):c.895G>C (p.Val299Leu)	ClinVar	362996	ABL1	c	895G>C
V299L	JAX	3783	ABL1	c	895G>C

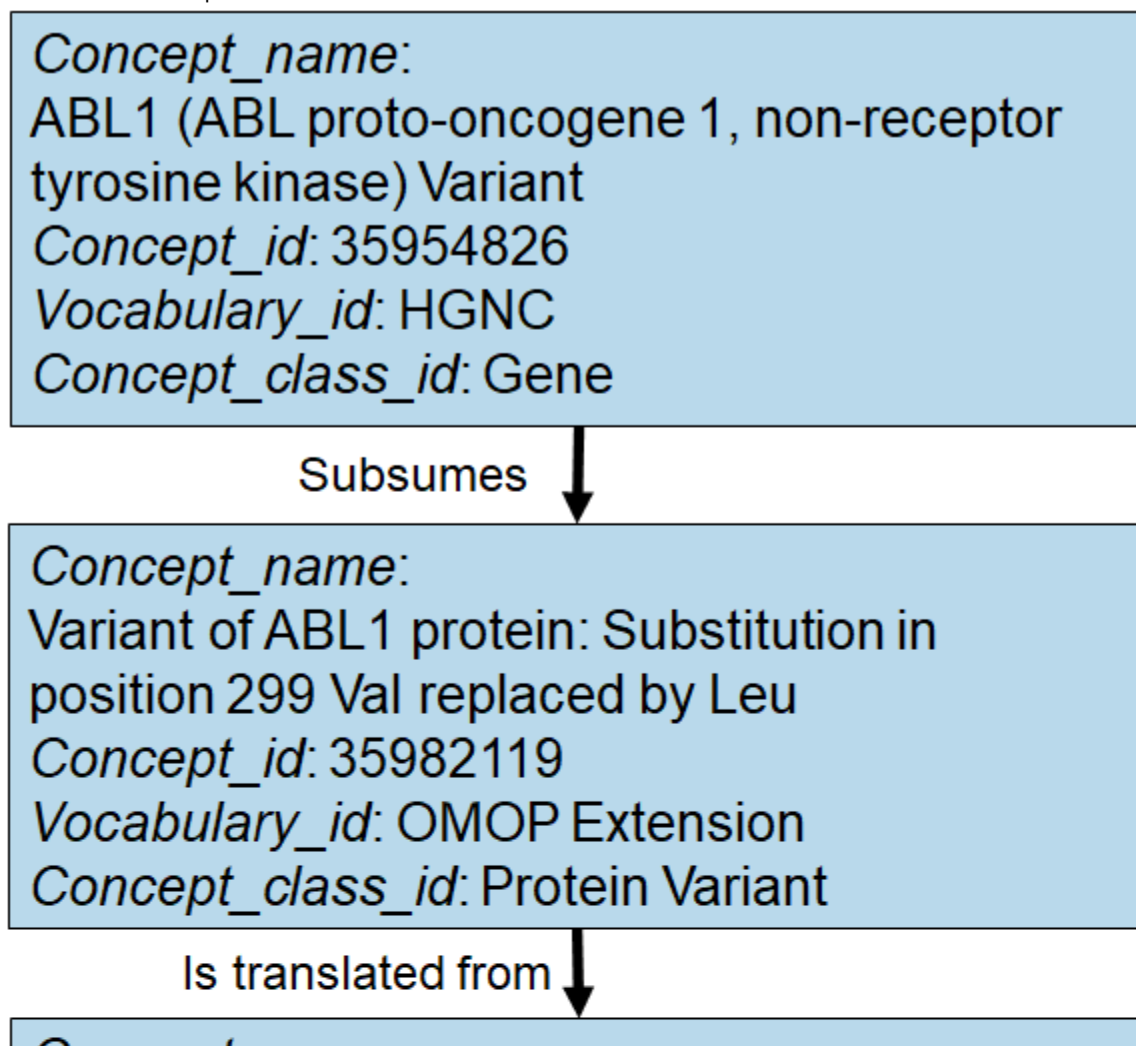
Genomic, Transcript and Protein variants were created as Standard concept in vocabulary OMOP Extension, where *concept\_code* represented as HGVS expression and *concept\_name* created from HGVS expression as human readable format.

<b>Concept_id</b>	35982287
<b>Concept_name</b>	Variant of ABL1 transcript: Substitution in position 895 of G replaced by C
<b>Domain_id</b>	Measurement
<b>Vocabulary_id</b>	OMOP Extension
<b>Concept_class_id</b>	Transcript Variant
<b>Standard_concept</b>	S
<b>Concept_code</b>	NM_005157:c.895G>C
<b>Valid_start_date</b>	22/09/2020
<b>Valid_end_date</b>	12/31/2099
<b>Invalid_reason</b>	

All terms from ontologies are mapped to Standard concepts



Standard Variants are placed into ancestor



*Concept\_name:*

Variant of ABL1 transcript: Substitution in position 895 of G replaced by C

*Concept\_id:* 35982287

*Vocabulary\_id:* OMOP Extension

*Concept\_class\_id:* Transcript Variant

Is transcribed from



*Concept\_name:*

Variant in Chromosome 9 Substitution in position 133747588 of G replaced by C

*Concept\_id:* 35981735

*Vocabulary\_id:* OMOP Extension

*Concept\_class\_id:* Genomic Variant