Names for (ISBT_102) GATA1 Alleles

General description: This gene encodes a hematopoietic-specific transcription

factor that induces high-level expression of adult betaglobin and other erythroid genes. The zinc-finger protein binds to the DNA sequence CCACACCCT found in the beta hemoglobin promoter. Heterozygous loss-of-function mutations in this gene result in the dominant In(Lu) blood phenotype. *GATA1* does not represent a blood group system, but a mutation in this gene is responsible for an X-

linked form of Lutheran-mod phenotype, XS2.

General Description (from GeneCards)

GATA1 (GATA Binding Protein 1) is a Protein Coding gene. Diseases associated with GATA1 include

Thrombocytopenia With Beta-Thalassemia, X-Linked and

Thrombocytopenia, X-Linked, With Or Without

Dyserythropoietic Anemia. Among its related pathways are

C-MYB transcription factor network and DREAM Repression and Dynorphin Expression. Gene Ontology (GO) annotations related to this gene include DNAbinding transcription factor activity and chromatin

binding.

An important paralog of this gene is *GATA2*.

Gene name: *GATA1*

LRG Ref LRG 559

Number of exons: 6

Initiation codon: within exon 2 Stop codon: within exon 6

Entrez Gene ID: 2623

LRG sequence: NG 008846.2 (genomic)

NM_002049.3 (transcript)

Antigens on reference allele n.a.

Phenotype	Allele name		Exon Intron	Predicted amino acid change	(Reference No.) PMID	Accession number	rs number
Common	GATA1*01						
XS2 Lu-mod	GATA1*BGM01	c.1240T>C	6	p.Ter414Argext*41	PMID: 23278136	none	rs587776456

References

PMID 23278136

Singleton BK, Roxby DJ, Stirling JW, Spring FA, Wilson C, Poole J and DJ Anstee. A novel GATA1 mutation (Stop414Arg) in a family with the rare X-linked blood group Lu(a-b-) phenotype and mild macrothrombocytic thrombocytopenia. Br J Haematol. 2013. 161(1):139-142.

Track of changes from version to version

1	Version			v1.0 30-JUN-2021
2 3	Author Review	created: reviewed:	n.a. n.a.	Margaret Keller unreviewed
4	General		n.a.	First Excel map version. Spread-sheets "Intro", "Allele Table", "References", and "Versioning" created.
5	End Version			v1.0 30-JUN-2021